# Estrogenomic Profile



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Patient: SAMPLE

**PATIENT** 

Age: 44 Sex: F MRN:

#### **Order Number:**

Completed: January 31, 2008 Received: January 16, 2008 Collected: January 14, 2008

### Apo E

#### Location:

Chromosome 19

# APOE

APO E2: cys / cys APO E3: cys / arg APO E4: arg / arg

Your Genotype:



The two SNPs lead to 3 possible variants for each chromosome, known as ApoE2, E3, & E4.

# Apolipoprotein E: CHOLESTEROL REGULATION

Apolipoprotein E (Apo E) plays a key role in lipid metabolism by helping to remove dietary cholesterol (chylomicrons and VLDL) from the bloodstream.

#### **Health Implications**

- $\cdot$  The APO E2/3 genotype is common, accounting for 10-15% of most populations
- · APO E2 is associated with lower LDL cholesterol and higher HDL-C, but higher triglycerides (as found in Metabolic Syndrome) compared to the other genotypes
- · APO E2 also confers a lower risk of atherosclerosis, myocardial infarction, stroke, and osteoporosis, and higher antioxidant activity

#### **Treatment Options**

- The cholesterol-lowering effect of a low saturated fat and low cholesterol diet is least effective with E2 individuals
- · Minimize high-glycemic index foods, which produce the largest triglyceride (TG) response in E2 carriers
- · Dietary fiber, fish oils, and exercise generally improve the lipid profile in this genotype; fish oils reduce TGs most effectively in E2 individuals
- · Alcohol may reduce LDL-C in men (neutral in women)
- E2 individuals generally respond the most favorably to statins and would therefore likely respond to statin mimetics such as inositol hexaniacinate, red rice yeast, and policosanol
- · Gemfibrozil may be particularly effective at lowering TGs and total cholesterol
- · HRT improves the lipid profile in this genotype, although oral estrogen may significantly increase TGs

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Neither chromosome carries the genetic variation.

**+** - (

One chromosome (of two) carries the genetic variation.

**+ +** Both chromosomes carry the genetic variation.

(You inherit one chromosome from each parent)

+ 1

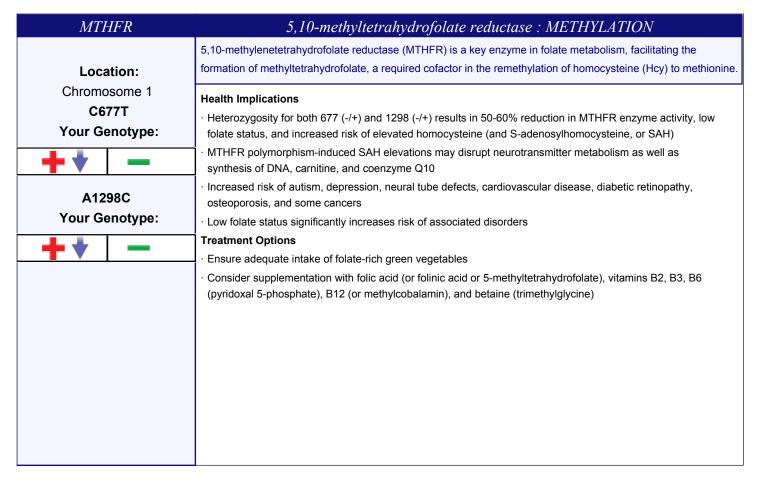
Gene activity increased



Gene activity decreased



# CYP1B1 Cytochrome p450 1B1: DETOXIFICATION CYP1B1 is a Phase I detoxification enzyme responsible for the 4-hydroxylation of estrogen as well as the activation of environmental toxins such as polycyclic aromatic hydrocarbons, PCBs, and aflatoxin B1. Location: Chromosome 2 **Health Implications** L432V $\cdot$ Hyper-induction of CYP1B1 upon exposure to substrates or inducers · Increased production of 4-hydroxyestrogens and potentially carcinogenic compounds Your Genotype: Tendency for lower 2:16α-hydroxyestrone ratio (higher risk of breast cancer) · Increased risk of breast cancer, especially if xenobiotic exposure (e.g., PAHs), high body mass index, estrogen therapy >= 4 yrs, or coexisting CYP1A1 polymorphism (I462V) N453S Possible increased risk of cancer of the ovary, uterus, prostate, and lung (esp. if exposed to second-hand Your Genotype: smoke) **Treatment Options** Minimize exposure to xenobiotics (e.g., PAHs) and xenoestrogens (e.g., organochlorines), which increase CYP1B1 activity Maintain a diet rich in antioxidants (colorful fruits and vegetables), consider supplementation Consider redirecting estrogen metabolism away from 4-hydroxylation with cruciferous vegetables and/or agents such as indole 3-carbinol (I3C), diindolylmethane (DIM), fish oils, or rosemary Caution using long-term estrogen therapy, especially conjugated equine estrogens, which are preferentially 4-hydroxylated. Combined estrogen/progestin therapy produces the greatest breast density in carriers of the Carcinogen-induced DNA damage may be minimized by agents such as curcumin, black cohosh, genistein, and DHEA



