

DEMO DEMO

Name: DEMO DEMO
Date of Birth: 05-29-1968
Biological Sex: Male
Age: 57
Height:
Weight:
Fasting: UNKNOWN

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FINAL REPORT

Accession ID: 2692051854

Provider Information

Practice Name: DEMO CLIENT, MD
Provider Name: DEMO CLIENT, MD
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Report Information

Current Result Previous Result In Control Moderate Risk

Specimen Information

Sample Type	Collection Time	Received Time	Report	Final Report Date
Metal Free Urine	2024-04-02 13:27 (UTC)	2024-04-03 22:22 (UTC)	Oxidative Stress Profile - P2	2024-04-16 18:06 (UTC)
Saliva	2024-04-02 13:25 (UTC)	2024-04-03 22:22 (UTC)	Oxidative Stress Profile - P2	2024-04-16 18:06 (UTC)

INTRODUCTION

Vibrant Wellness is pleased to present to you, 'Oxidative Stress Profile', to help you make healthy lifestyle, dietary and treatment choices in consultation with your healthcare provider. It is intended to be used as a tool to encourage a general state of health and well-being. The Vibrant 'Oxidative Stress Profile' is a test to identify and quantify the level of a large set of oxidative damage markers and to identify antioxidant genetics variations. The panel is designed to evaluate oxidative stress by measuring the levels of damage caused by oxidative species resulting from the impact of ROS and RNS on lipids, DNA, RNA and proteins and to give a complete picture of genetic predispositions that code for enzymes and antioxidants which can significantly impact oxidative stress response.

Methodology:

The Vibrant Oxidative Damage Markers panel uses tandem mass spectrometry methodology (LC-MS/MS) for quantitative detection of damage markers in urine samples. Urine creatinine is measured using a kinetic colorimetric assay based on the Jaffé method. All damage markers are reported as the quantitative result normalized to urine creatinine to account for urine dilution variations.

The Vibrant Antioxidant Genetics panel uses real-time PCR methodology. DNA is extracted and purified from saliva samples and a SNP (single nucleotide polymorphism) genotyping assay is performed using real-time PCR to detect the specific allele targets of each assay performed.

Interpretation of Report:

The report begins with the summary page which displays a summary flowchart of all antioxidant genetic variations of the human body's defense against oxidative stress and indicates the areas of concern from the genetic results observed. The set of analytes with risk associated variants are also summarized. The summary also includes the damage markers whose levels are high or moderate based on the reference range. This is followed by a graphical representation of the overall oxidative damage score which is calculated using the results from all urine damage markers tested applied to a linear regression model and displayed with respect to your age group. The score in green represents a normal score based on 50th percentile population, the score in yellow represents a moderate score based on 90th percentile and the score in red represent a high score based on the relatively healthy population. Reference ranges were determined using urine samples from 1000 apparently healthy individuals. Additionally, the previous value is also indicated to help check for improvements every time the test is ordered.

Following this section is the complete list of the genetic markers measured in the panel. Elevated risk associated variants are indicated with red, partially elevated risk associated variants are indicated with yellow and alleles with no risk are indicated with green. This is followed by a list of all damage marker results and their absolute levels are normalized with respect to Creatinine in a histogram format to enable a full overview along with the reference ranges. The level of the analyte with reference range is shown with three shades of color – Green, Yellow and Red. The result in green corresponds to 0th to 75th percentile indicates mild detection of the analyte. The result in yellow corresponds to 75th to 95th percentile indicates moderate detection of the analyte whereas the result in red corresponding to greater than 95th percentile indicates high detection of the analyte.

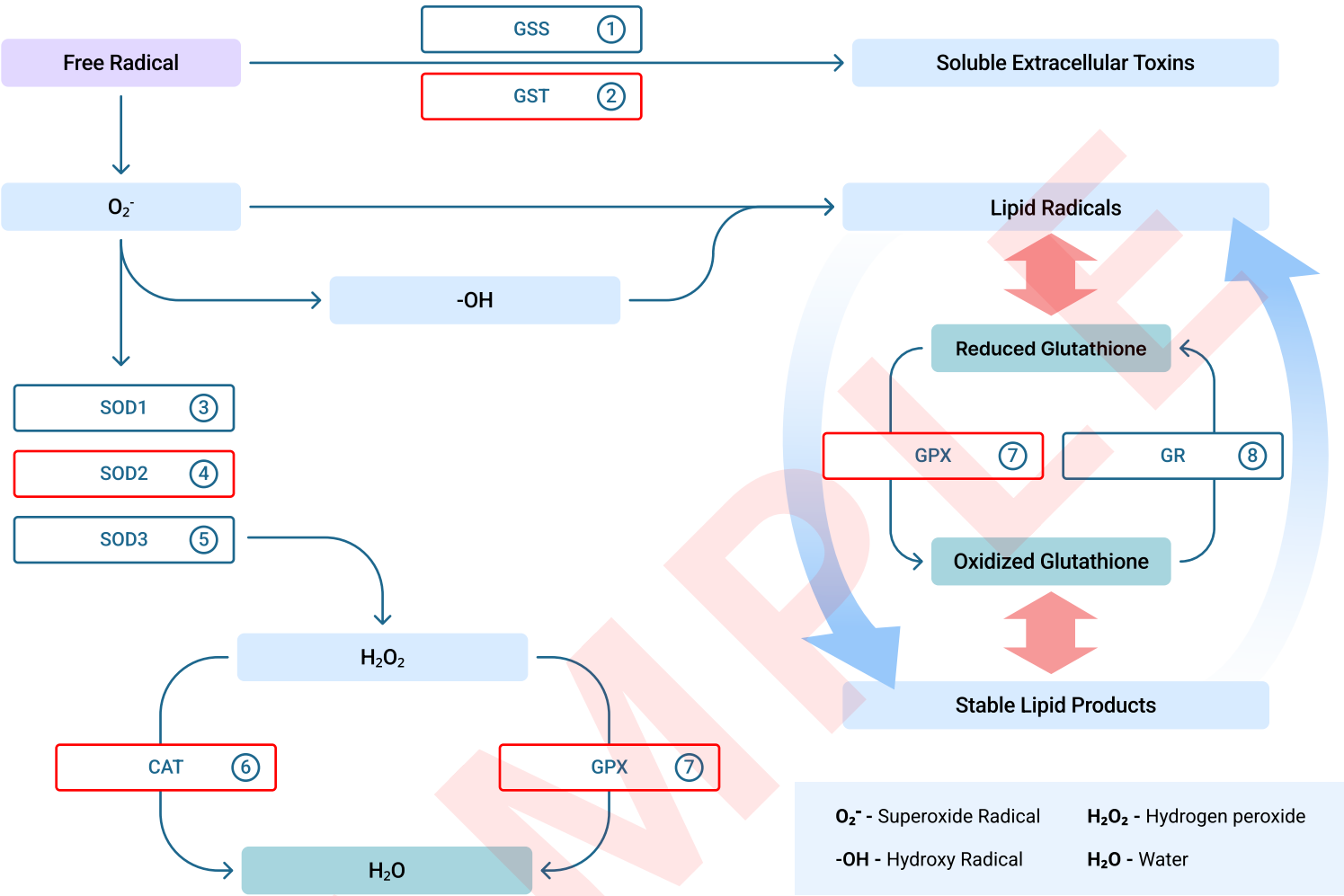
All contents provided in the report are purely for informational purposes only and should not be considered medical advice. Any changes based on the information should be made in consultation with the healthcare provider.

