NutraHacker

Complete Gene Mutation Report for Customer: 9e1cf8ea-ebe6-4cf8-865a-65775fc3bfa3

Instructions:

NutraHacker reports mutations (single nucleotide polymorphisms) in this uploaded genome. Genes not reported in this report are either normal, not actionable, or not currently detected by NutraHacker. The expected allele is the one seen in a normally functioning gene. The high risk alleles reported are the ones measured from the uploaded genome. NutraHacker reports the effects of these mutations as discovered by published empirical data and suggests nutritional supplements that can mitigate potential issues caused by these mutations.

This report is meant to serve as a guide for nutritional supplementation for the owner of the genome and is not applicable to any other individual. Supplement quantities and dosages are not included as they are indicated on the purchased product. Multiple recommendations for the same supplement does not mean that the dosage should be multiplied. In the case of a conflict (such as a particular vitamin being both encouraged and discouraged), the owner of the genome should assess his/her own personal biology to decide whether to include or discard that particular supplement.

NOTICE:

State law allows any person to provide nutritional advice or give advice concerning proper nutrition--which is the giving of advice as to the role of food and food ingredients, including dietary supplements. This state law does NOT confer authority to practice medicine or to undertake the diagnosis, prevention, treatment, or cure of any disease, pain, deformity, injury, or physical or mental condition and specifically does not authorize any person other than one who is a licensed health practitioner to state that any product might cure any disease, disorder, or condition.

NutraHacker reports are for scientific, educational and nutritional information only and are not intended to diagnose, cure, treat or prevent any disease, disorder or condition.

Thank you for using NutraHacker. To your health!

Gender of customer: Male

A total of 41 mutations were detected at this time for your genome out of the 195 polymorphisms assessed.

There were 12 homozygous mutations.

There were 29 heterozygous mutations.



Category	RSID	Gene	Expected	Genotype: Risk	Genotype Freq	Gene Function	Consequences	Encourage	Avoid
Detoxification	rs1131857	CPOX4	А	GT: 1/2	36.1087%	Coproporphyrinogen oxidase	N272H, his variant, referred to	NAC, Glutathione,	Sources of mercury
							herein as 'CPOX4', both increases	Possibly EDTA for	exposure
							sensitivity to the neurobehavioral	chelation	
							effects of Hg (Echeverria et al.		
							2006) and modifies urinary		
							porphyrin excretion as a potential		
							biomarker of this effect (Woods et		
							al. 2005, Li and Woods 2009).		
							The population frequencies of the		
							homozygous wildtype (A/A),		
							heterozygous (A/C) and		
							homozygous mutant (C/C)		
							genotypes within this cohort were		
							0.72, 0.25, and 0.03, respectively,		
							and were equally prevalent		
							among males and females,		
							suggesting substantial exposure		
							to the CPOX4 variant.		
Detoxification	rs2606345	CYP1A1	С	AC: 1/2	44.4935%	Phase I xenobiotic metabolism,	Reduced function of enzyme,	Diindolylmethane	
						PAH's, metabolize E2 to	effects vary with race		
						2-hydroxyestradiol			
Detoxification	rs762551	CYP1A2	А	AC: 1/2	46.8902%	Hydroxylation or dealkylation of	Slow to metabolize caffeine, Main	Induce with broccoli,	Curcumin, Cumin,
						xenobiotics, Phase I, metabolize	liver pathway	Cabbage,	Grapefruit
						E2 to 2-hydroxyestradiol		Diindolylmethane,	
								Glucarate, NAC,	
								Cardamom,	
								Sulforaphane	
Detoxification	rs1799853	CYP2C9	С	CT: 1/2	16.3208%	Metabolizes coumadin, NSAID's,	20% reduction in activity		Substrates of this
						aspirin, phenytoin and	heterozygous, 40% reduction		enzyme
						sulfonylureas	homozygous		

