

# GI Effects Stool Profiles

## **Interpretive Guide**

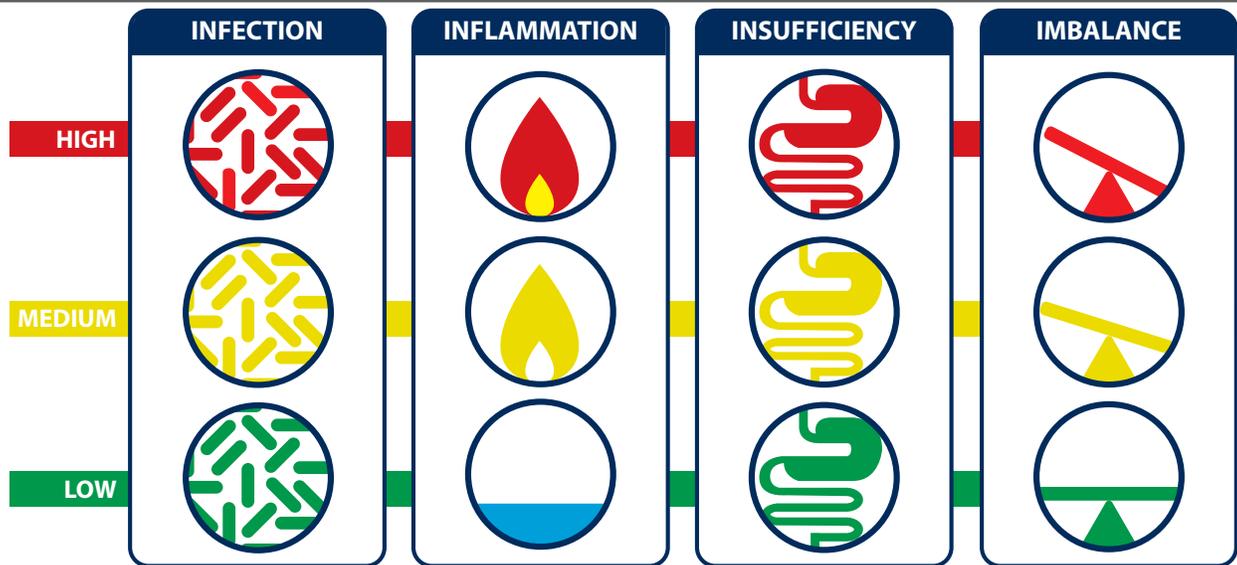


# GI Effects Interpretive Guide

The **GI Effects Interpretive Guide** has been created to provide a high-level approach to the GI Effects profile, biomarker interpretation, and therapeutic considerations. It is divided into three major sections: an overview of the GI Effects Interpretation At-a-Glance pages; a more in-depth review of the biomarkers comprising each of the Four Functional Pillars; and overall pattern recognition.

## Interpretation At-a-Glance Overview

Using evidence-based rules and weighted algorithms, the **Interpretation At-a-Glance** section on the first page of the GI Effects report synthesizes patient test results into key functional areas of clinical significance and provides a directional indication of potential next steps in patient management.



Four Functional Pillars Biomarker Map			
Infection Box	Inflammation Box	Insufficiency Box	Imbalance Box
<ul style="list-style-type: none"> <li>Any pathogenic organism present</li> </ul>	<ul style="list-style-type: none"> <li>Calprotectin</li> <li>Eosinophil Protein X (EPX)</li> <li>Fecal Secretory IgA</li> <li>Fecal Occult Blood</li> <li>Fecal Lactoferrin (if ordered)</li> </ul>	<ul style="list-style-type: none"> <li>Pancreatic Elastase 1 (PE1)</li> <li>Total Fecal Fats</li> <li>Products of Protein Breakdown (Total)</li> </ul>	<ul style="list-style-type: none"> <li>n-Butyrate</li> <li>Total SCFA</li> <li>Beta-glucuronidase</li> <li>Beneficial Bacteria <i>Lactobacillus</i>, <i>Bifidobacterium</i>, <i>E. coli</i> (PCR)</li> <li>Any potential pathogen (PP)</li> </ul>

## Four Functional Pillars

In this section, pertinent biomarkers have been grouped into four clinically actionable areas: Infection, Inflammation, Insufficiency, and Imbalance. The four functional pillars utilize a proprietary algorithm to evaluate key clinical markers in each of these four functional areas. The algorithm takes into account the level of each individual biomarker and its degree of clinical impact. As a result, an overall score of high, medium, or low is provided for each functional pillar. The score is represented by color-coded icons and informational graphics.

The specific biomarkers of concern that are utilized to establish the results for each functional pillar are listed above.

