-.ALEXANDRA MIDDLETON



P: 1300 688 522 E: info@nutripath.com.au A: PO Box 442 Ashburton VIC 3142 Date of Birth: 29-Jan-1958

Sex: F

Collected: 21/Feb/2022 Received: 23-Feb-2022 18 HILL STREET QUEENSCLIFF NSW 2096

Lab id: **3798976** UR#: 6600234

6 EDWARDS BAY ROAD MOSMAN NSW 2088

COMPLETE MICROBIOME MAPPING

General Macroscopic Description

Result		Range	Markers				
Stool Colour	Brown		Colour - Brown is the colour of normal stool. Other colours may indicate abnormal GIT conditions.				
Stool Form	Formed		Form -A formed stool is considered normal. Variations to this may indicate abnormal GIT conditions.				
Mucous	NEG	<+	Mucous - Mucous production may indcate the presence of an infection, inflammation or malignancy.				
Occult Blood	NEG	<+	Blood (Macro) - The presence of blood in the stool may indicate possible GIT ulcer, and must always be investigated immediately.				

GIT Functional Markers	Result	Range	Units	
Calprotectin.	7.2	0.0 - 50.0	ug/g	
Pancreatic Elastase	>500.0	> 200.0	ug/g	•
Faecal Secretory IgA	<i>165.9</i> *L	510.0 - 2010).0 ug/g	•
Faecal Zonulin	<i>107.1</i> *H	0.0 - 107.0	ng/g	•
Faecal B-Glucuronidase	1866.8	337.0 - 4433	3.0 U/g	
Steatocrit	1.0	0.0 - 15.0	%	
anti-Gliadin IgA	<20	0.0 - 100.0	units/L	•

Microbiome Mapping Summary

Parasites & Worms

Blastocystis hominis.

Bacteria & Viruses

Streptococcus species Klebsiella pneumoniae.

Fungi and Yeasts

Key Phyla Microbiota

-, ,				
Bacteroidetes	10.50	8.61 - 33.10	x10^11 org/g	•
Firmicutes	13.40	5.70 - 30.40	x10^10 org/g	•
Firmicutes:Bacteroidetes Ratio	0.13	< 1.00	RATIO	

