

GI Microbial Assay Plus



Who Should Have the GI-MAP Comprehensive Stool Analysis Done? YOU!

Almost every patient can benefit from a GI-MAP gut health assessment. Some patients are looking to achieve optimal health, while other patients have been chronically ill and frustrated without a diagnosis for years.

Some conditions that warrant testing are:

- Autoimmune diseases
- IBS/IBD
- Digestive complaints, diarrhea or constipation
- Brain fog
- Skin problems, like acne and psoriasis
- Mood disorders, depression, and anxiety
- Diabetes and weight loss issues



Pathogens

The GI-MAP includes pathogens (bacterial, parasitic, and viral) commonly known to cause intestinal gastroenteritis. It's important to note that not all individuals with positive findings for pathogens will present with symptoms. Many factors, including the health of the individual, the transient nature of some pathogens, and the presence and expression of virulence factors all contribute to an individual's symptoms.

Toxins are a type of virulence factor produced by certain pathogens. Since GI-MAP is a DNA-based test, results reflect the levels of pathogenic strains carrying the toxin genes, not the levels of any toxins that may be produced.

BACTERIAL PATHOGENS

- Campylobacter
- C. difficile Toxin A
- C. difficile Toxin B
- Enterohemorrhagic E. coli
- E. coli O157
- Enteroinvasive E. coli/Shigella
- Enterotoxigenic E. coli LT/ST
- Shiga-like Toxin E. coli stx1
- Shiga-like Toxin E. coli stx2
- Salmonella
- Vibro cholerae
- Yersinia enterocolitica

PARASITIC PATHOGENS

- Cryptosporidium
- Entamoeba histolytica
- Giardia

VIRAL PATHOGENS

- Adrenovirus 40/41
- Norovirus GI/II



H. pylori

Recent studies have shown that nearly 50% of the world's population may harbor H. pylori. And, although many carriers are asymptomatic, H. pylori is known to have a causative role in ulcers, chronic gastritis, and stomach cancer.

Additionally, in early phases of colonization, patients may experience hypochlorhydria followed by a change to hyper aciduria. Over time, additional H. pylori strains may colonize, including those with Virulence Factors and increased disease potential.

PYLORI & VIRULENCE FACTORS

Helicobacter pylori

- Virulence Factor, babA
- Virulence Factor, cagA
- Virulence Factor, dupA
- Virulence Factor, iceA
- Virulence Factor, oipA
- Virulence Factor, vacA
- Virulence Factor, virB
- Virulence Factor, virD



COMMENSAL/KEYSTONE BACTERIA

Trillions of microorganisms inhabit the human intestine to make up a complex ecosystem that plays an important role in human health.

Commensal bacteria extract nutrients and energy from our diets, maintain gut barrier function, produce vitamins (biotin and vitamin K), and protect against colonization by potential pathogens.

COMMENSAL BACTERIA

- Bacteroides fragilis
- Bifidobacterium spp.
- Enterococcus spp.
- Escherichia spp.
- Lactobacillus spp.
- Enterobacter spp.
- Akkermansia muciniphilia
- Faecalbacterium prausnitzii
- Roseburia spp.

BACTERIAL PHYLA

- Bacteroidetes
- Firmicutes
- Firmicutes/Bacteroidetes Ratio



OPPORTUNISTIC/OVERGROWTH MICROBES

Many bacteria measured on the GI-MAP are considered opportunistic pathogens, as they only cause disease and illness in some individuals, particularly the immune-compromised. Many individuals come into contact with opportunistic bacteria and experience no symptoms. Most sources consider these microbes to be normal in the stool. However, they can cause gastroenteritis and inflammation at high levels in vulnerable patients. Symptoms may include diarrhea, loose stools, abdominal pain, or even constipation.

Overgrowth and excessive colonization by opportunistic bacteria may occur when the commensal bacteria are impaired by poor diet, antibiotic use, parasitic infection, or a weakened immune system. When intestinal permeability is present (see zonulin), these microbes could escape the lumen of the gut and infect extraintestinal sites.

OPPORTUNISTIC/OVERGROWTH MICROBES

- Bicillus spp.
- Enterococcus faecalis
- Enterococcus faecium
- Morganella spp.
- Pseudomonas spp.
- Pseudomonas aeruginosa
- Staphylococcus spp.
- Staphylococcus aureus
- Streptococcus spp.

COMMENSAL OVERGROWTH MICROBES

- Desulfovibrio spp.
- Methanobacteriaceae (family)

INFLAMMATORY & AUTOIMMUNE-RELATED BACTERIA

- Citrobacter spp.
- Citrobacter freundii
- Klebsiella spp.
- Klebsiella pneumoniae
- M. avium subsp. paratuberculosis
- Proteus spp.
- Proteus mirabilis

COMMENSAL INFLAMMATORY & AUTOIMMINE-RELATED BACTERIA

- Enterobacter spp.
- Escherichia spp.
- Fusobacterium spp.
- Prevotella copri



FUNGI/YEAST

Fungal organisms are commonly found in the human digestive tract, but fungal overgrowth can cause illness in susceptible individuals. Fungal growth may be localized in the body. For instance, Candida spp. may be high in the large intestine but normal in the small intestine, and vice versa. In a patient with suspected fungal overgrowth, additional tests may be necessary to understand the complete picture of fungal overgrowth. Urinary D-arabinitol or antibodies to Candida are sometimes used.

