

P: 1300 688 522 E: info@nutripath.com.au A: PO Box 442 Ashburton VIC 3142 Date of Birth: 20-Oct-1986

Sex: F

Collected: 9/Aug/2021 Received: 12-Aug-2021 11/51 MCDONALD STREET FRESHWATER NSW 2096

Lab id: 3759996 UR#: 6589449

6 EDWARDS BAY ROAD MOSMAN NSW 2088

# COMPLETE MICROBIOME MAPPING

# General Macroscopic Description

Result		Range	Markers
Stool Colour	Brown		<b>Colour</b> - Brown is the colour of normal stool. Other colours may indicate abnormal GIT conditions.
Stool Form	Unformed		<b>Form</b> -A formed stool is considered normal. Variations to this may indicate abnormal GIT conditions.
Mucous	NEG	<+	<b>Mucous</b> - Mucous production may indcate the presence of an infection, inflammation or malignancy.
Occult Blood	+	<+	<b>Blood (Macro)</b> - 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.	9.6	0.0 - 50.0	ug/g	
Pancreatic Elastase	>500.0	> 200.0	ug/g	•
Faecal Secretory IgA	798.2	510.0 - 2010	0.0 ug/g	
Faecal Zonulin	<i>108.2</i> *H	0.0 - 107.0	ng/g	
Faecal B-Glucuronidase	3639.9	337.0 - 4433	3.0 U/g	•
Steatocrit	<i>22.0</i> *H	0.0 - 15.0	%	
anti-Gliadin IgA	<20	0.0 - 157.0	units/L	

# Microbiome Mapping Summary

# **Parasites & Worms**

# **Bacteria & Viruses**

Streptococcus species Proteus species

# **Fungi and Yeasts**

Key Phyla Microbiota

x10^11 org/g **Bacteroidetes 6.90** \*L 8.61 - 33.10 x10^10 org/g 5.70 - 30.40 **Firmicutes** 30.33 Firmicutes:Bacteroidetes Ratio 0.44 < 1.00 **RATIO** 





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Parasites and Worms.	Result	Range	Units		
Parasitic Organisms					
Cryptosporidium.	<dl< th=""><th>&lt; 1.0</th><th>x10^6 org/g</th><th></th></dl<>	< 1.0	x10^6 org/g		
Entamoeba histolytica.	<dl< th=""><th>&lt; 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Giardia lamblia.	<dl< th=""><th>&lt; 5.0</th><th>x10^3 org/g</th><th></th></dl<>	< 5.0	x10^3 org/g		
Blastocystis hominis.	<dl< th=""><th>&lt; 2.0</th><th>x10^3 org/g</th><th></th></dl<>	< 2.0	x10^3 org/g		
Dientamoeba fragilis.	<dl< th=""><th>&lt; 1.0</th><th>x10^5 org/g</th><th></th></dl<>	< 1.0	x10^5 org/g		
Endolimax nana	<dl< th=""><th>&lt; 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Entamoeba coli.	<dl< th=""><th>&lt; 5.0</th><th>x10^6 org/g</th><th></th></dl<>	< 5.0	x10^6 org/g		
Pentatrichomonas hominis	<dl< th=""><th>&lt; 1.0</th><th>x10^2 org/g</th><th></th></dl<>	< 1.0	x10^2 org/g		
Worms					
Ancylostoma duodenale, Roundworr	n Not De	etected		Comment: Not Detected results indicate	
Ascaris lumbricoides, Roundworm	Not Detected		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>&lt; 1.5</th><th>x10^5 org/g</th><th>•</th></dl<>	< 1.5	x10^5 org/g	•	
Enterococcus faecalis	<dl< th=""><th>&lt; 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Enterococcus faecium	<dl< th=""><th>&lt; 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Morganella species	<dl< th=""><th>&lt; 1.0</th><th>x10^3 org/g</th><th></th></dl<>	< 1.0	x10^3 org/g		
Pseudomonas species	<dl< th=""><th>&lt; 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Pseudomonas aeruginosa.	<dl< th=""><th>&lt; 5.0</th><th>x10^2 org/g</th><th></th></dl<>	< 5.0	x10^2 org/g		
Staphylococcus species	<dl< th=""><th>&lt; 1.0</th><th>x10^4 org/g</th><th></th></dl<>	< 1.0	x10^4 org/g		
Staphylococcus aureus	<dl< th=""><th>&lt; 5.0</th><th>x10^2 org/g</th><th></th></dl<>	< 5.0	x10^2 org/g		
Streptococcus species	<i>8.5</i> *H	< 1.0	x10^3 org/g	•	
Methanobacteriaceae	0.52	< 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	21.0	> 15.0	x10^7 org/g		
Potential Autoimmune Triggers					
Citrobacter species.	<dl< th=""><th>&lt; 5.0</th><th>x10^5 org/g</th><th></th></dl<>	< 5.0	x10^5 org/g		
Citrobacter freundii.	<dl< th=""><th>&lt; 5.0</th><th>x10^5 org/g</th><th></th></dl<>	< 5.0	x10^5 org/g		
Klebsiella species	<dl< th=""><th>&lt; 5.0</th><th>x10^3 org/g</th><th></th></dl<>	< 5.0	x10^3 org/g		
Klebsiella pneumoniae.	<dl< th=""><th>&lt; 5.0</th><th>x10^4 org/g</th><th></th></dl<>	< 5.0	x10^4 org/g		
Prevotella copri	<dl< th=""><th>&lt; 1.0</th><th>x10^7 org/g</th><th></th></dl<>	< 1.0	x10^7 org/g		
Proteus species	<i>345.5</i> *H	< 5.0	x10^4 org/g	•	
Proteus mirabilis.	<dl< th=""><th>&lt; 1.0</th><th>x10^3 org/g</th><th></th></dl<>	< 1.0	x10^3 org/g		
Fusobacterium species	0.19	< 10.00	x10^7 org/g		
Fungi & Yeast	Result	Range	Units		
Candida species.	0.9	< 5.0	x10^3 org/g		
Candida albicans.	<dl< th=""><th>&lt; 5.0</th><th>x10^2 org/g</th><th></th></dl<>	< 5.0	x10^2 org/g		
Geotrichum species.	<dl< th=""><th>&lt; 3.0</th><th>x10^2 org/g</th><th></th></dl<>	< 3.0	x10^2 org/g		
Microsporidium species	<dl< th=""><th>&lt; 5.0</th><th>x10^3 org/g</th><th></th></dl<>	< 5.0	x10^3 org/g		
Rhodotorula species.	<dl< th=""><th>&lt; 1.0</th><th>x10^3 org/g</th><th>•</th></dl<>	< 1.0	x10^3 org/g	•	



