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**GI-MAP™** DNA Stool Analysis

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## Pathogens

### Bacterial Pathogens

|                                      | Result          | Expected |
|--------------------------------------|-----------------|----------|
| <i>Campylobacter</i>                 | <b>Negative</b> | Negative |
| <i>C. difficile</i> , Toxin A        | <b>Negative</b> | Negative |
| <i>C. difficile</i> , Toxin B        | <b>Negative</b> | Negative |
| <i>E. coli</i> O157                  | <b>Negative</b> | Negative |
| Enterotoxigenic <i>E. coli</i> LT    | <b>Negative</b> | Negative |
| Enterotoxigenic <i>E. coli</i> ST    | <b>Negative</b> | Negative |
| Shiga-like Toxin <i>E. coli</i> stx1 | <b>Negative</b> | Negative |
| Shiga-like Toxin <i>E. coli</i> stx2 | <b>Negative</b> | Negative |
| <i>Shigella</i>                      | <b>Negative</b> | Negative |
| <i>Salmonella</i>                    | <b>Negative</b> | Negative |
| <i>Vibrio cholerae</i>               | <b>Negative</b> | Negative |
| <i>Yersinia enterocolitica</i>       | <b>Negative</b> | Negative |

### Viral Pathogens

|                      | Result          | Expected |
|----------------------|-----------------|----------|
| Adenovirus 40/41     | <b>Negative</b> | Negative |
| <i>Norovirus</i> GI  | <b>Negative</b> | Negative |
| <i>Norovirus</i> GII | <b>Negative</b> | Negative |
| <i>Rotavirus</i> A   | <b>Negative</b> | Negative |

### Parasitic Pathogens

|                              | Result          | Expected |
|------------------------------|-----------------|----------|
| <i>Cryptosporidium</i>       | <b>Negative</b> | Negative |
| <i>Entamoeba histolytica</i> | <b>Negative</b> | Negative |
| <i>Giardia</i>               | <b>Negative</b> | Negative |

## H. pylori

|                            |                 |             | Range CFU/g |
|----------------------------|-----------------|-------------|-------------|
| <i>Helicobacter pylori</i> | <b>1.1e3</b>    | <b>High</b> | <1.0e3      |
| Virulence Factor, cagA     | <b>Negative</b> |             | Negative    |
| Virulence Factor, vacA     | <b>Positive</b> |             | Negative    |

## Normal Bacterial Flora

|                             |               |            | Range org/g     |
|-----------------------------|---------------|------------|-----------------|
| <i>Bacteroides fragilis</i> | <b>1.3e7</b>  |            | 5.00e5 - 3.00e9 |
| <i>Bifidobacterium spp.</i> | <b>1.0e10</b> |            | >1.10e8         |
| <i>Enterococcus spp.</i>    | <b>2.9e3</b>  | <b>Low</b> | 1.2e4 - 3.1e6   |
| <i>Escherichia spp.</i>     | <b>3.5e6</b>  |            | 1.00e4 - 7.60e7 |
| <i>Lactobacillus spp.</i>   | <b>5.8e7</b>  |            | 1.0e6 - 5.80e9  |

## Opportunistic Bacteria

|                              |               |             | Range CFU/g |
|------------------------------|---------------|-------------|-------------|
| <i>Citrobacter freundii</i>  | <b>2.0e4</b>  | <b>High</b> | <1.0e4      |
| <i>Klebsiella pneumoniae</i> | <b>&lt;dl</b> |             | <7.2e3      |
| <i>Proteus spp.</i>          | <b>7.4e3</b>  | <b>High</b> | <6.2e3      |
| <i>Proteus mirabilis</i>     | <b>&lt;dl</b> |             | <1.0e3      |

## Additional Dysbiotic/Overgrowth Bacteria

|                               |               | Range org/g |
|-------------------------------|---------------|-------------|
| <i>Morganella spp.</i>        | <b>&lt;dl</b> | <1.0e3      |
| <i>Pseudomonas spp.</i>       | <b>&lt;dl</b> | <2.5e3      |
| <i>Pseudomonas aeruginosa</i> | <b>&lt;dl</b> | <1.0e3      |
| <i>Staphylococcus spp.</i>    | <b>&lt;dl</b> | <1.0e4      |
| <i>Streptococcus spp.</i>     | <b>&lt;dl</b> | <1.0e3      |

## Parasites

|                                 |                 | Range |
|---------------------------------|-----------------|-------|
| <i>Blastocystis hominis</i>     | <b>Negative</b> | Neg   |
| <i>Dientamoeba fragilis</i>     | <b>Negative</b> | Neg   |
| <i>Endolimax nana</i>           | <b>Negative</b> | Neg   |
| <i>Entamoeba coli</i>           | <b>Negative</b> | Neg   |
| <i>Chilomastix mesnelli</i>     | <b>Negative</b> | Neg   |
| <i>Cyclospora spp.</i>          | <b>Negative</b> | Neg   |
| <i>Pentatrichomonas hominis</i> | <b>Positive</b> | Neg   |

## Fungi/Yeast

|                            |                 | Range  |
|----------------------------|-----------------|--------|
| <i>Candida albicans</i>    | <b>&lt;dl</b>   | <5.0e3 |
| <i>Candida spp.</i>        | <b>Low</b>      | Neg    |
| <i>Geotrichum spp.</i>     | <b>Negative</b> | Neg    |
| <i>Microsporidium spp.</i> | <b>Mod</b>      | Neg    |
| <i>Trichosporon spp.</i>   | <b>Negative</b> | Neg    |

## Additional Tests

|                    | Result          |             | Range ug/g |
|--------------------|-----------------|-------------|------------|
| Secretory IgA      | <b>1570</b>     |             | 510 - 2010 |
| Anti-gliadin IgA   | <b>298</b>      | <b>High</b> | 0 - 100    |
| Elastase-1         | <b>421</b>      |             | >200       |
| Calprotectin       | <b>84</b>       | <b>High</b> | <50        |
| Fecal Occult Blood | <b>Negative</b> |             | Negative   |