



Gene Comprehensive Nutrigenomic Report

Accession Number: #####

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Specimen Received: ##/##/####

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Male



Do not make any decisions about your health solely based on the information contained in this report.
Always consult with a licensed and experienced health practitioner when you receive this report.

– 10 – Male

(-/-) No clinical abnormality

(+/-) Heterozygous result

(+/-) Heterozygous result (+/+) Homozygous result

| rsID | Gene | Genetic Result | Therapeutics Associated With Positive Result | Highly Recommended Therapeutics / Neurobiologix Formulas | Provider Discretion: As Needed Formula Recommendations | Lifestyle Recommendations | Laboratory Recommendations |
|-----------------------------------|-------------|----------------|--|---|--|---|--|
| Gastrointestinal | | | | | | | |
| Inflammation Environmental | | | | | | | |
| rs10156191 | AOC1 | +/- | Poor Ability To Break Down External Histamine | | GI Hist Support™ | Avoid Foods Containing Histamine | |
| rs11558538 | HNMT | -/- | | | | | |
| rs12995000 | HNMT | -/- | | | | | |
| rs492602 | FUT2 | +/+ | Probiotics Needed | Biotic Blend Pro™, Biotic Multi Blend Pro™, Probiotic Boost Daily™, Biotic Boost Chews For Kids™ | | | Consider Microbiome Testing If GI Inflammation Present |
| rs2248814 | NOS2 | +/- | Anti-Infectives, Beta Glucans | | Neuro-Immune Infection Control™ | | |
| rs2187668 | HLA DQA1 | -/- | High Risk of Gluten Based Issues | | | | |
| rs2858331 | HLA DQA2 | +/- | | | | | |
| Lactose Intolerance | | | | | | | |
| rs4988235 | MCM6 | +/+ | High incidence of lactose Intolerance | | | Avoid Lactose (milk products) | |
| Autophagy Consideration | | | | | | | |
| rs510432 | ATG5 | +/- | Curcumin, Lithium Orotate, D-Chiro-Inositol, Catechins, Resveratrol, Caffeine, 12 Hour Fasting | N.A.S. Enhancer™ (NRF2 Autophagy SOD Support), Metabolic Stimulator™ | | Calorie Restriction, 12-15 Hour Fasting | Routine Blood Sugar, Insulin and Hb A1c |
| rs10210302 | ATG16L1 | +/- | | | | | |
| rs2066845 | NOD2 CARD15 | -/- | Increase susceptibility to bacterial GI infections and Crohn's | | Neuro-Immune Infection Control™, N.A.S. Enhancer™ (NRF2 Autophagy SOD Support) | | |
| rs2241880 | ATG16L1 | +/- | | | | | |

Summary for Gastrointestinal

Highly Recommended Therapeutics / Neurobiologix Formulas

- **Biotic Blend Pro™, Biotic Multi Blend Pro™, Probiotic Boost Daily™, Biotic Boost Chews For Kids™**
- **N.A.S. Enhancer™ (NRF2|Autophagy|SOD Support), Metabolic Stimulator™**

Lifestyle Recommendations

- Avoid Foods Containing Histamine
- Avoid Lactose (milk products)
- Calorie Restriction, 12-15 Hour Fasting

Laboratory Recommendations

- Consider Microbiome Testing If GI Inflammation Present
- Routine Blood Sugar, Insulin and Hb A1c

Gene Information Key

| rsID | Gene | "-" variant | "+" variant |
|------------|-------------|----------------|----------------|
| rs10156191 | AOC1 | C | T |
| rs10210302 | ATG16L1 | C | T |
| rs11558538 | HNMT | C | T |
| rs12995000 | HNMT | C | T |
| rs2066845 | NOD2/CARD15 | G | C |
| rs2187668 | HLA-DQA1 | C | T |
| rs2241880 | ATG16L1 | A | G |
| rs2241880 | ATG16L1 | A | G |
| rs2248814 | NOS2 | G | A |
| rs2858331 | HLA-DQA2 | A | G |
| rs492602 | FUT2 | A | G |
| rs4988235 | MCM6 | G | A |
| rs510432 | ATG5 | C | T |

