myDNA Comprehensive Health Report





Welcome to the future of health and human potential

ID:		
Name:		
DOB:		
Barcode:		
Date:	03/07/23	



TEST METHODOLOGY AND LIMITATIONS

Recommendations in this report apply to all ages, however for any patient under 18 years, a guardian must purchase the test and be present for the report recommendations. The information in this report is not intended to treat, diagnose or cure any medical condition or disease.

Gene By Gene, a wholly owned subsidiary of myDNA, Inc., is a College of American Pathologists (CAP) accredited and Clinical Laboratory Improvement Amendments (CLIA) certified clinical laboratory qualified to perform high-complexity testing. This test was developed and its performance characteristics determined by Gene by Gene. It has not been cleared or approved by the FDA. FDA does not require this test to go through premarket FDA review. This test is used for clinical purposes. It should not be regarded as investigational or for research. Only the genomic regions listed below were tested; there is a possibility that the tested individual is a carrier for additional, undetected mutations. Although molecular tests are highly accurate, rare diagnostic errors may occur that interfere with analysis. Sources of these errors include sample mix–up, trace contamination, and other technical errors. The presence of additional variants nearby may interfere with mutation detection. Genetic counseling is recommended to properly review and explain these results to the tested individual.



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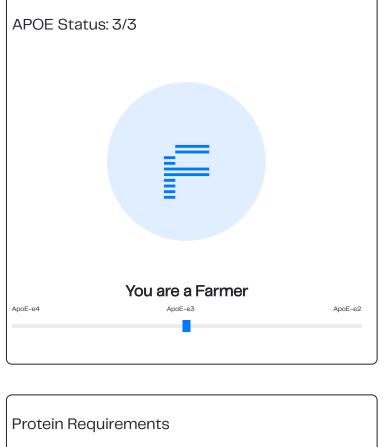
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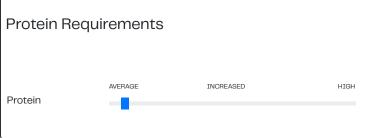
NUTRIENT METABOLISM &

DIGESTION



Apolipoprotein E (ApoE) is a lipid-binding protein that transports triglycerides and cholesterol in multiple tissues, including the brain. The e4 allele is common in hunter-gatherer communities, while the e3 and e2 alleles are most common in agricultural communities.

- ApoE 3/3 is the most common ApoE genotype found in agricultural communities and has numerous benefits
- Extended cognitive fitness and enhanced expression of anti-aging sirtuins
- Improved HDL and LDL profile
- · Improved ability to repair synapses and neural protection
- Higher viral protection
- · Higher response to plant bioactive compounds
- In the brain, e2 and e3 accumulate in neurons 2 to 4-fold higher than e4





Traditional protein intake ranges based on latitude from less than 18% of total calories to approximately 35% in the far northern climates. Recommended protein intake varies based on weight and exercise intensity.

 Genetically, your requirements fall on the average side of the spectrum, approximately 18–20% of total caloric intake

Your carbohydrate intake range is based on the latitude of your ancestors and whether a hunter–gatherer diet or modern agricultural diet made a larger imprint on your genes.

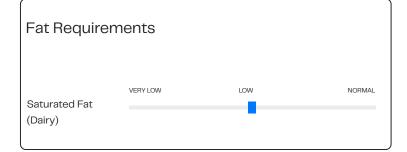
- Your genotype combination is associated with improved carbohydrate metabolism, allowing 40% to 55% of total calories from carbohydrates if desired
- For a 2,000 calorie diet, this comes to 200 to 275 grams of carbohydrates per day



The differences between responses in individuals to refined carbohydrates have been linked to a genetic adaptation occurring during the agricultural age.

Your genotype is associated with an adaptation for lowering the sensitivity to refined carbohydrates

Fat Requirements	INCREASED	нідн	The NIH has set the recommended intake of omega-3's from 1.1 to 1.6 grams per day from a combination of ALA, EPA and DHA. Omega-3 fatty acids are essential for brain, eye, and cardiovascular health. • Your genotype combinations are associated with a higher requirement of EPA and DHA
			 ApoE e2 and e3 carriers can benefit from non-phospholipid fish oil intake, however, e4 carriers should use phospholipid-based EPA and DHA as found in fish and fish roe For ApoE e4 carriers, fish oil supplements do not appear as effective as phospholipid-based EPA and DHA as found in fish and fish roe E4 carriers may have impaired transport of free DHA and require phospholipids for successful transport
Fat Requirements			Traditional total fat intake ranges based on latitude, with as low at 25% consumed in countries closer to the equator and up to 55% of total calories
			consumed in countries closer to the equator, and up to 55% of total calories from fat being consumed in northern latitudes.
AVERAGE Monounsaturated and Polyunsaturated	INCREASED	HIGH	 Genetic testing can show which fats to focus on, but total fat will range based on your climate and health goals Your genoypes are associated with a higher emphasis on monounsaturated and polyunsaturated fats from olive oil, avocados, poultry, nuts and seeds
Fat Requirements			The 2020 Dietary Guidelines in the U.S. recommends limiting calories from saturated fats to less than 10% of the total calories you eat and drink each day. That's about 200 calories for a 2,000 calorie diet. Traditionally, saturated fat intake from animal foods ranged based on the season and the geographical location, with higher latitudes and more mountainous regions consuming more.
VERY LOW Saturated Fat	LOW	NORMAL	Based on your genotype combinations, you should aim to get less



The 2020 Dietary Guidelines in the U.S. recommends limiting calories from saturated fats to less than 10% of the total calories you eat and drink each day. That's 200 calories for a 2,000 calorie diet.

saturated fat in your diet from red meatLimit your red meat consumption to twice a week

• Your genotype combinations are associated with benefiting from getting less than 22 grams of saturated fat in your diet, especially from dairy

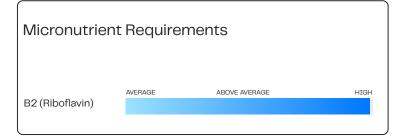
(Red Meat)



Celiac disease is an immune reaction to eating gluten, a protein found in wheat, barley and rye. Published research shows that approximately 30 percent of the general population have variants in the celiac disease risk genes HLA-DQA1 through HLA-DQB, yet only 3% of these individuals develop celiac disease.

- Your genotype combination is associated with a slight risk for celiac disease
- The diagnosis for celiac disease is more than two times higher in females than in males
- On a global level, the rates of celiac disease are not related either to the amount of wheat consumed by each country or to the prevalence of the HLA DR3-DQ2 and DR4-DQ8 genotypes worldwide
- First-degree relatives of people with celiac disease including parents, siblings and children have a 1 in 10 risk compared to 1 in 100 in the general population, which may be increased by existing autoimmune disorders
- Talk with your doctor about further testing if celiac disease is suspected





Micronutrient Requirements

B3 (Niacin)

AVERAGE

The recommended daily allowance (RDA) for thiamine is 1.2mg. Thiamine requirements are analyzed based on ethanol metabolism, however, chronic intake of alcohol depletes thiamine.

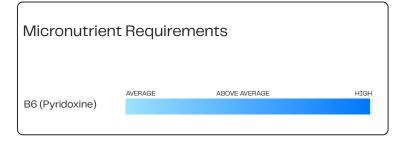
Your genotype is associated with an average need for B1

The recommended daily allowance (RDA) for riboflavin is 1.3mg. Riboflavin is used as a co-factor for numerous reactions associated with protein, fat, and carbohydrate metabolism. Riboflavin requirements are analyzed based on MTHFR gene function.

- Your genotype is associated with a higher than average need for B2
- B2 is high in liver (2.8mg), lamb (0.4mg), salmon (0.8mg), yogurt (0.6mg) and oyster mushrooms (0.3mg)

The recommended daily allowance (RDA) for niacin is 16mg. Niacin targets genes associated with cardiovascular and skin health, while also balancing methylation levels.

- Your genotype is associated with a higher sensitivity to low niacin intake
- Niacin in high in yellowfin tuna (37.5mg), canned tuna (21.9mg), wild salmon (17mg), ground turkey (20mg), chicken breast (16mg), liver (14.2mg), skirt steak (9.5mg), white button mushrooms (6.8mg), and brown rice (5.2mg)



ABOVE AVERAGE

HIGH

The recommended daily allowance (RDA) for B6 is 1.7mg. B6 deficiency can manifest as anorexia, irritability, anxiety, depression, muscle pain, bad PMS/low progesterone, nausea, seizures, migraines, dermatitis, age related macular degeneration (with low folate and B12) and lethargy.

- Your genotype is associated with a higher than average need for B6
- Women of reproductive age, especially current and former users of oral contraceptives, teenagers, male smokers, non–Hispanic African– American men, and men and women over age 65 are most at risk of B6 deficiency
- B6 is high in yellowfin tuna (6 oz., 1.8mg), wild salmon (6 oz., 1.2mg), liver (3oz., 0.8mg), chicken breast (6 oz., 1mg), unfiltered fermented drinks (16oz., 0.8mg), pistachios (1 oz., 0.5mg), avocado (1 whole, 0.5mg), sweet potatoes (1 whole, 0.3mg), and spinach (1/2 cup, 0.1mg)

Micronutrient	Requireme	nts	
B9 (Folate)	AVERAGE	ABOVE AVERAGE	HIGH

The recommended daily allowance (RDA) for folate is 400mcg. Folate is one of the – if not most – influential nutrigenomic micronutrient. It has a powerful influence on genes related to pregnancy, homocysteine, and cancer prevention.

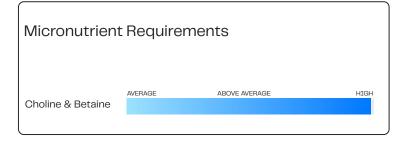
- · Your genotype is associated with a higher than average need for folate
- Folate is depleted by proton pump inhibitors, oral contraceptives, NSAIDs, anticonvulsants, antivirals, antibiotics, and antacids
- Folate is high in liver (3 oz., 215mg), collard greens (1 cup cooked, 177mcg), beets (1 cup raw, 148mcg), black-eyed peas (1/2 cup, 105mg), raw spinach (1 cup 58mg), asparagus (4 spears, 89mg), hummus (1/2 cup, 83mcg), broccoli (1/2 cup cooked, 52mg), romaine lettuce (1 cup, 64mg), strawberries (1 cup, 40mcg), orange (1 whole, 39mcg), sprouted lentils (1/2 cup, 38mcg), and parsley (1 sprig, 15.2mg)



The recommended daily allowance (RDA) for B12 is 2.4mcg. B12 influences genes related to homocysteine, brain health, pregnancy, and energy. B12 requirements are based on serum levels associated with the FUT2 gene.

Your genotype is associated with an average requirement for B12



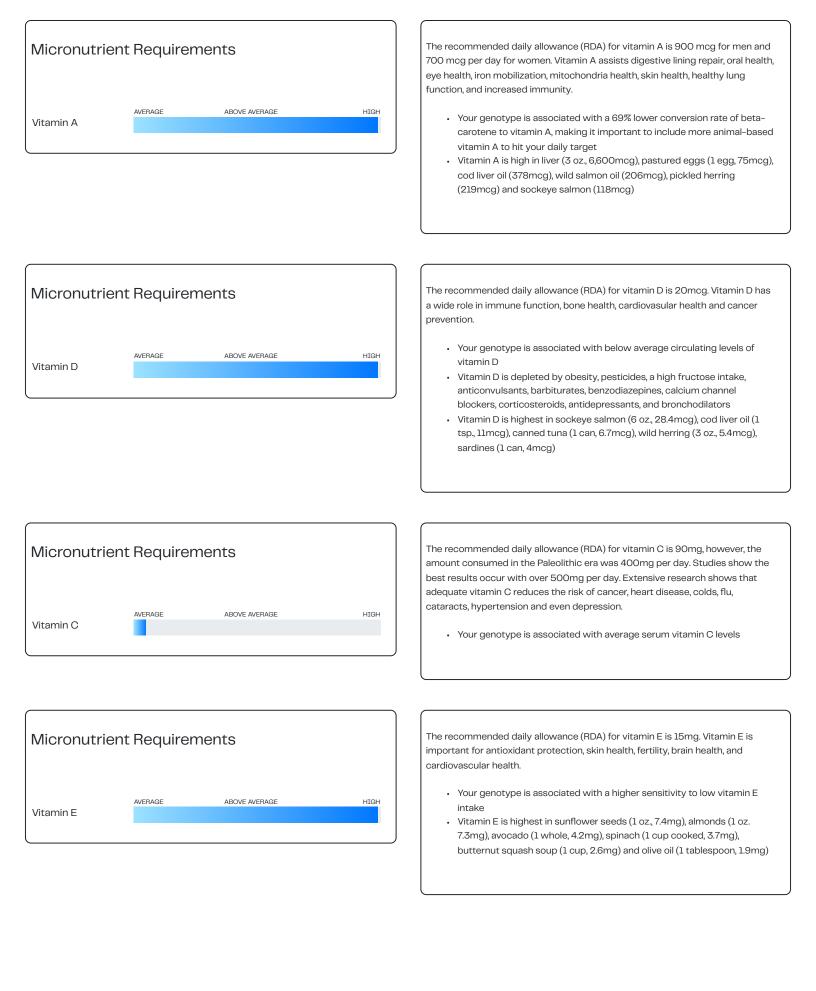


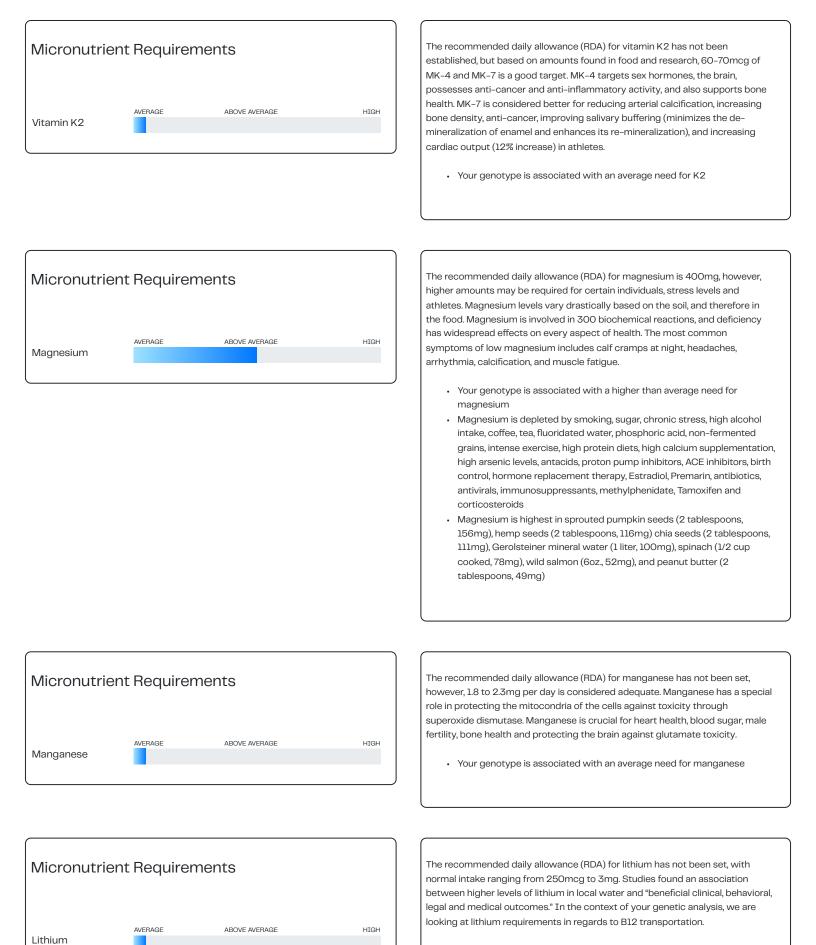
The recommended daily allowance (RDA) for boron has not been set, but 1–3mg is considered adequate. Boron is connected to bone health, hormone health and healthy SAMe levels for brain health. Men with low testosterone and women with osteoporosis or osteopenia will benefit from more boron.

• Your genotype is associated with an average need for boron

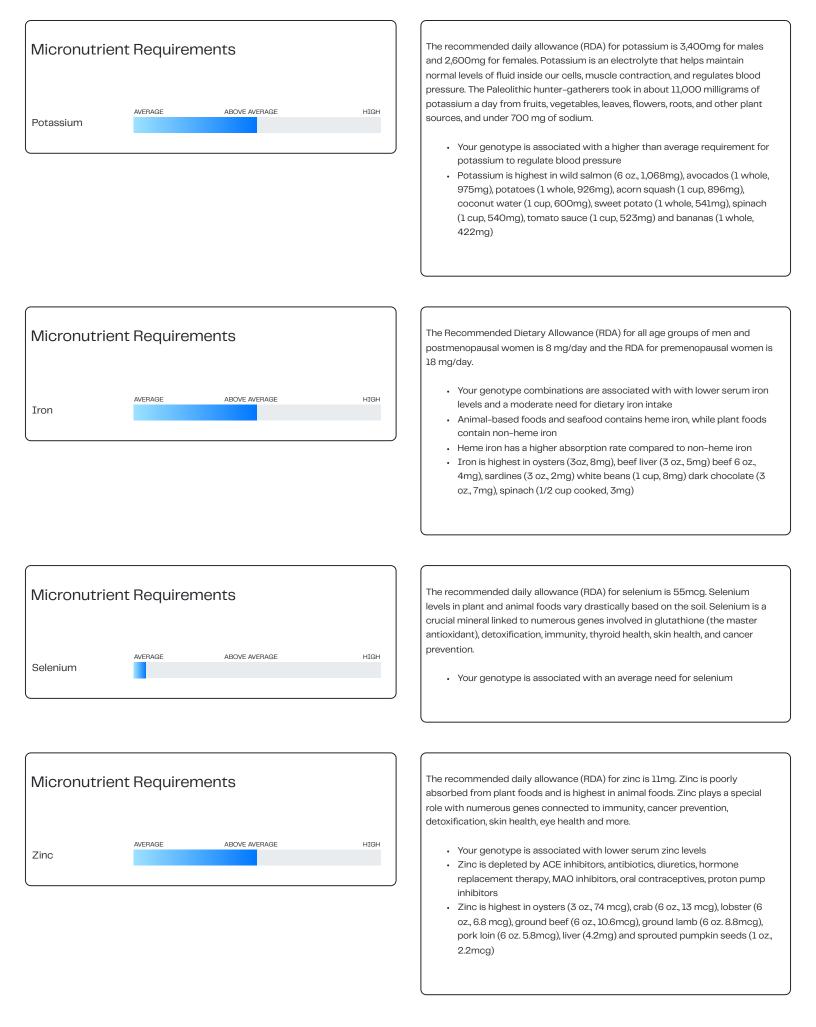
The recommended daily allowance (RDA) for choline is 550mg, while betaine hasn't been set. The more betaine you consume, the less choline you require. Choline is crucial for pregnancy, lowers anxiety, prevents fatty liver, assists detoxification, and improves memory.

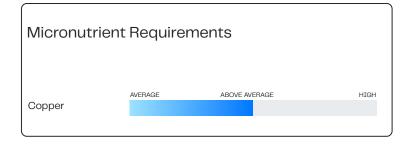
- Your genotype is associated with a higher than average need for choline
 and betaine
- Choline is depleted by nighttime pain relievers, antihistamines, sleep aids, antidepressants, incontinence drugs and narcotic pain relievers
- Intense endurance exercise depletes choline levels, and increasing
 phosphatidylcholine has been found to improve exercise capacity during
 high-intensity cycling and running, as well as reduce muscle soreness
- Choline is highest in liver (3 oz., 356mg), pastured eggs (2 eggs, 294mg), beef round (6 oz., 234mg), heart (3 oz., 194mg), chicken (6 oz., 144mg), wild cod (6 oz., 142mg), bacon (3.5 oz., 125mg), and edamame (1/2 cup, 107mg)
- Betaine is highest in spinach (3.5 oz., 645mg), shrimp (3.5 oz., 218mg), beets (3.5 oz., 200mg) and whole grain sourdough wheat bread (2 slices, 201mg)





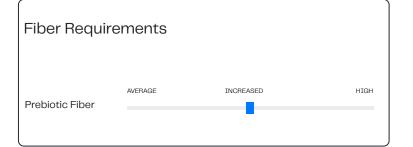
· Your genotype is associated with an average need for lithium





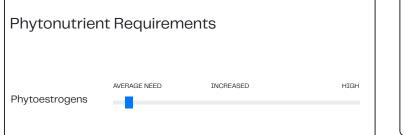
Typical diets meet or exceed the copper RDA and copper deficiency is rare. The RDA for copper is 900mcg. Copper is involved in the regulation of gene expression, brain development, neurotransmitters, cardiovascular health, and immune system functioning.

- Your genotype is associated with slightly lower serum copper levels
- Copper is highest in liver (3 oz. 12,400mcg), oysters (4,850mcg), potatoes (675mcg), shiitake mushrooms (1/2 cup, 650mcg), cashew (1 oz., 629mg), sunflower seeds (1/4 cup, 615mg) and dark chocolate (1 oz., 615mcg)



The recommended amount of fiber is up to 25 grams per day for women and up to 38 grams per day for men.

Your genotypes are associated with a slighter higher than average requirement for prebiotic fiber



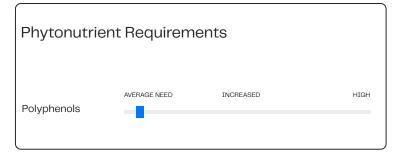


Phytoestrogens are plant derived compounds found in a wide variety of foods. There are pros and cons to phytoestrogen intake that appears to have a genetic, age, and gut health connection for determining optimal intake.

Your genotype combinations are associated with a low to average need for phytoestrogens for healthy hormones

A recommended daily intake of lutein and zeaxanthin hasn't been established. Lutein and zeaxanthin can help protect your eyes from harmful high-energy light waves like UV sunlight.

- Your genotype is associated with an increased need for foods high in lutein and zeaxanthin to support eye health
- Around 700 carotenoids have been discovered and only lutein and zeaxanthin are found in the eye
- American adults typically consume 1–3 mg/day of lutein and zeaxanthin, the Spanish consume 3.5 mg/day, the Germans consume 5.33 mg/day, and older Australians consume 0.9mg per day
- For reducing the risk of eye disorders, the estimated target is 6mg or more of lutein and zeaxanthin daily
- The foods highest in lutein and zeaxanthin include cooked spinach (1/2 cup, 12.64 mg lutein), raw spinach (1/2 cup, 6.6mg lutein), cooked kale (1/2 cup, 8.88mg lutein), egg yolks (1 egg, 237mcg lutein and 216mcg zeaxanthin), and orange peppers (208mcg lutein and 1665mcg zeaxanthin)



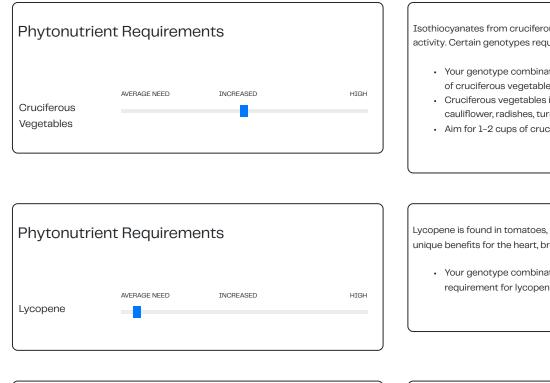
Research strongly suggests that long term consumption of diets rich in plant polyphenols offer protection against development of cancers, cardiovascular diseases, diabetes, osteoporosis and neurodegenerative diseases.

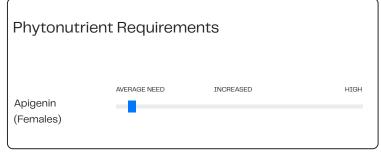
 Your genotype is associated with a slower metabolism of certain polyphenols, which means you have a higher benefit with a lower intake of green tea, coffee, berries, and chocolate

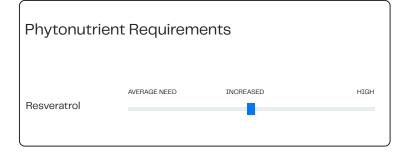
Phytonutrient Requirements

Cinnamon lowers blood glucose usually without causing hypoglycemia and increases satiety.

Your genotype is associated with an average need for cinnamon to control blood sugar







Isothiocyanates from cruciferous vegetables are known for their anti-cancer activity. Certain genotypes require higher levels of this anti-cancer activity.

- · Your genotype combinations are associated with a higher requirement of cruciferous vegetables
- · Cruciferous vegetables include broccoli, Brussels sprouts, cabbage, cauliflower, radishes, turnips, Bok choy, and watercress
- · Aim for 1-2 cups of cruciferous vegetables per day

Lycopene is found in tomatoes, watermelon, guava and pink grapefruit, and has unique benefits for the heart, breast, prostate and skin.

· Your genotype combinations are associated with a lower to average requirement for lycopene

Apigenin is a flavonoid that possess anti-inflammatory, antioxidant and anticancer properties. Certain genotypes require higher levels for breast health.

· Your genotype is associated with an average need for apigenin for breast health

Resveratrol is a plant chemical produced in response to stress from the elements, and has been found to protect against heart disease and potentially extend life.

- · Your genotype combinations are associated with a higher than average requirement for resveratrol and heart health
- · Resveratrol has been found to increase PON1 and ACE2 levels, decreasing the risk of LDL oxidation, the negative effects of a high fat intake, and improving blood pressure
- · Resveratrol is highest in grapes, peanuts, pistachios, blueberries, cranberries and dark chocolate



Lactose is the major carbohydrate in milk. The arrival of farming in Europe around 8,500 years ago necessitated adaptation to new environments, pathogens, diets, and social organizations. One of the best examples of genetic dietary changes to this is the lactase enzyme in northern Europeans that only dates to the last 4,000 years.