Page 1 of 8 Complete Microbiome Map Lab ID: 3798976 Patient Name: ANITA SCHUMANN



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Parasites and Worms.	Result	Range	Units		
Parasitic Organisms					
Cryptosporidium.	<dl< th=""><th>< 1.0</th><th>x10^6 org/g</th><th></th></dl<>	< 1.0	x10^6 org/g		
Entamoeba histolytica.	<dl< th=""><th>< 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Giardia lamblia.	<dl< th=""><th>< 5.0</th><th>x10^3 org/g</th><th></th></dl<>	< 5.0	x10^3 org/g		
Blastocystis hominis.	233.5 *H	< 2.0	x10^3 org/g	•	
Dientamoeba fragilis.	<dl< th=""><th>< 1.0</th><th>x10^5 org/g</th><th></th></dl<>	< 1.0	x10^5 org/g		
Endolimax nana	<dl< th=""><th>< 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Entamoeba coli.	<dl< th=""><th>< 5.0</th><th>x10^6 org/g</th><th></th></dl<>	< 5.0	x10^6 org/g		
Pentatrichomonas hominis	<dl< th=""><th>< 1.0</th><th>x10^2 org/g</th><th></th></dl<>	< 1.0	x10^2 org/g		
Worms					
Ancylostoma duodenale, Roundworn	n Not De	etected		Comment: Not Detected results indicate	
Ascaris lumbricoides, Roundworm	Not De	etected		the absence of detectable DNA in this	
Necator americanus, Hookworm	Not De	etected		sample for the worms reported.	
Trichuris trichiura, Whipworm	Not De	etected			
Taenia species, Tapeworm	Not De	etected			
Enterobius vermicularis,Pinworm	Not De	etected			
Opportunistic Bacteria/Overgr	Result	Range	Units		
Bacillus species.	<dl< th=""><th>< 1.5</th><th>x10^5 org/g</th><th></th></dl<>	< 1.5	x10^5 org/g		
Enterococcus faecalis	0.6	< 1.0	x10^4 org/g	•	
Enterococcus faecium	0.7	< 1.0	x10^4 org/g		
Morganella species	<dl< th=""><th>< 1.0</th><th>x10^3 org/g</th><th>•</th></dl<>	< 1.0	x10^3 org/g	•	
Pseudomonas species	<dl< th=""><th>< 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Pseudomonas aeruginosa.	<dl< th=""><th>< 5.0</th><th>x10^2 org/g</th><th>•</th></dl<>	< 5.0	x10^2 org/g	•	
Staphylococcus species	<dl< th=""><th>< 1.0</th><th>x10^4 org/g</th><th>•</th></dl<>	< 1.0	x10^4 org/g	•	
Staphylococcus aureus	<dl< th=""><th>< 5.0</th><th>x10^2 org/g</th><th></th></dl<>	< 5.0	x10^2 org/g		
Streptococcus species	1.1 *H	< 1.0	x10^3 org/g	•	
Methanobacteriaceae	0.12	< 5.00	x10^9 org/g		
Desulfovibrio piger	<dl< th=""><th>0.0 - 18.0</th><th>x10^7 org/g</th><th></th></dl<>	0.0 - 18.0	x10^7 org/g		
Oxalobacter formigenes	154.1	> 15.0	x10^7 org/g	•	
Potential Autoimmune Triggers					
Citrobacter species.	<dl< th=""><th>< 5.0</th><th>x10^5 org/g</th><th></th></dl<>	< 5.0	x10^5 org/g		
Citrobacter freundii.	<dl< th=""><th>< 5.0</th><th>x10^5 org/g</th><th></th></dl<>	< 5.0	x10^5 org/g		
Klebsiella species	<dl< th=""><th>< 5.0</th><th>x10^3 org/g</th><th></th></dl<>	< 5.0	x10^3 org/g		
			0.0		
Klebsiella pneumoniae.	<i>438.3</i> *H	< 5.0	x10^4 org/g	•	
Klebsiella pneumoniae. Prevotella copri	438.3 *H <dl< th=""><th>< 1.0</th><th></th><th></th></dl<>	< 1.0			
Prevotella copri Proteus species		< 1.0 < 5.0	x10^4 org/g x10^7 org/g x10^4 org/g	•	
Prevotella copri	<dl< th=""><th>< 1.0 < 5.0 < 1.0</th><th>x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g</th><th>•</th></dl<>	< 1.0 < 5.0 < 1.0	x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g	•	
Prevotella copri Proteus species	<dl< th=""><th>< 1.0 < 5.0</th><th>x10^4 org/g x10^7 org/g x10^4 org/g</th><th>•</th></dl<>	< 1.0 < 5.0	x10^4 org/g x10^7 org/g x10^4 org/g	•	
Prevotella copri Proteus species Proteus mirabilis.	<dl <dl <dl< th=""><th>< 1.0 < 5.0 < 1.0</th><th>x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g</th><th></th></dl<></dl </dl 	< 1.0 < 5.0 < 1.0	x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g		
Prevotella copri Proteus species Proteus mirabilis. Fusobacterium species Fungi & Yeast Candida species.	<dl <dl <dl 1.08</dl </dl </dl 	< 1.0 < 5.0 < 1.0 < 10.00 Range < 5.0	x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g x10^7 org/g		
Prevotella copri Proteus species Proteus mirabilis. Fusobacterium species Fungi & Yeast	<dl <dl <dl 1.08 Result</dl </dl </dl 	< 1.0 < 5.0 < 1.0 < 10.00 Range < 5.0 < 5.0	x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g x10^7 org/g Units x10^3 org/g x10^2 org/g		
Prevotella copri Proteus species Proteus mirabilis. Fusobacterium species Fungi & Yeast Candida species.	<dl< th=""><th>< 1.0 < 5.0 < 1.0 < 10.00 Range < 5.0 < 5.0 < 3.0</th><th>x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g x10^7 org/g Units x10^3 org/g</th><th></th></dl<>	< 1.0 < 5.0 < 1.0 < 10.00 Range < 5.0 < 5.0 < 3.0	x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g x10^7 org/g Units x10^3 org/g		
Prevotella copri Proteus species Proteus mirabilis. Fusobacterium species Fungi & Yeast Candida species. Candida albicans.	<dl< th=""><th>< 1.0 < 5.0 < 1.0 < 10.00 Range < 5.0 < 5.0</th><th>x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g x10^7 org/g Units x10^3 org/g x10^2 org/g</th><th></th></dl<>	< 1.0 < 5.0 < 1.0 < 10.00 Range < 5.0 < 5.0	x10^4 org/g x10^7 org/g x10^4 org/g x10^3 org/g x10^7 org/g Units x10^3 org/g x10^2 org/g		