The Vibrant Wellness platform provides tools for you to track and analyze your general wellness profile. Testing for the Oxidative Stress panel is performed by Vibrant America, a CLIA certified lab CLIA#:05D2078809 and Vibrant Genomics, a CLIA certified lab CLIA#: 05D2098445. Vibrant Wellness provides and makes available this report and any related services pursuant to the Terms of Use Agreement (the "Terms") on its website at www.vibrant-wellness.com. By accessing, browsing, or otherwise using the report or website or any services, you acknowledge that you have read, understood, and agree to be bound by these terms. If you do not agree to accept these terms, you shall not access, browse, or use the report or website. The statements in this report have not been evaluated by the Food and Drug Administration and are only meant to be lifestyle choices for potential risk mitigation. Please consult your healthcare provider for medication, treatment, or lifestyle management. This product is not intended to diagnose, treat, or cure any disease.

Please note:

Pediatric ranges have not been established for this test. It is important that you discuss any modifications to your diet, exercise, and nutritional supplementation with your healthcare provider before making any changes.

Oxidative Stress Profile Summary



Please note that the flowchart only includes only a subset of the genes tested. The complete set of the genes is listed in the tables below under "Antioxidant Genetics".

<div><div>①</div><div>GSS - Glutathione Synthetase (0/1) Involved in Glutathione synthesis</div></div> <div><div>②</div><div>GST - Glutathione S Transferase (3/3) Aids Glutathione in toxin removal Decreasing antioxidant activity leads to elevated oxidative stress, Decreased antioxidant activity</div></div> <div><div>③</div><div>SOD1 - Superoxide Dismutase (0/1) Aids in quenching superoxide free radical</div></div> <div><div>④</div><div>SOD2 - Superoxide Dismutase (1/1) Aids in quenching superoxide free radical Impaired anti-oxidant activity</div></div>	<div><div>⑤</div><div>SOD3 - Superoxide Dismutase (0/2) Aids in quenching superoxide free radical</div></div> <div><div>⑥</div><div>CAT - Catalase (2/3) Aids in quenching hydrogen peroxide Mitochondrial dysfunction</div></div> <div><div>⑦</div><div>GPX - Glutathione Peroxidase (2/5) Aids in reduction of hydrogen peroxide by glutathione Lower selenoprotein enzyme levels, Lower selenoprotein concentrations</div></div> <div><div>⑧</div><div>GR - Glutathione Reductase (0/1) Aids in recycling glutathione</div></div>
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Oxidative Stress Profile - Summary

Antioxidant Genetics					
⊕ ⊕ Homozygous Mutant ⊕ ⊖ Heterozygous ⊖ ⊖ Homozygous Wild					
Test Name	Gene Name	Risk Association	Your Mutation	Your Risk	Reference
rs4756146	CAT	Mitochondrial dysfunction	⊕ ⊕ T/T	Elevated	C/C, C/T
<p>The CAT gene encodes for the catalase enzyme, localized in mitochondria. Mitochondrial catalase was shown to protect cells from oxidative injury induced by hydrogen peroxide by degrading hydrogen peroxide generated by peroxisomal oxidases to water and oxygen, thereby protecting cells from the toxic effects of hydrogen peroxide. Thus, the enzyme participates in antioxidant functions in the body. Mutations in the gene lead to decreased catalase production resulting in excess ROS production. This induces mitochondrial dysfunction and elevated oxidative stress. Individuals with TT genotypes exhibit reduced catalase production leading to ROS buildup and increased risk of oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs7943316	CAT	Mitochondrial dysfunction	⊕ ⊕ T/T	Elevated	A/T, A/A
<p>The CAT gene encodes for the catalase enzyme, localized in mitochondria. Mitochondrial catalase was shown to protect cells from oxidative injury induced by hydrogen peroxide by degrading hydrogen peroxide generated by peroxisomal oxidases to water and oxygen, thereby protecting cells from the toxic effects of hydrogen peroxide. Thus, the enzyme participates in antioxidant functions in the body. Mutations in the gene lead to decreased catalase production resulting in excess ROS production. This induces mitochondrial dysfunction and elevated oxidative stress. Individuals with TT genotypes exhibit reduced catalase production leading to ROS buildup and increased oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs2071566	GPX2	Lower selenoprotein enzyme levels	⊖ ⊖ G/G	Elevated	A/G, A/A
<p>GPX2 is an endogenous selenium-dependent antioxidant encoding for the major antioxidant enzyme called Glutathione peroxidase 2. The GPX2 enzyme catalyzes the reduction of hydrogen peroxide to water and oxygen as well as catalyzing the reduction of peroxide radicals to alcohols and oxygen thus participating in the antioxidant defense system by protecting cells against reactive oxygen species. GPX2 modulates redox-dependent mitochondrial function where mitochondria generate reactive oxygen species (ROS) and respond to ROS-mediated changes in the cellular redox state. Mutations in the gene lead to higher selenoprotein enzyme levels and reduced oxidative damage. Individuals with GG genotypes have a reduced enzyme level and GPX activity leading to oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs4902346	GPX2	Lower selenoprotein concentrations	⊖ ⊖ T/T	Elevated	C/T, C/C
<p>GPX2 is an endogenous selenium-dependent antioxidant encoding for the major antioxidant enzyme called Glutathione peroxidase 2. The GPX2 enzyme catalyzes the reduction of hydrogen peroxide to water and oxygen as well as catalyzing the reduction of peroxide radicals to alcohols and oxygen thus participating in the antioxidant defense system by protecting cells against reactive oxygen species. GPX2 modulates redox-dependent mitochondrial function where mitochondria generate reactive oxygen species (ROS) and respond to ROS-mediated changes in the cellular redox state. Mutations in the gene cause higher selenoprotein enzyme levels and reduced oxidative damage. Individuals with TT genotypes have a reduced enzyme level and GPX activity leading to oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs366631	GSTM1	Decreased antioxidant activity	⊖ ⊖ T/T	Elevated	C/C
<p>GSTM1 gene encodes for an enzyme, glutathione S-transferase Muv 1. In addition to the cytoplasm, it is found in the mitochondria, lysosomes, and nuclear regions. Through the inhibition of cardiolipin (a lipid found in mitochondria) peroxidation and cytochrome c release, mitochondrial GST contributes to the protection of organelles from oxidative stress. The enzyme plays an important regulatory role in detoxification by catalyzing the modification of toxic compounds to glutathione, which is an antioxidant that helps combat Reactive oxygen species (ROS). The mutation leads to lead to significant damage in cells and mitochondrial function, which causes a build-up of ROS therefore, increasing the risk for oxidative stress. Individuals with TT genotypes who have impaired mitochondrial function with ROS build-up have increased oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					

