



CDSA 2.0

Patient: **JANE DOE**
DOB: September 28, 1940
Sex: F
MRN:

Order Number: **E1210572**
Completed: October 05, 2011
Received: September 21, 2011
Collected: September 20, 2011

Digestion/Absorption

Analyte	Result	Reference Range
1. Pancreatic Elastase 1 [♦]	18	>= 201 mcg/g
2. Putrefactive SCFAs (Total*)	<dl	1.3-8.6 micromol/g

*Total values equal the sum of all measurable parts.

Digestion/Absorption

Digestion encompasses the functional activities of: mastication, gastric acid production, pancreatic activity, bile production and brush border maintenance. Absorption depends on all of the above actions, as well as a healthy gut mucosal barrier.

Gut Immunology

Analyte	Result	Reference Range
3. Eosinophil Protein X	1.9	<= 7.0 mcg/g
4. Calprotectin [♦]	>500	<=50 mcg/g

Gut Immunology

Eosinophil Protein X (EPX) reflects IgE-mediated inflammation and tissue damage and can be elevated in celiac disease, collagenous colitis, helminthic/parasitic infection, and IgE mediated food allergies. Elevated EPX requires further diagnostic testing to determine the cause. Calprotectin is a neutrophilic marker specific for inflammation in the gastrointestinal tract. It is elevated with infection, post-infectious IBS, and NSAID enteropathy. Fecal calprotectin can be used to differentiate IBD vs. IBS, to monitor treatment in IBD, and to determine which patients should be referred for endoscopy and/or colonoscopy. Levels between 50-120 should be repeated at 4-6 weeks and confirmed.

Metabolic

Analyte	Result	Reference Range
5. Beneficial SCFAs (Total*)	39.7	>= 13.6 micromol/g
6. n-Butyrate	5.9	>= 2.5 micromol/g
7. pH [♦]	6.3	6.1-7.9
8. Beta-glucuronidase	<dl	337-4,433 U/g

Secondary Bile Acids

9. Lithocholic acid (LCA)	6.20	0.65-5.21 mg/g
10. Deoxycholic acid (DCA)	6.47	0.67-6.76 mg/g
11. LCA / DCA Ratio	0.96	0.39-2.07

*Total values equal the sum of all measurable parts.

Metabolic

Gut metabolism is representative of the bacterial milieu, primarily through the presence of commensal bacteria. Metabolic activities include: mucous production, vitamin synthesis and absorption, deconjugation of steroid hormones and bile acids, fat regulation, and SCFA metabolism. These metabolic activities require a normal population of commensal bacteria without active bacterial, viral, or parasitic infection.



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Microbiology

Bacteriology

12. Beneficial Bacteria

Lactobacillus species	(2+)
Escherichia coli	(4+)
Bifidobacterium	(1+)

14. Mycology

Candida albicans/dubliniensis	<div style="display: flex; align-items: center; gap: 10px;"> <div style="border: 1px solid black; padding: 2px;">NP</div> <div style="background-color: yellow; border: 1px solid black; border-radius: 10px; padding: 2px 10px;">(1+)</div> </div>
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Human microflora is influenced by environmental factors and the competitive ecosystem of the organisms in the GI tract. Pathological significance should be based upon clinical symptoms and reproducibility of bacterial recovery.

*NG	NP	PP	P
*NG	NP	PP	P
No Growth	Non-Pathogen	Potential Pathogen	Pathogen

Microbiology

The Markers in this section reflect the bacteriological status of the gut.

Beneficial bacteria Beneficial flora controls potentially pathogenic organisms, influences nutrient production, removes toxins from the gut and stimulates the intestinal immune system (GALT). The composition of the colonic flora is affected by diet, transit time, stool pH, age, microbial interactions, colonic availability of nutrients, bile acids, sulfate and the ability of the microbes to metabolize these substrates. Ideally, levels of Lactobacilli and E. coli should be 2+ or greater. Bifidobacteria being a predominate anaerobe should be recovered at levels of 4+.

Additional bacteria

Non-pathogen: Organisms that fall under this category are those that constitute normal, commensal flora, or have not been recognized as etiological agents of disease.

Potential Pathogen: Organisms that fall under this category are considered potential or opportunistic pathogens when present in heavy growth.

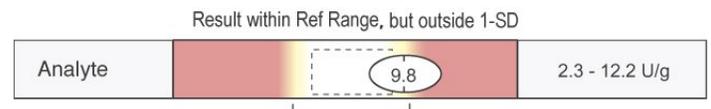
Pathogen: The organisms that fall under this category are well-recognized pathogens in clinical literature that have a clearly recognized mechanism of pathogenicity and are considered significant regardless of the quantity that appears in culture.

Mycology: Organisms that fall under this category constitute part of the normal colonic flora when present in small numbers. They may, however, become potential pathogens after disruption of the mucosal lining, which enables fungi to colonize and establish a local infection.

Lab Comments

Elastase repeated and confirmed. 09/29/2011 UL

The **Reference Range** is a statistical interval representing 95% or 2 Standard Deviations (2 S.D.) of the reference population. One Standard Deviation (1 S.D.) is a statistical interval representing 68% of the reference population. Values between 1 and 2 S.D. are not necessarily abnormal. Clinical correlation is suggested. (See example below)



Commentary is provided to the practitioner for educational purposes, and should not be interpreted as diagnostic or treatment recommendations. Diagnosis and treatment decisions are the responsibility of the practitioner.

The performance characteristics of all assays have been verified by Genova Diagnostics, Inc. Unless otherwise noted with as cleared by the U.S. Food and Drug Administration, assays are For Research Use Only.



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Parasitology

Microscopic Exam Results:

No Ova or Parasites seen
White Blood Cells: Rare

Parasitology

Optimized Parasite Recovery (OPR) is a technique used by Genova Diagnostics Inc. that involves combining multiple stool specimens submitted from the same patient for intestinal parasite examination as compared to individual sample evaluation. Research demonstrates that this method increases parasite recovery.

Data from analysis shows that parasites are detected in 22% of samples submitted to Genova Diagnostics Inc. This implies that a significant portion of the population suffers from infection with parasites, many of whom experience minimal gastrointestinal symptoms.

PARASITOLOGY EIA TESTS:

	In Range	Out of Range
Cryptosporidium	Negative	
Giardia lamblia	Negative	
Entamoeba histolytica/dispar	Negative	