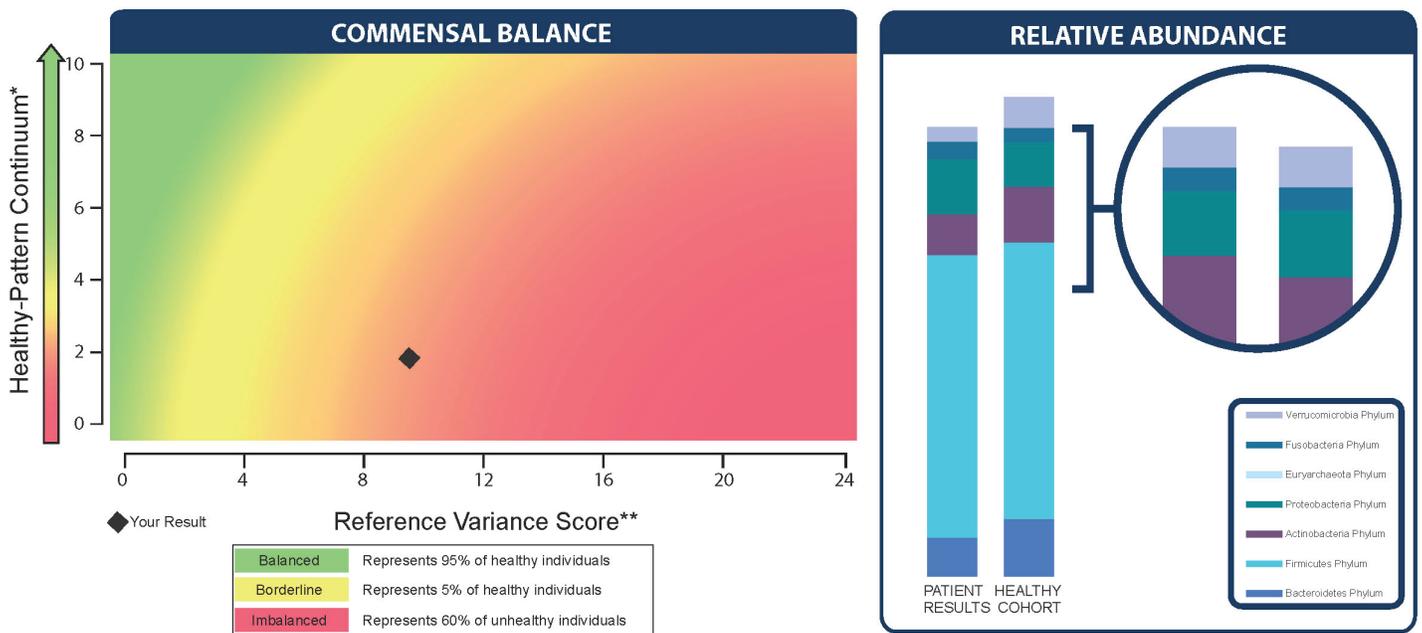
## Commensal Balance and Relative Abundance

The Commensal Balance infographic has been designed to provide a more precise view of an individual patient's commensal bacteria (PCR) results relative to a healthy cohort. It is a composite of two measures:

- The **Healthy-Pattern Continuum** (formerly known as the Diversity Association Index) is a progressive ranking scale based on a Genova proprietary algorithm that differentiates healthy and unhealthy commensal patterns. This algorithm is applied to an individual patient's GI Effects commensal bacteria (PCR) findings, and produces a numeric result ranging from 0 to 10 and is denoted by the 'y' axis of the Commensal Balance infographic.
- The **Reference Variance Score** reflects the total number of an individual patient's commensal bacteria (PCR) results that are out of reference range. This number ranges from zero to 24, and is denoted by the 'x' axis of the Commensal Balance infographic.

The patient's result on the Commensal Balance infographic is denoted by a black diamond against a color-coded gradient (green, yellow and red). The position of the patient's result against this background provides an At-a-Glance comparison of the patient's current commensal findings against those seen in healthy and diseased cohorts. Green suggests balanced commensal health status, yellow borderline, and red imbalanced.

The **Relative Abundance (RA)** graphic represents the proportional levels of selected phyla in an individual's microbiome and is represented relative to similar measures derived from a healthy cohort of individuals.



\* A progressive ranking scale based on a Genova proprietary algorithm that differentiates healthy and unhealthy commensal patterns.

\*\*The total number of Commensal Bacteria (PCR) that are out of reference range for this individual

## Commensal and Biomarker Clinical Associations

To create its Clinical Associations charts, Genova utilized the GI Effects test-results database which allowed comparison of commensal and biomarker results in patients with self-reported clinical conditions (IBD, Metabolic Syndrome, Chronic Fatigue, Autoimmune dysfunction, Type 2 Diabetes, High Blood Pressure, and Mood Disorders, and ROME III criteria (IBS)) to those found in the healthy cohort.

Differences between the healthy cohort and individuals with clinical conditions are denoted by the arrows in the Clinical Associations charts. In addition, these charts feature patient results marked as high (H) or low (L) compared to the reference range for each commensal bacteria or biomarker.

### Clinical Associations Interpretative Key

Statistically significant differences between patients with a clinical condition and the healthy cohort are denoted in the chart with either up or down arrows, or in some instances, a combination of both.