Oxidative Stress Profile - Summary

Antioxidant Genetics					
⊕ ⊕ Homozygous Mutant ⊕ ⊖ Heterozygous ⊖ ⊖ Homozygous Wild					
Test Name	Gene Name	Risk Association	Your Mutation	Your Risk	Reference
rs9932581	CYBA	Elevated ROS production	⊖ ⊖ G/G	Elevated	A/A, A/G
<p>The CYBA gene encodes the p22phox subunit of NADPH oxidase, an enzyme that plays an essential role in the immune system. Upon the detection of foreign invaders, phagocytes are stimulated, and NADPH oxidase is assembled. This enzyme catalyzes the conversion of oxygen to superoxide, a toxic molecule that is used to generate several other highly reactive and toxic substances collectively known as reactive oxygen species (ROS). Phagocytes use these ROS to kill foreign invaders, preventing them from reproducing in the body and causing illness. Mutations in the CYBA gene are associated with higher p22phox expression and increased levels of ROS. The accumulation of ROS can thus result in oxidative stress. Individuals with GG genotype have higher p22phox expression and higher ROS levels, increasing their susceptibility to oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs2796498	PRKAA2	Impaired antioxidant activity	⊕ ⊕ G/G	Elevated	A/G, A/A
<p>The PRKAA2 gene encodes for an enzyme AMP-activated protein kinase (AMPK). AMPK is an important energy-sensing enzyme that monitors cellular energy status. AMPK plays a role in cellular energy homeostasis, largely to activate glucose and fatty acid uptake and oxidation when cellular energy is low. AMPK is part of the antioxidant defense system and is needed to protect the cells from oxidative stress. AMPK promotes mitochondrial biogenesis (a process that occurs in response to increased energy expenditure to produce more ATP). Mutation reduces the expression of the PRKAA2 gene causing impaired AMPK synthesis and impaired antioxidant activity leading to oxidative stress. Individuals with GG genotypes have impaired AMPK synthesis and impaired antioxidant activity leading to oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs1548357	TXNRD2	Impaired mitochondrial oxygen radical scavenging activity	⊖ ⊖ T/T	Elevated	C/T, C/C
<p>The TXNRD2 gene encodes a member of the thioredoxin (Trx) system, a selenocysteine-containing enzyme essential for mitochondrial oxygen radical scavenging. This protein plays a role in antioxidant defenses by reducing thioredoxin 2 (TXN2), which in turn reduces oxidized cysteine in cellular proteins and scavenges peroxides by peroxiredoxins (PRDX), thus protecting cells against oxidative stress. Mutation in the gene decreases enzyme activity which can affect mitochondrial oxygen radical scavenging and antioxidant functions resulting in increased oxidative stress. Individuals with TT genotypes have reduced enzyme activity and mitochondrial oxygen radical scavenging activity leading to higher oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs4880	SOD2	Impaired anti-oxidant activity	⊕ ⊖ C/T	Partially elevated	C/C
<p>The SOD2 gene encodes the mitochondrial enzyme superoxide dismutase 2 (MnSOD), which protects cells from mitochondrial superoxide by converting it to hydrogen peroxide and molecular oxygen. This function allows SOD2 to clear mitochondrial reactive oxygen species (ROS), thereby protecting against oxidative stress. Individuals with CT genotypes exhibit moderate MnSOD activity, with partially impaired mitochondrial import, resulting in intermediate antioxidant defense and moderate oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs10911021	GLUL	Decreased levels of glutamine synthetase and glutathione	⊕ ⊖ C/T	Partially elevated	T/T
<p>The GLUL gene encodes for glutamate ammonia ligase (glutamine synthetase) enzyme. Glutamine synthetase plays a role in maintaining cellular levels of glutamine, an amino acid with multiple functions, including antioxidant properties. Glutamine serves as a precursor for the synthesis of glutathione, a key antioxidant molecule. Glutathione protects the cell from oxidative stress, its availability in reduced form is mandatory to control the redox status of the cell. The mutation leads to the downregulation of the gene leading to enzyme inefficiency that may cause a deficiency of glutamine required for the synthesis of glutathione. Thus increasing the risk of oxidative stress. Individuals with CT genotype exhibit reduced levels of glutamine synthetase enzyme and glutathione, leading to increased oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					