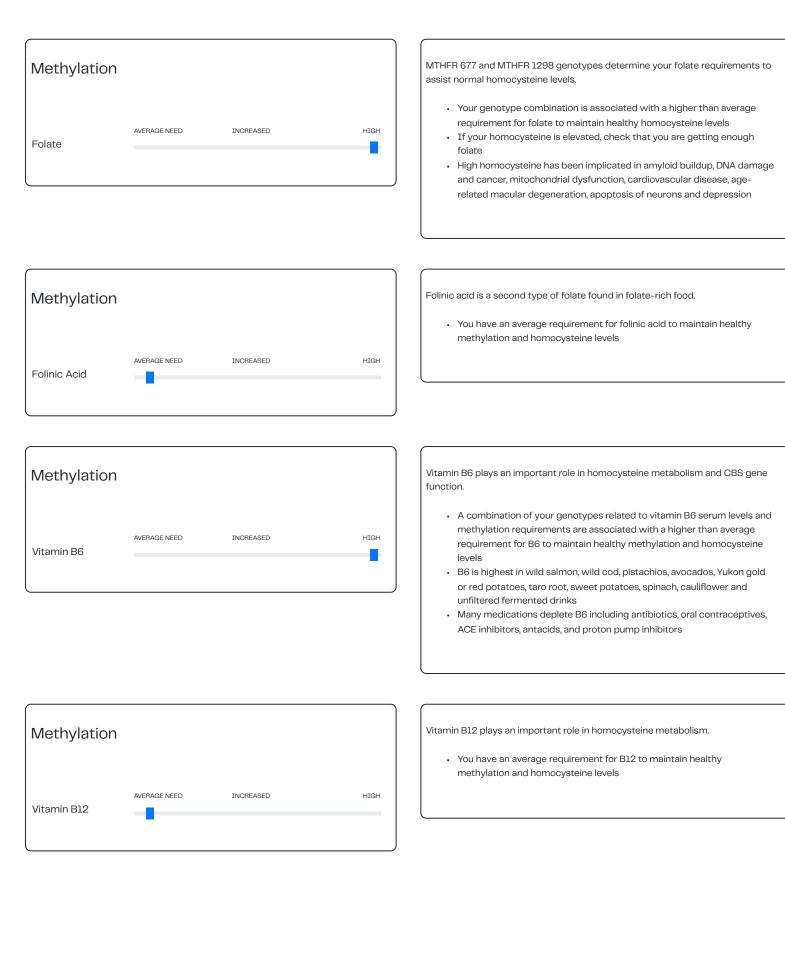
- · Your LCT genotype is associated with lactose tolerance
- The ability to digest lactose is much more common in people of European ancestry
- Approximately 32 percent of the world's population is lactose tolerant
- Since this gene only looks at lactose, sensitivities to dairy can still exist

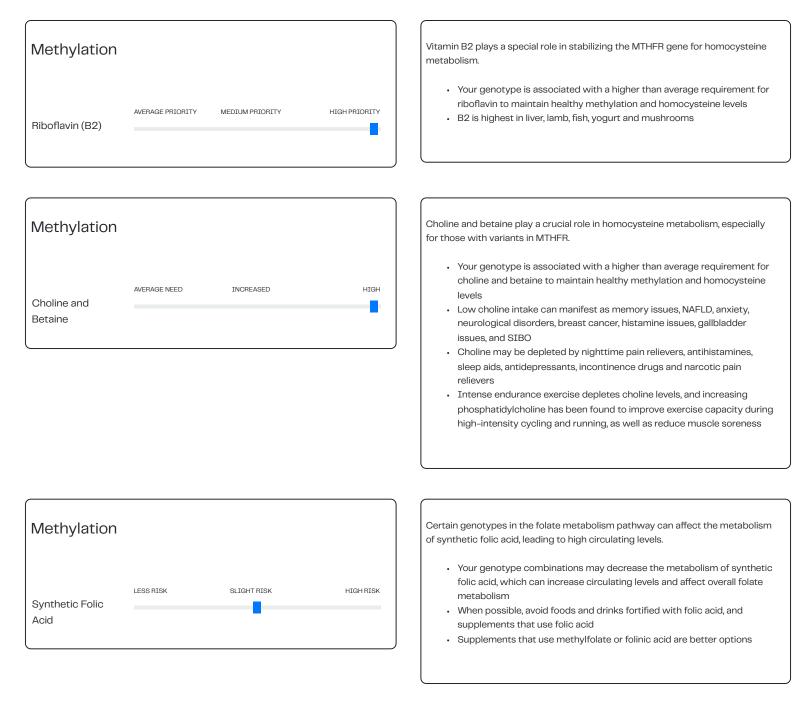


Variants in the CYP1A2 gene determine the rate at which you metabolize caffeine.

• You are a fast metabolizer of caffeine, meaning that you feel the effects of caffeine more quickly, but the effects also wear off quickly













Certain glutathione SNPs are associated with breast protection.

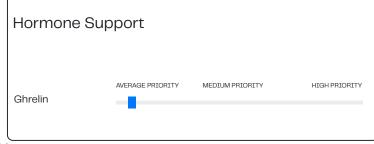
- Your genotypes for multiple genes are associated with slightly lower glutathione protection for breast health
- Boosting glutathione can be accomplished with selenium, glycine, cysteine, vitamin C, and cruciferous vegetables

There are multiple genes in the estrogen detoxification pathway that have a cumulative value on the ability to properly detoxify estrogen.

- Your combination of numerous genotypes in the estrogen pathway are
 associated with reduced estrogen detoxification
- To reduce the risk of harmful estrogen metabolites, you should avoid xenoestrogens, manage stress levels, and focus on gut health
- Increasing prebiotic fiber, polyphenols, magnesium and bifidobacteria may improve breast health by reducing the amount and activity of harmful estrogen metabolites







T3 and T4 level variations have been associated with variants in the DIO1 gene.

- · Your genotype is associated with normal T3 and T4 levels
- T3 and T4 can still be out of range based on other epigenetic factors

ADIPOQ encodes for adiponectin, a protein secreted by fat cells that affect insulin and glucose metabolism. Low levels of adiponectin play a role in obesity, insulin resistance and Type 2 diabetes.

 Your genotype is associated with normal adiponectin levels, which can increase the effect of insulin, improve glucose metabolism and assist a healthy body weight

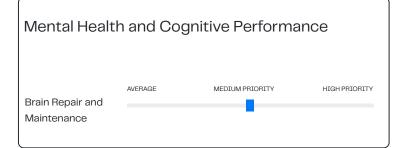
Variants in genes related to ghrelin levels and dopamine receptor density have been shown to create a larger appetite and the potential for overeating in multiple populations.

- Your genotypes are not associated with higher ghrelin levels
- You are at a decreased risk for overeating and abdominal weight gain



MENTAL HEALTH & COGNITIVE

PERFORMANCE



Mental Health and Cognitive Performance

MEDIUM PRIORITY

HIGH PRIORITY

AVERAGE

Episodic Memory

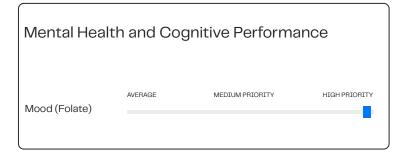
Multiple genes are responsible for daily neural repair and maintenance, and a combination of genotypes are associated with decreased neural repair.

- Your genotype combination is associated with slightly reduced neural repair, which can affect healing from brain injuries and amplify damage from poor sleep patterns
- · Limit or avoid activities with a high risk of concussions
- · Get eight hours of sleep per night for optimal repair
- Be proactive with neural repair by focusing on safe endurance exercise, DHA, B-vitamins, Lion's Mane mushroom, zinc, vitamin C, and vitamin E

The 5-HT2A gene is associated with episodic memory, which is the ability to recall details of an event.

- Your 5-HT2A genotype is associated with a reduced episodic memory
- Episodic memory can be enhanced with tryptophan, green or black tea, catecholamines, betaine, and choline
- Mindfulness training, increasing your VO2 max and reducing your reliance on smartphones can also enhance episodic memory

Mental Health and Cognitive Performance

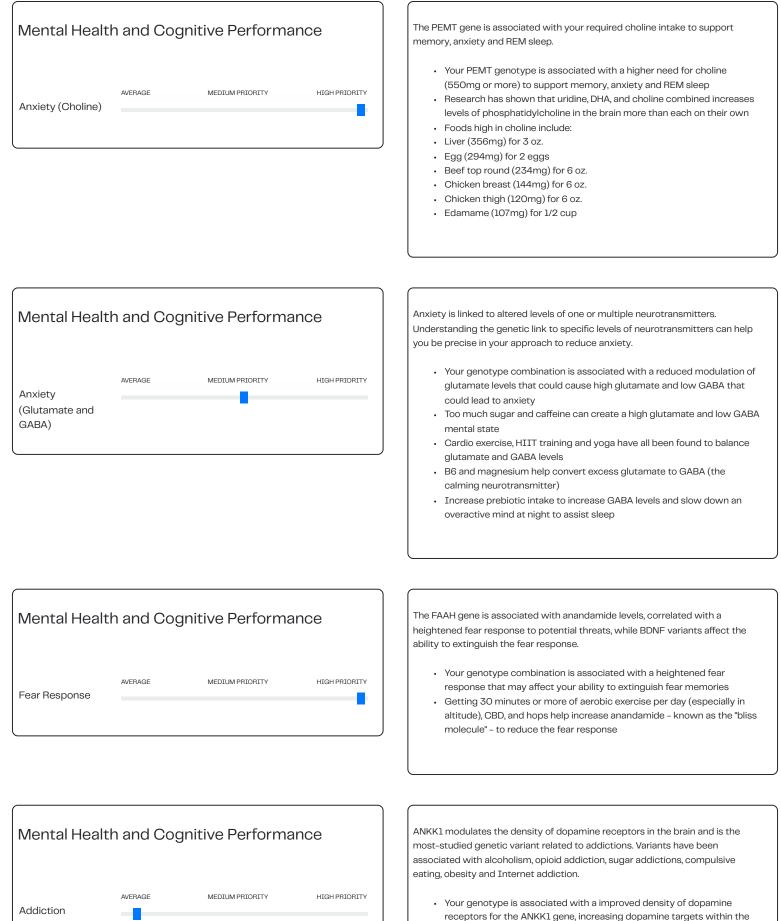


Cardiovascular exercise has a tremendous influence on neurotransmitter balance, memory and cognitive fitness.

- You have a higher than average requirement for cardiovascular exercise to improve mood and cognitive fitness
- Research shows that at least 30 minutes of cardio, 5 times a week, is the target to hit for improved mood and cognitive fitness

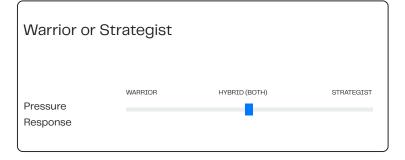
MTHFR genotypes determine folate requirements for healthy BH4 levels responsible for neurotransmitter balance. The current daily value for folate is 400mcg DFE.

- Your genotype may require 400–600mcg (or more) for healthy BH4 levels responsible for neurotransmitter balance
- Foods high in folate include:
- Liver (215mg) 3 oz.
- Spinach (131mg) 1/2 cup cooked
- Asparagus (89mg) 4 spears
- Brussels sprouts (78mg) 1/2 cup
- Broccoli (52mg) 1/2 cup



striatum of the brain and reducing addiction susceptibility

Mental Health and Cognitive Performance

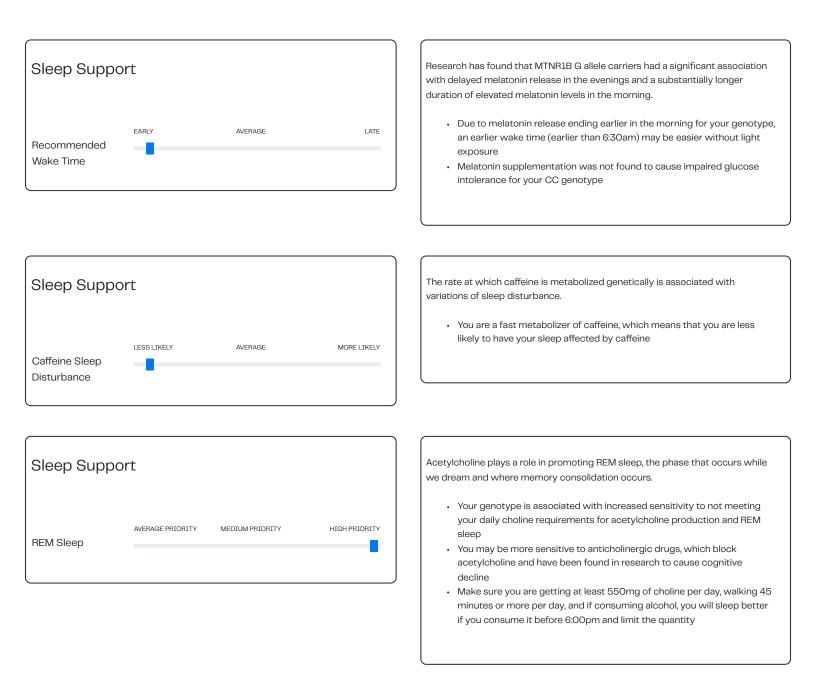


COMT rs4680 has been linked in a meta-analysis to variations in the reward response based on genotypes associated with low and high dopamine levels.

 Your genotype is associated with a slightly higher dopamine response to reward processing that could improve motivation as well as decision making

Your COMT genotype is associated with the "Hybrid" that has intermediate to high levels of dopamine and may be able to express the Warrior or Strategist traits depending on life experience. Higher dopamine is useful in complex environments that require maximal performance in terms of memory and attention for survival.

- If your levels of dopamine get too high and you find yourself irritable, impulsive, and stressed, add strength training 3–5 times a week and increase your magnesium and vitamin C intake for balance.
- Average to low catecholamine intake recommended (coffee, green tea, berries, chocolate)
- For men and premenopausal women, avoid IPA beers due to a higher
 estrogenic effect that can slow COMT down further







Toxin Sensitiv	vity		Mycotoxins are toxic compounds that are naturally produced by certain types of fungi. Research suggests that mycotoxins can decrease the formation of glutathione due to decreased gene expression of the enzymes needed to form glutathione.
Mycotoxins	AVERAGE PRIORITY MEDIUM PRIORITY	HIGH PRIORITY	 Your genotype is associated with lower glutathione levels which may cause glutathione depletion to occur at a faster rate and decrease mycotoxin detoxification The highest exposure to mycotoxins can be in foods grown or stored in damp conditions This may include grains, nuts, corn, coffee, wine, beer, grape juice, sorghum, rice, dried beans, apples, pulses, cacao products, and spices Boosting glutathione can be accomplished with selenium, glycine, cysteine, alpha lipoic acid, vitamin C, and cruciferous vegetables
Toxin Sensitiv	vity		Xenoestrogens are synthetic hormone disruptors found in plastics and pesticides.
Xenoestrogens	AVERAGE PRIORITY MEDIUM PRIORITY	HIGH PRIORITY	 Your genotype is associated with a slower metabolism of xenoestrogens, and therefore the damage may be greater from xenoestrogen exposure Increasing magnesium targets the enzyme responsible for assisting xenoestrogen detoxification
Toxin Sensitiv	vity		Workers exposed to certain chemicals over a long period in the metalworking, petroleum, agricultural industries and in glass factories are at increased risk for occupational skin cancers.
Occupational Toxins	AVERAGE PRIORITY MEDIUM PRIORITY	HIGH PRIORITY	 Your genotypes are associated with an increased sensitivity to these toxins Focus on zinc, selenium, niacin, and vitamin C to improve DNA protection for skin health Ellagic acid, lutein, zeaxanthin, cocoa polyphenols, chaga tea, green tea and citrus have all been found to help protect against skin damage and cancerous growth
Toxin Sensitiv	vity		ALDH2 encodes for aldehyde dehydrogenase, and variants can affect the levels of acetaldehyde and therefore the carcinogenic effect of alcohol.
Ethanol	AVERAGE PRIORITY MEDIUM PRIORITY	HIGH PRIORITY	 Your genotype is not associated with a higher risk of alcohol-related adverse reactions including flushing, palpitation, nausea, headache, drowsiness, breathlessness, and general discomfort

Toxin Sensitiv	ʻity		
Formaldehyde	AVERAGE PRIORITY	MEDIUM PRIORITY	

The International Agency for Research on Cancer has classified formaldehyde as carcinogenic to humans. Sources of formaldehyde in the home include building materials, smoking, household products, gas stoves, kerosene space heaters, as a food preservative, permanent–press clothes, and draperies, as a component of glues and adhesives, and as a preservative in some paints and coating products. Variants in the XRCC3 DNA repair gene have been associated with higher DNA damage from formaldehyde.

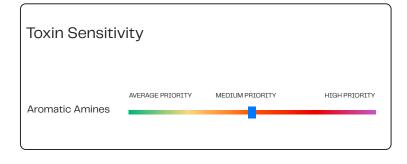
- Your genotype is associated with an increased sensitivity to formaldehyde
- Formaldehyde can cause watery eyes, burning sensations in the eyes and throat, nausea, and difficulty in breathing in some humans exposed at elevated levels
- Formaldehyde has also been shown to cause cancer in animals and may cause cancer in humans
- Average concentrations in older homes before the 1970's are generally well below 0.1 (ppm), but in homes with significant amounts of new pressed wood products, levels can be greater than 0.3 ppm
- The detoxification pathway for formaldehyde includes the thiol pathway, ribulose monophosphate pathway, and pterin dependent pathway
- B-vitamins, zinc, selenium, vitamin C, cysteine, glycine and glutamine all target these pathways and DNA repair to improve formaldehyde detoxification

 AVERAGE PRIORITY
 MEDIUM PRIORITY
 HIGH PRIORITY

 Benzo(a)pyrene
 Image: constrainty
 Image: constrainty

Benzo(a)pyrene is a carcinogenic compound produced from the burning of wood or trash, tobacco smoke, asphalt, coal, diesel exhaust, charred meat, and gas cooking.

- Your genotype combinations are associated with decreased detoxification of benzo(a)pyrene
- It is recommended to increase your intake of cruciferous vegetables, vitamin C, vitamin E, vitamin A, resveratrol, curcumin, green tea, and white tea to protect and detoxify benzo(a)pyrene



Toxin Sensitivity

 AVERAGE PRIORITY
 MEDIUM PRIORITY
 HIGH PRIORITY

 Benzene
 Image: Comparison of the priority
 HIGH PRIORITY

Aromatic amines are found in cigarettes, rubber factories, hair dyes that contain 4-aminobiphenyl, and meat cooked at high temperatures.

- Your genotype combinations are associated with a slightly below
 average detoxification ability of aromatic amines
- If your exposure is higher to aromatic amines, increase cruciferous
 vegetable intake, carotenoids, vitamin C, and use marinades for meat
 when barbecuing

Sources of benzene include cigarette smoke, gasoline, exhaust, fires, industrial emissions, paint, detergent, glue, furniture wax, building materials, paint, petrochemical manufacturing, oil storage tanks, urban-industrial areas, service stations, certain foods, certain creams, groundwater contamination, and underground gasoline leaks.

· Your genotype is associated with an average sensitivity to benzene

Toxin Sensitiv	vity		
Aspartame	AVERAGE PRIORITY	MEDIUM PRIORITY	HIGH PRIORITY

Aspartame is an artificial sweetener that has been linked to behavioral, neurological and cognitive problems, increased blood sugar, thyroid issues, and certain types of cancer.

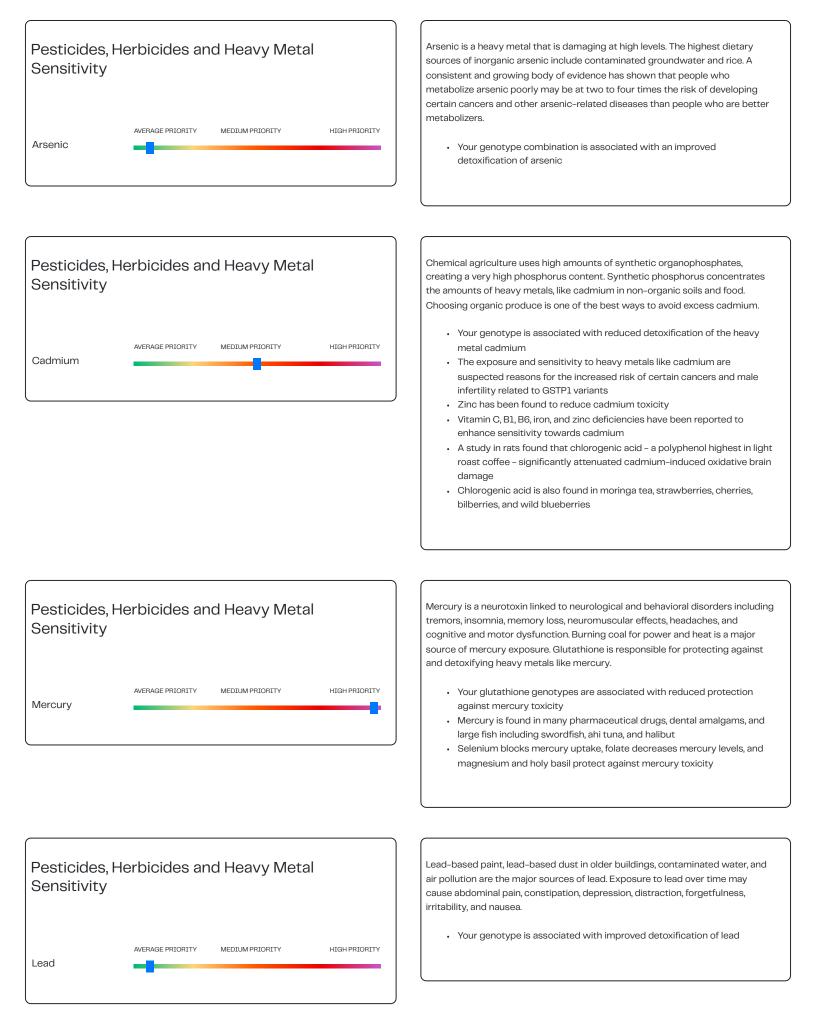
- Your genotype combinations may increase the sensitivity to aspartame
- Possible neurophysiological symptoms include learning problems, headache, seizure, migraines, irritable moods, anxiety, depression, and insomnia
- Artificial sweeteners in general increased waist circumference 500
 percent while aspartame increased blood sugar in diabetes-prone mice
- Aspartame has been found to contribute to the formation of tumors in the CNS such as gliomas, medulloblastomas, and meningiomas, increased lymphoma and leukemia and, is an excitotoxin to brain neurons
- Aspartame in the body further metabolizes to formaldehyde, and rat studies found that formaldehyde (as a metabolite of aspartame) caused increased TSH levels and worsens the capacity of the gland leading to thyroid failure



Food dyes have been found to inhibit mitochondrial respiration: the ability of the powerhouse of your cells to convert nutrients to energy. They have also been found to especially affect those with ADHD.

- Your genotype combination is associated with a higher sensitivity to food dyes
- Avoid foods and drinks that use food dyes when possible

Pesticides, Herbicides and Heavy Metal Sensitivity AVERAGE PRIORITY MEDIUM PRIORITY Glyphosate	 Glyphosate is an herbicide that has been found to be highly toxic. Your genotype is associated with average cellular protection against the herbicide glyphosate The highest glyphosate levels have been found in non-organic wheat and non-organic pulses like beans, lentils, and peas A meta-analysis of human epidemiological studies suggests a link between exposures to glyphosate and an increased risk for non-Hodgkin's lymphoma An association between glyphosate and thyroid disease comes from plots over time of the usage of glyphosate in the U.S. on corn and soy time-aligned with plots of the incidence rate of thyroid cancer in the U.S. Manganese deficiency and toxicity can occur simultaneously from glyphosate exposure due to a disruption in liver enzymes, causing transportation of manganese through the vagus nerve to the brainstem where excess manganese can lead to Parkinson's disease The gut bacterium Lactobacillus is negatively impacted by glyphosate and the depletion in associated with celiac disease Humic acid from Shilajit has been shown in vivo to reduce glyphosate concentration, inhibit the destructive effect of glyphosate on beneficial bacteria, and protect and repair against tight junction injury of the digestive system
Pesticides, Herbicides and Heavy Metal Sensitivity	Organochlorines are found in certain pesticides, PCBs and sucralose. • Your genotype is associated with improved protection against organochlorines
Organochlorines	PON1 plays a large role in removing pesticides and is also involved with
Pesticides, Herbicides and Heavy Metal Sensitivity	HIGH PRIORITY Supporting HDL function and LDL oxidation. Organophosphates are a class of insecticides, including parathion and chlorpyrifos, that were among the most widely used insecticides available until the 21st century. • Your PON1 genotype is associated with reduced PON1 levels and detoxification of organphosphate insecticides
Organophosphate Insecticides	 Organophosophate insectides work by damaging an enzyme in the body called acetylcholinesterase Residential proximity to agricultural organophosphate application is associated with faster cognitive and motor symptom decline among Parkinson's disease patients Reduce exposure to pesticides, get adequate calcium and magnesium, and consume pomegranates, broccoli sprouts, and high quality olive oil to increase PON1 levels





IMMUNE SUPPORT,

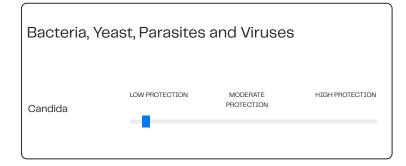
ANTIOXIDANTS AND

INFLAMMATION

Bacteria, Yeast, Parasites and Viruses

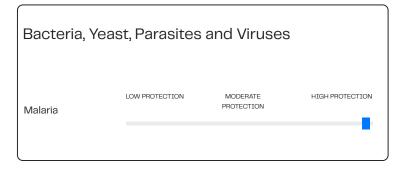
The inactive "non-secretor" genotype for FUT2 confers resistance to H. Pylori. H. Pylori is present in approximately 50% of the population in developed countries.

- You have the non-secretor genotype for FUT2, associated with a decreased susceptibility to H. Pylori
- H. Pylori inhibition has been demonstrated with alcohol extracts of the mushroom Lion's Mane



The inactive "non-secretor" genotype for $\ensuremath{\mathsf{FUT2}}$ decreases resistance to Candida overgrowth.

- You have the non-secretor genotype for FUT2, associated with an increased susceptibility to Candida overgrowth
- Approximately 70 percent of Candida overgrowth occurs in women
- Blood sugar control with a low glycemic load and balance of protein, fat, fiber and carbohydrates is key for both sexes
- In women, the focus should be on balancing estrogen and progesterone levels



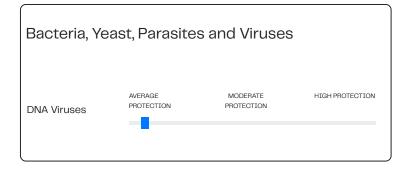
Research has shown that MTHFR genotypes influence T-lymphocytes, natural killer cells, and protection against malaria.

- Your genotype is associated with boosted levels of thymidine, which may increase lymphocyte replication and immune function in response to malaria
- The malaria parasite needs higher amounts of folate to survive and replicate
- For malaria-endemic regions, a homozygous MTHFR 677 genotype is superior due to lower folate levels

Bacteria, Yeast, Parasites and Viruses

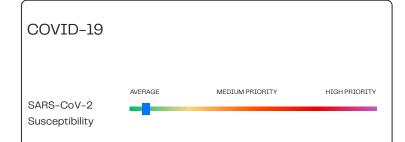
The inactive "non-secretor" genotype for FUT2 confers resistance to the Norovirus.

You have the non-secretor genotype for FUT2, associated with a decreased susceptibility to the Norovirus



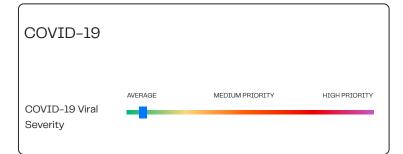
DNA viruses include HPV, Epstein Barre, herpes, and smallpox. Folate is a precursor to BH4 to produce nitric oxide. Nitric oxide acts as an antiviral that is more potent against DNA viruses.

- Your genotype combination is associated with lower BH4 levels with insufficent folate, lowering protection against DNA viruses
- Low BH4 affects the aggressiveness of DNA viruses
- To increase BH4, include foods high in folate, vitamin C, L-arginine, B6, magnesium, and selenium for healthy nitric oxide levels and DNA virus support
- BH4 is depleted by high blood sugar, high omega-6 intake, chronic stress, high levels of mercury, arsenic, lead and aluminum, aspartame, and oxidative stress



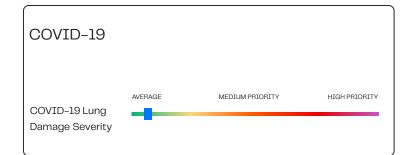
Genome-wide association studies have identified a region of chromosome 3p21.31 as the for conferring susceptibility to infection with LZTFL1 as the candidate gene. ApoE-e4, ACE2 and TMPRSS2 polymorphisms have been shown to be strongly associated with the susceptibility, severity, and clinical outcomes of COVID-19.

- Your genotype combination is associated with a reduced probability to SARS-CoV-2 infection
- Advanced age, obesity, and being male are considered the top risk factors for SARS-CoV-2 susceptibility, especially when combined with Type 2 diabetes, high blood pressure, and cardiovascular disease
- Research has shown that CBD, Chaga mushroom, birch bark and olive oil may stop SARS CoV-2 entry by helping block the "lock" for viral entry
- The flavonols kaempferol, quercetin, myricetin, fisetin and their derivatives were the most documented molecules with antiviral activities against SARS-CoV-2
- Propolis has antiviral activity and inhibitory effects on ACE2, TMPRSS2 and PAK1 signaling pathways used by SARS-CoV-2, while promoting immunoregulation of pro-inflammatory cytokines, and reducing the risk of cytokine storm syndrome



ACE2 (the receptor for SARS-CoV-2) is reduced in individuals that are carriers of ApoE4. Once the virus fuses with the cellular membranes, it takes control, shuts down more ACE2 receptors, closing the door behind it. Fewer ACE2 receptors lead to elevated angiotensin II levels, which increases the viral load. This leads to a more severe infection, NF-kb activation, lung damage, and viral replication.