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Bacterial Pathogens	Result	Range	Units		
Aeromonas species.	<dl< th=""><th>&lt; 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g		
Campylobacter.	<dl< th=""><th>&lt; 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>&lt; 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>&lt; 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>&lt; 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>&lt; 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>&lt; 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>&lt; 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>&lt; 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>&lt; 1.0</th><th>x10^3 CFU/g</th><th>•</th><th></th></dl<>	< 1.0	x10^3 CFU/g	•	
Salmonella.	<dl< th=""><th>&lt; 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>&lt; 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>&lt; 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>&lt; 1.0</th><th>x10^3 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^3 CFU/g		
Comment: Helico Pylori virulence	factors	will be liste	d below if o	letected POSITIVE	
H.pylori Virulence Factor, babA	Not Do	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 Do	etected	H.pylori Virulence Factor, vacA		Not Detected
H.pylori Virulence Factor, virB	Not Do	etected	H.pylori Viru	lence Factor, virD	<b>Not Detected</b>
Viral Pathogens	Result	Range	Units		
Adenovirus 40/41	<dl< th=""><th>&lt; 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>&lt; 1.0</th><th>x10^7 CFU/g</th><th></th><th></th></dl<>	< 1.0	x10^7 CFU/g		
Bocavirus	<dl< th=""><th>&lt; 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	1.2 *L	1.6 - 250.0	x10^9 CFU/g	•	
Bifidobacterium species	46.9	> 6.7	x10^7 CFU/g		
Bifidobacterium longum	42.5	> 5.2	x10^6 CFU/g		•
Enterococcus species	4.2	1.9 - 2000.0	x10^5 CFU/g	•	
Escherichia species	660.6	3.7 - 3800.0	x10^6 CFU/g	•	
Lactobacillus species	29.3	8.6 - 6200.0	x10^5 CFU/g		
Lactobacillus Rhamnosus	<i>5.1</i> *L	8.3 - 885.0	x10^4 CFU/g	•	
Clostridium species	31.7	5.0 - 50.0	x10^6 CFU/g		
Enterobacter species	3.1	1.0 - 50.0	x10^6 CFU/g	•	
Akkermansia muciniphila	2.63	0.01 - 50.00	x10 <sup>3</sup> CFU/g		
Faecalibacterium prausnitzii	930.4	1.0 - 500000	x10^3 CFU/g	•	
Short Chain Fatty Acids	Result	Range	Units		
Short Chain Fatty Acids, Beneficial	24.1	> 13.6	umol/g		
Butyrate	19.4	10.8 - 33.5	%		
Acetate	62.3	44.5 - 72.4	%		
Propionate	16.0	0.0 - 32.0	%		
Valerate	2.3	0.5 - 7.0	%		

## -.ALEXANDRA MIDDLETON



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## Pathogen Summary:

## **Macroscopy Comment**

BROWN coloured stool is considered normal in appearance.