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Bacterial Pathogens	Result	Range	Units			
Aeromonas species.	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th>_</th></dl<>	< 1.0	x10^3 CFU/g		_	
Campylobacter.	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
C. difficile, Toxin A	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
C. difficile, Toxin B	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
Enterohemorrhagic E. coli	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
E. coli O157	<dl< th=""><th>< 1.0</th><th>x10^2 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^2 CFU/g			
Enteroinvasive E. coli/Shigella	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th>•</th><th></th></dl<>	< 1.0	x10^3 CFU/g	•		
Enterotoxigenic E. coli LT/ST	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
Shiga-like Toxin E. coli stx1	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
Shiga-like Toxin E. coli stx2	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
Salmonella.	<dl< th=""><th>< 1.0</th><th>x10^4 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^4 CFU/g			
Vibrio cholerae	<dl< th=""><th>< 1.0</th><th>x10^5 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^5 CFU/g			
Yersinia enterocolitica.	<dl< th=""><th>< 1.0</th><th>x10^5 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^5 CFU/g			
Helicobacter pylori	<dl< th=""><th>< 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g			
Comment: Helico Pylori virulence	Comment: Helico Pylori virulence factors will be listed below if detected POSITIVE					
H.pylori Virulence Factor, babA	Not De	etected	H.pylori Viru	lence Factor, cagA	Not Detected	
H.pylori Virulence Factor, dupA	Not Detected		H.pylori Viru	lence Factor, iceA	Not Detected	
H.pylori Virulence Factor, oipA	Not De	etected		lence Factor, vacA	Not Detected	
H.pylori Virulence Factor, virB	Not De	etected	H.pylori Viru	lence Factor, virD	Not Detected	
Viral Pathogens	Result	Range	Units			
Adenovirus 40/41	<dl< th=""><th>< 1.0</th><th>x10^10 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^10 CFU/g			
Norovirus GI/II	<dl< th=""><th>< 1.0</th><th>x10^7 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^7 CFU/g			
Bocavirus	<dl< th=""><th>< 1.0</th><th>x10^10 CFU/g</th><th>•</th><th></th></dl<>	< 1.0	x10^10 CFU/g	•		
Normal Bacterial GUT Flora	Result	Range	Units			
Bacteroides fragilis	4.2	1.6 - 250.0	x10^9 CFU/g			
Bifidobacterium species	90.6	> 6.7	x10^7 CFU/g			
Bifidobacterium longum	82.4	> 5.2	x10^6 CFU/g			
Enterococcus species	539.2	1.9 - 2000.0	x10^5 CFU/g	•		
Escherichia species	1.2 *L	3.7 - 3800.0	x10^6 CFU/g	•		
Lactobacillus species	29.5	8.6 - 6200.0	x10^5 CFU/g	•		
Lactobacillus Rhamnosus		8.3 - 885.0	x10^4 CFU/g	•		
Clostridium species	10.6	5.0 - 50.0	x10^6 CFU/g	•		
Enterobacter species	6.9	1.0 - 50.0	x10^6 CFU/g	•		
Akkermansia muciniphila	4.37	0.01 - 50.00	_	•		
Faecalibacterium prausnitzii	830.5	1.0 - 500000	x10^3 CFU/g	•		
Short Chain Fatty Acids	Result	Range	Units			
Short Chain Fatty Acids, Beneficial	18.8	> 13.6	umol/g			
Butyrate	13.8	10.8 - 33.5	%	•		
Acetate	56.5	44.5 - 72.4	%			
Propionate	27.4	0.0 - 32.0	%		•	
Valerate	2.3	0.5 - 7.0	%			



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Pathogen Summary:

Macroscopy Comment

BROWN coloured stool is considered normal in appearance.

Metabolism Comment

In a healthy gut Short Chain Fatty Acids are exhibited in the following proportions; Butyrate, Acetate, Propionate (16%:60%:24%)

VALERATE:

Valerate is a short chain fatty acid that is important for gut health. Although Acetate, propionate, and butyrate make up the the most abundant SCFAs in gastrointestinal tract (95%), Valerate and other SCFA's make up the remaining and work optimally when within range.