Oxidative Stress Profile - Summary

Antioxidant Genetics					
⊕ ⊕ Homozygous Mutant ⊕ ⊖ Heterozygous ⊖ ⊖ Homozygous Wild					
Test Name	Gene Name	Risk Association	Your Mutation	Your Risk	Reference
rs3754446	GSTM5	Decreased antioxidant activity	⊕ ⊖ G/T	Partially elevated	T/T
<p>The glutathione S transferase Muv 5 (GSTM5) gene belongs to the GST gene family. In addition to the cytoplasm, it is found in the mitochondria, lysosomes, and nuclear regions. Through the inhibition of cardiolipin (a lipid found in mitochondria) peroxidation and cytochrome c release, mitochondrial GSTP contributes to the protection of organelles from oxidative stress. The enzyme plays an important regulatory role in detoxification by catalyzing the modification of toxic compounds to glutathione, which is an antioxidant that helps combat Reactive oxygen species (ROS). The mutation leads to lead to significant damage in cells and mitochondrial function, which causes a build-up of ROS therefore, increasing the risk for oxidative stress. Individuals with GT genotypes exhibit impaired mitochondrial function with ROS build-up and have increased oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs4485648	TrxR2	Impaired mitochondrial redox balance	⊕ ⊖ C/T	Partially elevated	C/C
<p>The TrxR2 gene encodes for the enzyme thioredoxin reductase 2. Thioredoxin reductases are a family of enzymes that maintain cellular redox balance and regulate various cellular processes. TrxR2 is primarily located in the mitochondria and plays a crucial role in maintaining the redox state of proteins and other molecules within the mitochondria. TrxR2 is responsible for reducing oxidized thioredoxin, an antioxidant protein, which allows thioredoxin to carry out its antioxidant and regulatory functions. Mutations or alterations in the TrxR2 gene can disrupt the normal functioning of the enzyme and impair mitochondrial redox balance, resulting in increased oxidative stress. Individuals with CT genotypes have reduced enzyme activity and impaired mitochondrial oxygen radical scavenging activity leading to higher oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs4673	CYBA	Elevated ROS production	⊕ ⊖ C/T	Partially elevated	C/C
<p>The CYBA gene encodes the p22phox subunit of NADPH oxidase, an enzyme that plays an essential role in the immune system. Upon the detection of foreign invaders, phagocytes are stimulated, and NADPH oxidase is assembled. This enzyme catalyzes the conversion of oxygen to superoxide, a toxic molecule that is used to generate several other highly reactive and toxic substances collectively known as reactive oxygen species (ROS). Phagocytes use these ROS to kill foreign invaders, preventing them from reproducing in the body and causing illness. Mutations in the CYBA gene are associated with higher p22phox expression and increased levels of ROS. The accumulation of ROS can thus result in oxidative stress. When present alongside other functional SNPs such as -930 A/G, this variant has been shown to synergistically enhance oxidative stress, as evidenced by increased malondialdehyde (MDA) levels. Individuals with the CT genotype have higher p22phox expression and higher ROS levels, increasing their susceptibility to oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					
rs206812	XDH	Elevated ROS production	⊕ ⊖ A/G	Partially elevated	G/G
<p>The XDH gene encodes the enzyme xanthine dehydrogenase, which is primarily involved in the metabolism of purine compounds. Xanthine dehydrogenase is responsible for the conversion of hypoxanthine to xanthine and xanthine to uric acid, which is antioxidant in nature. However, under certain conditions, xanthine dehydrogenase can undergo conversion to its other form called xanthine oxidase (XO). XO, the oxidized form of xanthine dehydrogenase, has the ability to produce superoxide radicals as a byproduct of its enzymatic activity. Superoxide radicals are reactive oxygen species (ROS) that can be generated during normal cellular processes. It is crucial for cells to efficiently break down these ROS to prevent cellular damage and oxidative stress. Mutations in the XDH gene can disrupt the normal regulation of xanthine dehydrogenase and promote the conversion to XO more readily. This increases the XO activity for higher production of ROS, including superoxide radicals. The accumulation of ROS can thus result in oxidative stress. Individuals with the AG genotype exhibit heightened XO activity and experience an elevation in ROS levels, increasing their susceptibility to oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.</p>					

