

invivo[®]

The Human Microbiome Company

TEST REPORTED: 17/06/2021
 TEST RECEIVED: 08/06/2021
 PATIENT NAME: SAMPLE PATIENT
 PATIENT DOB:
 GENDER: FEMALE

REPORT STATUS: COMPLETED
 CLINICIAN NAME: SAMPLE CLINICIAN
 ACCESSION NO:
 SAMPLE TYPE: STOOL

Lab Director:
 Eviatar Natan, PhD



Immune

RESULTS:

RANGE:

Beta Defensin 2	5ng/g	NORMAL		<68ng/g
Secretory IgA	159ug/g	NORMAL		<188ug/g

Inflammation

RESULTS:

RANGE:

Calprotectin	127ug/g	HIGH		<100ug/g
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Digestion

RESULTS:

RANGE:

Bile Acids	5466umol/L	HIGH		<3477umol/L
Pancreatic Elastase	162ug/g	LOW		>200ug/g

Other

RESULTS:

RANGE:

FIT (Occult Blood)	0ug/g	NORMAL		<10ug/g
Zonulin	2ng/g	NORMAL		<100ng/g



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Commensal Bacteria

RESULTS:

RANGE:

			0 - 4	5 - 8	9 - 12	13 - 16	17 - 20	
<i>Akkermansia muciniphila</i>	2.2							1.7-7.1
<i>Anaerostipes caccae</i>	5.9							3.0-7.3
<i>Bacteroides spp.</i>	16.4							14.8-17.5
<i>Bifidobacterium spp.</i>	12							7.3-16.3
<i>Blautia obeum</i>	9.4	LOW						15.5-17.9
<i>Coprococcus eutactus</i>	2.1	LOW						10.4-16.5
<i>Escherichia coli</i>	2.4	LOW						4.5-12.0
<i>Eubacterium rectale</i>	11.4							7.7-14.8
<i>Faecalibacterium prausnitzii</i>	17							14.2-18.3
<i>Lactobacillus spp.</i>	<DL	LOW						2.7-8.9
<i>Roseburia homini</i>	7							5.7-10.2
<i>Ruminococcus bromii</i>	15.6							14.2-17.7
<i>Subdoligranulum variabile</i>	6.3							6.3-12.5

Bacteroides Sub Group

RESULTS:

RANGE:

			0 - 4	5 - 8	9 - 12	13 - 16	17 - 20	
<i>Bacteroides dorei</i>	15.3							6.1-17.0
<i>Bacteroides fragilis</i>	13.3							10.7-14.8
<i>Bacteroides ovatus</i>	1.1	LOW						3.5-9.2
<i>Bacteroides thetaiotaomicron</i>	8.8							6.2-12.5
<i>Bacteroides uniformis</i>	10.1	LOW						10.9-16.5
<i>Bacteroides vulgatus</i>	16.1							14.6-17.4



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Gram Negative (-) Bacteria

	RESULTS:		RANGE:
<i>Bilophila wadsworthia</i>	7.4		2.4-9.5
<i>Citrobacter freundii</i>	<DL		<1.0
<i>Citrobacter koseri</i>	<DL		<DL
<i>Desulfovibrio spp.</i>	12.1	HIGH	<6.9
<i>Enterobacter cloacae</i>	<DL		<2.8
<i>Fusobacterium nucleatum</i>	0.9		<2.8
<i>Hafnia alvei</i>	<DL	LOW	0.8-9.0
<i>Klebsiella oxytoca</i>	<DL		<1.5
<i>Klebsiella pneumoniae</i>	<DL		<2.5
<i>Morganella morganii</i>	<DL		<0.5
<i>Oxalobacter formigenes</i>	<DL		<1.6
<i>Prevotella copri</i>	<DL		<11.4
<i>Proteus mirabilis</i>	<DL		<0.4
<i>Pseudomonas aeruginosa</i>	<DL		<0.7
<i>Serratia marcescens</i>	<DL		<0.4
<i>Veillonella spp.</i>	5.7		4.0-10.0



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Gram Positive (+) Bacteria

	RESULTS:	RANGE:
<i>Clostridium perfringens</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <4.0
<i>Clostridium sporogenes</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <DL
<i>Enterococcus faecalis</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <3.0
<i>Enterococcus faecium</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <2.6
<i>Enterococcus gallinarum</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <0.9
<i>Methanobrevibacter smithii</i>	12.6 HIGH	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <8.3
<i>Mycobacterium avium</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <0.2
<i>Ruminococcus gnavus</i>	5.8	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 4.1-10.7
<i>Ruminococcus torques</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <2.3
<i>Staphylococcus aureus</i>	1.3	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <3.5
<i>Streptococcus agalactiae</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <0.9
<i>Streptococcus pneumoniae</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <0.1
<i>Streptococcus pyogenes</i>	<DL	0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 <DL

Helicobacter pylori

	RESULTS:	RANGE:
<i>Helicobacter pylori</i> Stool Antigen	NEGATIVE	NEGATIVE



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Mycology

RESULTS:

RANGE:

			0 - 4	5 - 8	9 - 12	13 - 16	17 - 20	
<i>Aspergillus fumigatus</i>	<DL							<0.1
<i>Candida albicans</i>	<DL							<0.6
<i>Candida tropicalis</i>	<DL							<0.3
<i>Malassezia restricta</i>	<DL							<0.5

Parasitology

RESULTS:

RANGE:

			0 - 4	5 - 8	9 - 12	13 - 16	17 - 20	
<i>Blastocystis hominis</i>	9.7	HIGH						< 6.4
<i>Dientamoeba fragilis</i>	<DL							< 12.5
<i>Entamoeba histolytica</i>	<DL							< DL
<i>Giardia lamblia</i>	<DL							< DL

Bacterial Pathogens

RESULTS:

RANGE:

			0 - 4	5 - 8	9 - 12	13 - 16	17 - 20	
<i>B. fragilis</i> Enterotoxigenic	<DL							< DL
<i>Clostridium difficile</i>	<DL							< DL
<i>Clostridium difficile</i> (tox A)	<DL							< DL
<i>Clostridium difficile</i> (tox B)	<DL							< DL
<i>Yersinia enterocolitica</i>	<DL							< DL

The GI EcologiX™ profile utilises the highly sensitive quantitative PCR (qPCR) TaqMan technology for analysis of the gastrointestinal microbiota. Microbial genes of interest are quantified within a sample and their abundances are normalised to an endogenous and highly conserved gene. The qPCR results are therefore reported as the relative abundance of a microorganism as proportional to the whole microbial community.



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