## COMTCatechol-O-MethylTransferase: METHYLATION COMT is a key enzyme in the deactivation of catechol compounds such as catecholamines, estrogens, various chemicals, and toxins. COMT modulates the neurotransmitter functions of dopamine and norepinephrine. Location: Chromosome 22.11q **Health Implications** V158M $\cdot$ 3-4-fold reduction in COMT activity with increased bioavailability of catecholamines and impaired methylation Your Genotype: of catechol estrogens · Increased risk of nervousness, anxiety, or panic disorder Reduced pain threshold and increased risk of fibromyalgia **Treatment Options** · Ensure adequate B6, B12, folate, magnesium, and methionine to support formation of S-adenosylmethionine and prevent elevated homocysteine · Ensure adequate anti-oxidants to prevent oxidation of dopamine and pro-carcinogenic 4-hydroxyestrogens · Exercise caution using MAO inhibitors, tricyclics, mirtazapine (Remeron®) or stimulants - especially in patients with bipolar disorder · Inferior anti-depressant response to mirtazapine (Remeron®)

TNF-a	Tumor Necrosis Factor a : INFLAMMATION
Location: Chromosome 6 -308G-A Your Genotype:	Tumor Necrosis Factor a : INFLAMMATION  TNF-α is a pro-inflammatory cytokine secreted from activated macrophages that plays an important role in host defense. Excessive TNF-α release can lead to inflammatory reactions and oxidative stress.  Health Implications  Decreased production of TNF-α, decreased inflammatory tendency and oxidative stress  Decreased risk of autoimmune disease, osteoporosis, insulin resistance  May be associated with increased risk of some cancers because of TNF-α's anti-neoplastic properties  Treatment Options  Risk of inflammatory disorders is minimal  Diet and lifestyle associated with minimizing cancer risks is prudent

IL-6	InterLeukin-6 : INFLAMMATION
Location: Chromosome 7 -174G - C Your Genotype:	IL-6 is a TH-2 cytokine that promotes maturation of antibody-producing B-cells. IL-6 mediates inflammatory and stress-induced responses.  Health Implications  · Higher baseline levels of IL-6; increased risk of inflammation  · Increased stress response, with stimulation of HPA axis and cortisol  · Increased risk of osteoporosis, atherosclerosis, stroke, auto-immune disease, and Type II diabetes (in some populations)  Treatment Options
	<ul> <li>Stress management; support adrenal function</li> <li>Avoid trans fats, ensure adequate intake of Ω-3 fatty acids</li> <li>IL-6 release is reduced by a Mediterranean-style diet, N-acetyl cysteine, anti-oxidants, Siberian ginseng, curcumin, conjugated linoleic acid, estrogen, progesterone, DHEA, and COX-2 inhibitors</li> </ul>

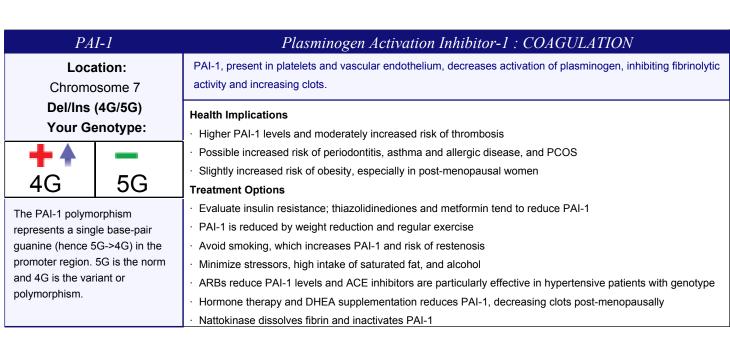
# Vitamin D Receptor: HORMONAL BONE FORMATION **VDR** VDR is an intracellular hormone receptor that specifically binds the active form of vitamin D and interacts with target-cell nuclei to produce effects. Location: Chromosome 12 **Health Implications Bsml RFLP** · Slight impairment of vitamin D receptor with resistance to vitamin D3 Your Genotype: · Slightly increased risk of impaired calcium absorption, increased bone loss, lower bone mineral density, and enhanced bone lead accumulation · Moderately reduced risk of prostate cancer **Treatment Options** · Carriers of the (+) allele benefit from vitamin D supplementation · Ensure adequate calcium (Ca) intake; studies suggest minimum of 900 mg/day · Vitamin K may help to compensate for the higher risk of bone loss $\cdot$ Caffeine intake >300 mg/day may accelerate bone loss, especially when low calcium intake · Favorable bone response to etidronate and raloxifene and HRT