- An analysis of your ApoE genotype and numerous genes involved in the viral inflammatory pathways show an association with a reduced risk of viral severity
- The risk of viral severity can still be increased by existing health issues including Type 2 diabetes, obesity, high blood pressure, and cardiovascular disease
- In September 2020, Boston University found that SARS-CoV-2 is highjacking the NF-kB pathway and should be a target for suppressing viral aggressiveness
- In addition to vaccination, NF-Kb inhibitors can be both preventative for inflammation and successful with slowing viral aggressiveness
- These include vitamin C, vitamin D, zinc, selenium, magnesium, cordyceps mushrooms, resveratrol, triterpenoids (Chaga, reishi, olive oil, holy basil), caffeic acid (coffee, Chaga, elderberry), and anthocyanins (elderberry, goji berries, cacao)
- The dietary flavonols kaempferol, quercetin, myricetin, fisetin were the most documented molecules with antiviral activities against SARS-CoV-2, and had a broad spectrum of biological activities that could reduce the severity of infection symptoms and enhance the immune response
- The strong association between air pollution exposure and COVID-19 mortality suggests that inhaled noxious particles influence COVID-19 outcomes



COVID-19

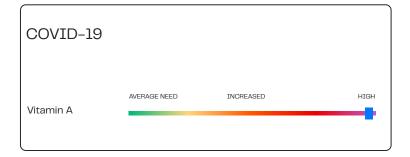
AVERAGE NEED INCREASED HIGH
Glutathione

Scientists at the University of Oxford published results in November 2021 that variants in the LZTFL1 gene doubles the risk of lung failure and death from COVID-19. Approximately 60% of people with South Asian ancestry, 15% of people with European ancestry, 2% of people with African-Caribbean ancestry and 1.8% of people with East Asian ancestry carry the high-risk variant. Scientists found that the LZTFL1 gene high-risk variant affects the lungs, but does not have an impact on the immune system.

Your genotype is not associated with an increased risk COVID-19
related lung damage severity

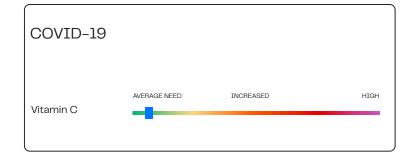
Glutathione is the master antioxidant system involved in oxidative stress, detoxification, and immunity. The functional capacity of immune cells and the ability to cope with oxidative stress has been proposed as one of the significant markers of health and longevity. In both animals and humans, those who reach exceptionally old age have immune markers the same as young adults.

- Your genotype combination is associated with decreased baseline
 glutathione levels
- Glutathione decreases with age, and low levels of glutathione are associated with chronic exposure to chemical toxins, heavy metals and excess alcohol, immunocompromised conditions, and neurodegenerative disorders
- Glutathione has been found to increase by 20% with deep breathing practices like Tai Chi or yoga
- For exercise, a combination of aerobic exercise and circuit weight training produced the highest glutathione effect
- Selenium, glycine, cysteine, vitamin C, and cruciferous vegetables all improve glutathione levels
- Chicken or bone broth, herbs, and spices are some of the best dietary
 ways to maintain higher levels of glutathione
- Some of the all-stars include cinnamon, anise, sage, and thyme due to
 also containing the antiviral compound caffeic acid



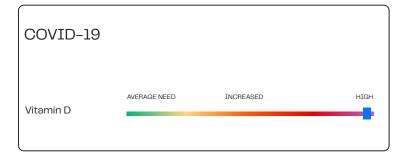
Vitamin A and some other retinoids show important immunomodulatory properties, including the ability to increase the efficiency of actions of type 1 interferons, an important antiviral cytokine released by the innate immune system against viral infections. Coronaviruses similar to SARS-CoV-2 can suppress the host IFN-I-based antiviral response as part of their infection mechanism.

- Your genotype is associated with a 69% lower conversion rate of betacarotene to vitamin A, making it important to include more animal-based vitamin A to hit your daily target
- Vitamin A intake by country shows that Spain is the country with the lowest population meeting nutritional requirements for vitamin A, followed by Belgium and Finland
- Germany and Portugal show the best, and with the exception of Finland, countries with suboptimal Vitamin A status are correlated (although not significantly) with their COVID-19 incidence and mortality



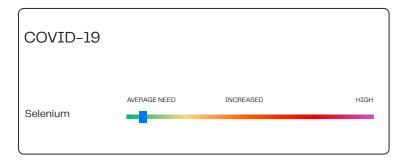
Optimal status of vitamin C plays an important role in the proper working of the immune system.

- · Your genotype is associated with average serum vitamin C levels
- Countries such as the UK, France, Netherlands, and Belgium do not reach
 optimal dietary intake of vitamin C
- Germany stands out for its level of vitamin C intake in comparison with
 other countries
- Despite suboptimal vitamin C intake correlating weakly with COVID-19 incidence, it correlates strongly with deaths percentage, which could suggest a positive effect to fight infection once the individual has already been infected with SARS-CoV-2



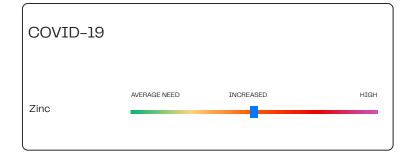
Vitamin D plays a key role in modulating the immune system, and suboptimal or deficient consumption of vitamin D is associated with various conditions related to a malfunction of the immune system and dysregulations in inflammatory status.

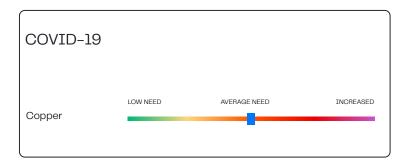
- Your genotype is associated with below average circulating levels of vitamin D
- A 2021 study found that those with the GG genotype for CYP2R1 were associated with 5.9 higher odds of experiencing severe COVID-19
- Vitamin D intake is deficient in all countries studied with COVID severity, with Spain, France, and Italy as the countries with the lowest intake
- A meta-analysis of the studies appears to show that vitamin D is only useful for those who are clinically low (below 20 ng/ml), with moderate doses daily or weekly to raise levels being more effective than periodic large doses



Suboptimal or deficient levels of selenium are associated with decreased cytotoxicity of NK cells, decreased antibody titers, and impaired cellular immunity. Supplementation is commonly related to improvements in cellular immunity and an improved optimal immune response against viruses, including an inhibitory effect on the development of the poliovirus and influenza.

- Your genotype is associated with an average need for selenium
- Glutathione peroxidase 1 (GPX1) is a selenoenzyme with described antioxidant and antiviral properties that depends on nutritional selenium status
- Spain is at the top for meeting selenium requirements while Denmark is
 at the bottom
- The only two populations above the median of the countries analyzed included Finland and France, while the rest of the countries are below the general median



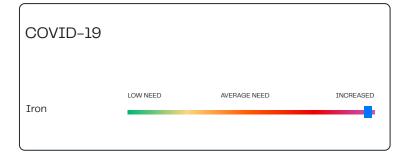


Optimal status of zinc is essential for the proper operation of the immune system and regulates NF-kb, where zinc deficiency in the setting of severe infection provokes a systemic increase in NF-kB activation. In vitro approaches have shown that zinc can inhibit SARS-CoV-1 replication.

- · Your genotype is associated with lower serum zinc
- Scientific evidence supports that optimal zinc intake or supplementation should be considered part of the strategy to reduce COVID-19 effects, with early reports finding that 15-23mg a day show significant improvement in symptoms

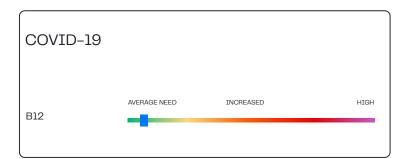
Copper plays a key role in optimal performance of relevant components of the immune system, such as NK cells, macrophages, neutrophils, and monocytes. A deficiency has been related to less effective immune responses against infections, vulnerability for the heart and blood vessels to damage, and increased virulence. Excessive intake is also associated with negative immune function.

· Your genotype is associated with lower serum copper



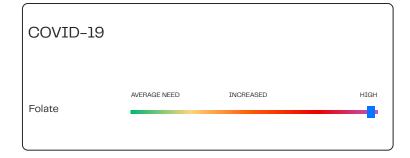
Suboptimal levels of iron are associated with decreased killer efficiency of NK cells and lymphocytes as well as with compromised cytokine production. Both iron uptake disturbances and metabolism are implicated in virulence of airway hospital-acquired infection and chronic respiratory infections. In contrast, excessive iron levels can generate harmful cellular toxicity, so their serum levels must be well regulated.

- Your genotype combination is associated with lower serum iron levels
- A retrospective study based on 50 hospitalized Chinese subjects with confirmed COVID-19 demonstrated that 90% of these subjects had abnormally low serum iron concentrations
- Populations with lower iron status could be more prone to suffer a mild to severe (or critical) symptomatology of COVID-19 and the fact of monitoring patient iron levels has been proposed as a potential early marker to predict COVID-19 severity and mortality
- · Iron levels should always be monitored by your practitioner



Sufficient vitamin B12 intake is essential for antibody production and a deficiency is related to a lower concentration of circulating lymphocytes and altered antibody-based responses. SARS CoV-2 infection is related to an aggravation of the cellular metabolism and the homocysteine pathway causing severe complications from COVID-19, and the correct supply of vitamin B12, folate and B6 may be crucial for COVID-19 patients.

- · Your genotype is associated with high B12 levels
- Some of the countries least affected by SARS-CoV-2 show the highest levels of vitamin B12 intake (Portugal and Finland)
- Some of the countries most affected by SARS-CoV-2 (Belgium and Spain) have intakes below the median



COVID-19

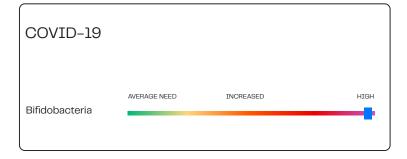
B6 (Pyridoxine)

Folate is crucial for optimal Th-1 mediated immune response and proper antibody production. Suboptimal levels of folate may trigger imbalances in T and NK cell mediated immune responses and decrease the amount of antibody production.

- Your genotype combination is associated with a higher than average need for folate
- The correct intake of vitamin B6, folate and B12 in patients affected by COVID-19 has been proposed as part of the disease treatment, even by supplementation formulas, in an attempt to regulate the disruption of cellular metabolism of the homocysteine pathway caused by the SARS-CoV-2 infection

Vitamin B6 is essential for maintaining cytotoxic activity of NK cells, lymphocyte development, and B-cell antibody production. Suboptimal intake is associated with lower concentrations of circulating lymphocytes, impaired lymphocyte maturation, and decreased antibody-based responses.

- Your genotype is associated with low serum B6 levels, requiring a higher than average intake of B6
- The correct intake of vitamin B6, folate and B12 in patients affected by COVID-19 has been proposed as part of the disease treatment, even by supplementation formulas, in an attempt to regulate the disruption of cellular metabolism of the homocysteine pathway caused by the SARS-CoV-2 infection



INCREASED

HIGH

AVERAGE NEED

Approximately 80% of your immune system is in your gut. The good bacteria bifidobacterium is highest in breast-fed infants and has been found to be lower in the higher-risk demographics for COVID-19 including those with diabetes, obesity, asthma and the elderly. Bifidobacteria populations have been found to vary based on the FUT2 genotype.

- Your genotype is associated with lower bifidobacteria levels in the gut that could increase the susceptibility to lower and upper respiratory infections
- Prebiotics found in foods like bananas, garlic, leeks, barley, asparagus, pistachios, onions, and polyphenol-rich foods – have been found in human trials to increase bifidobacteria levels

COVID-19			
Glycine	AVERAGE NEED	INCREASED	HIGH

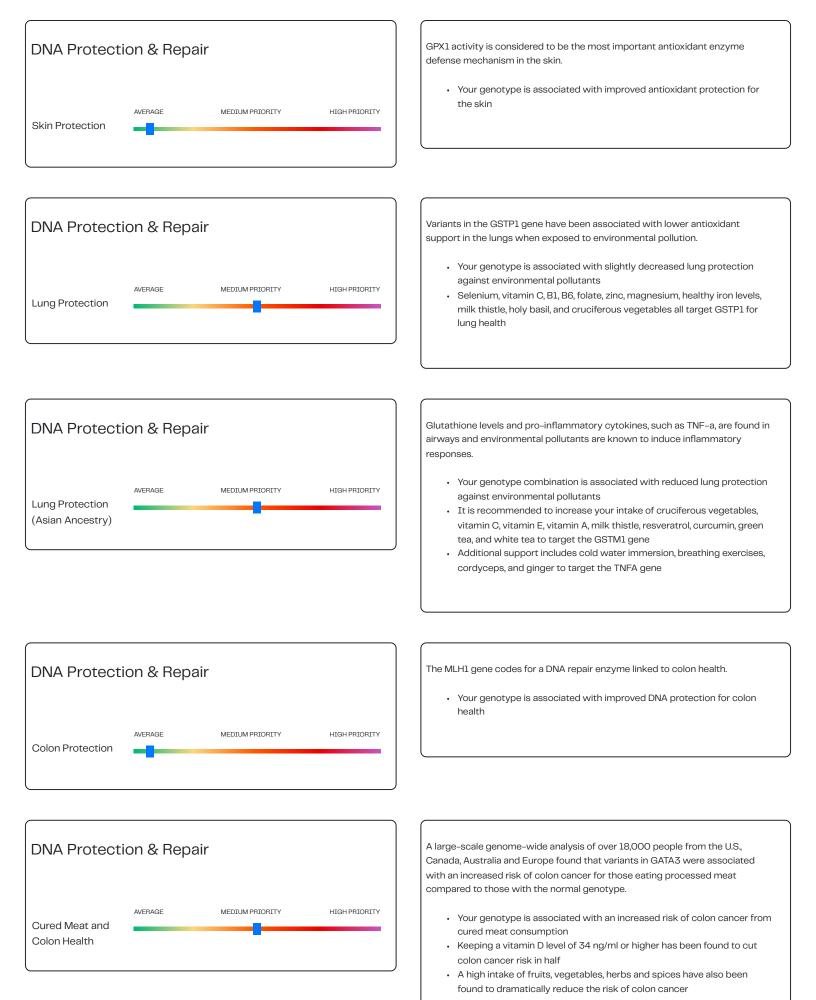
Glycine is one of the three major amino acids for glutathione production, protecting the body from oxidative damage during the immune response, and supporting T-cell proliferation.

- Your genotype is associated with low type 1 collagen production, increasing your glycine requirement
- Type I collage is a major structural protein in the lung and is stimulated during certain inflammatory reactions in the lung
- Collagen protein, bone, or chicken broth that gelatinizes, gelatin, meat with the skin, ribs, shanks, and drumsticks are all ways to increase dietary glycine
- Baobab is considered an exceptionally good source of plant-based glycine found in the hunter-gatherer Hazda diet



DNA PROTECTION & REPAIR

DNA Protec	tion & Repair			Glutathione is the master antioxidant system involved in oxidative stress, detoxification, and immunity. Glutathione status parallels telomerase activity, an important indicator of lifespan.
Glutathione Protection	AVERAGE	MEDIUM PRIORITY	HIGH PRIORITY	 Your genotype combinations are associated with decreased baseline glutathione levels Glutathione decreases with age, and low levels of glutathione are associated with chronic exposure to chemical toxins, heavy metals and excess alcohol, immunocompromised conditions, and neurodegenerative disorders Glutathione has been found to increase by 20% with deep breathing practices like Tai Chi or yoga For exercise, a combination of aerobic exercise and circuit weight training produced the highest glutathione effect Selenium, glycine, cysteine, vitamin C, and cruciferous vegetables all improve glutathione levels Chicken or bone broth, herbs, and spices are some of the best dietary ways to maintain higher levels of glutathione Some of the all-stars include cinnamon, anise, sage, and thyme due to also containing the antiviral compound caffeic acid
DNA Protec	tion & Repair			CAT makes an enzyme called catalase, which helps reduce oxidative stress. CAT is present in all aerobic cells while research has found the highest correlation to prostate, breast, liver, and blood health.
Catalase Protection	AVERAGE	MEDIUM PRIORITY	HIGH PRIORITY	Your genotype is associated with improved catalase levels
DNA Protec	tion & Repair			The SOD2 gene is responsible for superoxide dismutase levels, an important protector of the mitochondria, the powerhouse of the cell.
Mitochondrial		MEDIUM PRIORITY	HIGH PRIORITY	Your genotype is associated with increased mitochondrial protection
Protection				
DNA Protec	tion & Repair			One hypothesis for variants in MTHFR 677 is that they were selected based on higher folate intake and UV exposure, both common in Mediterranean climates. What happens in the body when MTHFR enzymatic function is reduced is that thymidine production increases. Thymidine enhances the repair of UV-induced
UV Protection	AVERAGE	MEDIUM PRIORITY	HIGH PRIORITY	 DNA damage to help quickly repair sun damage. Your MTHFR genotype is associated with increased UV protection from the sun

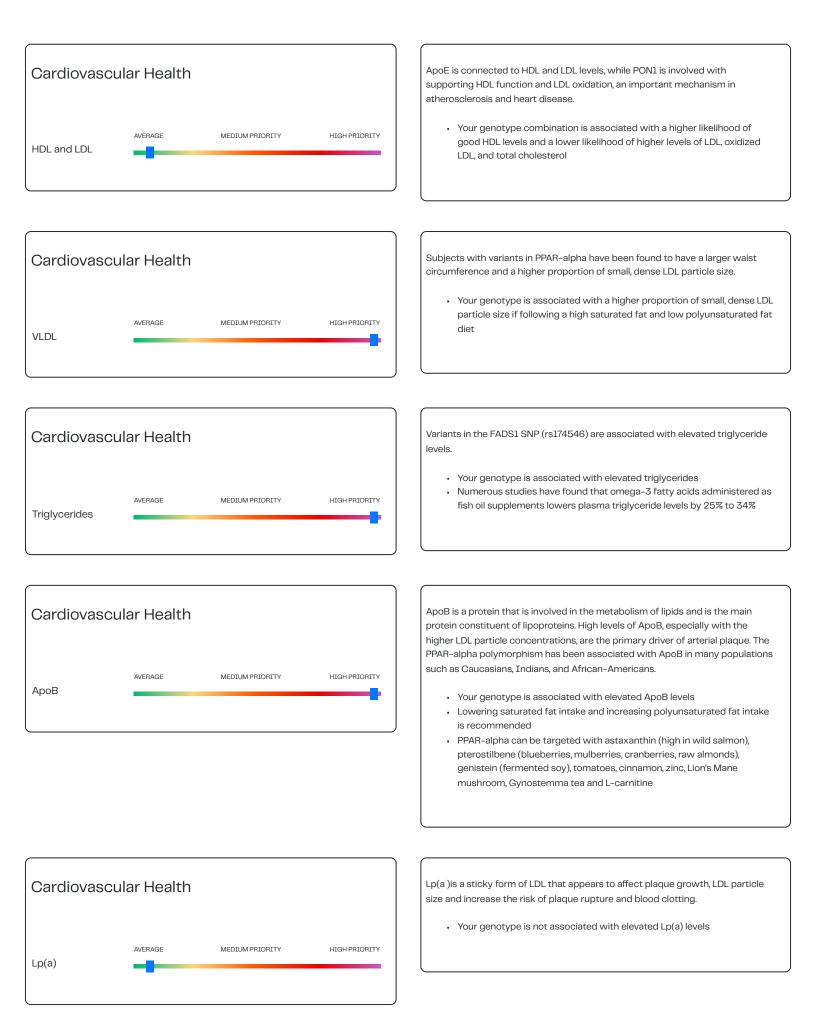






CARDIOVASCULAR HEALTH &

EXERCISE



Cardiovascu	lar Health			The risk of heart attacks and cardiovascular disease conferred by the 9p21 gene appears to be modified by a prudent diet high in raw vegetables and fruits for South Asian, Latin American, Arab, Chinese and European populations for variants in rs4977574.
Raw Plant Intake	AVERAGE	MEDIUM PRIORITY	HIGH PRIORITY	• You have a higher than average requirement for raw fruits and vegetables to maintain a healthy heart
Cardiovascu	lar Health			The uncoupling of nitric oxide has been linked to play an essential role in cardiovascular pathologies including dilated cardiomyopathy, ischemia-reperfusion injury, endothelial dysfunction, atherosclerosis, and hypertension.
Nitric Oxide	AVERAGE	MEDIUM PRIORITY		 Your genotype combinations are associated with a higher than average need for folate to produce adequate BH4, the precursor to nitric oxide BH4 is depleted by high blood sugar, high omega-6 intake, chronic stress, high levels of mercury, arsenic, lead and aluminum, aspartame, and oxidative stress Other strategies to increase BH4 include vitamin C, L-arginine, B6, magnesium, and selenium
(
Cardiovascu	lar Health			Homocysteine is a non-protein amino acid that is produced from methionine, can be recycled back into methionine and converted into cysteine in the methylation cycle. High homocysteine levels have been connected to depression, blood clots, inflammation, macular degeneration, dementia, and
Homocysteine	AVERAGE	MEDIUM PRIORITY	HIGH PRIORITY	 You have a higher than average need for folate to maintain healthy homocysteine levels



Cardiovascular Health
AVERAGE MEDIUM PRIORITY HIGH PRIORITY
Blood Clots

Age-related increases in blood pressure have been observed in almost every population, except among hunter-gatherers and farmers. High physical activity, low-stress levels, and potentially protective diets high in fruits, vegetables, potassium, lower in calories, salt, and alcohol are the major contributing factors to the stark differences.

Your genotype combination is associated with normal blood pressure levels

Deep vein thrombosis is a condition that occurs when a blood clot forms in a vein deep inside a part of the body and is most common for those over 60. Variants in F5 increase the risk of deep vein thrombosis.

· Your genotype is not associated with deep vein thrombosis



Cardiovascular Health
AVERAGE MEDIUM PRIORITY HIGH PRIORITY
Fibrinogen

Cardiovascular Health						
Hemochromatosis		SLIGHT RISK	MORE LIKELY			

Variants in the ABCG2 gene have been associated with elevated uric acid levels and an increased risk of gout in Asians, Europeans, African Americans, Mexican Americans, and American Indians. Epidemiological studies have shown that uric levels are positively correlated with gout, hypertension, atherosclerosis, atrial fibrillation, and heart failure.

· Your genotype is associated with average uric acid levels

Fibrinogen is a soluble protein in the plasma that is broken down to fibrin by the enzyme thrombin to form clots. Fibrinogen serves to stop excessive bleeding, but in high amounts can cause blot clots.

• Your genotype is not associated with elevated fibrinogen levels

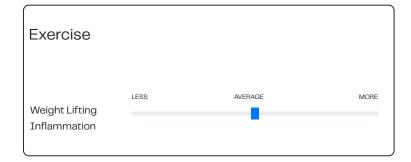
A homozygous HFE C282Y may lead to an iron overload due to increased iron absorption and disrupted metabolism. People who are homozygous for the HFE C282Y gene mutation comprise 85 to 90 percent of those with hemochromatosis in whites of western European descent.

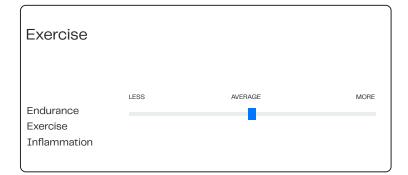
Your wild-type HFE genotype is associated with a reduced likelihood of genetically linked hemochromatosis



ACTN3 is currently the most promising gene for predicting the likelihood of becoming an Olympic level sprint and power athlete in males and females. The RR (CC) genotype expresses the ACTN3 protein found in Type II muscle fibers, which produces explosive and powerful contractions.

- You have the RX genotype for the ACTN3 gene associated with enhanced improvements in strength, power and muscle hypertrophy
 More powerful muscle contractions
- Higher muscle hypertrophy response
- Faster recovery







Exercise			
ACL and Shoulder Dislocation Risk	LESS RISK	AVERAGE	HIGH RISK

Weight lifting leads to a variation in muscle inflammatory markers based on genetics and intensity.

- Your genotype combination is associated with average levels of muscle inflammation (creatine kinase) for weight lifting
- To accelerate recovery, ice baths, whey protein, American ginseng, curcumin, vitamin C, and collagen protein have all been found to attenuate creatine kinase levels

Endurance training leads to a variation in creatine kinase levels based on genetics.

Your genotype combination is associated with average levels of muscle inflammation (creatine kinase) for endurance exercise

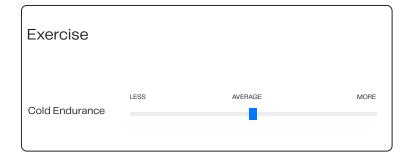
High-intensity exercise is defined as 70% to 85% of your maximum heart rate, and inflammation variation has been associated with the SOD2 gene.

Your SOD2 genotype is associated with less muscle inflammation in response to high-intensity exercise

The COL1A1 gene is associated with ACL and shoulder injury risk.

- Your COL1A1 genotype is associated with an increased need for dietary collagen to prevent ACL and shoulder injuries
- Vitamin C, zinc, copper, glycine, proline, lysine, and B6 are all precursors to collagen production







The ACTN3 gene is linked to increased or decreased risk of ankle and hamstring injuries.

 Your ACTN3 genotype is associated with an average risk of ankle and hamstring injuries

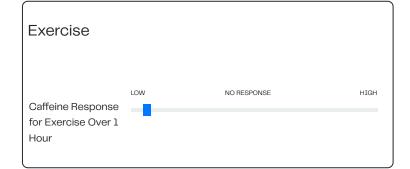
The ACTN3 gene is associated with a lower or higher adaptation rate to cold endurance.

 You have the ACTN3 RX genotype, associated with an average adaptation rate to cold endurance

VO2 max is measurement for oxygen capacity and endurance training. The most recent research has shown that 97 genes predicted VO2 max trainability. The phenotype is dependent on several of these variants combined together, which may contribute to approximately 50% of an individual's VO2 max trainability.

Research has shown that your genotypes for multiple genes are associated with a higher VO2 max training response





The CYP1A2 gene is associated with caffeine response for improving or decreasing exercise performance.

Fast metabolizers have been found to show an increase in exercise
performance for activities under 1 hour from caffeine consumption

The CYP1A2 gene is associated with caffeine response for improving or decreasing exercise performance.

• Your CYP1A2 genotype is associated with a decrease in exercise performance activities over 1 hour from caffeine consumption due to being a fast metabolizer

MY HEALTH REPORT: STRENGTHS

This section is a thorough overview of your individual gene function across the entire analysis in just a few pages. If you are looking for a brief summary of the most important parts of your report without doing a deep dive into the genotype tables and clinical research sections, this is the place to start. Be proud of your inherent genetic strengths!

NUTRIENT METABOLISM & DIGESTION

- Prebiotics, Probiotics and B12-FUT2 The "non-secretor" AA genotype in European, African, and Indian populations is associated with higher B12 levels and confers resistance to certain infections including the Norovirus, Rotavirus, and H. Pylori, but is more susceptible to other infections.
- — Vitamin C-SLC23A1 Your genotype is associated with improved whole-body vitamin C homeostasis through dietary absorption and renal reabsorption.
- Adiponectin-ADIPOQ Your genotype is associated with a higher probability of normal adiponectin levels, linked to improved bodyweight, insulin, and glucose levels.
- _ Iron Your genotype is associated with a lower risk of iron overload for the HFE C282Y gene.
- **_____ Saturated Fat-PPAR-alpha** While your heterozygous genotype is associated with abnormal lipid metabolism and a poor response to the ketogenic diet, it has also been found to have lower C-reactive protein levels, a marker of inflammation.
- **—** Ghrelin and Appetite-FTO Your genotype is associated with normal ghrelin levels (hunger hormone), decreasing the risk for overeating and abdominal weight gain.
- = Saturated Fat-APOA2 Your genotype is associated with a reduced likelihood of saturated fats causing weight gain.
- = Carbohydrates-TCF7L2 Your genotype is associated with an improved insulin response for grain-based carbohydrates.
- _ Lactose You have the homozygous AA genotype that is associated with a lower probability of lactose intolerance.
- **Histamines-APB1** You have the wild-type genotype that is associated with improved histamine breakdown in the digestive tract.
- _ Uric Acid-ABCG2 Your genotype is associated with a lower probability of chronically elevated uric acid levels.
- = Ethanol Metabolism-ALDH2 Your genotype is less likely to experience facial flushing from alcohol due to improved acetaldehyde metabolism.
- Stress and IBS-ADRB2 You have the ADRB2 CC wild-type genotype that is associated with a reduced percentage of digestive disorders, IBS, and anxiety related to adrenaline levels.