Oxidative Stress Profile - Summary

Antioxidant Genetics

⊕ ⊕ Homozygous Mutant ⊕ ⊖ Heterozygous ⊖ ⊖ Homozygous Wild

Test Name	Gene Name	Risk Association	Your Mutation	Your Risk	Reference
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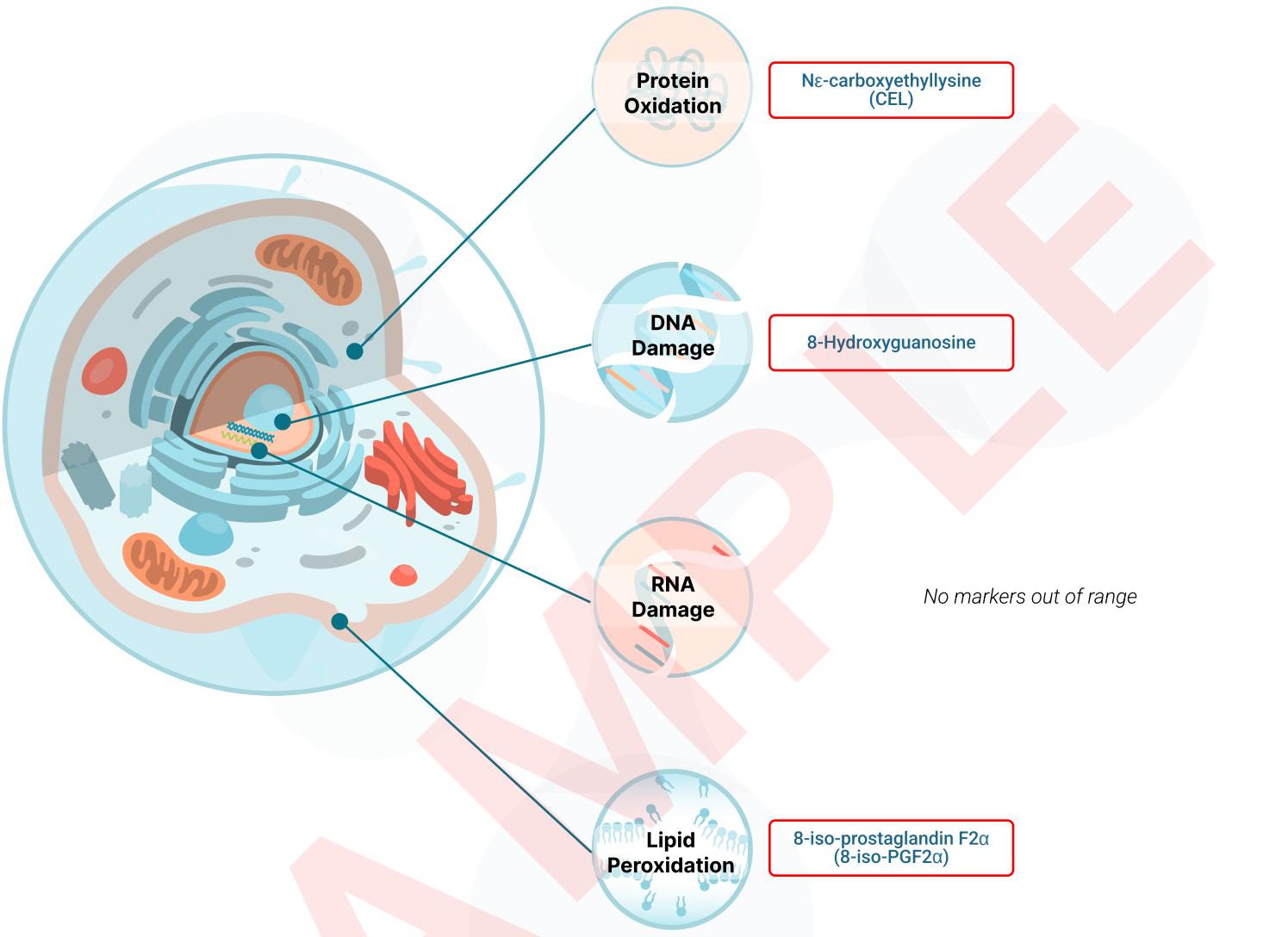
rs1695	GSTP1	Decreasing antioxidant activity leads to elevated oxidative stress	⊕ ⊖ A/G	Partially elevated	A/A
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Glutathione S-transferase P is an enzyme that in humans is encoded by the GSTP1 gene. In addition to the cytoplasm, it is found in the mitochondria, lysosomes, and nuclear regions. Through the inhibition of cardiolipin (a lipid found in mitochondria) peroxidation and cytochrome c release, mitochondrial GSTP contributes to the protection of organelles from oxidative stress. The enzyme plays an important regulatory role in detoxification by catalyzing the modification of toxic compounds to glutathione, which is an antioxidant that helps combat Reactive oxygen species (ROS). The mutation decreases enzyme activity and directly elicits mitochondrial dysfunction, resulting in the rapid generation of ROS, and thus leading to oxidative stress. Individuals with AG genotypes who have decreased gene activity have decreased antioxidant activity and thus increased oxidative stress. Consume antioxidant-rich foods like berries, nuts, and green tea while avoiding processed foods. Engaging in regular moderate exercise is recommended.

rs3877899	SELENOP	Impaired plasma selenium production	⊕ ⊖ C/T	Partially elevated	C/C
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SELENOP encodes a protein selenoprotein P that affects blood selenium or selenoprotein levels in response to supplementation. This selenoprotein accounts for most of the selenium in plasma. It has been implicated as an extracellular antioxidant, and in the transport of selenium to extra-hepatic tissues via apolipoprotein E receptor-2 (apoER2). Mutation in the gene reduces the gene activity and impairs plasma selenium production which leads to increased selenium deficiency, which has the potential of weakening an individual's capacity to respond to oxidative damage involved in the aging process and in most chronic diseases including cancer, cardiovascular disease, diabetes, and dementia. Individuals with CT genotypes who have gene deficiency have a slightly increased risk of selenium deficiency and oxidative stress. Individuals with selenium deficiency are advised to consume brazil nuts, pork, beef, turkey, chicken, fish, shellfish, and eggs. A diet such as breads, grains, meat, poultry, fish, and eggs can increase selenium levels.

Oxidative Stress Profile Summary



Oxidative Stress Biomarkers

Lipid Peroxidation	Current	Previous	Result	Reference
8-iso-prostaglandin F2α (8-iso-PGF2α) (ug/g)	0.12		<div><div></div></div>	≤0.26

8-iso-prostaglandin F2α (8-iso-PGF2α) is an isoprostane generated through the non-enzymatic peroxidation of arachidonic acid in membrane phospholipids. It is found in human plasma and excreted in urine. This biomarker serves as an indicator of oxidative stress and can reliably reflect lipid peroxidation in chronic diseases. Elevated levels of 8-iso PGF2α can lead to DNA oxidation and subsequent structural DNA damage. 8-iso PGF2α is thereby, valuable in assessing oxidative damage to DNA and understanding its implications for cellular health and disease development. Studies have shown that increased levels of 8-iso PGF2α contribute to heightened oxidative stress associated with aging, hypertension, diabetes mellitus, hypercholesterolemia, smoking, and coronary artery disease.

DNA Damage	Current	Previous	Result	Reference
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Free radicals produced either endogenously or exogenously can attack nucleic acid in living cells. Reactions of reactive oxygen species (ROS) and reactive nitrative species (RNS) with RNA yield 8-hydroxyguanosine (8-HdG). Among the known oxidative lesions in nucleic acids, 8-HdG is abundant and appears to be most deleterious due to its high mutagenic potential. This implies that 8-HdG is capable of inducing genetic mutation. RNA dysfunction caused by oxidative damage may contribute to the development of various degenerative diseases. Urinary levels of 8-HdG have risen as indicators of oxidative damage of RNA by ROS.