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GIT Markers Comment

PANCREATIC ELASTASE: Normal exocrine pancreatic function.

Pancreatic Elastase reflects trypsin, chymotrypsin, amylase and lipase activity.

This test is not affected by supplements of pancreatic enzymes.

Healthy individuals produce on average 500 ug/g of PE-1. Thus, levels below 500 ug/g and above 200 ug/g suggest a deviation from optimal pancreatic function.

The clinician should therefore consider digestive enzyme supplementation if one or more of the following conditions is present: Loose watery stools, Undigested food in the stools, Post-prandial abdominal pain, Nausea or colicky abdominal pain, Gastroesophageal reflux symptoms, Bloating or food intolerance.

CALPROTECTIN Normal:

Faecal calprotectin values <50 ug/g are not indicative of inflammation in the gastrointestinal tract. Subjects with low faecal calprotectin levels normally do not need to be further investigated by invasive procedures.

FAECAL SECRETORY IgA:

Production of sIgA is important to the normal function of the gastrointestinal mucosa as an immune barrier.

It represents the first line immune defense of the GIT.

Elevated levels are associated with an upregulated immune response.

LOW/LOW NORMAL sIgA LEVEL:

The primary function of secretory IgA (sIgA) is an antibody protein secreted into the gastrointestinal tract as a first line of immune defence against pathogenic microorganisms. sIgA binds to invading micro organisms and toxins and entrap them in the mucus layer or within the epithelial cells, so inhibiting microbial motility, agglutinating the organisms and neutralising their exotoxins and then assist in their harmless elimination from the body in the faecal flow. sIgA also 'tags' food as acceptable, so low sIgA leads to increased sensitivity to foods.

Several studies link stress and emotionality with levels of sIgA. Production is adversely affected by stress, which is mediated by cortisol levels.

**Reduced sIgA levels may be associated with sub optimal adrenal output. Consider an Adrenocortex Stress profile.

Treatment: Investigate the root cause of inflammation. Consider the use of probiotics (saccharomyces boulardii), choline, essential fatty acids, glutathione, glycine, glutamine, phosphatidylcholine, Vitamin C and Zinc which are all required for efficient production of sIqA.

ELEVATED ZONULIN LEVELS:

Zonulin is a protein that modulates intestinal barrier function. Zonulin release facilitates the opening of tight junctions between the cells of the intestinal lining to allow for passage of nutrients and fluids into the body. However, Zonulin release can be "overstimulated" by certain external factors to cause excessive opening of tight junctions, leading to intestinal hyperpermeability or "leaky gut", inflammation, liver overload, nutrient deficiencies, rheumatoid arthritis and autoimmune disorders.

Identify the possible cause/s (Gut microorganism imbalance or the presence of dietary Gluten/gliadin) and remove to reduce further damage.

If it's gluten for gluten sensitivity or celiac disease, remove gluten.

If bacterial overgrowth or dysbiosis, treat the bacterial overgrowth.

Treatment:

Firstly, fix the gut. Treat/repair the gut before before proceeding with other protocols; nutrients and other supplements can be damaging to the system if they get out of the gut

Follow a grain - free diet for at least 12 months.

Eliminate gluten, sugar, processed food, artificial flavorings, colors, trans fats.

Supplementation:

Caprylic acid, Probiotics, acidophylis and B complex, fish oil, Magnesium D3, CoQ10, Mg Citrate, Boswellia & Curcumin, Milk Thistle, Selenium

For patients with chronic digestive issue: Vitamin A, L-Glutamine, Probiotics

Further investigations to consider:

- SIBO Breath Test,
- IgG or IgA 96 Food Sensitivity

Parasites/Worms Comment

ELEVATED BLASTOCYSTIS HOMINIS LEVEL:

Significant copies per gene of Blastocystis hominis have been detected in this stool sample. The role of B. hominis in terms of colonisation and disease is still considered controversial. When this organism is present in the absence of any other parasites, enteric

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organisms or viruses, it may be considered the etiological agent of disease. Symptoms can include diarrhoea, cramps, nausea, fever, vomiting and abdominal pain. B. hominis has been associated with irritable bowel syndrome, infective arthritis and intestinal obstruction.