DNA DAMAGE, PROTECTION AND REPAIR

- _ DNA Repair-ATM Your genotype is associated with improved DNA repair function for pancreatic health.
- = Breast-ESR2 For women with the AA ESR2 rs2987983 genotype, your genotype is associated with improved tumor suppression function for breast health. All genes related to breast health should be analyzed to better determine the cumulative value for breast protection.
- _ DNA Repair-MDM2 Your MDM2 genotype is associated with improved DNA repair for sun damage if you are female.
- **DNA Repair-MLH1** Your genotype is associated with improved DNA repair for colon, endometrium, lung, and brain protection.
- Longevity-SIRT1 Your SIRT1 genotype is associated with normal SIRT1 activity for longevity. While not a weakness, you may want to increase SIRT1 activity epigenetically to increase the probability of longevity, especially if you have the APOE-e4 allele. A sedentary lifestyle, aging, poor diet, and obesity lowers SIRT1 activity. Exercise, fasting, 7-8 hours of sleep per night, saunas, polyphenols, vitamin D, omega-3 fatty acids, resveratrol, magnesium, and melatonin have all been found to increase SIRT1 activity.

METHYLATION

- **Folate-MTHFR 677** You have the homozygous genotype that is associated with improved protection against UVinduced DNA damage from the sun and a reduced risk of malaria.
- **Folate-MTHFR 1298** Your genotype is associated with improved BH4 levels and neurotransmitter function. Healthy BH4 levels assist in the management of cardiovascular health, mental health, and digestive health.
- **Folate-MTHFD1 G1958A** Your genotype is associated with improved metabolism for folinic acid, the second most common type of folate after methylfolate.
- **Folate-DHFR** Your genotype is associated with an improved breakdown of synthetic folic acid at the beginning of the folate cycle. However, variants in MTHFR 677 can also affect folic acid metabolism.
- ____ B12, B2 and Zinc-MTR You may have improved MTR function, assisting homocysteine metabolism.
- = B12-TCN2 Your genotype is associated with improved B12 transportation.
- = B6-CBS Your genotype is associated with improved homocysteine and hydrogen sulfide levels, assisting gut repair and brain health.
- _ Arsenic-CBS Your genotypes are associated with improved arsenic metabolism and detoxification for the CBS genes.

HORMONE SUPPORT

- **Sex Hormone Binding Globulin** If you are female, your genotype is associated with helping maintain normal estrogen and testosterone levels. Other epigenetic factors like obesity, fatty liver, and Type 2 diabetes should be considered when assessing SHBG levels.
- **Thyroid-DI01** Your genotype is associated with average DI01 gene function for T3 and T4 thyroid function, however other epigenetic factors should be assessed.
- **Thyroid-DI02** Your genotype is associated with average T3 and T4 thyroid function in the brain for the DI02 gene. However, other factors can affect T3 and T4 levels including thyroid surgeries.
- **Estrogen Metabolism-CYP1A1** Your CYP1A1 wild-type genotype is improved for the beginning phase of estrogen metabolism. Please review all genes involved in estrogen metabolism for a complete picture of the process.
- **—** Estrogen Metabolism-CYP1A2 For men and women with the CYP1A2 AA genotype, coffee intake was found to be more protective against estrogen receptor-positive breast cancer and prostate cancer.
- MTNR1B-Melatonin You have the CC MTNR1B genotype, which is associated with a normal circadian rhythm of melatonin production at night and in the morning. This gives you a wider time range for breakfast and dinner for glycemic control.

MENTAL HEALTH, NEUROTRANSMITTERS & COGNITIVE PERFORMANCE

- **Dopamine, Adrenaline and Estrogen-COMT** The heterozygous genotype for COMT V158M and H62H scored significantly higher on insight problem-solving tasks and had a greater ability for social facilitation and cooperativeness.
- **Dopamine Receptors-ANKK1** Your genotype is associated with an improved density of dopamine receptors for healthy dopamine levels.
- **Histamines and Migraines-DAO** The wild-type CC genotype for DAO rs1049793 is associated with a reduced risk of histamine-induced migraine headaches.
- Cholesterol-APOE You have the ApoE e3/e3 genotype, improving cholesterol transport and the maintenance of brain neurons. The ApoE e3 allele improves cognitive fitness, HDL and LDL profiles, viral protection, and the response to plant bioactive compounds.

IMMUNE SUPPORT, ANTIOXIDANTS AND INFLAMMATION

- _ Cell Protection-SOD2 You may have improved SOD2 function to protect to the mitochondria (powerhouse of the cell).
- _ Cell Protection-CAT Your genotype is associated with improved catalase levels, mitigating damage to your cells.
- **—** Glutathione-GSTM1 While the GSTM1 null genotype has been associated with a greater sensitivity to benzo(a)pyrene, there is also a benefit to this genotype. The benefit is that the null genotype may retain a higher level of isothiocyanates, the anti-cancer compounds found in cruciferous vegetables that may also be required in higher amounts for this genotype.
- Glutathione-GSTP1 While the homozygous GG genotype for GSTP1 rs1695 is associated with a higher sensitivity to heavy metals, one advantage may be an increased VO2 max response from endurance training compared to the wildtype genotype.
- Glutathione-GPX1 Your genotype is associated with improved selenium status and glutathione peroxidase to boost DNA protection, heat stress tolerance, skin protection and longevity.
- Glutathione-CTH Your genotype is associated with improved gene function, leading to adequate cysteine for glutathione production.
- Nitric Oxide-NOS1 Your genotype is associated with an average required intake of red, yellow, and orange vegetables to modulate the inflammatory process for NOS1.
- Nitric Oxide-NOS2 Your NOS2A gene is functioning optimally for reducing the probability of age-related macular degeneration from cigarette smoke.

DETOXIFICATION

- **____** Liver Enzyme-CYP1A1 Your genotype is associated with improved detoxification of benzopyrene from cigarette smoke and will assist the function of your GSTM1 gene.
- **____** Liver Enzyme-THC and CYP2C9 You have the wild-type genotype that is associated with improved metabolism of THC, the active psychoactive compound in cannabis.
- Liver Enzyme-CYP2D6 Your genotype is associated with improved metabolism of certain drugs associated with CYP2D6 rs1065852. However, more CYP2D6 SNPs are needed for a complete panel. Please talk to your doctor about further testing for CYP2D6 and drug metabolism.
- **____** Liver Enzyme-CYP2E1 Your genotype is associated with improved metabolism of benzene and acrylamide for colon health.
- **____** Liver Enzyme-CYP3A4 Your genotype is associated with normal metabolism of certain drugs that use this enzyme. We recommend further pharmacogenomic testing with your doctor for more information regarding CYP3A4.
- Aromatic Amines-NAT2 You have the fast acetylator genotype for NAT2, which is associated with a reduced risk of bladder cancer in smokers and may improve the detoxification of aromatic amines found in commercial hair dyes, industrial and manufacturing plants, meat cooked at high temperatures, and diesel exhaust.
- Vitamin K2-VOKRC1*2 Your genotype is associated with normal vitamin K2 levels unless gut function is compromised from antibiotics, SIBO, leaky gut syndrome, IBS, IBD, Crohn's disease or parasites.
- Statins-COQ2 Your genotype is associated with a lower likelihood of statin drug-induced muscle pain.

CARDIOVASCULAR HEALTH & EXERCISE

- **Power and Recovery-ACTN3** You have the RX genotype associated with enhanced improvements in strength, muscle hypertrophy, sprint times, protection from eccentric training-induced muscle damage, and a reduced risk of sports injury. The RX genotype may represent the best of both ACTN3 genotypes for strength training, maintaining lean muscle mass later in life, and longevity.
- Lung Cytokines-TNFA If you have Asian ancestry, your genotype is associated with improved TNF-a gene function for lower inflammation in the lungs.
- — Muscle Inflammation-SOD2 You have the wild-type genotype that is associated with improved SOD2 function in response to high-intensity exercise, reducing the inflammatory response in muscle tissue compared to the homozygous carriers.
- _ LDL-LPA Your genotype is associated with healthy Lp(a) levels, a sticky form of LDL that affects plaque levels.
- **Caffeine-CYP1A2** You have the homozygous AA genotype and are a "rapid metabolizer" of caffeine. This means that caffeine will quickly be metabolized from your body and the effects lasting a shorter period of time. Variants in COMT can increase the sensitivity to catecholamines in coffee, and oral contraceptives can slow down caffeine metabolism.
- **____** Fibrinogen-ESR2 Your genotype is associated with improved fibrinogen levels.
- = Blood Clots-F5 Your genotype is associated with improved gene function for a lower probability of deep vein thrombosis.
- = Blood Pressure-ACE1 Your genotype is associated with intermediate baseline ACE levels. If you are female, ACE levels may be lower. Depending on ACE2 levels, you may have a more balanced renin-angiotensin system for blood pressure.
- **Blood Pressure-AGTR1** You have the wild-type genotype, associated with a lower probability for high blood pressure, elevated triglycerides, elevated ApoB, and NAFLD from excess dietary fat and carbohydrate intake.
- Blood Pressure-ACE2 Your genotype is associated with higher baseline ACE2, improving the balance between ACE1 and ACE2 for blood pressure, and potentially lowering the risk of COVID-19 severity. Other dietary habits and health issues could affect this result.
- **Phytoestrogens-TMPRSS2** You have the AG genotype that is associated with a lower expression of TMPRSS2 and may decrease the susceptibility to viral infections and prostate cancer (men).

MY HEALTH REPORT: WEAKNESSES

Genes are not your destiny - they are your blueprint. Please understand that these weaknesses can be turned into strengths based on the personalized recommendations given below. Making strategic changes to diet, environment, stressors, and even relationships can have a profound effect on optimizing gene function. Aim to turn every weakness into a strength by giving attention to the proactive, customized dietary and lifestyle modification recommendations in this section!

NUTRIENT METABOLISM & DIGESTION

- = Beta Carotene to Vitamin A Conversion Rate-BCMO1 Your BCMO1 genotype combination is associated with a reduced conversion rate of plant-based beta carotene (squash, sweet potatoes, carrots) to vitamin A. This increases your need for foods higher in vitamin A like eggs, cod liver oil, wild salmon oil and organ meats for skin, digestion, healthy eyes, lungs, and immunity.
- ALA to EPA and DHA Conversion-FADS2 Your genotype is associated with a reduced conversion of plant-based omega-3 ALA (walnuts, flax seeds, and pumpkin seeds) to EPA and DHA. Increased EPA and DHA intake may be needed.
- — Prebiotics, Probiotics and B12-FUT2 The "non-secretor" AA genotype in European, African, and Indian populations is associated with lower bifidobacteria and increased susceptibility to pneumonia, meningitis, ear infections, bronchitis, bloodstream infections, throat swelling, skin infections, infectious arthritis, the mumps virus, Candida, and certain autoimmune disorders. Increasing bifidobacteria with prebiotic foods will assist with GABA production (the calming neurotransmitter) and immune function.
- **B6-NBPF3** You are more likely to have low B6 levels due to variants in the NBPF3 gene, increasing the sensitivity to medications that deplete B6 (oral contraceptives, antibiotics, ACE inhibitors, antacids, proton pump inhibitors and more). You need to focus on increasing foods high in B6 like wild salmon, pistachios, avocados and potatoes.
- — Saturated Fat-PPAR-alpha You have the heterozygous genotype that is associated with abnormal lipid metabolism and a poor response to the ketogenic diet. This is more pronounced if you are male and have multiple variants in the other fat metabolism genes. You may need to lower your saturated fat intake, increase your polyunsaturated fat intake, and focus on PPAR-alpha activators including astaxanthin, pterostilbene, cinnamon, tomatoes, zinc, Lion's Mane mushroom, Gynostemma tea, and increasing L-carnitine production.
- **—** Fat Metabolism-ACSL1 Your genotype is associated with higher fasting glucose levels from a higher saturated fat intake. If your fasting glucose is high and you have variants in the other fat metabolism genes, fatty red meat and dairy should be reduced and more focus should be on monounsaturated and polyunsaturated fats.

DNA DAMAGE, PROTECTION AND REPAIR

- DNA Repair-TP53 You have the homozygous CC genotype that may be advantageous for fertility in cold climates, but also increases the need for selenium, zinc, vitamin C, reishi, and niacin for DNA repair against chemical toxicity to the thyroid gland and skin.
- Processed Meat and Colon Cancer-GATA3 Your genotype is associated with a sensitivity to processed meats (hot dogs, salami, pepperoni) and colon cancer risk due to variants in GATA3. Reduce processed meat intake, optimize vitamin D levels and increase berries, apples, sauerkraut, broccoli, tomatoes, basil, rosemary, garlic, onions and leeks.

METHYLATION

- Folate-MTHFR 677 You have the homozygous genotype that is associated with a reduced function of approximately 50-70%. This increases the need for riboflavin and methylfolate for normal homocysteine levels.
- = B12-MTRR Your genotype is associated with a potentially higher sensitivity to B12 deficiency if there are variants in MTR as well.
- — Choline-PEMT Your genotype is associated with an increased need for dietary choline for liver health, normal homocysteine levels, breast health for women, and a healthy pregnancy for women.

HORMONE SUPPORT

- **Vitamin D-CYP2R1** Your genotype is associated with low circulating vitamin D levels that can affect immunity, breast health in women, and testosterone levels in men. Check your vitamin D levels and make sure you are in range.
- **Estrogen Metabolism-CYP2C19** Individuals with the CC genotype for CYP2C19*17 are considered the normal metabolizer phenotype, which may lack the estrogen metabolism benefits of the ultra-rapid metabolizer phenotype. Please review all genes involved in estrogen metabolism for a complete picture of the process.
- **Estrogen Metabolism-COMT** For estrogen metabolism and detoxification, those with the intermediate AG COMT V158M genotype may have an increase in harmful estrogen metabolites that can cause DNA damage. To reduce the risk of these metabolites, you should avoid xenoestrogens, manage stress levels, maintain gut health, increase magnesium intake, and consume green tea polyphenols.
- **Estrobolome-FUT2** Due to having the homozygous FUT2 genotype, the gut phase of estrogen detoxification may be affected. This could increase the sensitivity to depleted bifidobacteria from antibiotics, proton pump inhibitors, glyphosate, and sucralose that all deplete bifidobacteria. Increasing prebiotic fiber, polyphenols, and bifidobacteria may improve breast, prostate, bladder, liver, and digestive health.

MENTAL HEALTH, NEUROTRANSMITTERS & COGNITIVE PERFORMANCE

- — Serotonin Receptor-Memory You have the heterozygous genotype that is associated with reduced episodic memory, the ability to recall details regarding personal experiences, specific events, and what exactly occurred, depending on your diet and lifestyle. Episodic memory can be enhanced with tryptophan, green or black tea, mindfulness training, increasing VO2 max, reduced reliance on smartphones, and a higher catecholamine, betaine, and choline intake depending on COMT, BDNF, and PEMT gene status.
- Serotonin Receptor-Stress The 5-HT2A heterozygous genotype may be more impactful in females who also have variants in the BDNF gene. Chronic stress may increase the susceptibility to anxiety, depression, OCD, and IBS for these genotypes. If you experience higher perceived stress and chronic stress levels, you may require more aerobic exercise, cognitive behavioral therapy, mindfulness training, meditation, yoga, singing, prebiotics, lactobacillus helveticus, bifidobacterium longum, tryptophan, green or black tea, and B-vitamins.
- — Dopamine, Adrenaline and Estrogen-COMT The heterozygous AG COMT V158M genotype is associated with a slower breakdown of dopamine, adrenaline, and estrogen, creating higher circulating levels in response to stress due to variants in the COMT genes. This may increase your need for magnesium, vitamin C, strength training, and sprints to reduce stress levels.
- — Histamines and Hyperactivity-HNMT You have the AA genotype that is associated with increased hyperactivity in response to food dyes and sodium benzoate. Excess histamine is lowered by vitamin C, choline, folate, magnesium, chamomile, basil, stinging nettle, echinacea, fennel, ginger and wild oregano.
- — Anandamide-FAAH You have the common CC genotype that encodes for the fast activity of FAAH. This is associated with naturally lower anandamide levels that could increase anxiety, pain, pesticide sensitivity and a heightened stress response to threatening situations. You may benefit from aerobic exercise over 30 minutes (especially in altitude), CBD oil, red clover tea (women), kaempferol (raspberries, capers, cumin, cloves, almonds, cherry tomatoes, red wine), cacao, echinacea, rosemary, and hops to increase anandamide levels.
- = Brain Health-PEMT Your genotype is associated with an increased need for dietary choline and daily walks for memory, anxiety, and REM sleep.
- — Glutamate-BDNF Your genotype is associated with lower BDNF levels that can affect mood, head injury recovery, memory, and blood sugar levels. Research has shown that running, DHA, lithium, green tea, milk thistle, acetylcholine, sunlight, saunas, hot baths, the probiotic Bifidobacterium longum, intermittent fasting, turmeric, and optimal estrogen levels (women) all improve BDNF levels.
- **—** Glutamate Transport-SLC17A7 Your genotype is associated with delayed recovery from head injuries. We recommend also reviewing your APOE and BDNF genotype to determine cumulative impact. It is advised to be proactive with zinc, omega-3 fatty acids, Lion's Mane mushroom, magnesium and consistent exercise in case a head injury occurs.

IMMUNE SUPPORT, ANTIOXIDANTS AND INFLAMMATION

- — Glutathione-GSTM1 You have the null genotype that is associated with a higher sensitivity to benzo(a)pyrene from the burning of wood or trash, tobacco smoke, asphalt, coal, diesel exhaust, charred meat, and gas cooking. If you have the GSTM1 null and NAT2 slow acetylator combination, that may affect lung, breast, bladder, skin, colon, and kidney health. It is recommended to increase your intake of cruciferous vegetables, vitamin C, vitamin E, vitamin A, milk thistle, resveratrol, curcumin, green tea, and white tea.
- **—** Glutathione-GSTP1 You have the homozygous GG genotype for GSTP1 rs1695 that is associated with a higher sensitivity to mercury, cadmium, arsenic, pesticides, and air pollution for breast, prostate, urinary, esophagus, and skin health. Your GSTP1 rs1138272 genotype may increase or decrease this sensitivity. Selenium, vitamin C, vitamin E, milk thistle, and cruciferous vegetables all assist GSTP1 gene function.
- Heavy Metals-GSTP1 You have the heterozygous CT genotype for GSTP1 rs1138272 that is associated with a higher sensitivity to mercury, cadmium, arsenic, pesticides, and air pollution for colon, prostate, lung, throat, and fertility health. Your GSTP1 rs1695 genotype may increase or decrease this sensitivity. Selenium, vitamin C, B1, B6, folate, zinc, magnesium, healthy iron levels, milk thistle, holy basil, and cruciferous vegetables all assist GSTP1 rs1138272.
- **Eye Health-CFH** Your genotype is associated with an increased need for lutein, zeaxanthin, bilberry, lingonberry, vitamin C, and vitamin E for healthy eyes.
- **—** Eye Health-ARMS2 Your genotype is associated with a higher sensitivity to the negative effects of smoking on eye health.

DETOXIFICATION

- **—** Liver Enzyme-CYP1A2 You have the AA genotype for CYP1A2 that is associated with a higher sensitivity to heterocyclic amines (fried meat) depending on if you have the homozygous null GSTM1 genotype or the NAT2 slow acetylator genotype. Marinades, unfiltered fermented drinks (Kombucha, beer, wine), cruciferous vegetables, parsley, and spinach have all been found to reduce the carcinogenic effect of heterocyclic amines.
- **—** Liver Enzyme-CYP1B1 You have the GG genotype that is associated with reduced detoxification of polycyclic aromatic hydrocarbons (highest in vegetable oils), oral contraceptives, cigarette smoke, an increased sensitivity to excessive sun exposure, and high-dose biotin supplementation. You can assist CYP1B1 with seaweed, celery, berries, rooibos tea, red wine, and dark roast coffee.

CARDIOVASCULAR HEALTH & EXERCISE

- — VO2 Max-PPARGC1A Your genotype is associated with a higher need for more strategies to increase oxygen capacity for aerobic exercise, including a structured endurance program, cold exposure, and adaptogens. Your genotype in the GSTP1 rs1695 gene can also influence this result.
- — Muscle Recovery-IL6 You have the CG genotype that is associated with higher levels of creatine kinase a marker of muscle damage from workouts. To accelerate recovery, whey protein, cold water immersion, American ginseng, curcumin, allicin, optimal testosterone levels, vitamin C, and collagen protein have all been found to attenuate creatine kinase levels.
- — Muscle Injury-COL1A1 You have the wild-type CC genotype that is associated with an increased need for dietary collagen for healthy skin, tendons, corneas, lungs, and bones. Vitamin C, zinc, copper, glycine, proline, lysine, and B6 are all precursors to collagen production.
- — Pesticides, HDL and LDL-PON1 Your genotype is associated with decreased PON1 gene activity and reduced pesticide detoxification that could affect LDL oxidation. Elevated mercury levels and high homocysteine can further negatively affect PON1. There are numerous strategies to improve PON1 including choosing organic foods, adequate calcium and magnesium, pomegranates, broccoli sprouts, high-quality olive oil, and a glass of red wine.
- **—** Raw Fruit and Vegetable Intake-9p21 You have the homozygous genotype that is associated with an increased need for phytonutrients from a higher raw fruit and vegetable intake for a healthy heart.
- **Triglycerides-FADS1** Your genotype is associated with a higher need for EPA and DHA omega-3 fatty acids to maintain healthy triglyceride levels.
- **Stress-ADRB2** You have the heterozygous AG genotype for ADRB2 rs104271 that is associated with a higher sensitivity to chronic stress on your heart, especially with variants in COMT. Optimize COMT function and ADRB2 with foods that contain magnesium and vitamin C, deep breathing, and consider adaptogens to lower the stress response.
- — Potassium and Magnesium-ADD1 If you have Asian ancestry, your genotype is associated with an increased risk of a higher sodium intake causing elevated blood pressure. Increasing potassium, vitamin D, magnesium, calcium, garlic, and omega-3's all lower blood pressure.

YOUR PERSONALIZED DNA-BASED GROCERY LIST

This section of the report represents the most expansive, actionable summary of what you can do, right now, to dramatically up-regulate gene function, building a happier, healthier you! No technical expertise is required - just make these recommendations non-negotiable when you visit the grocery store.

Your grocery list is generated based on a combination of unique gene variants that require an increased intake of the following vitamins, minerals, phytonutrients, amino acids, fiber and more. This list generates the foods and drinks based on the highest levels for each section and does not take into account any food allergies or sensitivities.





















Glucosinolates

Brussels sprouts, mustard greens, turnips, savoy cabbage, kale, watercress, red cabbage, broccoli cauliflower, and Bok Choy

Glycine

Broth, collagen powder, meat with the skin, ribs, shanks, drumsticks, and baobob

Cooked spinach, cooked kale, raw spinach, raw parsley, goji berries, asparagus, and

eggs

Sprouted pumpkin seeds, hemp seeds, chia seeds, Gerolsteiner mineral water,

spinach, wild salmon, and peanut butter

Lutein and Zeaxanthin

Magnesium

Niacin

Omega-3's

Potassium

Prebiotics

Pistachios, leeks, asparagus, radicchio, bananas, garlic, kiwi, onions, artichokes, Tiger nuts, chicory root, yacon syrup and foods high in polyphenols

Probiotics

Fermented drinks like Kombucha, fermented veggies like sauerkraut, yogurt and kefir

Yellowfin tuna, canned tuna, wild salmon, ground turkey, chicken breast, liver, skirt steak, white button mushrooms, and brown rice

Wild salmon, avocados, potatoes, acorn squash, coconut water, sweet potato, spinach, tomato sauce, and bananas

Seafood and pastured eggs













Resveratrol

Tryptophan

Vitamin A

Vitamin C

Vitamin D

Vitamin E

Red grapes, peanut butter, Itadori tea, and dark chocolate

Wild salmon, oats, bananas, eggs, chicken, turkey, chocolate, peanuts and pumpkin seeds

Liver, pastured eggs, cod liver oil, wild salmon oil, eel, and sockeye salmon

Bell peppers, guava, black currants, strawberries, oranges, and broccoli

Sockeye salmon, cod liver oil, canned tuna, wild herring, and sardines

Sunflower seeds, almonds, avocado, spinach, butternut squash soup, and olive oil

Zinc

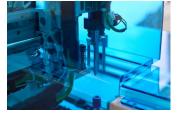
Oysters, crab, lobster, beef, lamb, pork loin, liver, and sprouted pumpkin seeds

PERSONALIZED BLOOD WORK

These results are generated based on a combination of gene variants unique to you. These biomarkers may not be out of range based on your diet and lifestyle habits, but they may be the ones for you to monitor to ensure you are making the right choices based on your genetic results (your predispositions).

For example, if vitamin D comes up in this section, it does not mean that your current levels of vitamin D are actually low. What we are saying is that based on a variety of genetic factors, your variants could make it more difficult to obtain recommended levels of circulating vitamin D, so it might be prudent to further monitor to ensure that you are taking the necessary steps to turn genetic weaknesses into strengths and maintain correct levels.





Triglycerides

Triglycerides should be <150



Vitamin D

Vitamin D should be between 35-50 ng/ml. Check both 25 and 1,25dihydroxyvitamin D.

MY CLINICAL RESEARCH SUMMARY: NUTRIENT METABOLISM & DIGESTION

Beta Carotene to Vitamin A Conversion Rate-BCMO1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
BCMO1 R267S rs12934922	Heterozygous AT
BCMO1 A379V rs7501331	Heterozygous CT

Recap

Improves BCMO1 Gene Function: Vitamin A in the form of retinol and zinc.

Decreases BCMO1 Gene Function: Relying on beta-carotene for vitamin A requirements.

BETA CAROTENE TO VITAMIN A CONVERSION RATE-BCMO1

Research: If you are heterozygous or homozygous for BCMO1 A379V or BCMO1 RS267S, you have a reduced conversion of beta-carotene to vitamin A. If you have a heterozygous or homozygous BCMO1 RS267S and BCMO1 RS267S, the reduction is even more dramatic. Many nutrition labels will have beta-carotene listed as vitamin A, however this is not true vitamin A.

The normal conversion for beta-carotene (carrots, sweet potatoes) to retinol is 1:6 and 1:12 for other carotenoids. Female volunteers carrying the T variant of rs7501331 (379V) had a 32% lower ability to convert beta-carotene, and those carrying at least one T in both SNPs (379V and R267S) show a 69% lower ability to convert beta-carotene into retinol.

In a cohort study of 48,400 US men and 75,170 US women, during a follow-up period of more than 26 years, a higher total vitamin A intake was associated with a reduction in cutaneous squamous cell carcinoma risk.

You want to make sure you consume animal based vitamin A (pastured egg yolks, wild salmon oil, cod liver oil, butter) along with zinc for digestive lining repair, oral health, eye health, iron mobilization, mitochondria health, skin health (sunburns deplete vitamin A in the skin, and acne responds to vitamin A), healthy lung function, and increased immunity.

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ALA to EPA and DHA Conversion-FADS2

Below is a summary of your most significant variant genotypes:

FADS2 rs1535 Homozygous GG
FADS2 rs174575 Homozygous GG

Recap

Improves FADS2 Gene Function: EPA and DHA omega-3 fatty acids.

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Decreases FADS2 Gene Function: Relying on plant-based omega-3 fatty acid ALA for those with the heterozygous or homozygous variant.

ALA TO EPA AND DHA CONVERSION-FADS2

Research: You may have a decreased conversion rate of the plant based omega-3 fatty acid ALA to DHA and should choose DHA sources for sufficient omega-3's.