Glycation is a spontaneous non-enzymatic reaction wherein free reducing sugars bind to free amino groups of proteins, DNA, and lipids. This results in the formation of advanced glycation end-products (AGE). Glycation and oxidative stress are closely linked, and they are together referred to as "glycooxidation". All steps of glycooxidation generate free radicals, some of them being common with the lipid peroxidation pathway. Owing to this, AGE has been considered a urinary biomarker of oxidative stress. The AGE product, Nε-carboxyethyllysine (CEL) is formed when methylglyoxal (formed from the oxidation of lipids and sugars) reacts with lysine. CEL interacts with AGE receptors (RAGEs) which may give rise to oxidative stress. This may even induce cellular dysfunction. Urinary levels of CEL can be used to monitor the degree of oxidative stress in the body system.

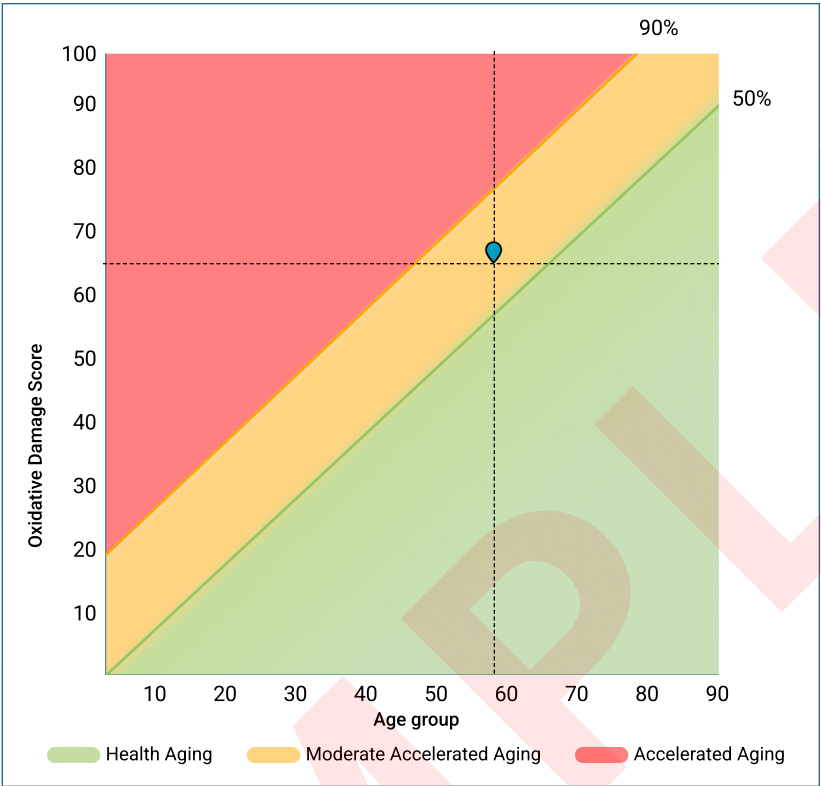
Test Name	Current	Previous	Result	Reference
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Oxidative Stress Profile - Summary

Oxidative Damage Score

Current Result

Previous Result



Result

Your Given Age: 57

Your Oxidative Stress Profile looks similar to a **66.02** year old.

Supplementation Suggestions

Nutrients	Dosage	Purpose
Vitamin C	90 mg/day	Vitamin C enhances GPX1 activity by donating electrons to reduce glutathione (GSH), which is then used by GPX1 to neutralize harmful reactive oxygen species (ROS), thereby protecting cells from oxidative damage. Vitamin C enhances GPX2 activity by donating electrons to reduce glutathione (GSH), which is then used by GPX2 to neutralize harmful reactive oxygen species (ROS), thereby protecting cells from oxidative damage. Vitamin C supplements enhance AMPK activity by promoting the phosphorylation of AMPK through activation of the upstream kinase, LKB1, leading to increased cellular energy sensing and metabolic regulation. Vitamin C supplements enhance catalase activity by donating electrons to the enzyme's active site, increasing its ability to break down hydrogen peroxide into water and oxygen, thus bolstering the antioxidant defense system. Vitamin C supplements enhance the activity of superoxide dismutase (SOD) enzymes by providing electrons necessary for SOD's antioxidant function, thereby increasing SOD activity and reducing oxidative stress markers. Vitamin C supplementation decreases 8-iso-prostaglandin F2α (8-iso-PGF2α) levels by acting as a powerful antioxidant, scavenging free radicals and inhibiting lipid peroxidation, thereby reducing oxidative stress.