Interpretation At-a-Glance									
Commensal Bacteria	Patient Results Out of Reference Range	Genova Diagnostics Commensal Bacteria Clinical Associations*							
		IBS	IBD	Metabolic Syndrome	Chronic Fatigue	Auto-immune	Type 2 Diabetes	High Blood Pressure	Mood Disorders
<b>Bacteroidetes Phylum</b>									
<i>Bacteroides-Prevotella</i> group		↑	↑	↑	↑	↑	↑	↑	↑
<i>Bacteroides vulgatus</i>		↑			↑	↑		↑	↑
<i>Barnesiella</i> spp.									
<i>Odoribacter</i> spp.									
<i>Prevotella</i> spp.	H	↑		↑	↑	↑		↑	↑
<b>Firmicutes Phylum</b>									
<i>Anaerotruncus colihominis</i>	H	↑	↑	↑	↑	↑	↑	↑	↑
<i>Butyrivibrio crossotus</i>									
<i>Clostridium</i> spp.									
<i>Coprococcus eutactus</i>		↑			↑	↑		↑	↑
<i>Faecalibacterium prausnitzii</i>	H	↑				↑			↑
<i>Lactobacillus</i> spp.									
<i>Pseudoflavonifractor</i> spp.	H	↑	↑	↑	↑	↑	↑	↑	↑
<i>Roseburia</i> spp.			↓						
<i>Ruminococcus</i> spp.	L	↓↑	↓	↓	↓	↓↑	↓↑	↓↑	↓↑
<i>Veillonella</i> spp.		↑	↑	↑	↑	↑	↑		↑
<b>Actinobacteria Phylum</b>									
<i>Bifidobacterium</i> spp.	H								
<i>Bifidobacterium longum</i>									
<i>Collinsella aerofaciens</i>	L	↓↑	↓↑	↓	↓↑	↓↑	↓↑	↓↑	↓↑
<b>Proteobacteria Phylum</b>									
<i>Desulfovibrio piger</i>									↑
<i>Escherichia coli</i>	H	↑	↑	↑	↑	↑	↑	↑	↑
<i>Oxalobacter formigenes</i>	H	↑		↑	↑				↑
<b>Euryarchaeota Phylum</b>									
<i>Methanobrevibacter smithii</i>		↑				↑			↑
<b>Fusobacteria Phylum</b>									
<i>Fusobacterium</i> spp.		↑	↑	↑	↑	↑	↑	↑	↑
<b>Verrucomicrobia Phylum</b>									
<i>Akkermansia muciniphila</i>		↓	↓	↓	↓	↓	↓	↓	↓

\*Information derived from GDx results data comparing a healthy cohort to various clinical condition cohorts. The chart above showing a comparison of patient results to clinical conditions is meant for informational purposes only; it is not diagnostic, nor does it imply that the patient has a specific clinical diagnosis or condition.

The arrows indicate Genova's clinical condition cohort test results falling below ↓ or above ↑ the reference range that is greater than that of Genova's healthy cohort.

↑↓ Indicates Genova's clinical condition cohort test results falling below and above the reference range that are greater than that of Genova's healthy cohort.

Cells with bolded arrows indicate Genova's clinical condition cohort had more test results falling above versus below ↓↑ or more below versus above ↑↓ the reference range compared to that of Genova's healthy cohort.

## Interpretation At-a-Glance

Biomarker	Patient Results Out of Reference Range	Genova Diagnostics Biomarker Clinical Associations*							
		IBS	IBD	Metabolic Syndrome	Chronic Fatigue	Auto-immune	Type 2 Diabetes	High Blood Pressure	Mood Disorders
Pancreatic Elastase		↓	↓	↓	↓	↓	↓	↓	↓
Products of Protein Breakdown (Total)							↓↑		
Fecal Fat (Total)		↑		↑	↑	↑	↓↑	↑	↑
Triglycerides	<b>H</b>	↑			↑	↑	↑	↑	↑
Long Chain Fatty Acids		↑			↑	↑	↓↑	↑	↑
Cholesterol							↓↑	↑	
Phospholipids		↑	↑	↑	↑	↑	↑	↑	↑
Calprotectin			↑					↑	
Eosinophil Protein X (EPX)			↑						
Fecal sIgA	<b>H</b>	↑	↑	↑	↑	↑	↑	↑	↑
Short Chain Fatty Acids (SCFA) (Total)					↓	↓			
n-Butyrate Concentration				↓					
n-Butyrate %									
Acetate%					↓↑		↓↑		
Propionate %				↑			↑	↑	
Beta-glucuronidase						↓↑			↓↑

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Cells with bolded arrows indicate Genova's clinical condition cohort had more test results falling above versus below ↑↑ or more below versus above ↓↓ the reference range compared to that of Genova's healthy cohort.