Treatment:

Metronidazole (Flagyl) is considered the most effective drug. Iodoquinol and paromomycin are also effective medications. Recommended therapy can also eliminate G. lamblia, E. histolytica and D. fragilis, all of which may be concomitant undetected pathogens and part of patient symptomology.

Using a combination of herbs that contain berberine (e.g. Golden seal, Coptis chinensis, Barberry, Oregon grape and Phellodendron) is desirable for the treatment of certain organisms. It is important to investigate the percentage of berberine contained in the dry weight extract of the berberine containing herb and then dose accordingly for the therapeutic dose of berberine.

Total therapeutic dose of berberine: 200mg four times daily.

Further Investigation:

PCR stool analysis should be considered in 4 weeks' time to ensure infection has cleared.

Opportunistic Bacteria Comment

STREPTOCOCCUS SPECIES:

Description:

Streptococcus is a gram-positive bacteria in the Firmicutes phylum. Streptococcus is generally a common isolate from gut flora. However, emerging research suggests that high levels in the intestine may result from low stomach acid, PPI use, reduced digestive capacity, SIBO or constipation; Elevated levels may also be indicative of intestinal inflammatory activity, and may cause loose stools.

Sources:

Recent infections with streptococcus pyogenes or scarlet fever can be linked to the presence of this species in faeces.

Treatment:

Treatment of streptococcus in gut flora is not always recommended. A practitioner may take into consideration a range of patient factors and symptoms to determine if treatment is necessary. In this case please refer to the 4R treatment protocol located at the end of this report.

METHANOBACTERIACEAE:

Family of bacteria-like microbes that produce methane. Facilitates carbohydrate fermentation and short-chain fatty acid production by beneficial bacteria.

LOW levels may indicate reduced production of short-chain fatty acids and may be associated with inflammation.

HIGH levels linked to chronic constipation, as well as some types of SIBO and IBS.

DESULFOVIBRIO COMMENT:

Sulfate is present in different concentrations in the intestine dependent on diet. Remnants not absorbed, alongside the presence of lactate, promote the growth of Sulfate reducing bacteria (SRB). Desulfovibrio Piger is the dominant SRB genus and has been implicated in gastrointestinal disorders such as ulcerative colitis via the reduction of sulfate to hydrogen sulphide in the gut. High Delsulfovibrio piger levels serves as an indicator of inflammatory bowel disease.

Treatment options include lowering the intake of sulfate rich foods such as some breads, dried fruits, beers, ciders and wines. Reference: Kushkevych et. Al., J. Clin. Med. 2019, 8, 1054; doi:10.3390/jcm8071054

OXALOBACTER COMMENT:

Oxolate is formed in the liver by amino acid catabolism as well as present in a wide range of foods including tea, coffee, chocolate and certain fruits and vegetables. High concentration of oxalate in the urine is related to the potential formation of calcium oxalate kidney stones. Oxolobacter Formigenes is the main known bacterial species involved in oxalate degradation in the gut. Levels of O. Formigenes tends to decrease with age as well as with the use of antibiotics or other drugs, with low levels identified as a risk factor for calcium oxide stone formation. Treatment options include probiotic treatment and low oxalate diet modification. Urinary oxalate levels can also be monitored by test code 4025 (oxalate urinary).

Reference: Duncan et. al., Applied and Environmental Microbiology, Aug. 2002, p. 3841-3847

Kaufman et. al., J Am Soc Nephrol. 2008 Jun; 19(6): 1197-1203.

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Potential Autoimmune Comments

ELEVATED KLEBSIELLA PNEUMONIAE LEVEL:

Sources:

Klebsiella is part of the Enterobacteriaceae family and as such is a gram-negative bacteria. Klebsiella is Isolated from foods and environmental sources. Klebsiella appears to thrive in individuals on a high starch diet.

Pathogenicity:

Part of the normal gut flora in small numbers, but can be an opportunistic pathogen.

Klebsiella is capable of translocating from the gut when in high numbers.

Certain strains of K. oxytoca have demonstrated cytotoxin production.

Symptoms:

When Klebsiella is found in considerable amounts, symptoms may include abdominal pain, bloating, loose stools, anxiety, insomnia, food allergies.