Supplementation Suggestions

Nutrients	Dosage	Purpose
Selenium	55 mcg/day	Selenium supplements increase GPX1 activity by incorporating selenium atoms into the GPX1 enzyme's active site, enhancing its ability to catalyze the reduction of harmful reactive oxygen species. Selenium supplements increase GPX2 activity by incorporating selenium atoms into the GPX2 enzyme's active site, enhancing its ability to catalyze the reduction of harmful reactive oxygen species. Selenium, when incorporated into selenoproteins, enhances the activity of catalase by serving as a cofactor, facilitating the breakdown of hydrogen peroxide into water and oxygen, thus increasing catalase's antioxidant function. Selenium supplements enhance the synthesis of selenoproteins, including selenium-dependent superoxide dismutase (SOD), which, in turn, increases SOD activity and levels, boosting cellular antioxidant defenses against superoxide radicals. Selenium reduces 8-iso-prostaglandin F2α (8-iso-PGF2α) levels by acting as a cofactor for the enzyme glutathione peroxidase, which helps neutralize reactive oxygen species responsible for the formation of 8-iso-PGF2α.
Quercetin	500 mg/day	Quercetin suppresses Xanthine Dehydrogenase (XDH) activity by inhibiting its conversion to Xanthine Oxidase (XO), thus reducing the production of reactive oxygen species and preventing oxidative stress. Quercetin supplements downregulate p22phox expression by inhibiting NF-κB activation, thereby reducing oxidative stress through decreased NADPH oxidase activity. Quercetin supplements may increase thioredoxin 2 (Trx2) levels by acting as an antioxidant, scavenging reactive oxygen species (ROS) and reducing oxidative stress, which in turn upregulates Trx2 expression through redox-sensitive pathways.
Vitamin E	22 IU/day	Vitamin E supplements enhance cellular antioxidant defenses by reducing lipid peroxidation, indirectly leading to increased catalase enzyme activity, which helps neutralize harmful reactive oxygen species (ROS). Vitamin E supplements enhance the activity of superoxide dismutase (SOD) enzymes by reducing lipid peroxidation and protecting cellular membranes, which indirectly increases SOD antioxidant markers, helping to neutralize harmful superoxide radicals. Vitamin E supplements reduce 8-iso-prostaglandin F2α (8-iso-PGF2α) levels by acting as an antioxidant, neutralizing reactive oxygen species that would otherwise promote the formation of 8-iso-PGF2α.
Vitamin D	600 IU/day	Vitamin D increases GPX1 activity by binding to vitamin D receptors (VDRs) in cells, which in turn promotes the transcription of GPX1 gene, leading to higher GPX1 enzyme levels and enhanced antioxidant defense. Vitamin D increases GPX2 activity by binding to vitamin D receptors (VDRs) in cells, which in turn promotes the transcription of GPX2 gene, leading to higher GPX2 enzyme levels and enhanced antioxidant defense.

Supplementation Suggestions

Nutrients	Dosage	Purpose
Lutein	10mg/day	Lutein supplements increase GPX1 activity by enhancing the antioxidant defense system through their ability to scavenge free radicals, reducing oxidative stress and thereby promoting GPX1 enzyme function. Lutein supplements increase GPX2 activity by enhancing the antioxidant defense system through their ability to scavenge free radicals, reducing oxidative stress and thereby promoting GPX2 enzyme function.

Oxidative Stress Profile

Antioxidant Genetics					
⊕ ⊕ Homozygous Mutant ⊕ ⊖ Heterozygous ⊖ ⊖ Homozygous Wild					
Test Name	Gene Name	Risk Association	Your Mutation	Your Risk	Reference
rs2234694	SOD1	Increased superoxide levels	⊖ ⊖ A/A	Normal	A/A
rs4880	SOD2	Impaired anti-oxidant activity	⊕ ⊖ C/T	Partially elevated	C/C
rs1799895	SOD3	Increased risk of impaired EC-SOD tissue protection	⊖ ⊖ C/C	Normal	C/G, C/C
rs8192287	SOD3	Disrupted EC-SOD activity	⊕ ⊕ T/T	Normal	T/T, G/T
rs1001179	CAT	Mitochondrial dysfunction	⊖ ⊖ C/C	Normal	C/C
rs4756146	CAT	Mitochondrial dysfunction	⊕ ⊕ T/T	Elevated	C/C, C/T
rs7943316	CAT	Mitochondrial dysfunction	⊕ ⊕ T/T	Elevated	A/T, A/A
rs10911021	GLUL	Decreased levels of glutamine synthetase and glutathione	⊕ ⊖ C/T	Partially elevated	T/T
rs1050450	GPX1	Aberrant redox signaling	⊖ ⊖ C/C	Normal	C/C
rs1987628	GPX1	Reduced antioxidant enzyme leads to selenium deficiency	⊕ ⊕ C/C	Normal	C/C
rs2071566	GPX2	Lower selenoprotein enzyme levels	⊖ ⊖ G/G	Elevated	A/G, A/A
rs4902346	GPX2	Lower selenoprotein concentrations	⊖ ⊖ T/T	Elevated	C/T, C/C
rs713041	GPX4	Lower selenoprotein concentrations	⊖ ⊖ C/C	Normal	C/T, C/C
rs121909307	GSS	Lower glutathione levels	⊕ ⊕ C/C	Normal	C/C
rs2071746	HMOX1	Decreased heme oxygenase 1 activity	⊕ ⊕ T/T	Normal	T/T
rs366631	GSTM1	Decreased antioxidant activity	⊖ ⊖ T/T	Elevated	C/C
rs3754446	GSTM5	Decreased antioxidant activity	⊕ ⊖ G/T	Partially elevated	T/T
rs4485648	TrxR2	Impaired mitochondrial redox balance	⊕ ⊖ C/T	Partially elevated	C/C
rs4673	CYBA	Elevated ROS production	⊕ ⊖ C/T	Partially elevated	C/C
rs9932581	CYBA	Elevated ROS production	⊖ ⊖ G/G	Elevated	A/A, A/G
rs10789038	PRKAA2	Impaired antioxidant activity	⊖ ⊖ A/A	Normal	A/A
rs2796498	PRKAA2	Impaired antioxidant activity	⊕ ⊕ G/G	Elevated	A/G, A/A
rs206812	XDH	Elevated ROS production	⊕ ⊖ A/G	Partially elevated	G/G

Oxidative Stress Profile

Antioxidant Genetics					
⊕ ⊕ Homozygous Mutant ⊕ ⊖ Heterozygous ⊖ ⊖ Homozygous Wild					
Test Name	Gene Name	Risk Association	Your Mutation	Your Risk	Reference
rs2073316	XDH	Elevated ROS production	⊕ ⊖ C/T	Normal	C/T, T/T
rs7310505	TXNRD1	Poor antioxidant activity	⊕ ⊖ A/C	Normal	C/C, A/C
rs1048943	CYP1A1	Elevated ROS production	⊕ ⊕ A/A	Normal	A/A
rs1548357	TXNRD2	Impaired mitochondrial oxygen radical scavenging activity	⊖ ⊖ T/T	Elevated	C/T, C/C
rs1695	GSTP1	Decreasing antioxidant activity leads to elevated oxidative stress	⊕ ⊖ A/G	Partially elevated	A/A
rs20417	COX-2	Elevated ROS production	⊕ ⊖ C/G	Normal	C/C, C/G
rs3877899	SELENOP	Impaired plasma selenium production	⊕ ⊖ C/T	Partially elevated	C/C
rs8190955	GSR	Increased oxidative stress in red blood cells	⊕ ⊕ C/C	Normal	C/C
rs916321	CYB5R3	Elevated ROS production	⊕ ⊖ A/G	Normal	G/G, A/G