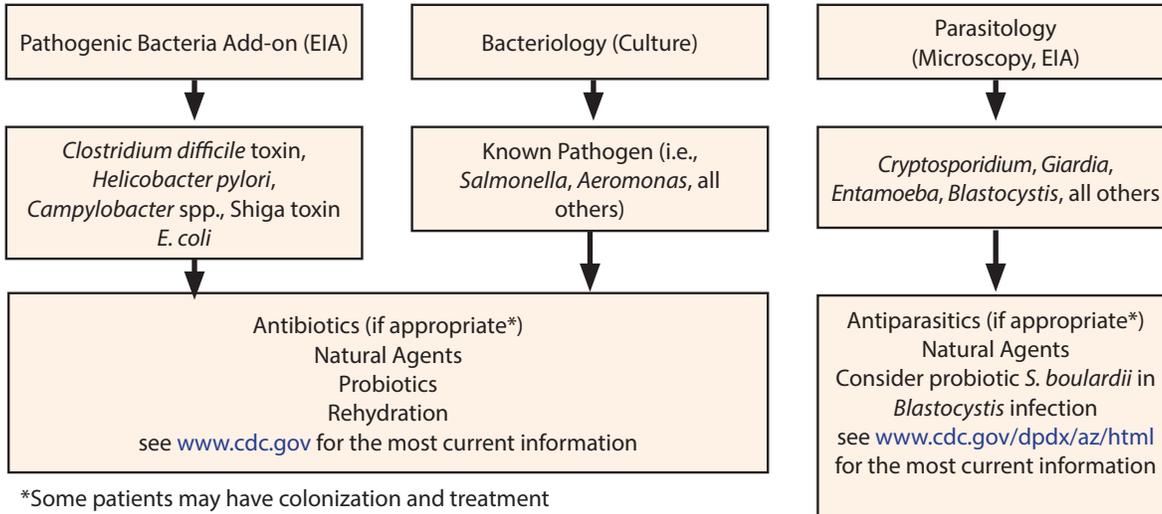
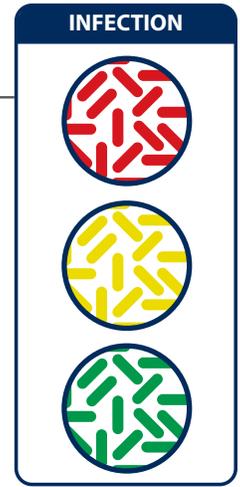
Displayed in an expanded Interpretation At-a-Glance section, the new report enhancements provide informative data derived from Genova Diagnostics patient results archive. All results were obtained with the same Genova Diagnostics technology platform(s); this allows for an equivalent or 'like to like' comparison of an individual patient's results to various clinical conditions.

**The remainder of this document will review biomarker interpretation and treatment considerations for each of the Four Pillars in more detail.**

# FOUR FUNCTIONAL PILLARS BIOMARKER DETAIL

## INFECTION

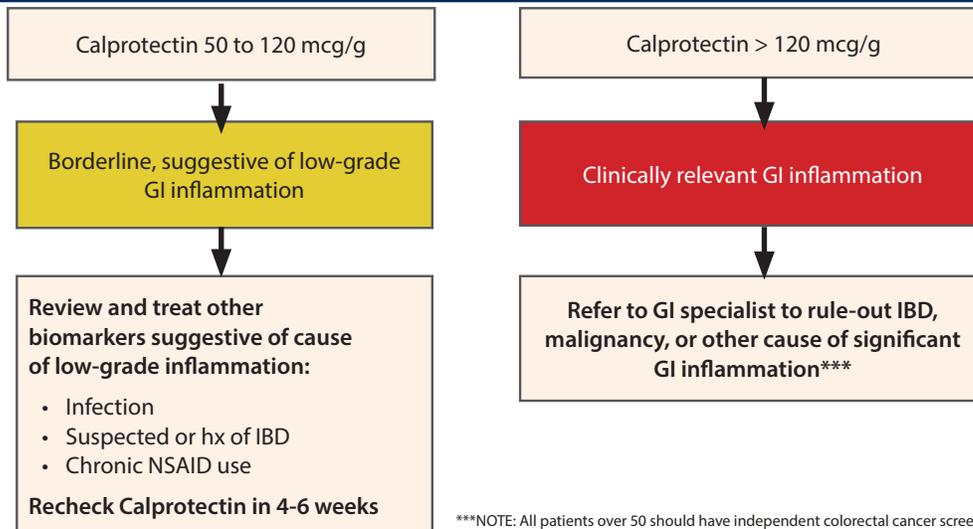
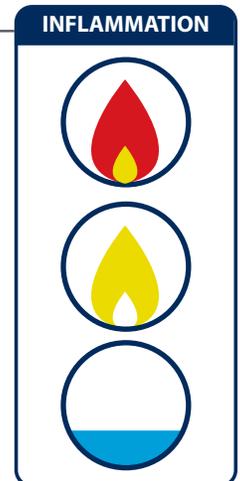
This pillar is where common infectious microorganisms are reported and includes **pathogenic bacteria and intestinal parasites**.



\*Some patients may have colonization and treatment decisions must be considered on an individual basis.