EYP1A1 is a Phase I detoxification enzyme responsible for the 2-hydroxylation of estrogen as well as the activation of common environmental toxins, such as polycyclic aromatic hydrocarbons (PAHs).  Bealth Implications Baseline "normal" CYP1A1 enzyme activity  Normal" degree of procarcinogen activation upon exposures to substrates
Baseline "normal" CYP1A1 enzyme activity  'Normal" degree of procarcinogen activation upon exposures to substrates
Possible decreased risk of endometriosis
reatment Options Regardless of CYP1A1 genotype, it is recommended to minimize exposure to PAHs (e.g. smoke and well-done meats), PCBs (e.g., contaminated fish or waste), and dioxins (e.g., chlorine bleaching, PVC plastics, incineration)
Maintain a diet rich in antioxidants (colorful fruits and vegetables)
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GSTM1	Glutathione S-Transferase mu-1 : DETOXIFICATION
Location: Chromosome 1	GST is responsible for Phase II detoxification of xenobiotics, carcinogens, and products of oxidative stress. GSTM1 is located primarily in the liver.
Your Genotype:	Health Implications  · GSTM1 enzyme activity is absent, with reduced detoxification capacity
The GSTM1 gene is either PRESENT or ABSENT (also called Null). If either copy is present, it is termed PRESENT. If both copies are absent, it is termed ABSENT.	<ul> <li>Increased risk of toxic burden, oxidative stress, atopic asthma, lung problems, cancer, chemical sensitivit and coronary artery disease</li> <li>Decreased risk of cancer, only with high intake of cruciferous vegetables</li> <li>Treatment Options</li> <li>Eat cruciferous vegetables and allium foods to reduce cancer risk</li> <li>Eat a diet rich in antioxidants (colorful foods), consider supplementation</li> <li>Ensure availability of glutathione precursors and cofactors</li> <li>Limit glutathione depletion with α-lipoic acid, milk thistle, or taurine</li> <li>Minimize exposure to xenobiotics, including PAHs and toxic metals</li> </ul>

### GSTP1 Glutathione S-Transferase pi-1: DETOXIFICATION GST is responsible for Phase II detoxification of xenobiotics, carcinogens, steroids, heavy metals, and products Location: of oxidative stress. GSTP1 is located primarily in the brain and lungs. Chromosome 11 A114V **Health Implications** Your Genotype: Polymorphisms are associated with either higher or lower enzyme activity, depending on specific environmental exposures; therefore, the (-/-) genotype may still increase risk for some disorders. The I105V snp is the more significant of the two. The I105V genotype (-/-) is associated with slightly increased risk of some cancers (especially if exposed **I105V** to cigarette smoke), also atopy, xenobiotic-induced asthma, and COPD Your Genotype: **Treatment Options** Ensure availability of glutathione precursors and cofactors, e.g., methionine-rich foods, NAC, L-glutamine, glycine, Mg, B6 Eat a diet rich in antioxidants (colorful foods), consider supplementation Minimize exposure to xenobiotics, including polycyclic aromatic hydrocarbons (e.g., cigarette smoke) and toxic metals

### Platelet Glycoprotein IIIa : COAGULATION GP3A PL(A) GP3A is a protein component of the platelet fibrinogen receptor IIbIIIa, playing a pivotal role in platelet Location: aggregation and thrombus formation. Chromosome 17 **PL(A1)/ PL(A2) Health Implications** Your Genotype: · Decreased platelet aggregability and decreased risk of clot formation · Greater risk of perioperative bleeding due to longer bleeding time **Treatment Options** · Aspirin and oral platelet antagonists are most effective in this genotype · This genotype may be less sensitive to platelet - inhibiting effects of estrogen The GP3A polymorphism is a L33P change that results from the substitution of cytosine for thymidine at position 1565. Clinical studies commonly refer to this change as PL(A1) -> PL(A2).



FACTOR II	Factor II (Prothrombin) : COAGULATION
Location: Chromosome 11 G20210A Your Genotype:	Factor II (Prothrombin): COAGULATION  Factor II is also known as prothrombin, which is converted to its active form, thrombin, and forms the essential part of a blood clot.  Health Implications  · Normal levels of prothrombin  · No increased risk of venous thromboembolism  Treatment Options  · None indicated

FACTOR V	Factor V (Leiden): COAGULATION
Location:	Factor V combines with Factor X to convert prothrombin to thrombin, the essential part of a blood clot. Factor Va is held in check by Protein C.
Chromosome 1  R506Q  Your Genotype:	Health Implications  · Normal inactivation of Factor V by activated Protein C  · No increased risk of venous thromboembolism  Treatment Options
	· None indicated

This test has been developed and its performance characteristics determined by Genova Diagnostics, Inc. It has not been cleared or approved by the U.S. Food and Drug Administration.

Commentary is provided to the practitioner for educational purposes, and should not be interpreted as diagnostic or treatment recommendations. Diagnosis and treatment decisions are the responsibility of the practitioner.

The accuracy of genetic testing is not 100%. Results of genetic tests should be taken in the context of clinical representation and familial risk. The prevalence and significance of some allelic variations may be population specific.

Any positive findings in your patient's test indicate genetic predisposition that could affect physiologic function and risk of disease. We do not measure every possible genetic variation. Your patient may have additional risk that is not measured by this test. Negative findings do not imply that your patient is risk-free.

DNA sequencing is used to detect polymorphisms in the patient's DNA sample. The sensitivity and specificity of this assay is <100%.