



Patient: Jane Doe

DOB: September 16, 1960

Sex: F

MRN:

2205 GI Effects™ Microbial Ecology Profile – Stool





Methodology: DNA by PCR

Gastrointestinal Microbiome

	Result CFU/g stool	QUINTILE DISTRIBUTION					Reference Range CFU/g stool
		1st	2nd	3rd	4th	5th	
Commensal Bacteria (PCR)							
Bacteroidetes Phylum							
<i>Bacteroides-Prevotella</i> group	4.3E7						7.3E6 - 2.3E9
<i>Bacteroides vulgatus</i>	1.2E8						<4.6E9
<i>Barnesiella</i> spp.	<DL						<3.3E8
<i>Odoribacter</i> spp.	5.6E7						<2.0E8
<i>Prevotella</i> spp.	8.6E5						2.4E5 - 3.0E7
Firmicutes Phylum							
<i>Anaerotruncus colihominis</i>	6.4E6						<6.1E7
<i>Butyrivibrio crossotus</i>	1.5E5						7.8E3 - 8.6E5
<i>Clostridium</i> spp.	2.7E9						3.1E8 - 3.2E10
<i>Coprococcus eutactus</i>	2.7E7						<2.0E8
<i>Faecalibacterium prausnitzii</i>	8.2E8						1.2E5 - 7.1E7
<i>Lactobacillus</i> spp.	6.9E8						1.5E7 - 7.6E9
<i>Pseudoflavonifractor</i> spp.	2.9E7						1.2E5 - 2.1E8
<i>Roseburia</i> spp.	2.8E9						1.7E8 - 4.1E9
<i>Ruminococcus</i> spp.	8.9E8						1.2E8 - 6.9E11
<i>Veillonella</i> spp.	1.4E6						2.6E5 - 1.0E8
Actinobacteria Phylum							
<i>Bifidobacterium</i> spp.	<DL						<1.5E10
<i>Bifidobacterium longum</i>	<DL						<1.3E9
<i>Collinsella aerofaciens</i>	1.4E8						1.5E7 - 3.7E9
Proteobacteria Phylum							
<i>Desulfovibrio piger</i>	<DL						<2.8E7
<i>Escherichia coli</i>	6.0E7						5.5E4 - 7.9E8
<i>Oxalobacter formigenes</i>	3.9E6						<2.8E7
Euryarchaeota Phylum							
<i>Methanobrevibacter smithii</i>	<DL						<1.9E8
Fusobacteria Phylum							
<i>Fusobacterium</i> spp.	1.9E4						<4.8E5
Verrucomicrobia Phylum							
<i>Akkermansia muciniphila</i>	1.8E7						>1.7E6
Firmicutes/Bacteroidetes Ratio							
<i>Firmicutes/Bacteroidetes</i> (F/B Ratio)	53						21 - 620

The gray-shaded portion of a quintile reporting bar represents the proportion of the reference population with results below detection limit.

Commensal results and reference range values are displayed in a computer version of scientific notation, where the capital letter "E" indicates the exponent value (e.g., 7.3E6 equates to 7.3 x 10⁶ or 7,300,000).

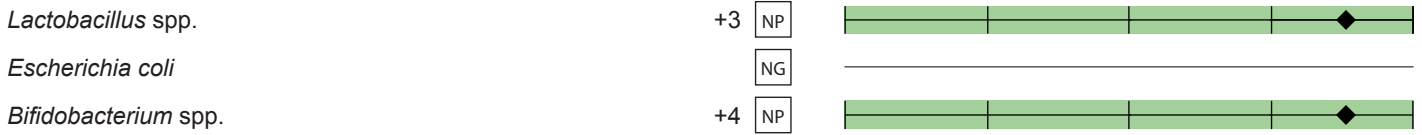
The Firmicutes/Bacteroidetes ratio (F/B Ratio) is estimated by utilizing the lowest and highest values of the reference range for individual organisms when patient results are reported as <DL or >UL.