Oxidative Stress Biomarkers					
Lipid Peroxidation	Current	Previous	Result	95th	Reference
11-β-Prostaglandin F2α (ug/g)	0.11		<div><div></div><div></div><div></div></div>	0.4	≤0.4
15(R)-Prostaglandin F2α (ug/g)	<0.05		<div><div></div><div></div><div></div></div>	0.22	≤0.22
8-iso-prostaglandin F2α (8-iso-PGF2α) (ug/g)	0.12		<div><div></div><div></div><div></div></div>	0.26	≤0.26
Glutathione 4-hydroxynonenal (GS-HNE) (ug/g)	0.23		<div><div></div><div></div><div></div></div>	2.5	≤2.5
Malondialdehyde (ug/g)	60.15		<div><div></div><div></div><div></div></div>	163.53	≤163.53
DNA Damage	Current	Previous	Result		Reference
8-Hydroxy-2-deoxyguanosine (ug/g)	0.73		<div><div></div><div></div><div></div></div>	4	≤4
8-Hydroxyguanine (ug/g)	10.87		<div><div></div><div></div><div></div></div>	49.4	≤49.4
8-Hydroxyguanosine (ug/g)	65.23		<div><div></div><div></div><div></div></div>	95.3	≤95.3
RNA Damage	Current	Previous	Result		Reference
8-Nitroguanine (ug/g)	22.01		<div><div></div><div></div><div></div></div>	107	≤107.47
8-Nitroguanosine (ug/g)	454.21		<div><div></div><div></div><div></div></div>	2608	≤2608.9

Oxidative Stress Profile

Oxidative Stress Biomarkers					
Protein Oxidation Products	Current	Previous	Result		Reference
			75th	95th	
3-Bromotyrosine (ug/g)	12.69		<div><div></div><div></div><div></div></div> <div>167.53349.6</div>		≤349.6
3-Chlorotyrosine (ug/g)	3.10		<div><div></div><div></div><div></div></div> <div>3.439.92</div>		≤9.92
Dityrosine (ug/g)	1.15		<div><div></div><div></div><div></div></div> <div>1.315</div>		≤5
Nitrotyrosine (ug/g)	29.99		<div><div></div><div></div><div></div></div> <div>91.32285.69</div>		≤285.69
Advanced Glycation Products	Current	Previous	Result		Reference
Nε-(carboxymethyl)lysine (CML) (ug/g)	15.31		<div><div></div><div></div><div></div></div> <div>015.870.3</div>		≤70.3
Nε-carboxyethyllysine (CEL) (ug/g)	28.61		<div><div></div><div></div><div></div></div> <div>019.591</div>		≤91

Risk and Limitations

This test has been developed and its performance characteristics determined and validated by Vibrant America and Vibrant Genomics LLC., CLIA certified laboratories. These assays have not been cleared or approved by the U.S. Food and Drug Administration. Vibrant Wellness provides additional contextual information on these tests and provides the report in a more descriptive fashion.

The Vibrant Oxidative Stress Profile does not demonstrate absolute positive and negative predictive values for any condition. Its clinical utility has not been fully established. Clinical history and current symptoms of the individual must be considered by the healthcare provider prior to any interventions. Test results should be used as one component of a healthcare provider's clinical assessment.

Vibrant has effective procedures in place to protect against technical and operational problems. However, such problems may still occur. Examples include failure to obtain the result for a specific test due to circumstances beyond Vibrant's control. Vibrant may re-test a sample to obtain these results but upon re-testing the results may still not be obtained. As with all medical laboratory testing, there is a small chance that the laboratory could report incorrect results. A tested individual may wish to pursue further testing to verify any results.

Genetic testing is helpful in analyzing risk to various diseases. However, it is important to note that Genetic risk determinants are neither necessary nor sufficient for the development of diseases. Environmental and lifestyle risk factors could also affect the risk of disease development. Results from genetic analysis should always be interpreted along with clinical findings on the individual. Genetic testing evaluates only for the particular genotypes indicated; it does not test for other genetic abnormalities found elsewhere in the genome. Different genetic variants can be tested by different genetic labs to evaluate the risk for a particular disease, depending on what is tested, genetic risk may not be comparable between labs. It should be realized that there are possible sources of error similar to any lab testing which include sample misidentification, trace contamination of PCR reactions, technical errors and rare genetic variants that may interfere with analysis.

Some individuals may feel anxious about getting their genetic test health results. If the potential user feels very anxious, such user should speak to his or her doctor or other health care professional prior to collection of a sample for testing. Users should consult with their doctor or other health care professional if they have any questions or concerns about the results of their test or their current state of health. Users of the test are also encouraged to discuss their test results with a genetic counselor, board-certified clinical molecular geneticist, or equivalent health care professional.

The information in this report is intended for educational purposes only. While every attempt has been made to provide current and accurate information, neither the author nor the publisher can be held accountable for any errors or omissions. Tested individuals may find their experience is not consistent with Vibrant's selected peer reviewed scientific research findings of relative improvement for study groups. The science in this area is still developing and many personal health factors affect diet and health. Since subjects in the scientific studies referenced in this report may have had personal health and other factors different from those of tested individuals, results from these studies may not be representative of the results experienced by tested individuals. Further, some recommendations may or may not be attainable, depending on the tested individual's physical ability or other personal health factors. A limitation of this testing is that many of these scientific studies may have been performed in selected populations only. The interpretations and recommendations are done in the context of these studies, but the results may or may not be relevant to tested individuals of different or mixed ethnicities.

Vibrant Wellness makes no claims as to the diagnostic or therapeutic use of its tests or other informational materials. Vibrant Wellness reports and other information do not constitute medical advice and are not a substitute for professional medical advice. Please consult your healthcare practitioner for questions regarding test results, or before beginning any course of medication, supplementation, or dietary changes.