UNFORMED/LIQUID stools may indicate the presence of infection and/or inflammation. Consider dysbiosis, food sensitivity, high dose vitamin C and magnesium, infection, intestinal permeability, laxative use,

malabsorption, maldigestion, stress. Other causes: bacterial, fungal, viral and other parasitic infections. Treatment:

- Investigate and treat possible underlying cause.
- Assess other CDSA markers such as pH, pancreatic elastase 1 & microbiology markers."

BLOOD PRESENT: Consider blood vessel injury, inflammation, infection, ulceration, hemorrhoids, severe constipation & other injury. Treatment:

- Investigate the cause of bleeding using other diagnostic tools such as endoscopy
- Assess other CDSA markers such as calprotectin, H. pylori, M2PK & microbiology markers.

## **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.

#### -.ALEXANDRA MIDDLETON



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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.

## **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

## **ELEVATED STEATOCRIT:**

The presence of steatorrhea is an indirect indicator of incomplete fat digestion. Consider high dietary fat intake, cholestasis, malabsorption and digestion (diarrhoea, pancreatic or bile salt insufficiency), intestinal dysbiosis, parasites, NSAIDs use, short bowel syndrome, whipple disease, crohn's disease, food allergies & sensitivities.

Treatment:

# o Prebiotic and probiotic supplementation

- o Supplement hydrochloride, digestive enzymes or other digestive aids
- o Investigate underlying causes
- o Investigate food sensitivities and allergies
- o Remove potential irritants
- o Review markers such as pancreatic elastase 1 and calprotectin

## **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.

#### -. ALEXANDRA MIDDLETON



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#### 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.

# **Potential Autoimmune Comments**

## **ELEVATED PROTEUS SPECIES LEVEL:**

Sources:

Food has been implicated as a vehicle of infection.

## Pathogenicity;

Part of the normal flora of the GI tract, though has been shown to be an independent causative agent of intestinal disorders. May also play a role as an opportunistic organism in enteric infection due to other pathogens.

## Symptoms

Occasionally implicated in diarrheal disorders.

Recently, it has been suggested that P. mirabilis may be an etiological agent in rheumatoid arthritis.

The mechanism may be related to the molecular cross reactivity between P. mirabilis and the HLA antigens, specifically HLA-DR4.

## Treatment

Currently, standard texts provide no specific antimicrobial guidelines for GI overgrowth of Proteus.

Ampicillin is recommended for extra-intestinal infections of P. mirabilis, followed by trimethoprim/sulfamethoxazole.

For further treatment suggestions, 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.



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## **Phyla Microbiota Comment**

#### LOW BACTEROIDETES LEVEL:

Gram-negative Bacteroidetes are a bacterial phyla that make up a large proportion of the human digestive tract, including the mouth, nose, throat, and colon. A low result in bacteroidetes may suggest imbalanced normal microbes in the GI tract. A lower level of bacteroidetes is considered an unfavourable outcome which allows for the potential of elevated firmicutes leading to a possible imbalanced firmicutes:bacteroidetes ratio.

Treatment:

It is suggested to eat a diverse range of foods including polyphenols. It is further suggested to decrease foods rich in fat and sugar as they encourage firmicute levels to rise. Investigate other causes relating to a low bacteroidetes level.

## **Normal Bacterial Flora Comment**

## LOW BACTEROIDES FRAGILIS LEVEL:

Organism of the Bacteroidetes phylum. Immune-modulating normal gut species believed to be involved in microbial balance, barrier integrity, and neuroimmune health.

Low levels may contribute to reduced anti- inflammatory activity in the intestine.

## 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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# The Four "R" Treatment Protocol

REMOVE	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	ANTIMICROBIAL	Oil of oregano, berberine, caprylic acid
		ANTIBACTERIAL	Liquorice, zinc camosine, mastic gum, tribulus, berberine, black walnut, caprylic acid, oil of oregano
		ANTIFUNGAL	Oil of oregano, caprylic acid, berberine, black walnut
	medication that may be acting as antagonists.	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
beneficial bacteria. Supplementation with probiotics, along with the us of prebioticshelpsre-establi	Recolonisation with healthy,	PREBIOTICS	Sippery elm, pectin, larch arabinogalactans
		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
REPAIR & REBALANCE	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
		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