Category	RSID	Gene	Expected	Genotype: Risk	Genotype Freq	Gene Function	Consequences	Encourage	Avoid
Detoxification rs1695	rs1695	GSTP1	G	AA: 2/2	48.1693%	Conjugation toxins to glutathione	Persons having the alleles AA or	NAC, Whey	Vitamin E
							AG 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.		
Detoxification	rs1041983	NAT2	С	TT: 2/2	N/A	This gene encodes an enzyme	Decreased activity	NAC, Vitamin B2,	
					1	that functions to both activate and		Vitamin B3, Vitamin	
						deactivate arylamine and		B5, Molybdenum	
						hydrazine drugs and carcinogens.			
Detoxification	rs1799930	NAT2	G	AA: 2/2	6.76530%	This gene encodes an enzyme	Slow metabolizer	NAC, Vitamin B2,	
					1	that functions to both activate and		Vitamin B3, Vitamin	
						deactivate arylamine and		B5, Molybdenum	
						hydrazine drugs and carcinogens.			
Detoxification	rs1800566	NQO1	С	AG: 1/2	35.5447%	Reduces quinones to	This is a null mutation and		
						hydroquinones (vitamin E	removal of carcinogenic quinones		
						alpha-tocopherol quinone,	is affected negatively		
						menadione, benzene quinones)			
Detoxification	rs182420	SULT2A1	А	CT: 1/2	25.5034%	Catalyze the sulfate conjugation	Decreased enzyme function	NAC, MSM, Taurine	
						of many hormones,			
						neurotransmitters, drugs, and			
						xenobiotic compounds			
Detoxification	rs2910397	SULT2A1	G	CT: 1/2	39.7265%	Catalyze the sulfate conjugation	Decreased enzyme function	NAC, MSM, Taurine	
						of many hormones,			
						neurotransmitters, drugs, and			
						xenobiotic compounds			
Neurotransmitter	rs578776	CHRNA5	Т	AG: 1/2	48.4664%	Neuronal acetylcholine receptor	Increased nicotine intake		Nicotine
Levels					1	subunit alpha-5			
Neurotransmitter	rs4633	COMT	С	TT: 2/2	17.6329%	Degrades catecholamines, Phase	Same amino acid sequence,	Hydroxy B12	Methyl B12, Methyl
Levels						II, inactivates hydroxy-estrogens	lower expression of gene, less	(hydroxycobalamin)	donors
							breakdown of catecholamines		

Category	RSID	Gene	Expected	Genotype: Risk	Genotype Freq	Gene Function	Consequences	Encourage	Avoid
Neurotransmitter	rs4646312	COMT	G	TT: 2/2	50.2164%	Degrades catecholamines, Phase	Decreased COMT activity	Hydroxy B12	Methyl B12, Methyl
Levels						II, inactivates hydroxy-estrogens		(hydroxycobalamin)	donors
Neurotransmitter	rs4680	COMT	G	AA: 2/2	16.5958%	Degrades catecholamines, Phase	Slower breakdown dopamine,	Hydroxy B12	Methyl B12, Methyl
Levels						II, inactivates hydroxy-estrogens	oestrogen, worrier, prone to	(hydroxycobalamin)	donors, Cannabis
							anxiety, more sensitive to green		
							tea		
Neurotransmitter	rs933271	COMT	Т	CT: 1/2	47.5934%	Degrades catecholamines, Phase	Decreased COMT activity	Hydroxy B12	Methyl B12, Methyl
Levels						II, inactivates hydroxy-estrogens		(hydroxycobalamin)	donors
Neurotransmitter	rs2391191	DAOA	G	AG: 1/2	49.3000%	D-amino acid oxidase activator,	Associated with cognitive manic	Idebenone,	
Levels						which degrades D-serine, a potent	symptoms	Piracetam,	
						activator of NMDA receptors		Magnesium, Taurine,	
								Lithium orotate	
Neurotransmitter	rs2241165	GAD1	А	CT: 1/2	49.1201%	Catalyzes production of GABA	High glutamate, low GABA	Taurine, Theanine,	MSG
Levels						from glutamate		NAC, Glycine	
Neurotransmitter	rs2769605	NTRK2	С	CT: 1/2	43.9018%	Neurotrophic tyrosine kinase	Decreased BDNF	Theanine, Curcumin,	
Levels						receptor type 2		Beta-alanine, Lithium	
								orotate,	
								Phosphatidylserine	
Folate One-Carbon	rs3758149	GGH	С	AG: 1/2	37.7989%	This gene catalyzes the hydrolysis	May have functional relevance		
Metabolism /						of folylpoly-gamma-glutamates	and result in alterations in plasma		
Methylation (FOCM)						and	homocysteine levels.		
						antifolylpoly-gamma-glutamates			
						by the removal of gamma-linked			
						polyglutamates and glutamate			
Folate One-Carbon	rs2236225	MTHFD1	G	AG: 1/2	46.1659%	Three distinct enzymatic activities	Increased requirement for choline	Choline	
Metabolism /						related to folate			
Methylation (FOCM)									
Folate One-Carbon	rs1801394	MTRR	А	AG: 1/2	49.3785%	Methylates, recycles vitamin b12	Poor methylation of Vitamin B12	Methyl B12,	
Metabolism /							leading to higher homocysteine	L-methylfolate	
Methylation (FOCM)							levels.		
Folate One-Carbon	rs7946	PEMT	С	TT: 2/2	16.8872%	Converts	Fatty liver due to low choline	Phosphatidylcholine	
Metabolism /					1	phosphatidylethanolamine to			
Methylation (FOCM)						phosphatidylcholine			