VIRUSES

FUNGI/YEAST

- Candida spp.
- Candida albicans
- Geotricum spp.
- Microsporidia spp.
- Rhodoturula spp.

VIRUSES

- Cytomegalovirus
- Epstein Bar Virus



PARASITES

A parasite is an organism that lives and feeds on a host organism at the expense of the host. The GI-MAP tests for pathogenic parasites and protozoa (some of which are non-pathogenic) most commonly occurring in the GI tract. Sources of exposure should be identified and eliminated to prevent reinfection.

INTESTINAL HEALTH MARKERS

PARASITES

PROTOZOA

- Blastocystis hominis
- Chilomastix mesnelli
- Cyclospora spp.
- Dientamoeba fragilis
- Endolimax nana
- Entamoeba coli
- Pentatrichomonas hominis

WORMS

- Ancyclostroma duodenale
- Ascaris lumbricoides
- Necator americanis
- Trichuris trichiura
- Taenia spp.

INTESTINAL HEALTH MARKERS

DIGESTION

- Steatocrit
- Elastase-1

GI MARKERS

- β-Glucuronidase
- Occult Blood FIT

IMMUNE RESPONSE

- Secretory IgA
- Anti-gliadin IgA
- Eosinophil Activation Protein (EDN)

INFLAMMATION

Calprotectin

ADD-ON TESTS

- Zonulin
- Gluten Peptide
- Universal Antibiotic Resistance Genes



H. PYLORI ANTIBIOTIC RESISTANCE GENES

The GI-MAP includes results for detection of antibiotic resistance genes in the microbiome. If an antibiotic resistance gene is present, then that class of antibiotics is designated POSITIVE for antibiotic resistance. A positive result for the presence of resistance genes for a given antibiotic indicates that the antibiotic is not an ideal choice for an antibiotic protocol.

Antibiotic resistance genes apply to all of the microorganisms found in the fecal sample. Since microbes can rapidly share DNA under stress, the presence of antibiotic resistance in any organism is reason enough to avoid that drug class.

Phenotypes | HELOBACTER

- Amoxicillen
- Clarithromycin
- Fluroquinolines
- Tetracycline



UNIVERSAL ANTIBIOTIC RESISTANCE GENES

The Universal Antibiotic Resistance (AR) Genes panel can be ordered as an add-on to the GI-MAP or to the GI-Pathogens panel. It detects the presence of 55 genetic elements associated with resistance to 10 different classes of antibiotics.

This AR Genes add-on panel detects genes associated with resistance to the most commonly prescribed antibiotics for gastrointestinal infections. The mobile genetic elements in the panel can be found in a variety of different microbes. The presence of these genes in a bacterial population have been associated with moderate to high levels of antibiotic resistance in human gastrointestinal infections.

UNIVERSAL ANTIBIOTIC RESISTANCE GENES

- β-lactamase
- Fluoroquinolones
- Vancomycin
- Macrolides
- Ciprofloxacin
- 5-Nitroimidazoles (non-Helicobacter pylori)
- Trimethoprim
- Sulfonamides
- Methicillin
- Chloramphenicol



Why Is Quantification Using qPCR Technology So Important?

Unlike other comprehensive stool tests on the market, the GI-MAP can provide practitioners with truly quantitative results. qPCR offers a much more accurate way to detect and quantify clinically-relevant organisms than standard PCR, culture, microscopy, or DNA sequencing-based methods. Accurately assessing how much of an organism's DNA is present in a patient's stool sample is essential for helping practitioners to determine the clinical significance of pathogenic organisms and dysbiosis patterns.

qPCR's Reliability, Reproducibility, and Use in Clinical Research

Although qPCR is becoming more commonplace in in-vitro diagnostics (IVD), we are the only laboratory in the United States exclusively using qPCR technology for advanced comprehensive stool testing. This technology is used routinely in clinical and academic research because it provides highly-accurate quantification, as well as high levels of sensitivity and specificity. Standard PCR technology doesn't offer the same level of sensitivity, or the ability to express precise numerical results.

The GI-MAP also provides consistently reproducible results. Reproducibility is of crucial importance to the practitioners and patients that rely on the efficacy of the GI-MAP. To achieve it, we perform rigorous quality control, and have validated all molecular target quantification assays to meet or exceed FDA standards.

GI-MAP Allows for the Personalized Treatment Plans and Informative Retests

The GI-MAP's accuracy and reliability allows practitioners to create personalized treatment protocols to address gut dysfunction based on which infections are urgent, which areas of the gut are already optimized, and which areas should be addressed after an infection is resolved.

Additionally, the quantification offers a remarkable ability to see how treatment modalities are working because a retest after treatment can show whether a parasite has resolved, dysbiosis has improved, and more.



Call Today To Order or Schedule An Appointment

920-737-1625

Cost For Labs & Results \$585

For A Detailed Discussion & Treatment Plan, Schedule An Appointment With A Provider

You Do Not Need To Be An Established Patient To Order a GI Map