FADS1 and FADS2 are enzymes that are involved in converting omega-3 and omega-6 fatty acids for brain development and inflammation control. Like the lactase gene, FADS1 is likely to be a critical gene of adaptation. In this case, it was in response to a plant-based diet versus a meat and fish based diet depending on migration routes and food availability.

It has been hypothesized that populations that began to rely more on plant-based diets adapted with the selected allele in FADS2 to synthesize more EPA and DHA from plants. The Inuit populations of Greenland, for example, who rely heavily on seafood with very little plant intake, have a deleted allele showing an opposite adaptation to a diet without plants.

A meta-analysis has found an association between variants in FADS2 in European heritage and a low conversion rate of ALA (plant-based omega-3) to DHA. There is also evidence for gene variants in those with African, Chinese, and Hispanic ancestry having a reduced conversion rate.

Children who had a higher dietary ratio of omega-6 to omega-3 were vulnerable for developing colitis if they also presented specific variants in FADS2.

A higher need of animal-based EPA and DHA may be needed for those with variants in FADS2.

Prebiotics, Probiotics and B12-FUT2

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
FUT2 rs601338	Homozygous AA

Recap

Improves FUT2 Gene Function: Prebiotics, probiotics, and gelatin.

Decreases FUT2 Gene Function: Antibiotics, proton pump inhibitors, glyphosate, and sucralose.

PREBIOTICS, PROBIOTICS AND B12-FUT2

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Research: Approximately 30 percent of people with European ancestry and 20 percent of those with African ancestry carry the homozygous AA version of SNP rs601338 in the FUT2 gene. It is also found in India, while a different FUT2 gene has been discovered in East Asia. While not conclusive, it appears that specific environmental pathogens caused selective pressure for each FUT2 genotype.

Approximately 80% of your immune system is in your gut. The good bacteria bifidobacterium is highest in breast-fed infants and has been found to be lower in the higher-risk demographics for COVID-19 including those with diabetes, obesity, asthma and the elderly. Those with celiac disease and dermatitis may also have lower levels of bifidobacteria.

Bifidobacteria is highly sensitive to glyphosate (a potent herbicide used on GMO corn and soy), and therefore choosing organic is important. Antibiotics, glyphosate and sucralose (Splenda) severely disrupt gut flora and increase the risk of salmonella and C-diff.

The inactive "non-secretor" homozygous AA genotype confers resistance to certain infections including the Norovirus, Rotavirus and H. Pylori, but is more susceptible to other infections. The presence of the A allele is also associated with higher vitamin B12 concentrations, compared to G allele carriers.

The AA genotype also has been found to increase the susceptibility to Haemophilus influenzae (pneumonia, meningitis, ear infections, bronchitis, bloodstream infections, throat swelling, skin infections, and infectious arthritis), the mumps virus, Candida, and certain autoimmune disorders.

A fascinating find is that the microbiome of the AA genotype has been found to be different than the GG or AG genotype. The AA individuals lacked or were rarely colonized by several strains of the probiotic Bifidobacteria ((B. bifidum, B. adolescentis and B. catenulatum/pseudocatenulatum), while several other bacterial genotypes were more common and dominant. This could also affect folate levels because the highest extracellular folate levels were produced by four strains of B. adolescentis and two of B. pseudocatenulatum.

Studies have found that red wine consumption can significantly modulate the growth of select gut microbiota in humans, which suggests possible prebiotic benefits associated with the inclusion of red wine polyphenols in the diet. This study also found that changes in cholesterol and C-reactive protein concentrations were linked to increases in bifidobacteria.

B6-NBPF3

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
NBPF3 rs4654748	Homozygous CC

Recap



Decreases NBPF3 Gene Function: Sugar, stress, high intake of alcohol and refined flour based carbohydrates, antibiotics, oral contraceptives, ACE inhibitors, antacids, proton pump inhibitors, Phenytoin, bronchodilators, Digoxin, diuretics, hormone replacement therapy, Estradiol, MAO inhibitors, St. John's Wort and Parnate.

B6-NBPF3

Research: You may require a higher intake of B6. Homozygotes have approximately a 2.90 ng/mL lower vitamin B6 blood concentration than the wild-type genotype.

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Vitamin B6 plays a major role in neurotransmitter health. B6 deficiency can manifest as anorexia, irritability, anxiety, depression, muscle pain, bad PMS/low progesterone, nausea, seizures, migraines, dermatitis, age related macular degeneration (with low folate and B12) and lethargy.

Researchers have found an inverse association between ovarian cancer risk and vitamin B6 intake. Subjects with the highest vitamin B6 intake showed a 24 percent decrease in the likelihood of developing ovarian cancer compared to the individuals with the lowest intake.

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Women of reproductive age, especially current and former users of oral contraceptives, teenagers, male smokers, non-Hispanic African-American men, and men and women over age 65 are most at risk of B6 deficiency. Data suggests that oral contraceptive users have extremely low plasma PLP levels. Three quarters of the women who reported using oral contraceptives, but not vitamin B6 supplements, were vitamin B6 deficient.

Saturated Fat-PPAR-alpha

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE		
PPAR-alpha rs1800206	Heterozygous CG		
		_	
ecap]	

SATURATED FAT-PPAR-ALPHA

Research: The PPAR-alpha gene is mainly expressed in tissues with extensive fatty acid metabolism and its activation leads to changes in the transcription of multiple genes that regulate lipid and lipoprotein metabolism. PPAR-alpha also regulates the metabolism of amino acids in the liver, influences the apolipoprotein (Apo) genes, HDL genes, LDL genes and ketone bodies during fasting.

The PPAR-alpha polymorphism (rs1800206) has been associated with increased levels of triglycerides, total cholesterol levels, and ApoB in many populations such as Caucasian, Indian and African-Americans, however variants may be more relevant in men. The positive trade-off is that variants in the PPAR-alpha genotype may also have lower C-reactive protein levels (CRP), a marker of inflammation.

Subjects with variants in PPAR-alpha have been found to have a larger waist circumference and a higher proportion of small, dense LDL particle size (risk factor for heart health) with a higher saturated fat intake and low polyunsaturated fat intake.

In mice missing the PPAR-alpha gene, fasting decreased the plasma level of the amino acid alanine and the ketogenic amino acid tyrosine. Those with variants in this gene have been found to have a poor response to fasting due to low ketone bodies and the research has stated that this genotype may explain diverse reactions to the ketogenic diet.

An agonist is a chemical that binds to a receptor and activates the receptor to produce a biological response. PPAR-I agonists activate the gene, promoting the uptake, utilization, and breakdown of fatty acids. The activation of PPAR-I has also been demonstrated to inhibit tumor growth and angiogenesis.

Animal models have shown that PPAR-alpha activation reduces body mass, and treats insulin resistance and non-alcoholic fatty liver. Research has found that polyunsaturated fats (fish, nuts, seeds), astaxanthin (high in wild salmon), pterostilbene (blueberries, mulberries, cranberries, raw almonds), genistein (fermented soy), tomatoes, cinnamon, zinc, Lion's Mane mushroom, Gynostemma tea and L-carnitine all activate PPAR-alpha gene expression. Human studies have shown that increasing polyunsaturated fat and decreasing saturated fat in those with variants in PPAR-alpha improved triglycerides levels, LDL particle size, and body weight.

One exception for saturated fats may be coconut oil. Rat studies found that virgin coconut oil decreased tissue lipid levels, reduced the activity of the enzymes involved in lipogenesis and enhanced the rate of fatty acid catabolism, which was mediated at least in part via the PPAR-alpha dependent pathways.

Fat Metabolism-ACSL1

Below is a summary of your most significant variant genotypes:

)	GENE	GENOTYPE
	ACSL1 rs9997745	Wild Type GG

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Improves ACSL1 Gene Function: Total fat intake under 35%, lower saturated fat intake and increased PUFA intake

FAT METABOLISM-ACSL1

Research: If you have the GG genotype, it may be beneficial for fat intake to be below 35% of your total calories or have a higher intake of polyunsaturated fat from fish, nuts and seeds if you struggle with weight and high glucose.

The GG genotype had higher fasting glucose and insulin concentrations compared with the minor A allele carriers from saturated fat intake, with the result that the GG genotype was more insulin resistant. Among individuals within the top 50th percentile of PUFA intake, the metabolic syndrome risk associated with GG genotype was eliminated.

Foods that are higher on the insulin index include dairy and red meat, and insulin inhibits fat breakdown. Fat should come primarily from nuts, seeds, olive oil, avocados, poultry and fish if there are issues with fasting glucose, insulin or weight.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
HLA DQ2.5	HLA DQ2.5- rs2187668		СТ	
HLA-DQ8	HLA-DQ8- rs7454108	TT		
BCMO1 R267S BCMO1 encodes the	BCMO1 R267S- rs12934922		AT	
conversion rate from beta- carotene to vitamin A.	BCMO1 A379V- rs7501331		СТ	
FADS2 The FADS2 gene encodes the	FADS2-rs1535			GG
conversion of plant based omega-3 fatty acid alpha linolenic acid (ALA) to EPA.	FADS2-rs174575			GG
FUT2 The FUT2 gene controls prebiotic production, B12 absorption and how much bifidobacteria you carry in your digestive tract. The rs601338 SNP is found in European, African and Indian populations.	FUT2-rs601338			AA
NBPF3 NBPF3 has been associated with vitamin B6 levels.	NBPF3-rs4654748			СС

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
SLC23A1 Solute carrier family 23 member 1 (SLC23A1) is one of the two transporters which aids in the absorption of vitamin C into the body. Polymorphisms in the gene are associated with reduced plasma vitamin C levels in the body.	SLC23A1- rs33972313	СС		
ACAT1-02 The ACAT gene converts protein and fat to ATP (energy) in the mitochondria, and plays an important role in cellular cholesterol homeostasis.	ACAT1-02- rs3741049	GG		
ADIPOQ ADIPOQ encodes for adiponectin, a protein secreted by fat cells that affect insulin and glucose metabolism. Low levels of adiponectin play a role in obesity, insulin resistance and Type 2 diabetes.	ADIPOQ- rs2241766	ΤΤ		
HFE-C282Y A homozygous HFE C282Y may lead to an iron overload due to increased iron absorption and disrupted metabolism. Compound heterozygotes C282Y/H63D and single C282Y heterozygotes carry a very low risk of hemochromatosis, making the homozygous C282Y the most clinically relevant.	HFE-C282Y- rs1800562	GG		
PPAR-alpha The PPAR-alpha gene plays a vital role in fatty acid metabolism and ketosis, and is considered one of the most critical targets for ameliorating abnormalities with triglycerides, HDL, LDL, VLDL, and ApoB.	PPAR-alpha- rs1800206		CG	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
ACSL1 Long-chain acyl CoA synthetase 1 (ACSL1) plays an important role in fatty acid metabolism and triglyceride synthesis. Disturbance of these pathways may result in dyslipidemia and insulin resistance, hallmarks of the metabolic syndrome.	ACSL1-rs9997745	GG		
FTO Polymorphisms in the FTO genes have been shown to cause higher ghrelin levels	FTO-rs9939609		AT	
(hunger hormone) in many populations, which can create a larger appetite and the potential for overeating.	FTO-rs17817449		GT	
APOA2 The APOA2 gene contains instructions for making a protein called apolipoprotein A- II, which is found in HDL cholesterol particles. The homozygous genotype has been linked to saturated fat intake and weight gain.	APOA2-rs5082	AA		
TCF7L2 TCF7L2 polymorphisms have been associated with low incretin hormones and impaired insulin secretion.	TCF7L2-rs7903146	СС		
LCT LCT is the gene connected with the ability to breakdown lactose in dairy.	LCT-rs4988235			AA
APB1 APB1 is encodes for the DAO enzyme to breakdown histamines primarily in the digestive tract. The homozygous genotype may increase the risk of migraines from histamines in women or a hypersensitivity to Aspirin in men.	APB1-rs10156191	CC		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
ABCG2 (Q141K) The ABCG2 (Q141K) gene is located at the membrane of kidney proximal tubule cells, where it mediates renal urate secretion. Variants in this gene are linked to reduced uric acid excretion.	ABCG2 (Q141K)- rs2231142	GG		
ALDH2 Alcohol metabolism in the liver most commonly involves the enzymes alcohol dehydrogenase and aldehyde dehydrogenase, metabolizing alcohol to acetaldehyde, and then to acetate. ALDH2 encodes for aldehyde dehydrogenase, and variants can affect the levels of acetaldehyde and therefore the carcinogenic effect of alcohol.	ALDH2-rs671	GG		
ADRB2 The beta-2-adrenergic receptor (ADRB2) is the main target of the catecholamine epinephrine, and a primary mediator of the stress response. ADRB2 is widely expressed both in the gastrointestinal tract and in the CNS.	ADRB2-rs1042714	СС		
PPCDC PPCDC is necessary for the biosynthesis of coenzyme A and variants in this SNP are associated with serum zinc levels.	PPCDC-rs2120019		СТ	
SELENBP1 The Protein Selenium Binding 1 gene codes for an integral membrane protein involved in antigen presentation and serum copper levels.	SELENBP1- rs2769264		GT	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
TFR2 The TFR2 gene provides instructions for making a protein called transferrin receptor 2 to help iron enter liver cells. The receptor on the surface of liver cells binds to transferrin, which transports iron through the blood to tissues throughout the body. When transferrin binds to transferrin receptor 2, iron is allowed to enter the cell.	TFR2-rs7385804			СС

MY CLINICAL RESEARCH SUMMARY: METHYLATION CYCLE

Folate-MTHFR 677

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
MTHFR 677 rs1801133	Homozygous AA

Recap

Improves MTHFR 677 Gene Function: Riboflavin and methylfolate.

	Decreases MTHFR 677 Gene Function: Proton pump inhibitors, oral contraceptives, NSAIDs, anticonvulsants, antivirals, antibiotics, acid blockers/antacids and hypothyroidism.
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FOLATE-MTHFR 677

The highest cluster of the homozygous MTHFR 677 genotype is found in Mexico and Hispanics in the US, Italy, Northern China, Spain, and France. The lowest frequency is found in black people (within and outside Africa), Inuit, Finland, Canada, the Netherlands, Germany, and Russia.

The homozygous MTHFR 677 gene has been found to have a 50-70% reduced enzymatic function. One hypothesis is that the homozygous MTHFR genotype was selected based on higher folate intake and UV exposure, both common in Mediterranean climates. What also happens in the body when MTHFR enzymatic function is reduced is that thymidine production increases. Thymidine enhances the repair of UV-induced DNA damage to help quickly repair sun damage.

The sun also depletes folate due to UV radiation. However, darker skin - common in the Mediterranean - contains higher melanin levels, which helps protect against folate loss. Increased thymidine and darker skin protected against the hot sun of the Mediterranean. At the same time, the environment provided more folate-rich fruits and vegetables to supply more dietary folate for other biochemical functions.

Another hypothesis is that malaria exposure - caused by a parasite through mosquito bites and has been prevalent in the Eastern Mediterranean and Southeast Asia - altered the MTHFR genotype selection. The malaria parasite needs higher amounts of folate to survive and replicate. Reduced MTHFR function lowered folate levels and boosted levels of thymidine, which may increase lymphocyte replication and immune function in response to malaria.

One concern with the homozygous MTHFR 677 genotype is high homocysteine. High homocysteine has been implicated in amyloid buildup, DNA damage and cancer, mitochondrial dysfunction, cardiovascular disease, age-related macular degeneration, apoptosis of neurons and depression. Multiple studies have found that riboflavin is essential for MTHFR 677 and that the homozygous genotype is associated with high homocysteine when riboflavin status is low. Reduced levels of methylfolate can also lead to decreased production of neurotransmitters, reduced conversion of homocysteine to methionine, and reduced s-adenosylmethionine (SAMe) concentrations.

One study in 259 post-menopausal women found that for those with variants in CYP1B1 (rs1056836), KRAS (rs61764370) and MTHFR (rs1801133 and rs1801131), oral contraceptives and hormone replacement therapy was associated with shorter leukocyte telomere length. Shorter leukocyte telomeres are connected to premature aging and may increase the risk of cancer, cardiovascular disease, obesity, diabetes, chronic pain, and sensitivity to perceived psychological stress.

A large meta-analysis showed the lack of statistically significant association between MTHFR mutations and coronary heart disease except in the Middle East and Japan, where it portrayed statistical significance.

It is essential to consider riboflavin intake, PEMT, MTR/MTRR, BHMT and CBS activity to assess overall homocysteine metabolism. Too high or too low levels of B12, B6, folate or their co-factors may cause dysregulation of methyl donor activity.

B12-MTRR

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
MTRR A66G rs1801394	Homozygous GG
Recap	
Improves MTRR Gene Function: B12, B6	and folate.

B12-MTRR

Research: Methionine synthase reductase (MTRR) is a vital enzyme of homocysteine/methionine metabolic pathway and is required for the conversion of inactive form of methionine synthase (MTR) to its active form. MTRR helps recycle B12. If MTR is working 100%, MTRR variants may be less pronounced. Variants in both MTR and MTRR may lead to issues with B12.

A clinically important allelic variant of MTRR A66G, with less enzymatic activity is reported with worldwide prevalence rate of 30%. The very high frequency of the homozygous genotype (greater than normal allele) was reported from Italy, France, Ireland, Netherlands and India.

Several epidemiological and case control studies have reported that the GG genotype may be a risk factor for several disease/disorders like neural tube defects, Down syndrome, coronary artery disease, male infertility and cancer through sustained hypomethylation if not addressed. An additional effect is a decreased availability of folate. Several studies show that DNA hypomethylation is the main causative factor of defective gene expression.

Caucasions with late life depression who were homozygous (GG) were more likely to still be depressed after a course of SSRI antidepressant treatment compared to individuals who are homozygous at the A allele or who were AG heterozygotes. This points toward the hypothesis that factors which elevate homocysteine concentrations have an adverse effect on depression treatment response.

Heterozygous or homozygous variants in MTRR may require more B6, folate and B12 if combined with MTHFR 677 and MTR variants. If you suffer from migraines, B6, folate and B12 may be effective in reducing migraines. Research has also found B6, folate and B12 to help prevent macular degeneration.

Choline-PEMT

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
PEMT rs7946	Heterozygous CT
PEMT rs12325817	Homozygous GG

77 Recap



Improves PEMT Gene Function: Choline, vitamin C and estrogen.

Decreases Gene Function: Nighttime pain relievers, antihistamines, anti-seizure medications, sleep aids, antidepressants, incontinence drugs and narcotic pain relievers.

Research: Phosphatidylethanolamine-N methyltransferase (PEMT) catalyzes the synthesis of phosphatidylcholine.

Choline is responsible for shuttling fat out of the liver, aiding the gallbladder, healthy cell membranes to protect against inflammation, lowering anxiety, preventing damage from glutamate spikes, deep sleep, healthy DNA, healthy pregnancy and breast health. Non-Alcoholic Fatty Liver Syndrome occurs mainly from a choline deficiency. Choline deficiency also increases sensitivity to carcinogenic chemicals, mycotoxins and vegetable oils due to poor cell membrane health.

Research shows that the highest dietary intake of choline is found from people in the Northern countries, whereas Mediterranean countries had the lowest intake. Worldwide, total choline intake in adults ranges from 284 mg/day to 468 mg/day for men, from Taiwan and Sweden, respectively; and from 263 mg/day to 374 mg/day for women, from Mexico and Sweden. Major food sources of dietary choline vary by country. For example, eggs, meat, and dairy are the major sources of total dietary choline in New Zealand, while eggs, seafood, meats, and soy products are the predominant sources in Japan and China.

Having one or more T alleles at rs7946 is associated with having lower phosphatidylcholine production in the liver.

More than 40% of women have a genetic polymorphism in PEMT (rs12325817) that makes this gene unresponsive to estrogen, which creates the same high choline requirement as men. These women may be especially sensitive to dietary choline variations during pregnancy. One study found that the highest quintile of choline consumption was associated with a lower risk of breast cancer compared with the lowest quintile.

Eighty percent of the women who were homozygous for the rs12325817 SNP manifested signs of choline depletion (liver or muscle dysfunction), relative to 43% of subjects carrying one copy of the variant allele and 13% of subjects without the SNP. Almost 75% of the North Carolina population in the United States has one variant allele.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
MTHFR 677 The MTHFR 677 gene encodes the MTHFR gene to convert folate into the active form, methylfolate. Variants in this gene slow down enzymatic function.	MTHFR 677- rs1801133			AA
MTHFR 1298 MTHFR 1298 is involved in converting 5-methylfolate (5MTHF) to tetrahydrofolate (THF). Unlike MTHFR 677, the 1298 variant does not lead to elevated homocysteine levels unless paired with a heterozygous MTHFR 677.	MTHFR 1298- rs1801131	тт		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
MTHFD1 G1958A (Methylenetetrahydrofolate dehydrogenase 1) encodes a protein that possesses three distinct enzymatic activities in the interconversion of 1- carbon derivatives of tetrahydrofolate.	MTHFD1 G1958A- rs2236225	GG		
DHFR A20965G Dihydrofolate reductase (DHFR) catalyzes the	DHFR A20965G- rs1643659	тт		
reduction of dihydrofolate to tetrahydrofolate (THF) and affect synthetic folic acid metabolism.	DHFR C19483A- rs1677693	GG		
MTR A2756G MTR (methionine synthase) combines folate, methyl B12 and homocysteine into methionine.	MTR A2756G- rs1805087	AA		
MTRR A66G MTRR attaches a methyl group to B12 and variants here will slow the process. When both MTR and MTRR exist, dysfunction can occur.	MTRR A66G- rs1801394			GG
TCN2 C766G Transcobalamin II (TCN2, or holotranscobalamin when bound) transports B12 to peripheral tissues. Variants in this gene may affect B12 transport.	TCN2 C766G- rs1801198		CG	
PEMT Variants in PEMT may increase the need for choline and increase the sensitivity to anticholinergic drugs.	PEMT-rs7946		СТ	
	PEMT-rs12325817			GG
CBS A13637G The Cystathione Beta- Synthase (CBS) enzyme pulls homocysteine to hydrogen sulfide (H2S) and glutathione, requiring B6 and SAMe as a modulator.	CBS A13637G- rs2851391	сс		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
CBS The Cystathione Beta- Synthase (CBS) enzyme pulls homocysteine to hydrogen	CBS-rs234709	СС		
sulfide (H2S) and glutathione, requiring B6 and SAMe as a modulator. CBS rs234709 and rs4920037 assists in arsenic detoxification.	CBS 191150T- rs4920037	GG		

MY CLINICAL RESEARCH SUMMARY: HORMONE SUPPORT

Vitamin D-CYP2R1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
CYP2R1 rs10741657	Homozygous GG

Recap

Improves CYP2R1 Gene Function: Sun exposure, adequate vitamin D intake and vitamin D co-factors.

Decre

Decreases CYP2R1 Gene Function: Lack of sun exposure, high fructose intake and lack of vitamin D co-factors.

VITAMIN D-CYP2R1

Research: Studies confirm that CYP2R1 is the principal 25-hydroxylase in humans and demonstrates that CYP2R1 alleles have dosage-dependent effects on vitamin D homeostasis.

A 2018 meta-analysis of sixteen articles with a total of 52,417 participants was reviewed for rs10741657. The GG genotype was associated with a clear descending trend of 25(OH)D levels when compared with the AA genotype in Caucasian and Asian populations.

Research has shown that oral administration of vitamin D led to negligible increases in serum 25-hydroxy-vitamin D for homozygotes, and significantly lower increases in serum 25-hydroxy-vitamin D in heterozygous subjects than in control subjects. The heterozygous effect may only be relevant in Caucasian populations.

Vitamin D can influence the expression of more than 1,000 genes and vitamin D deficiency has been linked to fatty liver, seizures, infertility, osteoporosis, cancer, autism (mother deficient), depression, heart attacks, Alzheimer's, dementia, high blood pressure, low testosterone in men, autoimmune disorders and more.

The literature is mixed on optimal vitamin D levels, which most likely vary based on your heritage, skin color and current health issues. The most well documented cause of Vitamin D deficiency is inadequate sunlight exposure such as high latitude countries. Paradoxically, despite its high sunlight hours, vitamin D deficiency is well recognized in Middle Eastern women, inner city young adults in America, athletes and dancers in Israel, elite gymnasts in Australia, young Hawaiian surfers, and adolescent girls in England.

For athletes, vitamin D deficiency has long been associated with muscle weakness and suboptimal muscle function. A positive relationship between serum vitamin D level and jump height, jump velocity and power was found in young women.

Clinical vitamin D deficiency is below 20 ng/ml. There is little evidence to prove there is a benefit for levels above 50 ng/ml. The latest cancer research has found that women with 25(OH)D concentrations greater than 40 ng/ml had a 67% lower risk of cancer than women with concentrations less than 20 ng/ml. Pesticides have been linked to suppressing vitamin D levels and creating a vitamin D deficiency. Your PON1 gene function should also be assessed.

Research has found that sunlight is the optimal way to optimize vitamin D levels along with exercise, vitamin D rich foods and vitamin D cofactors, however supplementation may be necessary.

Estrogen Metabolism-CYP1A2

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
CYP1A2 rs762551	Homozygous AA

Recap

Improves CYP1A2 Gene Function: Unfiltered fermented drinks (Kombucha, beer, wine), hops, marinades, cruciferous vegetables, blueberries, blackberries, red grapes, kiwi, watermelon, parsley, and spinach.



Decreases CYP1A2 Gene Function: Heterocyclic amines, nitrosamines, aflatoxin B1, polycyclic aromatic hydrocarbons, dioxins, and I-naphthoflavone. Omeprazole and primaquine are inducers. Caffeine and Tylenol combined with these compounds can make the effect worse.

ESTROGEN METABOLISM-CYP1A2

CYP1A2 is a key enzyme in caffeine metabolism and the 2-hydroxylation of the main estrogens, estrone, and estradiol. 2hydroxylation and 16a-hydroxylation are two mutually exclusive pathways in estrogen metabolism. 2-hydroxyestrone acts as a weak estrogen or anti-estrogen. 16I-OHE1 acts as a procarcinogen.

Coffee may protect against breast cancer by altering estrogen metabolism. Women with higher coffee intake and the CYP1A2 homozygous AA fast metabolizer genotype have a ratio of high 2-hydroxyestrone to low 16^{II}-OHE1. Researchers found that higher coffee intake was more protective against ER-positive breast cancer.

In men, a 2019 study found that low to moderate coffee intake and the AA fast caffeine metabolizer genotype were less likely to experience prostate grade cancer progression than non-consumers. In a large, pooled cohort of men with prostate cancer, coffee intake of more than 2.5 cups per day was associated with longer survival with the AA fast metabolizer genotype.

Estrogen Metabolism-COMT

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
COMT rs4680	Heterozygous AG

Recap



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Decreases Gene Function: Chronic stress, sugar, proton pump inhibitors, aspartame, low magnesium levels, low vitamin C levels, low and high copper levels, constipation, xenoestrogens, high homocysteine levels, high SAH levels, estrogenbased medications, and mercury toxicity.

ESTROGEN METABOLISM-COMT

COMT is a phase II enzyme involved in the inactivation of catechol estrogens that can otherwise lead to cancerous growth, while also increasing 2-methoxyestradiol, a metabolite that has been shown to inhibit the growth of breast cancer cells.

Variants in COMT V158M have been shown to decrease enzymatic activity and consequently increases the risk of carcinogenesis due to the accumulation of estrogen metabolites. COMT has been extensively investigated for correlation with different cancer risks including esophageal cancer, colorectal cancer, hepatocellular, carcinoma, lung cancer, breast cancer, ovarian cancer, endometrial cancer, testicular germ cell tumor, and bladder cancer with mixed results.

82 Due to the COMT V158M heterozygous and homozygous genotypes potentially having reduced estrogen clearance, slowing this pathway down further with chronic stress and a high catecholamine intake combined with poor gut health and low magnesium intake may affect the level of harmful estrogen metabolites.

However, this doesn't mean catecholamines should be avoided. It simply means that the dosage should be altered. For

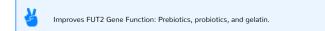
example, green tea has been found to be beneficial for breast cancer prevention in the COMT heterozygous and homozygous genotype because these individuals retained the polyphenols the longest. The wild type may need more to achieve the same benefit. Less is more for COMT variants.