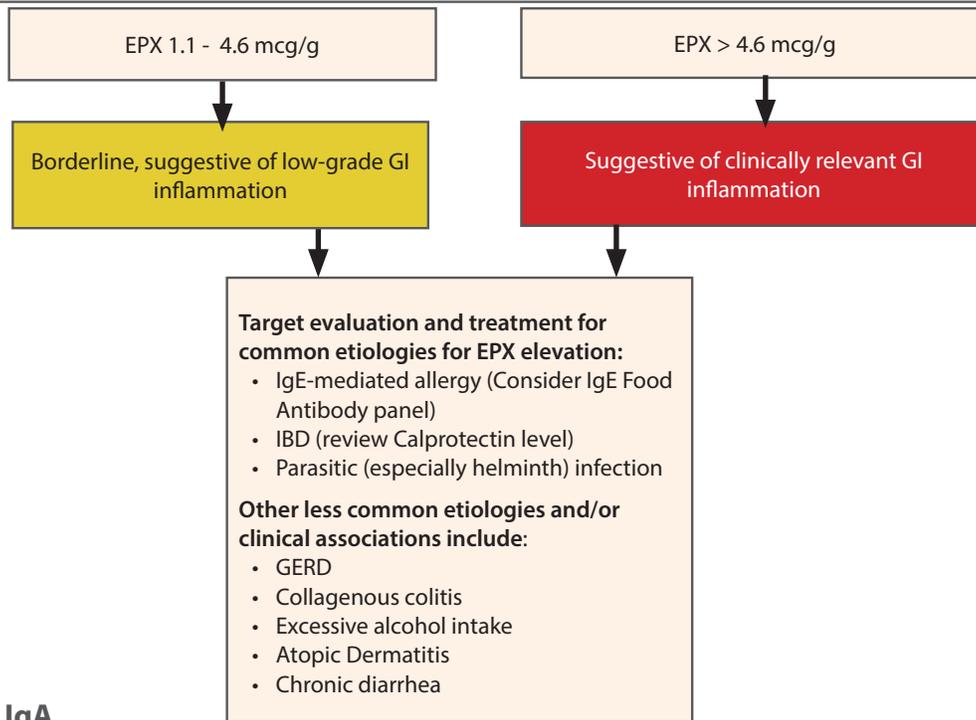
## INFLAMMATION

This pillar is where biomarkers that indicate inflammatory changes in the GI tract are reported. Biomarkers of GI inflammation and immunology provide information about the GI tract's interactions with, and responses to, the outside world. They indicate how well the GI tract is maintaining its role as a barrier, as well as whether the GI tract is undergoing pathological responses to external or internal challenges. The biomarkers are **Calprotectin**, a marker of neutrophil-driven inflammation; **Eosinophil Protein X (EPX)**, a marker of eosinophil-driven inflammation and allergic response; **Fecal Secretory IgA**, a marker of gut secretory immunity and barrier function; and **Fecal Occult Blood**.

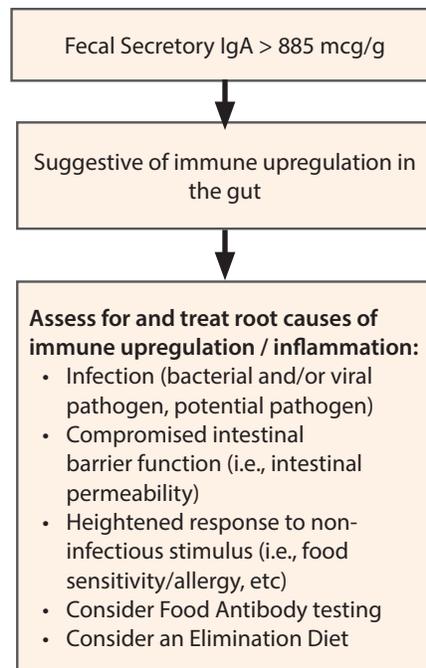


\*\*\*NOTE: All patients over 50 should have independent colorectal cancer screening per USPSTF recommendations. Although a normal fecal calprotectin does have a high negative predictive value for colorectal cancer, no single biomarker on the GI Effects panel is intended to exclusively rule out or to diagnose cancer.