Treatment:

Avoiding carbohydrates such as rice, potatoes, flour products and sugary foods reduces the amount of Klebsiella in the gut Other Herbal antimicrobials include:

Lemon and clove, Burr marigold, Thyme, Licorice, euphobia, cordyceps. For further treatment options, refer to the 4R treatment protocol located at the end of this report.

FUSOBACTERIUM SPECIES:

Fusobacterium species is a gram-negative bacteria in the Fusobacteria phylum. The bacteria is a common member of the human oral microbiome, this pro-inflammatory bacterium can also be found in the human gut. In the mouth, high levels are strongly linked to oral hygiene. In the gut, high levels have been observed in individuals with colon cancer and appendicitis.

Sources:

It primarily uses protein as its main source. However, research also shows that it can thrive from sugar.

Treatment:

Antimicrobial botanicals such as berberine, oregano, quercetin, curcumin, green and black tea extracts, blueberry extract, cinnamon and rosemary have shown to decrease levels.

Normal Bacterial Flora Comment

LOW ESCHERICHIA SPECIES LEVEL:

Organism of the Proteobacteria phylum. Escherichia coli (E. coli) is the primary species in this genus, and most are nonpathogenic. Low levels may indicate reduced mucosal health and decreased protection against pathogenic E. coli.

LOW LACTOBACILLUS RHAMNOSUS LEVEL:

Lactobacullus Rhamnosus is a bacteria in the Firmicutes phylum. Lactobacillus rhamnosus is one of the most widely used probiotic strains. Various health effects are well documented including the prevention and treatment of gastro-intestinal infections and diarrhea, and stimulation of immune responsesL. Low levels may be linked to poor digestive health, diarrhea and IBS symptoms.



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Sex : F

Collected: 21/Feb/2022 Received: 23-Feb-2022 18 HILL STREET QUEENSCLIFF NSW 2096

Lab id: 3798976 UR#: 6600234

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The Four "R" Treatment Protocol

	Using a course of antimicrobial, antibacterial, antiviral or anti parastic therapies in cases where organisms are present. It may also be necessary to remove offending foods, gluten, or medication that may be acting as antagonists.	ANTIMICROBIAL	Oil of oregano, berberine, caprylic acid
		ANTIBACTERIAL	Liquorice, zinc camosine, mastic gum, tribulus, berberine, black walnut, caprylic acid, oil of oregano
OVE		ANTIFUNGAL	Oil of oregano, caprylic acid, berberine, black walnut
HEM		ANTIPARASTIC	Artemesia, black walnut, berberine, oil of oregano
	Consider testing IgG96 foods as a tool for removing offending foods.	ANTIVIRAL	Cat's claw, berberine, echinacea, vitamin C, vitamin D3, zinc, reishi mushrooms
		BIOFILM	Oil of oregano, protease
REPLACE	In cases of maldigestion or malabsorption, it may be necessary to restore proper digestion by supplementing with digestive enzymes.	DIGESTIVE SUPPORT	Betaine hydrochloride, tilactase, amylase, lipase, protease, apple cider vinegar, herbal bitters
ш	Pecolonisation with healthy, beneficial bacteria. Supplementation with probiotics, along with the use of prebiotics helps re-establish the proper microbial balance.	PREBIOTICS	Sippery elm, pectin, larch arabinogalactans
RENOCULAT		PROBIOTICS	Bifidobacterium animalis sup lactise, lactobacillus acidophilus, lactobacillus plantarum, lactobacillus casei, bifidobacterium breve, bifidobacterium bifidum, bifidobacterium longum, lactobacillus salivarius sep salivarius, lactobacillus paracasei, lactobacillus rhamnosus, Saccaromyces boulardii
SALANCE	Restore the integrity of the gut mucosa by giving support to healthy mucosal cells, as well as immune support. Address whole body health and lifestyle factors so as to prevent future GI dysfunction.	INTESTINAL MUCOSA IMMUNE SUPPORT	Saccaromyces boulardii, lauric acid
REPAIR & REBAL		INTESTINAL BARRIER REPAIR	L-Glutamine, a loe vera, liquorice, marshmallow root, okra, quercetin, slippery elm, zinc camosine, Saccaromyces boulardii, omega 3 essential fatty acids, B vitamins
REP/		SUPPORT CONSIDERATION	Seep, diet, exercise, and stress management