Estrobolome-FUT2

Below is a summary of your most significant variant genotypes:



Recap



Decreases FUT2 Gene Function: Antibiotics, proton pump inhibitors, glyphosate, and sucralose.

ESTROBOLOME-FUT2

The estrobolome involves the bacterial genes capable of metabolizing estrogens. Gut dysbiosis can lead to increased colonies of bacteria that produce beta-glucuronidase, which deconjugates estrogen, and is reabsorbed into circulation. These deconjugated estrogens have carcinogenic potential.

The non-secretor FUT2 homozygous genotype has been found to have low bifidobacteria levels. Lactic acid bacteria, including bifidobacteria, were revealed to exert chemopreventative effects on colon, bladder, liver, breast, and gastric cancer types. Bifidobacteria is one of the species responsible for the metabolism of glucosinolates to isothiocyanates, the anti-cancer compound found in cruciferous vegetables.

Research has proposed that the microbiome contributes to breast carcinogenesis by modifying systemic estrogen levels. The Cleveland Clinic recently identified differences in the microbiome of breast cancer patients, showing a profound difference in bacteria. The microbiome's ability to affect systemic hormone levels may also be important in prostate cancer that is dually affected by estrogen and androgen levels.

In a North American case-control study involving 2,266 women with breast cancer and 7,953 healthy controls, all classes of antibiotics were associated with increased breast cancer risk, and the association remained after adjustment for factors including premenopausal or postmenopausal status, age at menarche, and family history.

Rat studies found that the consumption of sucralose decreased the total number of anaerobic and aerobic bacteria, bifidobacteria, lactobacilli, Bacteroides, and clostridium.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
SHBG Sex Hormone Binding Globulin (SHBG) is synthesized in the	SHBG-rs1799941	GG		
liver, and in the blood it transports and regulates the access of sex steroids to their target tissues.	SHBG-rs12150660	GG		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
DI01 DI01 is connected to thyroid health and is responsible for the deiodination of T4 into T3.	DI01-rs2235544		AC	
DI02 DI02 is connected to thyroid health and is responsible for the deiodination of T4 into T3. D2 is the only activating deiodinase in the brain.	DI02-rs225014	Π		
CYP2R1 Vitamin D is technically a hormone, and CYP2R1 is connected to circulating vitamin D levels.	CYP2R1- rs10741657			GG
CYP1A1 CYP1A1 is in the estrogen metabolism pathway along with CYP1B1, CYP1A2, CYP31A, SULT's and COMT.	CYP1A1-rs1048943	Π		
CYP2C19*17 Genetic variability impacts expression and activity of CYP2C19 and therefore can influence drug metabolism and catabolism of estrogens.	CYP2C19*17- rs12248560	сс		
CYP1A2 CYP1A2 is a key enzyme in caffeine metabolism and the 2- hydroxylation of the main estrogens, estrone, and estradiol.	CYP1A2-rs762551			AA
COMT COMT is involved in catecholamine, dopamine, adrenaline, and estrogen metabolism through the inactivation of the catechol estrogens.	COMT-rs4680		AG	
FUT2 The FUT2 gene controls prebiotic production, B12 absorption, and how much bifidobacteria you carry in your digestive tract.	FUT2-rs601338			AA

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
MTNR1B The MTNR1B gene encodes for the melatonin receptor 1B.	MTNR1B- rs10830963	сс		

MY CLINICAL RESEARCH SUMMARY: MENTAL HEALTH & COGNITIVE PERFORMANCE

MAO-Serotonin

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
MAO-A rs6323	Wild Type TT

Recap

Improves MAO-A Gene Function: Vitamin B6, folate, B12, B2, magnesium, vitamin C and probiotics.



Decreases Gene Function: Antibiotics, aspartame, oral contraceptives, proton pump inhibitors, high estrogen levels, constipation and deficiencies in the vitamins and minerals above.

MAO-SEROTONIN

Research: MAO-A (Monoamine oxidase A) is a critical enzyme involved in breaking down important neurotransmitters such as serotonin, estrogen, norepinephrine, and dopamine.

You have the TT genotype that encodes for the slow activity of the MAO-A enzyme, which helps preserve serotonin. The TT genotype, constipation, and high estrogen cause a very slow MAO-A enzyme in females, which can increase sensitivity to stress due to high levels of estrogen, serotonin, and poor detoxification. Fiber, B6, B2, and magnesium are crucial for this genotype. The TT genotype may also make you more sensitive to MAO inhibitors, SSRIs, antibiotics and oral contraceptives depending on your estrogen status.

Research has found that the transport of tryptophan (precursor to serotonin) is lower in children with ADHD, and one study found that the rs6323 TT genotype was a protective factor against ADHD in Korean children.

Serotonin levels are more complicated than assessing just MAO-A, including gender, estrogen fluctuations, chronic stress, antibiotic use and general gut health, COMT function, and serotonin transportation and receptor genes. Serotonin is responsible for well-being, happiness, memory, and appetite. When serotonin is too low, it can cause depression, lack of ambition, and a struggle to derive pleasure from life. When it is dysregulated, it can cause IBS, mania, OCD, and drug-induced serotonin syndrome.

To modulate healthy serotonin levels, research has found that aerobic exercise to fatigue, strength training, yoga, and nature walks all are effective. Fermented foods and probiotics (90% of serotonin is made in the gut), getting more sunlight, or taking vitamin D, dark chocolate, fish oil, and a weekly massage are also excellent strategies. However, both extremes of a sedentary lifestyle and excessive exercise negatively affect MAO-A.

Serotonin Receptor-Memory

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
5-HT2A rs6314	Heterozygous AG

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Improves Gene Function: Mindfulness training, meditation, yoga, exercise to fatigue, time in nature, strength training, green or black tea, tryptophan, B2, B6, B12, folate, and ginkgo biloba.

SEROTONIN RECEPTOR-MEMORY

Research: Serotonin plays an important role in memory formation and results suggest a possible role of 5-HT2A receptors in memory consolidation. Episodic memory is the ability to recall details regarding personal experiences, names of people, specific events, and what exactly occurred. It is this type of memory that is most affected by the serotonin gene 5-HT2A rs6314 and takes place mainly in the hippocampus of the brain.

Research has demonstrated a functional influence of the rs6314 SNP on human memory, with the A carriers showing 21% poorer episodic memory performance than the GG genotype. The A allele has been found with a frequency of 9.3% in Caucasian populations.

In one study, healthy young volunteers viewed six sets of semantically unrelated words and performed an incidental episodic memory task with immediate and delayed recall after 5 min and 24 hours. In both conditions, A carriers showed lower memory performance compared to the GG genotype. Moreover, it has been shown that this effect is modified by age in that it can be detected in young to middle-aged, but not in older populations. In another study, A carriers exhibited poor verbal delayed recall and recognition, but performed equivalent to controls on tests of immediate recall, attentional, and executive function. Further analysis showed A carries had reduced grey matter volume in the left hippocampus and left inferior temporal gyrus and, bilaterally, in the middle and superior temporal gyrus. The white matter was reduced in the left temporal regions as well.

Training in mindfulness, classically described as a receptive attentiveness to present events and experiences, has been shown to improve attention and episodic working memory. This training may have relevance for potential educational and occupational performance. An example of not being mindful and attentive would be getting distracted by a smartphone during a conversation or event, consequently missing visual and auditory cues for later storage. Harvard researchers found that after eight weeks, subjects spending 27 minutes per day on mindfulness exercises (like meditation) increased grey matter volume in the hippocampus, one of the most important sections of the brain for learning and memory.

Research has shown that the reliance on technology may be decreasing our ability to store long term memories. Researchers have argued that the reliance on smartphones and related technologies is not aiding mental functioning, but rather, is having a negative impact on our ability to think, remember, pay attention, and regulate emotion. Recent studies on memory and technology use found that participants who took pictures of an object vs. those who did not had a more difficult time recognizing these objects at a later date. Auditory memory was also impaired. Due to ubiquitous use of cameras on smartphones for day to day events, the consequences on long-term memory have yet to be researched.

Due to 5-HT2A receptor's role in BDNF regulation, the A allele for rs6314 was also found to cause a lower expression of BDNF in hippocampal neurons and potentially a decrease in neural plasticity. Theanine, a component of green tea and black tea, has been shown to increase BDNF levels, modulate serotonin and dopamine levels, and improve learning and memory. A human study that looked at matcha tea consumed in a realistic dose found that it can induce slight effects on speed of attention and episodic memory.

Cannabis use has been associated with episodic memory impairments and abnormal hippocampus morphology among both healthy individuals and schizophrenia subjects.

In a parallel, randomized and controlled trial, 88 patients diagnosed with exhaustion disorder participated in a 24-week multimodal rehabilitation program. After 12 weeks in the program, the patients were randomized to either a 12-week aerobic training intervention or to a control group with no additional training. The aerobic training group significantly improved in maximal oxygen uptake (VO2 max) and episodic memory performance.

Ginkgo biloba has been found to target the 5-HT2A serotonin receptor and improved episodic memory in patients with mildcognitive impairment. However, it was not found to improve memory in young adults.

Episodic memory is believed to be affected by both the hippocampus and prefrontal cortex function. Due to acetylcholine's role in memory in the hippocampal region, researchers found that increasing acetylcholine levels improved visual episodic memory.

Research has shown that tryptophan depletion impairs episodic memory and verbal short-term memory function in human subjects, while rodent studies have also shown that tryptophan-free protein-carbohydrate mixture to rats significantly lowered hippocampus tryptophan levels and produced impaired performance in visual working memory. However, dietary tryptophan

was found to increase brain serotonin levels and improve memory in rats.

5-HT2A receptors are the most abundant serotonin receptor in prefrontal cortex and regulate the dopamine pathway. Researchers found that for those with the GG COMT V158M genotype (low dopamine), a COMT inhibitor improved verbal episodic memory and prefrontal cortex function, while the AA COMT V158M homozygous genotype (high dopamine) got worse. If you have the GG COMT V158M genotype, you may improve episodic memory through coffee, chocolate, green tea, black tea, citrus, and bananas by slowing down the COMT enzyme and increasing dopamine.

Serotonin Receptor-Stress

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
5-HT2A rs6311	Heterozygous CT
5-HT2A rs6313	Heterozygous AG

Recap



Improves Gene Function: Moderate intensity aerobic exercise, cognitive behavioral therapy, mindfulness training, meditation, yoga, tryptophan, green or black tea, prebiotics, probiotics, B2, B6, B12, and folate.

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Decreases 5-HT2A Gene Function: Chronic stress, poor gut flora, high-dose lithium, cannabis abuse, and excessive smartphone use.

SEROTONIN RECEPTOR-STRESS

Research: The serotonin 2A receptor (5-HT2A) has been implicated in mental disorders with complex etiologies that are still not clearly understood, in processes such as learning and memory, and also in neurogenesis. Although the functional significance of 5-HT2A polymorphisms are not entirely understood, there is evidence that rs6311 modulates transcription factor binding and promoter methylation, affecting gene transcription (the first step of gene expression).

The T allele of the 5-HT2A gene rs6311 has been shown to increase the 5-HT2A expression in vitro and is associated with anxiety, IBS and depressive disorders. It has also been hypothesized that 5-HT2A variants may influence resting vagal activity among persons with chronically high levels of perceived stress.

One meta-analysis showed that the T allele of rs6311 or the linked A allele of rs6313 was significantly associated with obsessive compulsive disorder (OCD). This result was confirmed in the author's subsequent comprehensive meta-analysis in 2016 with a larger dataset. Multiple studies in this analysis indicated that the rs6311 T allele was more abundant in females with OCD compared to control females.

Another meta-analysis of 37 twin samples suggests that obsessions and compulsions arise from a combination of genetic factors and non-shared environment. OCD might be shaped by a large number of genes of modest impact, which combine to influence the risk for developing OCD. Polymorphisms in genes related to BDNF, GABA, glutamate, serotonin, acetylcholine, glycine, ubiquitin, bradykinin, myelinization, TNFA, gender and environmental trauma may all have a cumulative effect on whether or not someone develops OCD.

Psoriasis is a chronic inflammatory skin disease affecting about 2-4% of the population worldwide, and is thought to be a multifactorial disease with both genetic and immunogenic backgrounds. Psoriasis occurs in connection with stress and mood disorders and is apparently induced in patients who have been treated with antidepressants. The serotonergic system, which consists of serotonin-producing cells, serotonin receptors and serotonin transporters, may play a significant role in psoriasis.

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Theanine, a component of green tea and black tea, has been shown to increase BDNF levels, modulate serotonin and dopamine levels, and improve learning and memory. It has shown promise as an adjunct therapy for schizophrenia and depression, and researchers believe there may also be an application for anxiety disorders, panic disorder, OCD, and bipolar disorder.

Vagus nerve stimulation may be a promising add-on treatment for anxiety, depression, PTSD, seizures, and inflammatory bowel disease. Natural ways to stimulate the vagus nerve and increase vagal tone include singing, deep breathing, meditation

and yoga. Another way is to make a dietary shift towards good gut bacteria, shown to influence the activity of the vagus nerve.

In human volunteers as well as in a rat model, administration of a probiotic formulation consisting of Lactobacillus helveticus R0052 (traditionally used in the manufacture of Swiss-type cheeses and long-ripened Italian cheeses such as Emmental, Gruyere, Grana Padano and Parmigiano Reggiano) and Bifidobacterium longum R0175A (colonizes at birth, but levels vary genetically) significantly attenuated psychological distress and reduced anxiety-like behavior. Research has also found that prebiotics can improve non-REM sleep as well as REM sleep after a stressful event.

One pilot study found that a 12-week moderate intensity aerobic exercise program reduced OCD symptoms and the reductions lasted 6 months later.

Another study combined cognitive behavioral therapy and a 12-week moderate intensity aerobic exercise program with tremendous results, exceeding effects typically observed with individual and group-based cognitive behavioral therapy for OCD based on leading meta-analytic reviews.

Dopamine, Adrenaline and Estrogen-COMT

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
COMT V158M rs4680	Heterozygous AG
COMT rs4633	Heterozygous CT

Recap

Improves COMT Gene Function: Vitamin C, magnesium, and copper (copper should not be too low or too high).

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Decreases Gene Function: Chronic stress, sugar, proton pump inhibitors, aspartame, low magnesium levels, low vitamin C levels, low and high copper levels, constipation, xenoestrogens, high homocysteine levels, high SAH levels, estrogenbased medications and mercury toxicity.

DOPAMINE, ADRENALINE AND ESTROGEN-COMT

Research: COMT (catecholamine methyltransferase) shares a pathway with MAO-A and is the gene for dopamine, estrogen, adrenaline and catecholamine metabolism. This pathway requires magnesium, vitamin C and copper as co-factors.

Studies have found that the AG allele in COMT V158M (rs4680) results in an intermediate enzymatic function, while the wild type GG has fast activity, and the AA homozygous genotype has 4-5 times lower COMT activity. This means that dopamine and adrenaline levels should be more level in the AG genotype. However, multiple studies have shown that the AG genotype may fall on the higher end of the dopamine spectrum with cognitive tests.

Research has shown that individuals carrying the A allele of rs4680 or T allele of rs4633 scored significantly higher on insight problem-solving tasks, and for the COMT H62H rs4633 gene, the homozygous TT and heterozygous TC carriers had higher insight problem-solving scores than those with wild-type CC genotype.

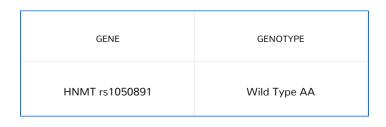
A small study found that Caucasian carriers of at least one G allele showed a greater effect for social facilitation and cooperativeness (working together in a group) than the AA homozygous group for COMT V158M.

There are both benefits and detrimental aspects to variants in COMT. The downside of the A allele in COMT V158 is that the body overreacts to stress and pressure that can lead to anxiety, depression, impulsiveness, obsessive behavior, irritability, ADHD and abnormal behavior. It can also create a sensitivity to a higher intake of catecholamines (coffee, black tea, green tea, red wine, chocolate), especially in a stressed state, leading to high dopamine and adrenaline levels making the stress response worse. However, green tea has been found to be beneficial for breast cancer prevention in the AG and AA genotype because these individuals retained the polyphenols the longest. Other genetic variants involved in dopamine transport and receptor function also influence this magnitude.

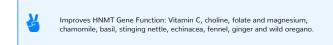
Having a heterozygous variant in COMT V158M may increase your need for magnesium, vitamin C, and healthy copper levels (not too high or low). Compound weight lifting (squats, bench press, deadlift), sprints, and chopping wood can assist a slow COMT enzyme by increasing testosterone levels, which speed up the pathway and lower the stress response. Supplementation of magnesium and vitamin C may be essential to modulate COMT due to low magnesium levels in the water and soil, or lack of freshly picked fruits and vegetables for vitamin C, and chronic stress levels.

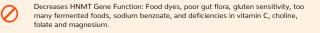
Histamines and Hyperactivity-HNMT

Below is a summary of your most significant variant genotypes:



Recap





HISTAMINES AND HYPERACTIVITY-HNMT

Research: HNMT stands for histamine methyltransferase. HNMT is the primary enzyme responsible for histamine metabolism in the brain. Dysfunction of the histaminergic nervous system is associated with various neuropsychiatric disorders including narcolepsy, Alzhemer's disease, Tourette's syndrome, eating disorders, and depression.

This gene requires adequate methyl donors from methionine and choline. If you do not have enough methyl groups available, you may be more prone to high histamine levels. HNMT polymorphisms differ considerably between Chinese and American populations.

In a 2019 mice study, results demonstrated that HNMT played an essential role in regulating brain histamine concentration, controlling aggression and sleep-wake cycles. HNMT disruption did not affect histamine concentration of the skin and stomach.

In children with ADHD, the adverse effect of food dyes and sodium benzoate on ADHD symptoms was determined by histamine degradation in the rs1050891 AA HNMT polymorphism. Histamine is lowered by vitamin C, choline, folate, magnesium, chamomile, basil, stinging nettle, echinacea, fennel, ginger and wild oregano.

Anandamide-FAAH

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
FAAH rs324420	Wild Type CC

Recap



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Improves FAAH Gene Function: Exercise over 30 minutes, red clover tea (women), kaempferol, cacao, genistein (fermented soy), Echinacea, 7-hydroxyflavone (parsley, onions, berries, tea, and citrus fruits), Il-caryophyllene (cloves, rosemary, hops).

Decreases FAAH Gene Function: Pesticides and phthalates

ANANDAMIDE-FAAH

Anandamide is a neurotransmitter and endogenous cannabinoid, and is known as the "bliss" molecule that targets the

endocannabinoid system.

The endocannabinoid system is involved in many physiological processes including reward, addiction, fertility, pain and energy regulation. This system was named from the cannabis plant, such as marijuana and hemp. THC closely resembles anandamide.

The endocannabinoids play a significant role in pain modulation and inflammation, and have been demonstrated to relieve pain by activating the CB1 and CB2 receptors.

The wild-type genotype (CC) encodes for the fast activity of FAAH, and therefore naturally leads to lower anandamide levels. Those with the homozygous genotype (AA), have the slow-activity of FAAH and naturally higher levels of anandamide. This means that the CC individuals may have more anxiety and have to work harder to achieve higher levels of happiness, while the AA individuals have less anxiety and naturally higher levels of the "bliss" molecule that stimulate feelings of happiness.

Low levels of anandamide have been linked to slower extinction of fear memories and a heightened stress response to threatening situations than those with higher anandamide levels. Healthy volunteers who carried the rs324420 "A" allele (low FAAH activity, high anandamide levels) had much less amygdala activation when placed in a threatening situation. They also had a weaker correlation between amygdala activation and trait anxiety, which is a general tendency to perceive situations to be threatening and to respond to such situations with subjective feelings of apprehension and tension.

Pesticides such as chlorpyrifos and diazinon alter the endocannabinod system and researchers have hypothesized that eating organic foods lacking pesticide residues may promote endocannabinoid balance. Phthalates are plasticizers added to water bottles, tin cans, food packaging, and even the enteric coating of pharmaceutical pills. Phthalates may act as endocrine disruptors and carcinogens, and have been found to block CB1 receptors, found in the brain.

However, there are also ways for people to lower excessive levels of chronic stress and anxiety by increasing anandamide levels in the body. One of best ways to do this is with exercise. Endorphins (endogenous opioids) enhance the effects of cannabinoids and what has been known as the "runner's high" may in fact be the increase of anandamide. Research found that running and biking over 30 minutes, along with strenuous hiking at high altitude significantly increased anandamide.

Clinical anecdotes suggest that stress-reduction techniques, such as meditation, yoga, and deep breathing exercises impart mild cannabimimetic effects.

Brain Health-PEMT

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
PEMT rs7946	Heterozygous CT
PEMT rs12325817	Homozygous GG

Recap



BRAIN HEALTH-PEMT

Research: Choline is required for acetylcholine, a neurotransmitter of the vagus nerve that innervates multiple organs including the lungs, heart, liver, stomach, ovaries, and temporal lobe of the brain. A deficiency could affect all of these, especially memory. Acetylcholine also plays a role in promoting REM sleep.

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Having one or more T alleles at rs7946 is associated with having lower phosphatidylcholine production in the liver.

Eighty percent of the women who were homozygous for the rs12325817 SNP manifested signs of choline depletion (liver or

muscle dysfunction), relative to 43% of subjects carrying one copy of the variant allele and 13% of subjects without the SNP.

Vitamin C has been shown to induce the release of acetylcholine from synaptic vesicles of neurons and increase acetylcholine levels in the brain.

Possible drugs that can cause memory loss include antidepressants, antihistamines, anti-anxiety medications, anti-seizure drugs, muscle relaxants, tranquilizers, sleeping pills, and pain medications given after surgery. Why? The majority of these are in a class called anticholinergic drugs and block acetylcholine.

A French study looking at 4,128 women and 2,784 men that reported taking anticholinergic drugs showed a greater decline over four years in verbal fluency scores and in global cognitive functioning than women not using anticholinergic drugs. In men, an association was found with a decline in visual memory and to a lesser extent in executive function. Significant interactions were observed in women between anticholinergic use and age, APOE genotype, or hormone replacement therapy. A significantly 1.4–2 fold higher risk of cognitive decline was observed for continuous anticholinergic users.

These drugs could be especially theoretically problematic for those with poor PEMT function, low estrogen (in women) and a family history of dementia and Alzheimer's disease.

Research shows that only 15% of women get enough choline, and one study found that those with lowest choline have the highest anxiety.

Glutamate-BDNF

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
BDNF rs6265	Heterozygous CT

Recap

Improves BDNF Gene Function: Running, DHA, milk thistle, green tea, low glycemic diet, normal levels of glutamate, lithium, acetylcholine, sunlight and heat exposure, bifidobacterium longum, intermittent fasting, turmeric, testosterone and estradiol (women).

Decreases BDNF Gene Function: Chronic stress, high blood sugar and head injuries

GLUTAMATE-BDNF

Research: BDNF (brain-derived neurotrophic factor) is the most important protein abundantly expressed in brain functions related to repair, spatial learning, episodic memory, and adaptability.

BDNF is a synaptic modulator of glutamate while GABA synapses are also regulated by BDNF. Therefore, BDNF regulates glutamate release and regulates glutamate receptor function. Chronic stress leads to decreased BDNF expression in the hippocampus, leading to enhanced anxiety-related behaviors.

There is building evidence that shows impaired glutamate synapses where compromised BDNF function has been observed such as Alzheimer's, Parkinson's, Huntington's disease, depression, bipolar disorder, migraines, and anxiety with the BDNF polymorphism Val66Met. It should be noted that a 2015 meta-analysis did not find a correlation with BDNF val66met and Alzheimer's disease.

BDNF is present in many regions of the central nervous system, including the hippocampus, cerebral cortex, cerebellum, hypothalamus, substantia nigra, amygdala, and spinal cord. The CT and TT genotypes have lower secretion and blood levels of BDNF and research suggests that upregulating BDNF-activated pathways may be therapeutically relevant.

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Evidence suggests that a decrease in hippocampal BDNF may account for the cognitive deficits and the impairment of memory in depression and anxiety disorders. Another study with depressed patients with BDNF polymorphisms found that the individuals with heterozygous or homozygous genotypes were significantly associated with an increased risk of suicidal behavior.

A 2012 and 2017 study found that episodic memory improves as maximal oxygen capacity increases. Aerobic activity induces a structural change in hippocampal volume and vasculature, responsible for episodic memory. It is the hippocampus that displays dramatic volume changes in disease states such as Alzheimer's disease and depression.

Mice studies have shown that the hormone irisin is generated by the muscles during endurance exercise and is responsible for producing BDNF. Further research has found that running produces a higher level of neurogenesis compared to resistance training and high-intensity training.

Exercise has been shown to cause a rise in serum BDNF and is especially enhanced in the heat. Since permeability of the blood-brain barrier increases with exercise in the heat, the hypothesis was raised that this causes a higher cerebral output of BDNF.

Research also found that BDNF levels are equally decreased in bipolar disorder during the occurrence of manic and depressive episodes. An interesting connection here is that when glutamate plummets, depression occurs. When glutamate spikes, mania occurs. If BDNF is suppressed, glutamate modulation is compromised.

An example of low glutamate can be seen in female patients with depression that have been found to have abnormally high expression levels of many genes that regulate the glutamate system. Recent studies found that a low dose of the drug ketamine, which alters glutamate system activity, can rapidly eliminate depression in two-thirds of patients who do not respond to conventional antidepressants. Conventional antidepressants target the monoamine (MAO) systems, which secrete the neurotransmitters dopamine, serotonin or norepinephrine.

In a 2017 study of 458 soldiers, those with the BDNF Met/Met genotype (homozygous TT), 57.9 percent had a history of one or more prior concussions, compared with 35.6 percent of those with other BDNF genotypes. Those with the BDNF Met/Met genotype also reported greater aggression and hostile personality characteristics. When combined in a predictive model, prior military deployments, being male, and having the BDNF Met/Met genotype were independently associated with an increased lifetime history of concussions in active-duty soldiers.

Low levels of BDNF have been shown in research to cause impaired glucose metabolism, highlighting the blood sugar connection of Type 2 diabetes to dementia and depression in epidemiological studies. Other studies have found high BDNF levels in those who already have Type 2 diabetes, with researchers hypothesizing that BDNF tries to overcompensate to reduce insulin and glucose levels, as has been found when BDNF is injected into diabetic rats.

When reviewing your genetic analysis, it is important to also look at the genes 5-HT2A, PEMT, CYP2R1, APOE, GAD1, SLC17A7, TCF7L2, FADS1, FADS2 and TCN2 to see how BDNF is most affected, and where you need to focus most nutritionally.

Glutamate Transport-SLC17A7

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
SLC17A7 rs74174284	Homozygous GG

Recap Improves SLC17A7 Gene Function: Zinc, omega-3 fatty acids (EPA and DHA), Iion's mane mushroom, B6, Iithium, magnesium, B2, folate, B12, vitamin C, melatonin, choline, vitamin D and exercise. Decreases SLC17A7 Gene Function: Head injuries

GLUTAMATE TRANSPORT-SLC17A7

93 **Research**: Polymorphisms in SLC17A7 are associated with delayed recovery time from head injuries. The hypothesis for this is that variants in this gene reduce glutamate transport, which leads to high concentrations of glutamate within the synaptic cleft after trauma.

Glutamate is the primary excitatory neurotransmitter in the brain, while GABA is the principal inhibitory (relaxing)

neurotransmitter. Following a head injury, high glutamate release is responsible for excitotoxicity that leads to neuronal injury, mitochondrial dysfunction and dysfunction of surviving neurons. The loss of GABA producing cells disrupts the balance of excitation and inhibition leading to further cell injury. This glutamate toxicity seen in brain injuries is also the process shared in epilepsy and neurodegenerative disorders. Therefore, a goal should be to restore normal glutamate and GABA function for a head injury recovery protocol.