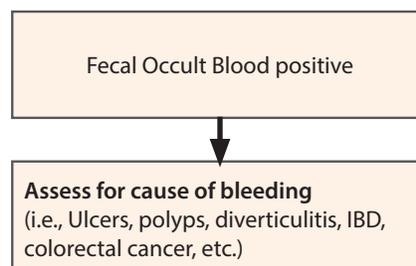
## EPX



## Fecal Secretory IgA



## Fecal Occult Blood



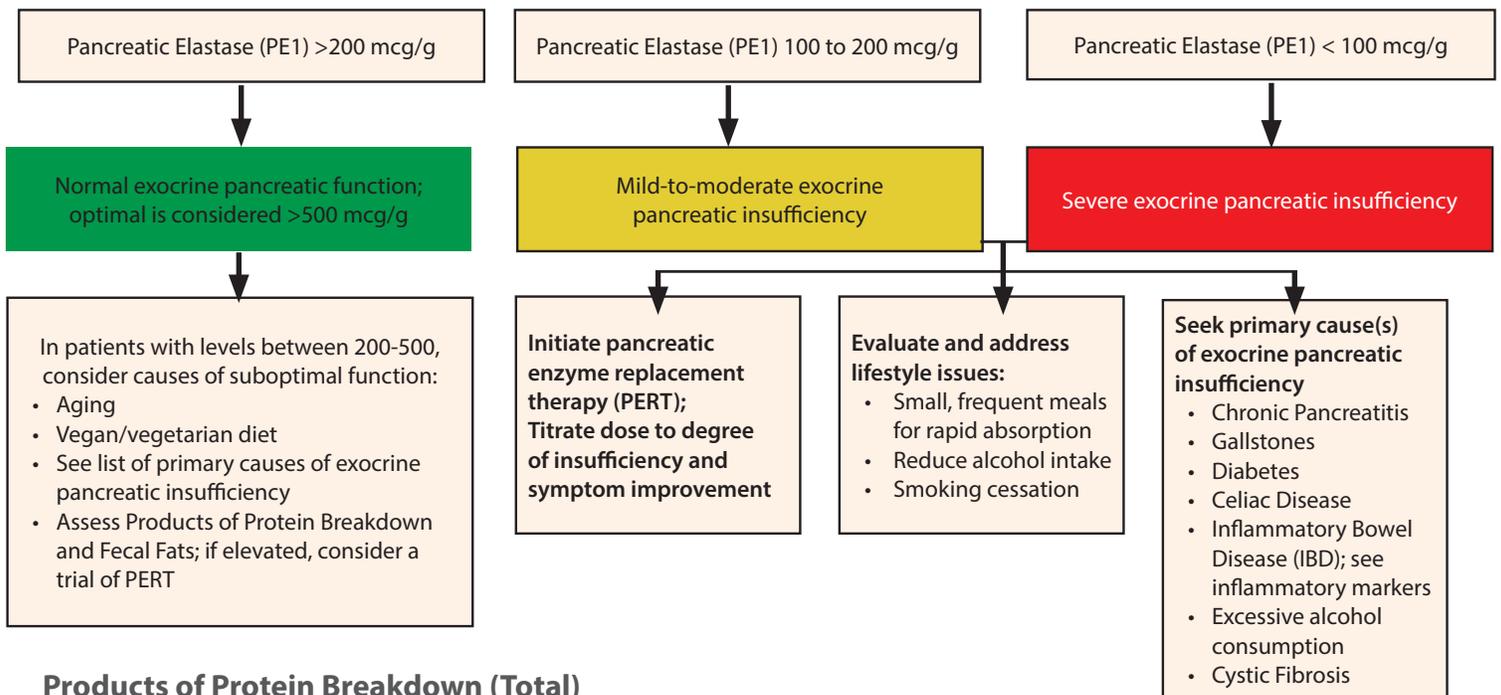
## INSUFFICIENCY

This pillar is where biomarkers that indicate digestive function are noted. Biomarkers of digestion and absorption provide information about nutrient breakdown and entry into the circulation. They ultimately indicate how well the GI tract is performing its basic digestive functions. The biomarkers are **Pancreatic Elastase 1**, a marker of exocrine pancreatic function; **Products of Protein Breakdown**, markers of undigested protein reaching the colon; and **Fecal Fat (Total)**, a marker of fat breakdown.