Definitions

| GASTROINTESTINAL | |
|---|--|
| MCM6 | A mutation in a DNA control region located in the MCM6 gene is associated with expression of the lactase gene. Individuals homozygous for this polymorphism are more likely to have hypolactasia, or lactose intolerance. |
| INFLAMMATORY | |
| This enzyme category has significant effects on the inflammatory state of a person's body. Polymorphisms in these specific enzymes will significantly increase the levels of inflammation in the body. By supplementing these enzyme deficiencies, the patient will effectively reduce inflammatory damage to the body. | |
| AOC1 | The SNP rs10156191 encodes a weaker form of the histamine degradation enzyme Amine Oxidase, Copper Containing 1 (AOC1). This mutation, Thr16Met, is predicted to produce an enzyme with less catalytic activity and associated higher levels of pro-inflammatory amines like histamine and putrescine. |
| ATG16L1 | The ATG16L1 gene encodes a protein that is a vital component of a protein complex necessary for the cellular phenomena known as autophagy. Autophagy is the process of degrading and cleaning of inert debris of the cell. Weakness in autophagy leads to abnormal accumulation of cellular "garbage" that will eventually affect the cellular function and lead to autophagy related disease states in including many neurological and immunological diseases, DM Type 2 and fatty liver disease. |
| ATG5 | Autophagy-related 5 protein (ATG5) is an important intracellular mediator of the autophagy response. ATG5 is involved in a wide range of "quality control" features inside the cell: autophagy vesicle formation, innate immune system signaling, consumption of damaged mitochondria, and apoptosis. Mutations in the ATG5 gene are associated with numerous neurological, immunological and endocrine syndromes. |
| FUT2 | Fucosyltransferase 2 (FUT2) is responsible for producing specific sugar groups that are secreted by the intestinal cells into the bowel to attract "good bacteria" . Polymorphisms in this gene produce "poor secreter" status. Lack of these sugars allows for gut dysbiosis and a higher risk of inflammatory bowel disease. |
| HLA-DQA1 | Major histocompatibility complex, DQ alpha 1 (HLA-DQA1) is a human gene responsible for a cell surface receptor essential to the function of the immune system. Patients with a polymorphism in this gene are at higher risk for auto-immune based inflammatory disease including Celiac disease, Crohn's, Ulcerative Colitis, and gluten sensitivity. |
| HLA-DQA2 | Major histocompatibility complex, DQ alpha 2 (HLA-DQA2) is a human gene responsible for a cell surface receptor essential to the function of the immune system. Patients with a polymorphism in this gene are at higher risk for auto-immune based inflammatory disease including Celiac disease, Crohn's, Ulcerative Colitis, and gluten sensitivity. |
| HNMT | The HNMT gene encodes the histamine degradative enzyme, histamine N-methyltransferase. HNMT, in contrast to AOC1, requires the methyl donor S-adenosylmethionine and a complete methylation pathway for normal function. Polymorphisms in HNMT gene expression or protein coding are predicted to prolong the pro-inflammatory effects of histamine signaling. |
| HNMT:Thr105Ile | The HNMT gene encodes the histamine degradative enzyme, histamine N-methyltransferase. HNMT, in contrast to AOC1, requires the methyl donor S-adenosylmethionine and a complete methylation pathway for normal function. Polymorphisms in HNMT gene expression or protein coding are predicted to prolong the pro-inflammatory effects of histamine signaling. |
| NOD2/CARD15 | NOD2 is a protein that integrates extracellular sensing of bacterial cell wall components with intracellular pro-inflammatory signaling to the nucleus. Mutations in the NOD2 gene are associated with poor ability to recognize abnormal bacterial pathogens in the intestinal tract and increased risk of intestinal dysbiosis. |
| NOS2 | Nitric Oxide Synthase 2 (NOS2) is responsible for producing nitric oxide, a biologic mediator used by the nervous system, immune system and in blood vessel function. Polymorphisms in this enzyme can cause reduced immune system function, exercise intolerance and fatigue. |

Disclaimers

METHODOLOGY AND LIMITATIONS:

Testing for genetic variation/mutation on listed genes was performed using ProFlex PCR and Real-Time PCR with TaqMan® allele-specific probes on the QuantStudio 12K Flex. All genetic testing is performed by GX Sciences, 4150 Freidrich Lane, Ste H, Austin, TX. 78744. This test will not detect all the known alleles that result in altered or inactive tested genes. This test does not account for all individual variations in the individual tested. Test results do not rule out the possibility that this individual could be a carrier of other mutations/variations not detected by this gene mutation/variation panel. Rare mutations surrounding these alleles may also affect our detection of genetic variations. Thus, the interpretation is given as a probability. Therefore, this genetic information shall be interpreted in conjunction with other clinical findings and familial history for the administration of specific nutrients. Patients should receive appropriate genetic counseling to explain the implications of these test results. Details of assay performance and algorithms leading to clinical recommendations are available upon request. The analytical and performance characteristics of this laboratory developed test (LDT) were determined by GX Sciences' laboratory pursuant to Clinical Laboratory Improvement Amendments (CLIA) requirements.

CLIA #: 45D2144988

DISCLAIMER:

This test was developed and its performance characteristics determined by GX Sciences. It has not been cleared or approved by the FDA. The laboratory is regulated under CLIA and qualified to perform high-complexity testing. This test is used for clinical purposes. It should not be regarded as investigational or for research. rsIDs for the alleles being tested were obtained from the dbSNP database (Build 142).

DISCLAIMER:

UND Result: If you have received the result Variant undetermined (UND) this indicates that we were not able to determine your carrier status based on your raw data. Please refer to the GX Sciences genetic knowledge database for more information: https://www.gxsciences.com/kb_results.asp

DISCLAIMER:

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DISCLAIMER:

These products are not approved by the Food and Drug Administration and are not intended to diagnose, treat, cure or prevent a disease. These recommendations are for report purposes only and an individual is not required to use such products. These are recommendations only and do not replace the advisement of your own healthcare practitioner.

GX Sciences SNP References

GASTROINTESTINAL SNP References

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