One study in 2016 took saliva samples from 40 athletes diagnosed with a sport-related concussion by a physician. An association was found between the normal genotype of SLC17A7 and recovery, where those carrying the minor G allele were 6.33-times more likely to experience prolonged recovery rates exceeding 20 days. Those carrying the GG genotype had worse motor speed scores upon initial assessment compared to both heterozygous (CG) and homozygous (CC) genotypes. Based upon these findings, rs74174284 is a potential predictive genetic marker for identifying athletes who are more susceptible for altered recovery times and worse motor speed scores after sport-related concussion.

The majority of traumatic brain injuries (TBI) cases can be attributed to motor vehicle accidents, motorcycle accidents, bicycle accidents, and pedestrian injuries. It is also a major concern in contact sports.

In football and hockey, the number of actual concussions is six or seven times higher than the number diagnosed. Approximately 70 percent of football players and 62 percent of soccer players get at least one concussion per year.

In a study of Norwegian soccer players, 81 percent had an impairment of attention, concentration, memory, and judgment ranging from mild to severe.

A study from the Archives of Pediatrics & Adolescent Medicine found that children who suffer concussions may experience lingering problems with memory and attention, even 12 months after the injury.

Human clinical data suggests that supplemental zinc can be used during recovery to improve cognitive and behavioral deficits associated with brain injury. Additionally, pre-clinical models suggest that zinc may increase resilience to traumatic brain injury, making it potentially useful in populations at risk for injury. It would appear that this is especially true for injuries to the temporal lobe.

A July publication of The Journal of Neurosurgery found that supplementing rats with EPA/DHA fish oil after head injuries reduced the observed issues with a concussion; "Animals receiving the daily fish oil supplement for 30 days post-concussion had a greater than 98 percent reduction in brain damage compared with the animals that did not receive the supplement. It is hypothesized that the omega-3 fatty acids in the fish oil reduced the neural inflammation induced by the concussion injury."

Current studies suggest that oxidative stress lasts at least 24 hours after a traumatic brain injury and that antioxidant reserves like vitamin C are severely compromised. Vitamin C has been shown to prevent excitotoxic damage caused by excessive extracellular glutamate and increase GABA receptor function.

An animal study found that that vitamin D3 may play a role in mechanisms relevant to protective properties against the neurotoxicity of glutamate through upregulation of VDR expression.

Studies show that brain magnesium levels fall 50% for 5 days after injury to the CNS. Studies of both animal and human brain trauma victims suggest higher magnesium levels are associated with better recovery. Post-traumatic administration of magnesium to restore normal magnesium homeostasis reduces neuronal cell death and increases the likelihood of recovery.

Melatonin has been evaluated to be effective in TBI where it improves mood and behavior, decreases brain edema, decreases intracranial pressure and significantly increased superoxide dismutase and glutathione peroxidase (both reduce inflammation).

University of Buffalo researchers published a study in the Clinical Journal of Sports Medicine that individualized exercise programs just below the onset of symptoms is safe and can relieve nearly all post- concussion symptoms. The athletes who exercised returned to normal within 11 to 36 days, while those who did not exercise required 41 to 112 days of intervention.

Please review BDNF and APOE genotypes for a more detailed assessment of head injury recovery.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
MAO-A MAO-A (Monoamine oxidase A) is a critical enzyme involved in breaking down important neurotransmitters such as serotonin, estrogen, norepinephrine, and dopamine.	MAO-A-rs6323	тт		
5-HT2A The 5-HT2A gene encodes for serotonin receptors found in the brain and central nervous system and is concentrated in the brain region essential for learning and cognition. Polymorphisms in rs6314 may result in reduced episodic memory in young and middle- aged individuals.	5-HT2A-rs6314		AG	
5-HT2A The 5-HT2A gene encodes for serotonin receptors found in the central nervous system. Polymorphisms in rs6311 and	5-HT2A-rs6311		СТ	
rs6313 may contribute to a reduced capacity to regulate stress, low vagal tone, anxiety, depression, OCD, and IBS, especially in females.	5-HT2A-rs6313		AG	
COMT V158M COMT is connected to dopamine, adrenaline,	COMT V158M- rs4680		AG	
estrogen and catecholamine metabolism.	COMT-rs4633		СТ	
ANKK1 ANKK1 modulates the density of dopamine receptors in the brain.	ANKK1-rs1800497	GG		
DAO C2029G DAO participates in the degradation of extracellular histamine. This gene is connected to migraines.	DAO C2029G- rs1049793	CC		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
HNMT C314T Histamine N- methyltransferase (HNMT) is a histamine-metabolising enzyme expressed in the brain. This gene is connected to migraines.	HNMT C314T- rs11558538	СС		
HNMT Histamine N- methyltransferase (HNMT) is a histamine-metabolising enzyme expressed in the brain. This gene is connected to hyperactivity and food dyes.	HNMT-rs1050891	AA		
FAAH FAAH (fatty acid amide hydrolase) is a gene that encodes for anandamide breakdown, a neurotransmitter and endogenous cannabinoid.	FAAH-rs324420	СС		
PEMT Choline is required for acetylcholine, a	PEMT-rs7946		СТ	
neurotransmitter of the vagus nerve that enervates numerous organs.	PEMT-rs12325817			GG
GAD1 GAD1 stands for "Glutamate Decarboxylase 1" and is responsible for the conversion of glutamate to GABA.	GAD1-rs3749034	GG		
BDNF BDNF is a synaptic modulator of glutamate while GABA synapses are also regulated by BDNF.	BDNF-rs6265		СТ	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
SLC17A7 SLC17A7 mediates the uptake of glutamate into synaptic vesicles at presynaptic nerve terminals of excitatory neural cells in the brain. Polymorphisms are associated with delayed recovery time from head injuries.	SLC17A7- rs74174284			GG
APOE Apolipoprotein E (APOE) is a lipid binding protein that transports triglycerides and cholesterol in multiple tissues, including the brain.	APOE-rs429358	тт		
	APOE-rs7412			СС
GAD1 GAD1 stands for "Glutamate Decarboxylase 1" and is responsible for the conversion of glutamate to GABA.	GAD1-rs3791851	тт		
	GAD1-rs2241165			ΤΤ
	GAD1-rs3791850			AA
	GAD1-rs769407	GG		

MY CLINICAL RESEARCH SUMMARY: DETOXIFICATION

Liver Enzyme-CYP1A2

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
CYP1A2 C164A rs762551	Homozygous AA

Recap

Improves CYP1A2 Gene Function: Unfiltered fermented drinks (Kombucha, beer, wine), hops, marinades, cruciferous vegetables, blueberries, blackberries, red grapes, kiwi, watermelon, parsley, and spinach.

~	Decreases CYP1A2 Gene Function: Heterocyclic amines, nitrosamines, aflatoxin B1,
\oslash	polycyclic aromatic hydrocarbons, dioxins, and I-naphthoflavone. Omeprazole and
V	primaquine are inducers. Caffeine and Tylenol combined with these compounds can
	make the effect worse.

LIVER ENZYME-CYP1A2

Research: Approximately 200 polymorphisms exist in CYP1A2 gene region, with numerous studies focusing on rs762551. You have the homozygous (AA) rs762551 genotype, which is the rapid metabolizer.

The cytochromes P450 liver enzymes play an important role in the development of various cancers since they are involved in the metabolic transformation of numerous endogenous and exogenous compounds including carcinogens. CYP1A2 is a key factor in the metabolic activity of carcinogenic aromatic and heterocyclic amines, and researchers have found that the inhibition activity of this enzyme may represent a logical strategy for preventing the development of human cancers induced by the aromatic and heterocyclic amines. Further research has shown a cumulative value of phase I (CYP-450 enzymes) and phase II enzymes (GSTM1, GSTP1 and NAT2) in determining individual carcinogenic potential of compounds.

Heterocyclic amines (HCAs) are created by high heat reacting with the proteins. The way to reduce HCAs is to use marinades. Marinades reduce HCAs by up to 90 percent. For further protection, pair with cruciferous vegetables (especially fermented like sauerkraut) and an unfiltered beer or Kombucha due to the protection of the yeast. Red wine, blueberries, blackberries, red grapes, kiwi, watermelon, parsley, and spinach all inhibit the mutagenic activity of certain HCAs in vitro.

High antioxidant fruits, lemon juice, herbs, and spices help keep meat fresh and juicy while protecting against HCAs and reducing AGEs.

Grass-fed meat is higher in vitamin E, and in a study adding concentrations of vitamin E to the surface of ground beef reduced HCA production by 70%. Aim for medium to medium-rare for red meat, flip often and avoid burning. The darker the color the higher the HCA concentrations.

Nitrosamines are used in pesticides, created by frying meat, and from a conversion in the gut by nitrites from cured meats. Vitamin C prevents nitrites from becoming nitrosamines. Limit cured meat consumption using nitrites and take vitamin C when needed.

Aflatoxin B1 is the most common in food and amongst the most potent genotoxic and carcinogenic. It can occur in grain-fed milk, nuts/grains stored in hot conditions or bins, vegetable oils, cocoa or coffee beans stored in warm conditions, and dried fruit. We don't recommend Brazil nuts because they are prone to aflatoxin contamination. Choose nuts and seeds in sealed bags, preferably sprouted. You also want to minimize or avoid oats (unless tested free of ochratoxin). Low protein diets may increase the toxicity of aflatoxin and promote cancerous growth.

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Hops in beer contain a flavonoid called xanthohumol, which strongly inhibits CYP1A2. Xanthohumol has anti-carcinogenic properties and has been found to scavenge reactive oxygen species, including hydroxyl- and peroxyl radicals, and to inhibit superoxide anion radical and harmful nitric oxide production.

Liver Enzyme-CYP1B1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
CYP1B1*6 L432V rs1056836	Wild Type GG

Recap



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Decreases Gene Function: Heterocyclic amines, xenoestrogens, high biotin supplementation, oral contraceptives, hormone replacement therapy, excessive sun exposure, vegetable oils, grains, fried meat, excess of smoked foods, cigarette smoke exposure and exhaust.

LIVER ENZYME-CYP1B1

Research: Due to the carcinogenic activation of polycyclic aromatic hydrocarbons (cigarette smoke, burning coal, vegetable oils, grains) and estrogens to genotoxic catechol estrogens - both which cause DNA mutations - variants in the CYP1B1 gene are important for breast, ovarian, colon, lung and prostate health. This is especially true for those with variants in GSTM1 and GSTP1. CYP1B1 may also be important for skin health, with excessive sun exposure negatively affecting CYP1B1 expression.

CYP1B1 participates in the first step of estrogen metabolism, the conversion of estrogens to 2- or 4-hydroxyestrogens, and specifically catalyzes the 4-hydroxylation of estrogens. 4-hydroxyestradiol is inactivated by COMT.

According to NCBI, C encodes the Leucine and G the Valine. The CYP1B1 L432V rs1056836 GG (valine) is associated with increased CYP1B1 messenger ribonucleic acid (mRNA) expression with a subsequent elevation in 4-hydroxyestradiol formation resulting in increased estrogen-mediated carcinogenicity. However, this has not been proven in human studies.

Minimizing polycyclic aromatic hydrocarbons, xenoestrogens and high estrogen levels in the body are a priority for CYP1B1. Vegetable oils (soy, corn) have been found to be one of the highest sources of polycyclic aromatic hydrocarbons, while also being a high source of omega-6 fatty acids that can disturb the healthy omega-3 and omega-6 ratio needed to prevent skin cancer growth.

A meta-analysis of 12 studies found that coffee consumption decreased the risk of cutaneous melanoma, while another study found that 2 cups of dark roast coffee per day for one month caused a 23% reduction in DNA damage.

Research has shown that optimal levels of iodine can help modulate the estrogen pathway and help prevent cancerous growth by targeting CYP1A1 and CYP1B1. Iodine deficient breast tissue exhibits early markers of breast cancer, and 30% of iodine stores are in the breast tissue.

One study found that high-dose biotin supplementation (often used in isolation for hair growth) increased CYP1B1 expression and was associated with an increase in the occurrence of single-stranded DNA breaks compared with biotin-deficient cells; while inhibitors of CYP1B1 prevented DNA strand breaks.

Inhibition of CYP1B1 activity was observed for the flavonols quercetin, apigenin and myricetin, while resveratrol has shown to convert to piceatannol through CYP1B!, a tyrosine kinase inhibitor and a compound of known anticancer activity. Chrysoeriol, present in rooibos tea and celery, also acts selectively to inhibit CYP1B1 in vitro and may be especially relevant to patients with CYP1B1 overactivity.

One study in 259 post-menopausal women found that for those with certain genotypes in CYP1B1 (rs1056836), KRAS (rs61764370) and MTHFR (rs1801133 and rs1801131), oral contraceptives and hormone replacement therapy was associated with shorter leukocyte telomere length. Shorter leukocyte telomeres are connected to premature aging, and may increase the risk of cancer, cardiovascular disease, obesity, diabetes, chronic pain, and sensitivity to perceived psychological stress.

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In observational studies, higher levels of exercise are related to longer telomere lengths in various populations, and athletes tend to have longer telomere lengths than non-athletes. This relationship is particularly evident in older individuals and physical activity may confer protection against stress-related telomere length shortening.

Higher coffee consumption has been associated with longer telomeres among female nurses. Be aware that there is a compounding effect with caffeine on the slow metabolizer CYP1A2 CC genotype. Research has shown that oral contraceptives significantly prolong the half-life of caffeine from 6.2 hours to 10.7 hours, and therefore could theoretically cause more cardiovascular issues from caffeine for the CYP1A2 CC genotype.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
CYP1A1*2C 4889 CYP1A1 is in the estrogen metabolism pathway along with CYP1B1, CYP1A2, CYP31A, SULT's and COMT. CYP1A1 is involved in the metabolism of benzopyrene.	CYP1A1*2C 4889- rs1048943	Π		
CYP1A2 C164A CYP1A2 metabolizes various environmental procarcinogens, such as heterocyclic amines, nitrosamines, aflatoxin B1 and ochratoxin A.	CYP1A2 C164A- rs762551			AA
CYP1B1*6 L432V The CYP1B1 gene metabolizes pro-carcinogens such as polycyclic aromatic hydrocarbons and 17 beta-estradiol.	CYP1B1*6 L432V- rs1056836	GG		
CYP2C9*3 A1075C Variants in CYP2C9 rs1057910 may alter the metabolism of THC, the psychoactive compound found in cannabis.	CYP2C9*3 A1075C- rs1057910	АА		
CYP2D6 T100C CYP2D6 metabolizes approximately 50% of drugs in clinical use.	CYP2D6 T100C- rs1065852	GG		
CYP2E1 Research has identified CYP2E1 as the primary P450 isozyme responsible for benzene metabolism at low concentrations, acrylamide to glycidamide, alcohol, Tylenol, and nitrosamines.	CYP2E1-rs2031920	СС		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
CYP3A4*1B The CYP3A4 enzyme is involved in the metabolism of approximately 50% of drugs that are used today, cholesterol homeostasis, and the oxidative deactivation of testosterone.	CYP3A4*1B- rs2740574	Π		
CYP2C19*17 Genetic variability impacts expression and activity of CYP2C19 and therefore can influence drug metabolism and catabolism of estrogens.	CYP2C19*17- rs12248560	сс		
NAT2 The NAT2 gene encodes an enzyme that functions to activate and deactivate arylamine, hydrazine drugs, and carcinogens.	NAT2-rs1495741	GG		
VKORC1*2 Variants in VOKRC1*2 may increase the need for vitamin K2 and a sensitivity to dosing of the drug Warfarin.	VKORC1*2- rs9923231	СС		
COQ2 The COQ2 gene encodes an enzyme that functions in the final steps in the biosynthesis of CoQ10 and homozygous variants may increase the risk of statin induced myopathy.	COQ2-rs4693596		СТ	

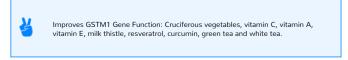
MY CLINICAL RESEARCH SUMMARY: IMMUNE SUPPORT, ANTIOXIDANTS AND INFLAMMATION

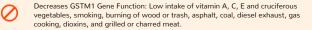
Glutathione-GSTM1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
GSTM1 rs366631	Wild Type AA

Recap





GLUTATHIONE-GSTM1

Research: GSTM1 rs366631 is a pseudo-SNP that can be used as a GSTM1 deletion marker. The deletion is also known as the null genotype and confers the absence of the GSTM1 protein. The frequency of the null genotype varies from 20% to 80%, depending on the ethnic group studied.

For example, the null genotype is less frequent in western and southern African populations, less frequent in South American populations, intermediate in the Japanese, but is higher in Egyptian, European, American, and Asian populations.

High frequencies of the GSTM1 null genotype have been found in patients with lung cancer (East Asians), breast cancer (over 50 age group and in Asians), bladder cancer (with NAT2 slow acetylator), colorectal cancer, skin cancer, gastric cancer (among Asians with H. Pylori), chronic bronchitis, kidney disease progression, acute myeloid leukemia, acute lymphoblastic leukaemia, head and neck cancer (combined with CYP1A1 variant), endometriosis, type 2 diabetes retinopathy, and recurrent pregnancy loss. All have been regarded as environmentally induced and the risk may change with ethnicity.

Of the major glutathione enzymes, GSTM1 appears to be the most effective at neutralizing cytotoxic and genotoxic reactive compounds. However, the research shows that the null genotype of GSTM1 on its own may not be able to determine carcinogen exposure cancer risk. Instead, a combination of genotypes in the other glutathione and antioxidant genes like GSTP1 and NFE2L2, detoxification genes like CYP1A1 and NAT2, and/or compounding epigenetic habits that appear to modify the effect.

GSTM1 catalyzes the detoxification of alkyl and polycyclic aromatic hydrocarbons, intermediate forms of many carcinogens, specifically metabolically generated epoxide intermediates of benzo(a)pyrene. Benzo(a)pyrene is part of a class of chemicals called polycyclic aromatic hydrocarbons. Sources of benzo(a)pyrene include the burning of wood or trash, tobacco smoke, asphalt, coal, diesel exhaust, and grilled or charred meat. There is evidence that it causes skin, lung, and bladder cancer in humans and in animals. Research has also shown that early markers of cardiovascular disease are associated with occupational exposure to polycyclic aromatic hydrocarbons.

A study also found sensitivity to gas cooking and the GSTM1 null genotype, increasing the sensitivity of the lungs to nitrogen dioxide. Nitrogen dioxide is also found in diesel exhaust. Exposure of human blood plasma to nitrogen dioxide caused rapid losses of ascorbic acid, uric acid, protein thiol groups, lipid peroxidation, and depletions of alpha-tocopherol, bilirubin, and ubiquinol leading to high levels of oxidative stress.

Animal studies and in vitro studies have shown that vitamin C, vitamin E, vitamin A, resveratrol, curcumin, green tea, and white tea can inhibit the carcinogenic effect of benzo(a)pyrene and nitrogen dioxide. In the Norwegian Mother and Child Cohort Study 50,651 women, a higher prenatal exposure to dietary benzo(a)pyrene was found to reduce birth weight. However, increasing dietary vitamin C intake during pregnancy helped reduce any adverse effects of benzo(a)pyrene on birth weight.

Isothiocyanates from cruciferous vegetables are known for their anti-cancer activity. They are stored as glucosinolates in

cruciferous vegetables and are hydrolyzed by myrosinase (an enzyme found in plants and intestinal microflora) to form isothiocyanates. Isothiocyanates from cruciferous vegetables are substrates and inducers of GSTM1.

GSTM1 variants may alter isothiocyanates clearance, with the null genotype retaining higher levels of isothiocyanates and therefore the benefits. In numerous studies, the GSTM1 null genotype was the most responsive to cruciferous vegetables for anti-cancer effects against lung cancer, colon cancer, breast cancer, and kidney disease.

The isothiocyanate levels in cruciferous vegetables will range based on growing conditions including sulfur and nitrogen levels, time after harvest and storage (cold transportation and storage of broccoli also cause a loss of glucosinolates up to 70-80%), plant genetics, and cooking preparation. Broccoli sprouts will yield the highest isothiocyanate levels.

Glutathione-GSTP1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
GSTP1 I105V rs1695	Homozygous GG

Recap



Improves GSTP1 Gene Function: Glycine, cysteine, selenium, vitamin C, B1, B6, zinc, magnesium, optimal iron levels, magnesium, alpha lipoic acid, milk thistle, holy basil and vitamin E supplementation for the homozygous GG genotype only.

0	Dec
V	pollu

eases GSTP1 Gene Function: Mercury, arsenic, cadmium, pesticides, and air

GLUTATHIONE-GSTP1

Research: Glutathione is the master antioxidant system involved in oxidative stress, detoxification and immunity. It requires the amino acids glycine, cysteine and glutamate. Selenium activates the glutathione system and works in concert with vitamin E as a potent antioxidant against plasma and LDL lipid peroxidation.

The functional polymorphism of the GSTP1 IIe105Val gene, which reduces enzymatic activity, involves an A-G substitution. Carriers of these mutations are less able to detoxify carcinogens, and epidemiological studies have suggested that individuals differing in the expression of allelic variants of GSTP1 gene differ in susceptibility to various chemical carcinogens.

A meta-analysis of 10,067 cancer cases and 12,276 controls in 41 independent case–control studies from 19 articles found a significant increase in risk in breast cancer in Caucasions with variants in GSTP1 rs1695. A second meta-analysis found the same results with Asians that had the GG genotype. A 2020 study found that the rs1695 homozygous GG genotype was associated with an increased risk of breast cancer, but not the AG genotype. Other research has shown the risk to be higher in premenopausal women vs. post-menopausal women.

An analysis of that included 3,035 breast cancer cases and 3,037 population controls in a Chinese population found that cruciferous vegetable intake helped offset the risk of the GG genotype, with a lower risk associated with a higher cruciferous vegetable intake.

A meta-analysis of 11,762 cases and 15,150 controls from 51 studies showed a statistically significant association between GSTP1 rs1695 polymorphism with prostate cancer risk and urinary system cancer among Asians.

GSTP1 rs1695 variants were reported to be associated with the risk of esophageal cancer and malignant melanoma in the Caucasian population, but not childhood acute lymphoblastic leukemia or bladder cancer.

Glutathione-related polymorphisms, such as GSTM1 and GSTP1 have also been found to increase the elevation and toxicity of mercury. Selenium blocks mercury uptake, folate decreases mercury levels and magnesium and holy basil protect against mercury toxicity.

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One benefit of the GSTP1 AG and GG genotype appears to be in athletic training. GSTP1 rs1695 AG and GG may be high responders to endurance training due to an impaired ability to remove excess reactive oxygen species. The hypothesis is that better activation of cell signaling pathways results in positive muscle adaptations. Women with at least one copy of the G allele showed a significantly greater increase in V?O2max in response to applied training.

In healthy control subjects, the effect of a-tocopherol supplementation on the production of inflammatory cytokines appears to be dependent on an individual's GSTP1 rs1695 genotype. These genotype-specific differences may help explain some of the discordant results in studies that used vitamin E. Persons having the alleles AA or AG in GSTP1 rs1695 had an increase in inflammatory interleukin-6 (IL-6) upon supplementing alpha-tocopherol (the most common form of Vitamin E in a North American diet) while those with GG saw a decrease.

Heavy Metals-GSTP1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
GSTP1 C341T rs1138272	Heterozygous CT

Recap





HEAVY METALS-GSTP1

Research: Glutathione is the master antioxidant system involved in oxidative stress, detoxification, and immunity. It requires the amino acids glycine, cysteine, and glutamate. Selenium activates the glutathione system and works in concert with vitamin E as a potent antioxidant against plasma and LDL lipid peroxidation.

Pesticides, cadmium, mercury, and arsenic have all been shown to lower GSTP1 expression, increasing the elevation and toxicity of these chemicals and heavy metals. The exposure and sensitivity to these chemicals and heavy metals are suspected reasons for the increased risk of certain cancers and male infertility related to GSTP1 variants.

A meta-analysis of 43 eligible case-control studies in 2019 found that the TT genotype of the GSTP1 rs1138272 polymorphism is likely related to the susceptibility to overall cancer in the Asian and African populations. Colorectal and head and neck cancers were increased in the Caucasian population, along with lung cancer in the CT genotype. Researchers stated that additional evidence is required to confirm these conclusions.

A 2020 study found that prostate cancer risk was increased 3.65-fold in the homozygous carriers of GSTP1 rs1138272, and increased incrementally depending on variants in GSTP1 rs1695 and GSTM1.

Children carrying GSTP1 rs1138272 or rs1695 minor alleles may constitute a susceptible population at increased risk of asthma associated with air pollution.

Men with the homozygous or heterozygous genotypes of GSTP1 rs1138272 were 3 times higher to experience a disturbance in sperm motility and 2.5 times higher for abnormalities in morphology of spermatozoa in a Russian population. Studies have also shown that men exposed to higher levels of air pollution were more likely to experience abnormal sperm morphology, decreased motility, and an increased chance of DNA fragmentation. The GSTP1 variants would increase this toxicity and require more antioxidant support.

Docetaxel is a chemotherapy drug used to treat breast, lung, prostate, stomach, and head and neck cancer. A 2015 case-control study of 150 women with early-stage breast cancer found that the GSTP1 rs1138272 polymorphism was associated with docetaxel-induced peripheral neuropathy, and the risk was five times higher for patients with a BMI \geq 30.

104 Choosing organic produce is one of the best ways to avoid excess cadmium. Chemical agriculture uses high amounts of synthetic organophosphates, creating a very high phosphorus content. Synthetic phosphorus concentrates the amounts of heavy metals, like cadmium and uranium in non-organic soils and food. Vitamin C, B1, B6, iron and zinc deficiencies have also been reported to enhance sensitivity towards cadmium.

Fish high in mercury including swordfish, ahi tuna, and halibut should be minimized or avoided. Selenium blocks mercury uptake, folate decreases mercury levels, and magnesium and holy basil protect against mercury toxicity.

Arsenic levels range throughout the world based on groundwater levels. Arsenic is also high in rice. Optimal selenium and folate intake have been found to improve arsenic detoxification and mitigate toxicity.

Eye Health-CFH

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
CFH rs1061170	Heterozygous CT

Recap



Decreases CFH Gene Function: Smoking, pesticides, benzene (found in certain laundry detergents, gasoline and paint), aspartame, oxidative stress, elevated TNFalpha, elevated IL-6, obesity, smoking, diabetes, hypertension, atherosclerosis and low intake of lutein and zeaxanthin.

EYE HEALTH-CFH

Research: Age related macular degeneration (AMD) is the leading cause of blindness in Western societies, but its etiology remains largely unknown.

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Variants in CFH confers a 2-fold higher risk of late AMD per copy in individuals of European descent. Research indicates that CFH (rs1061170) polymorphism impacts significantly on retinal function in early AMD patients, and supports the hypothesis that a dysfunctional CFH might result in early retinal function loss due to a reduction in the immune antioxidant defense mechanism. A study from 2005 found that variants in CFH likely explains approximately 43% of AMD in older adults.

Malondialdehyde (MDA) is a common lipid peroxidation product that accumulates in many pathophysiological processes, including AMD. In vivo studies in mice found CFH as a major MDA-binding protein that blocks MDA-modified proteins by macrophages and MDA-induced pro-inflammatory effects. The CFH polymorphism markedly reduces the ability of CFH to bind MDA, indicating a causal link to a cause of age related macular degeneration.

A recent meta-analysis found that the rates of myopia (nearsightedness) will increase 140% by 2050 due to our increased time in front of a screen. Myopia can increase the risk of numerous eye disorders. Blue light is a high-energy or short-wavelength visible light from your phone and computer that induces inflammation and retinal diseases such as age-related macular degeneration and retinitis pigmentosa. Research has found that bilberry and lingonberry exert protective effects against blue LED light-induced retinal photoreceptor cell damage due to their polyphenol content.

Lutein and zeaxanthin can inhibit oxidation of cell membranes and may be protective against UV-induced eye damage. Studies have demonstrated that people in the highest quintile of intake of dietary carotenoids, especially lutein and zeaxanthin concentrations have significantly lower risk of macular degeneration. Blue-eyed adults have far less lutein and zeaxanthin in their retinas.