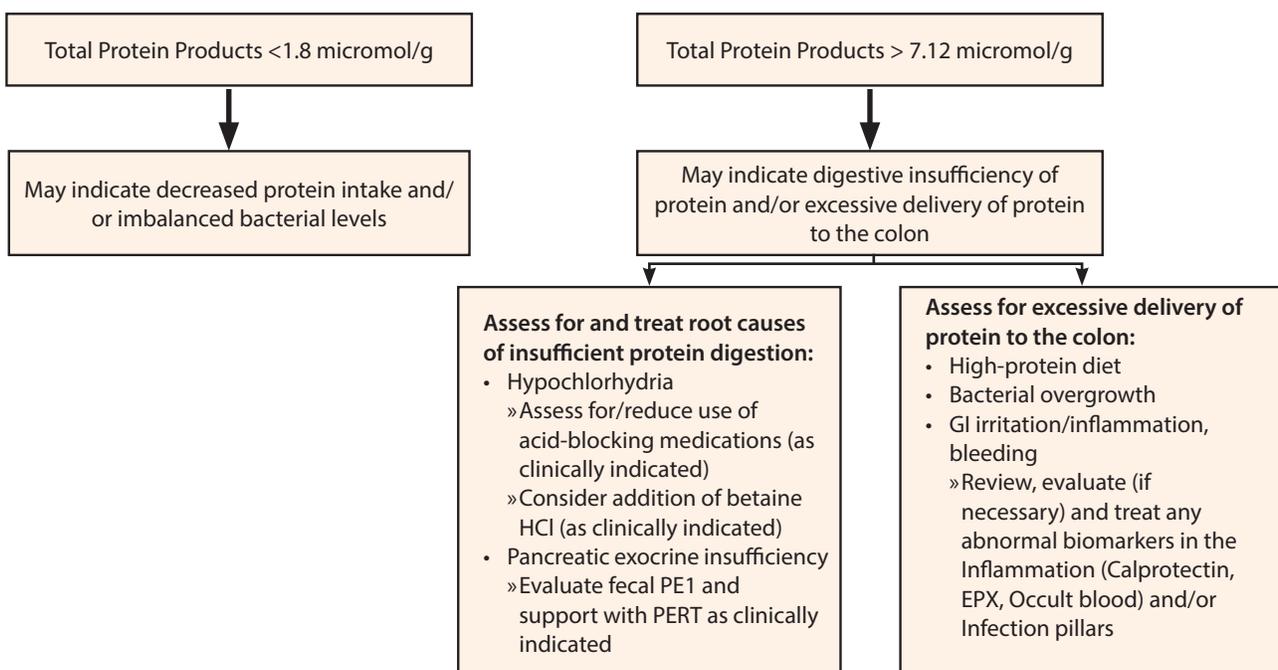
## INSUFFICIENCY



### Pancreatic Elastase 1 (PE1)

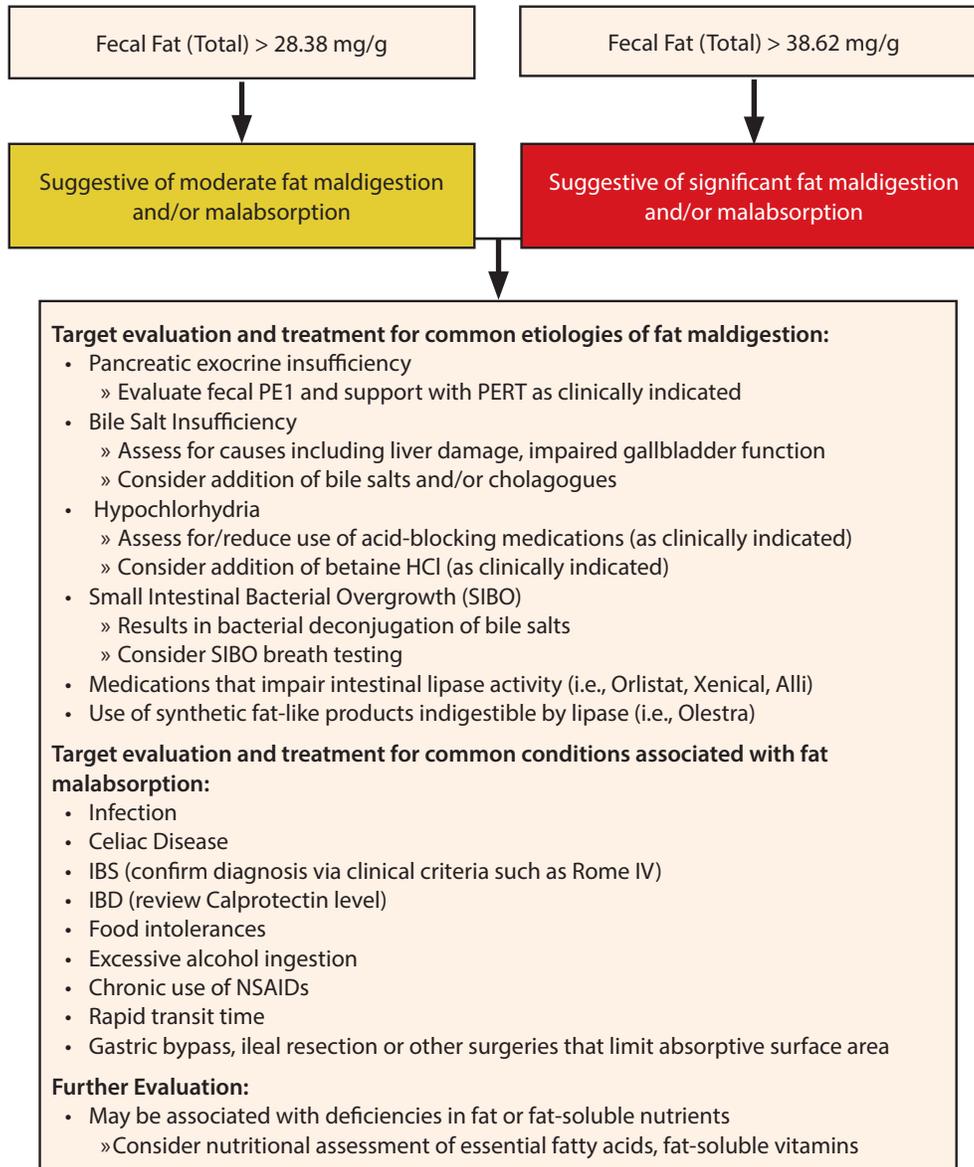


### Products of Protein Breakdown (Total)



## Fecal Fat (Total)

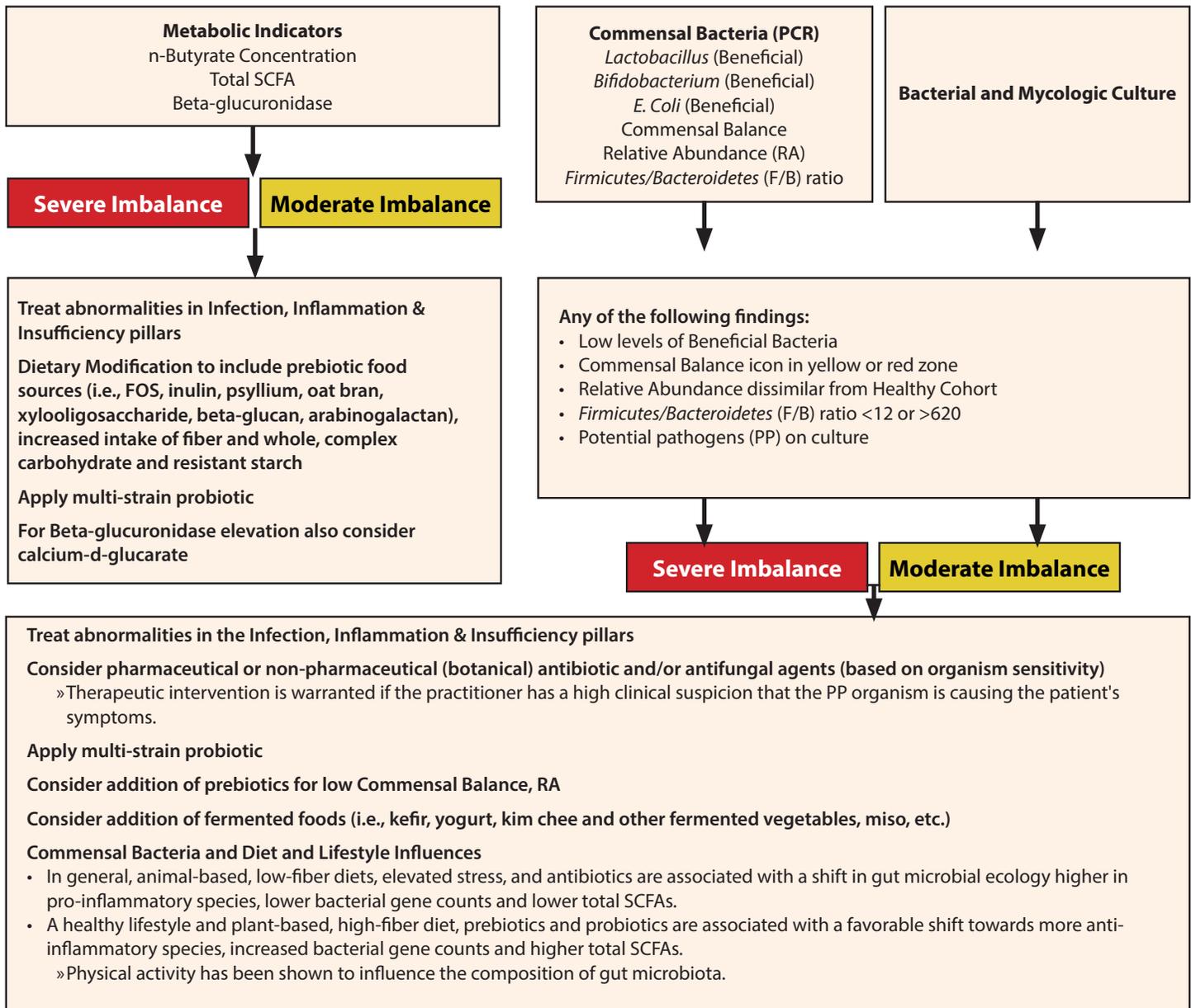
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# IMBALANCE

## IMBALANCE

This pillar is where imbalances in the population of GI organisms (i.e., dysbiosis) is noted. Biomarkers of the GI Microbiome provide information about the health, function, and abundance of the trillions of microbial cells in the GI tract. They indicate how well the microbiome is performing its shared metabolic functions with the human host. Abnormal results in this pillar may be associated with a large number of conditions and symptoms. The biomarkers which assess gut microbial imbalance are **Metabolic Indicators**, which demonstrate specific and vital metabolic functions performed by the microbiota; **Commensal Bacteria**, a PCR evaluation of 24 key bacterial groups/species; and **Bacterial and Mycologic Culture**, which identifies potentially pathogenic (PP) organisms.



## OVERALL PATTERN RECOGNITION

One of the benefits of ordering a comprehensive stool panel like GI Effects is the ability to observe trends across different groups of biomarkers. In addition to patterns discussed within the 4 functional pillars, clinicians commonly observe other patterns including Small Intestinal Bacterial Overgrowth (SIBO), Intestinal Permeability, and Immune Dysregulation/ Loss of Resilience. These patterns are not diagnostic, but rather suggestive of these common conditions and further workup may be appropriate.