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HPA axis / Endocrine	rs3774261	ADIPOQ	А	GG: 2/2	20.3505%	Important adipokine involved in	Decreased adiponectin	Omega-3 fatty acids	
						the control of fat metabolism and		like fish oil, Coffee,	
						insulin sensitivity, with direct		Leucine, Magnesium,	
						anti-diabetic, anti-atherogenic and		Fiber, Exercise	
						anti-inflammatory activities.			
HPA axis / Endocrine	rs1801260	CLOCK	А	AG: 1/2	34.2473%	Circadian Locomotor Cycles	Late sleeping time	Be mindful of sleep	
						Kaput		time	
HPA axis / Endocrine	rs1049353	CNR1	G	CT: 1/2	27.7148%	Cannabinoid Receptor 1	The rs1049353 polymorphism of	Omega-3 fatty acids	
							the CNR1 gene is associated with	like fish oil, Coffee,	
							decreased levels of adiponectin.	Leucine, Magnesium,	
								Fiber	
HPA axis / Endocrine	rs560887	G6PC2	Т	CC: 2/2	72.8756%	This gene encodes an enzyme	Fasting blood glucose level	Chromium, Vanadium	High carb diets
						belonging to the	higher. This is actually the more		
						glucose-6-phosphatase catalytic	common form		
						subunit family. These enzymes			
						are part of a multicomponent			
						integral membrane system that			
						catalyzes the hydrolysis of			
						glucose-6-phosphate, the terminal			
						step in gluconeogenic and			
						glycogenolytic pathways, allowing			
						the release of glucose into the			
						bloodstream. The family member			
						encoded by this gene is found in			
						pancreatic islets.			
HPA axis / Endocrine	rs852977	NR3C1	G	AA: 2/2	56.1304%	Glucocorticoid receptor	Mutation associated with	Phosphatidylserine,	
						· ·	generalized glucocorticoid	Possibly ketogenic	
							resistance, high cortisol, CFS	diet	
HPA axis / Endocrine	rs1544410	VDR	G	CT: 1/2	42.7506%	Vitamin D Receptor	Downregulated Vitamin D	Vitamin D3, Sage,	Methyl donors
						, i	receptor	Rosemary	·
HPA axis / Endocrine	rs731236	VDR	Α	AG: 1/2	43.3464%	Vitamin D Receptor	Downregulated Vitamin D	Vitamin D3, Sage,	Methyl donors
					-	1	receptor, can affect dopamine	Rosemary	,
							levels	,	
Cardiovascular	rs4654748	ALPL	С	CT: 1/2	45.9348%	alkaline phosphatase	Lower concentration b6	Vitamin B6	
- Cardio vasculai	10-100-17-10	1,,,,,,,	1~	01. I/L	13.30 70 /0	antanii o prioopriatase	201101 CONTROLLED TO	VIX.ATTIIT BO	<u> </u>