One study compared diets of 356 patients with macular degeneration with 520 patients with other eye diseases. The data revealed that beta carotene was not especially effective, but that lutein and zeaxanthin were. Another study found that the risk of macular degeneration was reduced 65 percent with high amounts of lutein and zeaxanthin.

Research has found that MDA levels are significantly increased in groups of subjects with deficient levels of vitamin C and vitamin E. Deficiency in these two antioxidants leads to insufficient defense against free radicals and increased MDA levels. Those with polymorphisms in CTH should increase vitamin C and vitamin E intake. In another study, the risk for macular degeneration was found to be 77% lower when vitamin C supplements and a low-glycemic diet was used.

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One study followed 3,600 people ages 55-80 years old for six years and found that those that took antioxidants plus zinc were less likely than those who took only antioxidants or only zinc to lose their vision.

Studies show that people who consume more fish, which is rich in DHA-fish fat, are less likely to develop macular degeneration. Eating fish one to three times a week has been associated with a 40 to 75 percent reduction in macular degeneration.

Eye Health-ARMS2

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
ARMS2 rs10490924	Heterozygous GT

Recap



Decreases ARMS2 Gene Function: Smoking, pesticides, benzene (found in certain laundry detergents, gasoline and paint), aspartame, oxidative stress, elevated TNFalpha, elevated IL-6, obesity, smoking, diabetes, hypertension, atherosclerosis and low intake of lutein and zeaxanthin.

EYE HEALTH-ARMS2

Research: ARMS2 is considered a second major AMD susceptibility gene next to CFH. CFH and ARMS2 share a common pathway in the pathogenesis of AMD, with ARMS2 polymorphisms disrupting mitochondrial function in the retina.

Research estimates that the risk of AMD can be attributed to 20% from smoking, 36% for variants in ARMS2, and 43% for variants in the CFH gene.

The overall effect of ARMS2 polymorphisms is driven primarily by a strong association in smokers. This gene-environment interaction is supported by statistically independent family-based and case-control analysis methods. Studies have shown that a genetic susceptibility coupled with a modifiable lifestyle factor such as cigarette smoking confers a significantly higher risk of AMD than either factor alone.

Polymorphisms in ARMS2 is also associated with polypoidal choroidal vasculopathy (PCV) and clinical severity in the subgroups of PCV in the Japanese population.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
SOD2 Superoxide dismutase (SOD2) is manganese dependent and protects against superoxide for the mitochondria of the cell. Variants here increase the need for intracellular antioxidant protection.	SOD2-rs4880	AA		

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
SOD3 Superoxide dismutase (SOD3) is zinc/copper dependent and protects against superoxide for the cell membrane. Variants here increase the need for intracellular and extracellular antioxidant protection.	SOD3-rs1799895	СС		
CAT C-262T CAT makes an enzyme called catalase, which helps reduce oxidative stress.	CAT C-262T- rs1001179	сс		
GSTM1 GSTM1 catalyzes the detoxification of alkyl and polycyclic aromatic hydrocarbons (PAHs), intermediate forms of many carcinogens, specifically metabolically generated epoxide intermediates of benzo(a)pyrene.	GSTM1-rs366631	AA		
GSTP1 I105V Glutathione S-Transferase (GSTP1) is linked to the metabolism of mutagens, carcinogens, and other poisonous chemicals. It plays a crucial role in the detoxification process, thereby protecting cells from these compounds. GSTP1 rs1695 is connected to breast, prostate, urinary, esophagus, and skin health.	GSTP1 I105V- rs1695			GG
GSTP1 C341T Glutathione S-Transferase (GSTP1) is linked to the metabolism of mutagens, carcinogens, and other poisonous chemicals. It plays a crucial role in the detoxification process, thereby protecting cells from these compounds. GSTP1 rs1138272 is connected to the colon, prostate, lung, throat, and fertility.	GSTP1 C341T- rs1138272		СТ	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
GPX1 The GPX1 (Glutathione peroxidase 1) gene encodes a protein responsible for the modulation and detoxification of hydroperoxides and hydrogen peroxide to protect the mitochondria and cytoplasm of cells against oxidative damage.	GPX1-rs1050450	GG		
CTH The CTH (Cystathionine Gamma-Lyase) gene encodes an enzyme in the trans- sulfuration pathway that converts cystathionine derived from methionine into cysteine. Glutathione synthesis in the liver is dependent upon the availability of cysteine.	CTH-rs1021737	GG		
NOS1 NOS1 (nNOS) codes for brain neural transmission, memory, learning, psychological stress, the peripheral nervous system and potentially the lymph nodes.	NOS1-rs3782218	СС		
NOS2 NOS2 (iNOS) encodes for wound, tissue damage, infection and hypoxia (low oxygen).	NOS2-rs2248814	GG		
CFH CFH (complement factor H) polymorphism is associated with increased risk of age related macular degeneration.	CFH-rs1061170		СТ	
ARMS2 ARMS2 polymorphism is associated with increased risk of age related macular degeneration (AMD).	ARMS2- rs10490924		GT	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
LZTFL1 The LZTFL1 gene influences the transition of specialized lung cells to less specialized lung cells during infection and inflammation.	LZTFL1- rs17713054	GG		

MY CLINICAL RESEARCH SUMMARY: DNA PROTECTION, DAMAGE & REPAIR

Processed Meat and Colon Cancer-GATA3

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE		
GATA3 rs4143094	Heterozygous GT		
Recap			
Improves GATA3 Gene Function: Avoidir	ng processed meats.	\oslash	Decreases GATA3 Gene Function: Processed meats like hot dogs, cu pepperoni and sausages.

PROCESSED MEAT AND COLON CANCER-GATA3

Research: A large-scale genome-wide analysis of over 18,000 people from the U.S., Canada, Australia and Europe found that variants in GATA3 (rs4143094) was associated with an increased risk of colon cancer for those eating processed meat compared to those with the normal genotype.

A meta-analysis revealed that by raising the serum level of vitamin D to 34 ng/ml, the incidence rates of colorectal cancer could be reduced by half. Researchers projected a two-thirds reduction in incidence with serum levels of 46 ng/ml, which corresponds to a daily intake of 2,000 IU of vitamin D3.

All of the foods below have been found in research studies to reduce the risk of colon cancer: black raspberries, blackberries, raspberries, cranberries, blueberries, apples, oranges, avocado, tomatoes, garlic, onions, shallots, leeks, cabbage, sauerkraut, broccoli, Brussels sprouts, sweet potatoes, beets, spinach, kale, asparagus, cauliflower, turmeric, rosemary, oregano, basil, thyme and parsley. Preventing constipation should be a priority.

Longevity-SIRT1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
SIRT1 rs7895833	Wild Type AA

Recap

Improves SIRT1 Gene Function: Exercise, fasting, 7-8 hours of sleep per night, sauna, polyphenols, vitamin D, omega-3 fatty acids, resveratrol, magnesium, and melatonin.

Decreases SIRT1 Gene Function: The APOE-e4 genotype, high blood sugar, and insulin resistance.

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LONGEVITY-SIRT1

Research: SIRT1 regulates numerous genes that accelerate the aging process, modulate DNA repair mechanisms and transcription factors like p53 (tumor suppressor gene), FOXOs (key regulators of lipid metabolism, stress resistance, and

apoptosis) and inhibits NF-kb, a pathway connected to viral inflammation.

SIRT1 activity goes down as we age, and DNA damage accumulates, and its activity is especially harmed by a sedentary lifestyle, poor diet, and obesity. Activation of sirtuins induces the growth of blood vessels, insulin sensitivity and better glucose control, and other health benefits in a wide range of age-related cardiovascular and metabolic disease models. Experimental models have shown that increasing the activity of the sirtuins is associated with the delay of age-related diseases and potentially increasing longevity.

Researchers have observed a significant increase in SIRT1 levels in longevity populations and found a significant positive correlation between SIRT1 levels and age in a Turkish population. The oldest people carrying AG genotypes for rs7895833 had the highest SIRT1 level compared to the AA genotype, suggesting an association between rs7895833 SNP and lifespan longevity.

The average age of older people carrying AG genotype (76.0 \pm 1.5 years) was significantly higher than the average age of older people carrying AA genotype (71.3 \pm 1.4 years).

Your APOE genotype may also affect SIRT1 activity for longevity. Research from the Buck Institute group found that APOE-e4 reduced expression of SIRT1. The reduced expression of SIRT1 was thought to impair beta-amyloid clearance observed in Alzheimer's. If you have the APOE-e4 allele, the AA SIRT1 genotype may require more SIRT1 activation.

Polyphenols are activators of SIRT1 and contain anti-inflammatory and apoptosis properties. These include piceatannol (a metabolite of resveratrol), olive oil, fisetin (strawberries, apples, grapes), quercetin (wine, peppers, berries, apples) and resveratrol (wine, blackberries, blueberries, pistachios and dark chocolate).

Other activators of SIRT1 that also benefit the APOE-e4 carriers include magnesium, melatonin, vitamin D, and omega-3 fatty acids. One study found that centenarians (those living over 100) have higher total body magnesium and lower calcium levels than most elderly people.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
XRCC3 XRCC3 participates in DNA double-strand break/recombination repair.	XRCC3-rs861539			AA
ATM D1853N ATM coordinates DNA repair by activating enzymes that fix double stranded DNA breaks.	ATM D1853N- rs1801516	GG		
ESR2 ESR2 acts as a tumor suppressor gene that codes for estrogen receptor beta (ER-beta), one of two main types of estrogen receptor activated by estrogen. ESR2 is strongly expressed in the breast.	ESR2-rs2987983	AA		
TP53 TP53 is a tumor suppressor gene responsible for DNA repair.	TP53-rs1042522			СС

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Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
MDM2 Variants in the MDM2 gene encode a protein that reduces cellular levels of the p53 tumor suppressor protein.	MDM2-rs2279744	тт		
MLH1 MLH1 codes for a DNA repair enzyme linked to colon health.	MLH1-rs1800734	GG		
GATA3 GATA3 factors are involved in cellular maturation with proliferation arrest and cell survival.	GATA3-rs4143094		GT	
SIRT1 SIRT1 senses changes in intracellular NAD+ levels and plays a role in DNA damage and repair.	SIRT1-rs7895833	AA		

MY CLINICAL RESEARCH SUMMARY: CARDIOVASCULAR HEALTH AND ATHLETIC PERFORMANCE

Power and Recovery-ACTN3

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
ACTN3 rs1815739	Heterozygous CT
Recap	
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POWER AND RECOVERY-ACTN3

The T (R) allele is associated with enhanced strength and training adaptation, improved protection from eccentric traininginduced muscle damage, lower risk of sports injury, and reduced frailty in the elderly. Testosterone levels were also higher in male and female athletes with at least one R allele compared to the XX genotypes. When stratified by race and gender for power athletes in a 2019 meta-analysis, Asian and male athletes benefited the most from the RX (TC) genotype.

The ACTN3 RR and ACTN3 RX groups have not been significantly different, indicating that the presence of one or two R alleles does not have a dose-dependent effect on 200-meter sprint speed in elite athletes. However, there was some evidence for a dose-effect of the ACTN3 R allele and 200-meter sprint speed in elite male African athletes. The ACTN3 RR individuals had (on average) a faster best personal sprint time than ACTN3 RX individuals.

Having the RX genotype may represent the best of both worlds for cold adaptation, longevity, strength training, and exercise recovery.

VO2 Max-PPARGC1A

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
PPARGC1A rs8192678	Heterozygous CT

Recap



Decreases PPARGC1A Gene Function: Sedentary lifestyle

113 VO2 MAX-PPARGC1A

Research: Peroxisome proliferator-activated receptor gamma coactivator 1-alpha (PPARGC1A) is a master regulator of mitochondrial biogenesis, mitochondrial respiration, skeletal muscle fiber transformation (from fast to slow twitch), glucose and fatty acid metabolism, and the anti-oxidation machinery. PPARGC1A is expressed in cell types with high oxidative function

(heart, skeletal muscle slow twitch fibers, liver, and pancreas) and in brown adipose tissue.

Several studies have shown that SNPs in PPARGC1A are associated with a significant lower level in aerobic power (i.e., VO2 max) in insulin resistant and untrained individuals as well as in athletes. Healthy untrained adults display a large individual variation in VO2 max that ranges from -20% to more than 50%.

Research indicates that the exercise-induced variation in VO2 max is 47% explained by genetics. If you have heterozygous or homozygous variants in PPARGC1A, you may have a naturally lower VO2 max for aerobic exercise and increased CRP (C-reactive protein) levels.

To increase VO2 max, consider cold exposure. Since mitochondria are what give us the ability to use oxygen in order to produce cellular energy, the more we have the more the aerobic potential.

Cold exposure activates the PPARGC1A gene and PGC1^{II} protein, which makes more mitochondria in the muscle. One study found that 15 minute exposure to cold water (50°F or 10°C) following high intensity running, increases PGC1^{II} in muscle tissue. Another study found that men that were immersed in cold water at 50°F (10°C) for 15 minutes, 3 times a week for four weeks after running were able to increase mitochondrial biogenesis occurring in their muscle tissue.

Adaptogens are another way to increase your VO2 max. One study found that ashwagandha increased velocity, power, VO2 max, lower limb muscular strength and neuromuscular coordination. A second study used elite Indian cyclists for 8 weeks. One group received 500mg of the root extract 2x a day, while the other group received a placebo. There was significant improvement in the experimental group in all parameters, namely, VO2 max and time for exhaustion on treadmill.

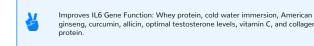
A study using eleuthero root found that using 800mg for 8 weeks increased VO2 max of by 12%, endurance time improved 23%, the highest heart rate increased 4%, and metabolism was altered which spared glycogen storage. The study concluded that "this was the first well-conducted study that shows that 8-week ES supplementation enhances endurance capacity, elevates cardiovascular functions and alters the metabolism for sparing glycogen in recreationally trained males."

Muscle Recovery-IL6

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
IL6 rs1800795	Heterozygous CG

Recap



Decreases Gene Function: Low testosterone (men), depression, obesity, bacterial overgrowth and workout routines without enough recovery days.

MUSCLE RECOVERY-IL6

Research: Exercise increases IL6 cytokines even when muscle damage hasn't occurred. It is produced in large amounts during heavy weight lifting and endurance races. The CG genotype is more common in sprint and power athletes compared to endurance and non-athletes.

C-allele carriers of the IL6 SNP have been found to have higher creatine kinase values (a marker of muscle damage) following exercise compared with GG homozygotes.

The highest post-exercise creatine kinase levels are found after prolonged exercise such as ultra distance marathon running, weight lifting and downhill running.

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To accelerate recovery, whey protein, cold water immersion, American ginseng, curcumin, optimal testosterone levels, vitamin C and collagen protein have all been found to attenuate creatine kinase levels.

Research has also found that purple sweet potatoes, cranberries, blueberries and beet root juice have verified health,

performance-enhancing, and exercise recovery benefits.

Perhaps the most promising results have come from two separate studies showing decreased muscle soreness and increased recovery from cherry juice and dehydrated cherry supplements. One of these studies had subjects perform ten sets of ten repetitions at 70% of a 1-RM back squat. The researchers found that Montmorency powdered tart cherry supplementation used daily and 48 hours post-workout significantly lowered muscle soreness strength decrement during recovery, and markers of muscle catabolism throughout the 48 hour post-lifting recovery period compared to placebo.

Muscle Injury-COL1A1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
COL1A1 rs1800012	Wild Type CC

Recap



Improves COL1A1 Gene Function: Vitamin C, zinc, copper, glycine, proline, lysine and B6 (all precursors to collagen production) and cryotherapy. Decreases COL1A1 Gene Function: Deficiencies in vitamin C, zinc, copper, glycine, proline, lysine, B6 and excessive NSAID use.

MUSCLE INJURY-COL1A1

Research: According to one study, the gene encoding for the alpha1 chain of type I collagen (COL1A1) has been shown to be associated with cruciate ligament ruptures and shoulder dislocations.

You have the CC genotype for COL1A1, which lowers the production of Type 1 collagen. Approximately 90% of collagen in the body is Type I. Type I collagen is found in the skin, tendons, corneas, lungs and in 95% of bone.

ACL ruptures are considered the most severe injury sustained in sports. The A variant produces more COL1A1. Two AA's reduced risk of ACL rupture by ten times, while only 5% of the population have two AA's.

Cryotherapy has been shown to inhibit harmful collagenase (activity on collagen enzyme that breaks down collagen) and also decreased the production of inflammatory E2 series prostaglandins. For athletes, cryotherapy post-training could be a useful tool to help prevent injuries.

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Pesticides, HDL and LDL-PON1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
PON1 rs662	Heterozygous CT

Recap

Improves PON1 Gene Function: Organic foods, calcium, magnesium, boron, lycopene, alpha-lipoic acid, gamma-linolenic acid (black cumin seed oil), broccoli sprouts, vitamins E, B1, B2, B5, B6, selenium, omega 3 fatty acids, high quality olive oil, polyphenols, naringenin, quercetin, pomegranates and alcohol in moderate amounts (1 drink for women and 1-2 drinks for men).

PESTICIDES, HDL AND LDL-PON1

Research: Paraoxonases (PON1) are a family of enzymes involved in breaking down chemicals including several types of pesticides and pharmaceutical drugs. They are involved in protecting both high and low-density lipoproteins from oxidation, an important mechanism in atherosclerosis and heart disease. The rs662 SNP is the most clinically relevant for PON1. The C allele is also known as the "R" allele in research studies and is connected to atherosclerosis and heart disease.

A 2018 meta-analysis found that carriers of the variant R allele had higher levels of oxidized LDL, triglycerides, total cholesterol, and low-density lipoprotein cholesterol than the non-carriers. This was most pronounced in Asians and coronary heart disease patients. The hypothesis is that decreased levels of PON1 activity may lead to increased circulating levels of oxidized LDL and reduce the capacity of PON1-mediated inhibition of LDL-C oxidation.

Mercury appears to decrease PON1 function and liver expression of the PON1 gene is down-regulated in mice with high homocysteine. The proatherogenic effects of homocysteine may involve decreased serum PON1 activity, leading to impaired antioxidant function and decreased capacity to degrade homocysteine thiolactone.

The availability and catalytic activity of PON1 are impaired in many children with Autism Spectrum Disorders, making them more susceptible to the toxic effects of pesticide residues which are most frequently found on grain.

The rs662 SNP is the most clinically relevant for PON1. You need to make sure you are focusing on foods and drinks that improve gene function.

All of the vitamins, minerals, and compounds in the "Improves PON1 Gene Function" section have been verified in research to improve PON1 function. One way that pomegranates protect cardiovascular health is by augmenting nitric oxide. In one study, pomegranates protected against atherosclerosis by reducing LDL's basal oxidative status by 90%.

Moderate drinkers can also rejoice. Research has found that alcohol in small amounts (1 drink for women, 1-2 for men based on weight), improved PON1 activity by 395%. However, too much alcohol decreased PON1 by 45%.

A recent study found that red wine induced significant increases in plasma total antioxidant status and significant decreases in plasma MDA (inflammation biomarker). The results show that the consumption of 400 mL/day (14 ounces) of red wine for two weeks, significantly increases antioxidant status and decreases oxidative stress in the circulation.

Non-organic wine in particular may have concentrated amounts of additives, pesticides, insecticides and fungicides, while beer that uses GMO crops may be high in glyphosate (RoundUp). Residual concentrations of many different pesticides that have been detected in bottled wine were similar to initial concentrations on the grapes. The US and France are heavier users of pesticides. Italy and Argentina have been found to have wine most likely free from pesticides and heavy metals.

Caffeine-CYP1A2

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
CYP1A2 C164A rs762551	Homozygous AA

Recap

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Increases CYP1A2 Gene Function: A higher cruciferous vegetable intake may help increase caffeine metabolism for those with the CC slow metabolizer genotype, along with exercise.

Decreases CYP1A2 Gene Function: Oral contraceptives

CAFFEINE-CYP1A2

You have the homozygous AA genotype and are a "rapid metabolizer" of caffeine. This means that caffeine will quickly be metabolized from your body and the effects lasting a shorter period of time. It is important to review your COMT gene function to better understand a sensitivity to coffee intake.

For the AA genotype, caffeine decreased 40-km time in cyclists by an average of 3.8 minutes in the AA homozygotes as compared to 1.3 minutes in the C allele carriers.

Triglycerides-FADS1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
FADS1 rs174546	Homozygous TT

Recap

Improves FADS1 Gene Function: Higher intake of the omega-3 fatty acids EPA and DHA.

Decreases FADS1 Gene Function: Low intake of EPA and DHA.

TRIGLYCERIDES-FADS1

Research: Variants in the FADS1 SNP (rs174546) are associated with elevated triglyceride levels, which appears to be due to a higher need for EPA and DHA from animal foods. Studies have found that plasma triglyceride levels were lower in wild-type CC genotype when compared to carriers of the minor T allele.

Population average triglyceride levels have increased since 1976 in parallel with the constant growing epidemic of obesity, insulin resistance and Type-2 diabetes. A meta-analysis of 17 population-based prospective trials including 46,413 men and 10,864 women identified plasma triglycerides levels as an independent risk factor for cardiovascular disease.

Triglycerides are essentially fat in the blood that are driven by excess sugar and carbohydrate consumption. They are the driving force behind lipoprotein particles that are potent causes of heart disease, such as small LDL and very low-density lipoprotein (VLDL).

Numerous studies have found that omega-3 fatty acids administered as fish oil supplements lowers plasma triglyceride levels by 25% to 34%. While fish oil is known to lower triglycerides, there doesn't appear to be a difference in the FADS1 genotype response to supplementation.

A meta-analysis of 13 randomized controlled trials found that 500mg of vitamin C resulted in a significant decrease in serum LDL cholesterol and triglyceride concentrations.

Stress-ADRB2

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE		
ADRB2 rs1042713	Heterozygous AG		
lecap			
Improves ADRB2 Gene Function: Deep breathing techniques, magnesium and vitamin C.		Decreases ADRB2 Gene Function: Chr	onic stress and shallow br

STRESS-ADRB2

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Research: ADRB2 activation regulates various biological functions, including heart rate, blood pressure or respiration, and it

may modulate the vasodilatation in normal coronary arteries.

In a meta-analysis of seven case-control studies with a total of 6,843 subjects, a higher frequency of polymorphisms in rs1042713 was found with heart attacks or coronary artery disease compared to healthy controls. A similar result was also obtained with polymorphisms in rs1042714. Ethnicity-stratified subgroup analysis suggested that the rs1042714 variants correlated with an increased risk of the two diseases in both Asians and Caucasians, while rs1042713 only contributed to the risk of two diseases in Asians.

If cardiovascular disease runs in your family, you may be more prone to anxiety, high blood pressure and arterial damage from stress. You may also be more sensitive to beta blockers, leading to high triglycerides.

Since ADRB2 is connected to the catecholamine epinephrine, it is also important to look at your COMT gene function. Magnesium, vitamin C and adaptogens should be considered while making lifestyle changes and strategies if chronic stress is present.

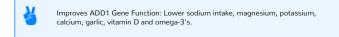
Relaxation techniques that involve slow, deep breathing have been found to be an effective therapeutic intervention that counteracts the adverse clinical effects of stress in disorders including hypertension, anxiety, insomnia and aging.

Potassium and Magnesium-ADD1

Below is a summary of your most significant variant genotypes:

GENE	GENOTYPE
ADD1 rs4961	Heterozygous GT

Recap



Decreases Gene Function: High sodium intake, excess weight, high sugar intake, sedentary lifestyle, smoking and stress.

POTASSIUM AND MAGNESIUM-ADD1

Research: A meta-analysis of 33 studies with 40,432 participants found that variants in rs4961 was significantly associated with hypertension in Asians. Other research found that carriers of the risk (T) allele responded better to diuretics and sodium-restricted diets, in that they tended to lower their blood pressure by ~10 mmHg points compared to rs4961(GG) homozygotes similarly treated.

Excess weight, high sugar intake, sedentary lifestyle, smoking, stress and high sodium intake all raise blood pressure. People living at higher latitudes throughout the world are at higher risk of hypertension, and patients with cardiovascular disease are often found to be deficient in vitamin D. Magnesium, potassium, calcium, vitamin D, garlic and omega-3's all lower blood pressure.

One study found that increasing potassium-rich foods to 4.7 grams was equivalent to cutting out 4 grams of sodium in terms of reducing blood pressure.

In another study, aged garlic extract given at a dose of 600-1500mg was just as effective as the drug atenolol in reducing blood pressure over a 24-week period.

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
ACTN3 ACTN3 encodes for the alpha- actin-3 protein found exclusively within type-II fast- twitch muscle fibers.	ACTN3-rs1815739		СТ	
PPARGC1A It has been demonstrated that variants in the PPARGC1A gene affect the exercise- induced change in maximal oxygen uptake (VO2).	PPARGC1A- rs8192678		СТ	
TNFA Tumor necrosis factor (TNF-a) is a pro-inflammatory cytokine. Variants may increase the risk of asthma in Asian populations.	TNFA-rs1800629	GG		
IL6 IL6 is an interleukin that acts as both a pro-inflammatory cytokine and an anti- inflammatory myokine.	IL6-rs1800795		CG	
SOD2 Superoxide dismutase (SOD2) is manganese dependent and protects against superoxide for the mitochondria of the cell. The homozygous genotype increases the need for antioxidant support in high- intensity athletes.	SOD2-rs4880	AA		
COL1A1 COL1A1 produces alpha 1 chain of type I collagen, a major protein in tendons and ligaments.	COL1A1- rs1800012	сс		
PON1 PON1 (Paraoxonase) plays a large role in removing pesticides. It is also involved with supporting HDL function and LDL oxidation.	PON1-rs662		СТ	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
LPA Lp(a)is a sticky form of LDL that appears to affect plaque growth, LDL particle size and increase the risk of plaque rupture and blood clotting.	LPA-rs3798220	ТТ		
CYP1A2 C164A Variants in CYP1A2 determine caffeine metabolism and effects on bone density and cardiovascular health.	CYP1A2 C164A- rs762551			AA
9p21 9p21 is considered an important genetic marker for cardiovascular health.	9p21-rs4977574			GG
FADS1 FADS1 is involved in fatty acid metabolism, and variants in this gene are associated with elevated triglyceride levels.	FADS1-rs174546			TT
ESR2 ESR2 codes for estrogen receptor beta (ER-II), one of two main types of estrogen receptor activated by estrogen and is linked to fibrinogen levels in post-menopausal women.	ESR2-rs4986938	СС		
F5 Variants in F5 increase the risk of deep vein thrombosis, especially if using oral contraceptives.	F5-rs6025	СС		
ADRB2 Beta-2 adrenergic receptor (ADRB2) is abundantly expressed in cardiac cells, and bronchial smooth muscle cells and is connected to stress levels and heart health.	ADRB2-rs1042713		AG	

Gene & Gene Function	Gene Rsid	Wild Type	Heterozygous	Homozygous
ACE1 G2350A ACE1 is part of the renin- angiotensin system responsible for the conversion of angiotensin I to angiotensin II, constricting blood vessels and elevating blood pressure.	ACE1 G2350A- rs4343		AG	
ADD1 Variants in ADD1 are associated with hypertension in Asians.	ADD1-rs4961		GT	
AGTR1 Angiotensin-II receptor type 1 (AGTR1) is a major component of the renin-angiotensin system for regulating blood pressure and is highly expressed in adipose tissue, liver, leukocytes and the intestine. The homozygous genotype may increase the risk of high blood pressure from excess dietary fat and carbohydrate intake.	AGTR1-rs5186	AA		
ACE2 A8790G ACE2 is part of the renin- angiotensin system, responsible for degrading angiotensin II and providing balance to ACE1 by dilating blood vessels and lowering blood pressure.	ACE2 A8790G- rs2106809		AG	
TMPRSS2 Transmembrane Serine Protease 2 is highly expressed in the prostate and lungs, and the expression is associated with viral susceptibility and prostate cancer.	TMPRSS2- rs2070788		AG	

Sources

Please click the link below if you'd like to browse peer-reviewed studies referenced by this analysis:

https://www.mydna.life/wp-content/uploads/myDNA-Comprehensive-Health-Report-references.pdf