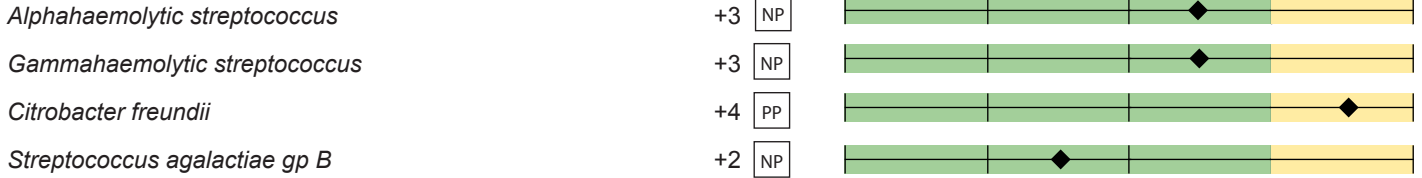
Methodology: culture/MALDI-TOF MS, Automated and Manual Biochemical Methods, Vitek 2® System Microbial identification and Antibiotic susceptibility

Gastrointestinal Microbiome

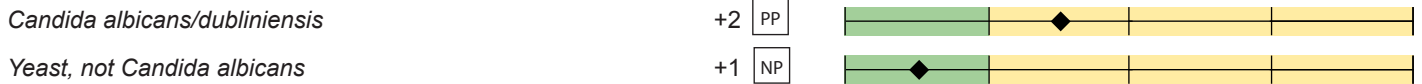
Bacteriology (Culture)



Additional Bacteria



Mycology (Culture)



Human microflora is influenced by environmental factors and the competitive ecosystem of the organisms in the GI tract. Pathogenic significance should be based upon clinical symptoms.

Microbiology Legend			
NG	NP	PP	P
No Growth	Non-Pathogen	Potential Pathogen	Pathogen

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.



Methodology: Direct Microscopic Examination, EIA

Parasitology

Microscopic Exam Results:

Blastocystis hominis: Many

Parasitology

Parasite Recovery: Literature suggests that >90% of enteric parasitic infections may be detected in a sample from a single stool collection. Increased sensitivity results from the collection of additional specimens on separate days.

Lab Comments

SENSI'S: All yeast, add'l bacteria

Parasitology EIA Tests:

	In Range	Out of Range
<i>Cryptosporidium</i> ◆	Negative	
<i>Giardia lamblia</i> ◆	Negative	
<i>Entamoeba histolytica</i> ◆	Negative	



Methodology: EIA, Fecal Immunochemical Testing (FIT)

Additional Results

	Result	Expected Value	
Fecal Occult Blood ♦	Negative	Negative	HpSA (<i>Helicobacter pylori</i> stool antigen) <i>Helicobacter pylori</i> is a bacterium which causes peptic ulcer disease and plays a role in the development of gastric cancer. Direct stool testing of the antigen (HpSA) is highly accurate and is appropriate for diagnosis and follow-up of infection.
Color††	Brown		
Consistency††	Formed/Normal		
HpSA - <i>H.pylori</i>	Negative	Negative	<i>Campylobacter</i> <i>Campylobacter jejuni</i> is the most frequent cause of bacterial-induced diarrhea. While transmission can occur via the fecal-oral route, infection is primarily associated with the ingestion of contaminated and poorly cooked foods of animal origin, notably, red meat and milk.
<i>Campylobacter</i> spp♦	Negative	Negative	
<i>Clostridium difficile</i> ♦	Negative	Negative	
<i>Shiga toxin E. coli</i> ♦	Negative	Negative	<i>Clostridium difficile</i> is an anaerobic, spore-forming gram-positive bacterium. After a disturbance of the gut flora (usually with antibiotics), colonization with <i>Clostridium difficile</i> can take place. <i>Clostridium difficile</i> infection is much more common than once thought.
Fecal Lactoferrin♦	Negative	Negative	

†† Results provided from patient input.

Shiga toxin E. coli is a group of bacterial strains that have been identified as worldwide causes of serious human gastrointestinal disease. *Enterohemorrhagic E. coli* includes over 100 different serotypes; 0157:H7 is the most significant, occurring in over 80% of all cases. Contaminated food continues to be the principal vehicle for transmission; foods associated with outbreaks include alfalfa sprouts, fresh produce, beef, and unpasteurized juices.

Tests were developed and their performance characteristics determined by Genova Diagnostics. Unless otherwise noted with ♦, the assays have not been cleared or approved by the U.S. Food and Drug Administration.