Category	RSID	Gene	Expected	Genotype: Risk	Genotype Freq	Gene Function	Consequences	Encourage	Avoid
Cardiovascular	rs174537	FADS1	G	GT: 1/2	35.4912%	Fatty Acid Desaturase 1	Regulation of saturation of fatty	Omega-3 fatty acids	
							acids, significantly associated with	like fish oil, Plant oils	
							lower concentrations of long-chain		
							PUFA. At baseline, men with the		
							rs174537T allele had lower		
							arachidonic acid (AA) and		
							AA/linoleic acid (LA), and higher		
							interleukin (IL)-6 levels than		
							rs174537GG counterparts. After 3		
							years, rs174537GG men had		
							significantly increased AA (P =		
							0.022),		
							AA/dihomo-gamma-linolenic acid		
							(DGLA) (P = 0.007),		
							docosapentaenoic acid (DPA),		
							low-density lipoprotein (LDL)		
							cholesterol, and oxidized LDL		
							(ox-LDL), but decreased		
							eicosatrienoic acid. The		
							rs174537T group showed		
							significantly increased		
							gamma-linolenic acid and ox-LDL,		
							and decreased eicosadienoic		
							acid, eicosapentaenoic acid		
							(EPA)/alpha-linolenic acid (ALA),		
							and IL-6.		
Cardiovascular	rs2070744	NOS3	Т	CT: 1/2	29.6035%	In a process dependent on BH4,	Elite power performance but	Omega-3 fatty acids	
						NOS converts arginine into nitric	certain risks to health	like fish oil	
						oxide and assists in ammonia			
						detoxification. In the absence of			
						BH4, NOS will convert Arginine			
						into peroxynitrite or superoxide			

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Cardiovascular	rs662	PON1	А	CT: 1/2	49.3911%	Major antiatherosclerotic component of HDL	Glutamine high activity, arginine low activity, position 192, Low serum PON1 activity in NIDDM	Omega-3 fatty acids like fish oil, Fat soluble antioxidants,	High fat diet
							may be related to an increased	Vitamin K	
							tendency to lipid peroxidation and	vitariiir ix	
							may also increase susceptibility to		
							toxicity from organophosphate		
							exposure.		
Cardiovascular	rs854571	PON1	С	CT: 1/2	48.5938%	Major antiatherosclerotic	Decreased function	Omega-3 fatty acids	Cholesterol, High fat
						component of HDL		like fish oil, Fat	diet
								soluble antioxidants,	
								Vitamin K	
Cardiovascular	rs9923231	VKORC1	С	TT: 2/2	15.9922%	Reduces vitamin K 2,3-epoxide to	Related to vitamin K recycling.	Vitamin K	
						the enzymatically activated form.			
Digestion / Elimination	rs492602	FUT2	Т	AG: 1/2	49.6357%	Fucosyltransferase 2 enzyme	Reduced intestinal microbiota	Probiotics	
						which determines 'secretor status'	diversity but higher vitamin B12		
							levels		
Digestion / Elimination	rs601338	FUT2	G	AG: 1/2	49.5914%	Fucosyltransferase 2 enzyme	Reduced intestinal microbiota	Probiotics	
						which determines 'secretor status'	diversity, non secretor		
Digestion / Elimination	rs602662	FUT2	G	AG: 1/2	49.7883%	Fucosyltransferase 2 enzyme	Reduced intestinal microbiota	Probiotics	
						which determines 'secretor status'	diversity. Interferes with		
							absorption of B12. Individuals on		
							vegetarian diet with GG		
							(homozygous major genotype)		
							have significantly lower levels of		
							vitamin B(12).		
Digestion / Elimination	rs10889677	IL-23R	С	AC: 1/2	47.5499%	Important part of the inflammatory	Affects intestinal health	Probiotics, Omega-3	
						response against infection. It		fatty acids like fish oil,	
						promotes upregulation of the		Vitamin D3	
						matrix metalloprotease MMP9,			
						increases angiogenesis and			
						reduces CD8+ T-cell infiltration.			

Category	RSID	Gene	Expected	Genotype: Risk	Genotype Freq	Gene Function	Consequences	Encourage	Avoid
Energy / Oxidation	rs4880	SOD2	А	AG: 1/2	48.9123%	Mitochondrial Superoxide	Decreased gene function. Noise	Manganese, Vitamin	Alcohol, Noise
						Dismutase 2	induced hearing loss, rs10370	E in tocotrienol form	(greater chance for
							'TT', rs4880 'GG' diplo-genotype		hearing loss)
							(diplotype) was associated with		
							more gray matter shrinkage in 76		
							individuals who report chronic		
							high levels of alcohol		
							consumption.		
Energy / Oxidation	rs2855262	SOD3	Т	CC: 2/2	14.9428%	Manganese superoxide dismutase	Decreased gene function	Vitamin E in	
								tocotrienol form,	
								Manganese	