Biomarker pattern, along with associated symptoms suggest	Biomarkers	Next steps
Small Intestinal Bacterial Overgrowth (SIBO)	<ul style="list-style-type: none"> <li>↑ Relative Abundance</li> <li>↑ Products of Protein Breakdown</li> <li>↑ SCFA</li> <li>↑ n-butyrate</li> <li>↑ Fecal Fat (total)</li> <li>↓ PE1 (limited evidence)</li> <li>↑ <i>Methanobrevibacter smithii</i></li> </ul>	<ul style="list-style-type: none"> <li>• Confirm with SIBO Breath Test</li> </ul>
Intestinal Permeability	<ul style="list-style-type: none"> <li>↑ Zonulin</li> <li>↑ sIgA</li> <li>↑ EPX</li> <li>↓ <i>Akkermansia muciniphila</i></li> </ul>	<ul style="list-style-type: none"> <li>• Confirm with Intestinal Permeability Assessment</li> <li>• Determine root cause of permeability</li> <li>• Consider GI repair/ support</li> <li>• Consider IgG and/or IgE Food Antibody testing</li> </ul>
Immune Dysregulation/ Loss of Resilience	<ul style="list-style-type: none"> <li>&lt;DL sIgA</li> <li>&lt;DL EPX</li> <li>↑ PP bacteria/yeast</li> <li>↓ Beneficial bacteria</li> <li>↓ <i>Akkermansia muciniphila</i></li> </ul>	<ul style="list-style-type: none"> <li>• Evaluate root cause of immune dysregulation (i.e., Adrenocortex Stress Profile)</li> <li>• Consider prebiotics/probiotics</li> </ul>

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