

## **Gut Case Studies, Part 4**

## CASE #7: 38-YEAR-OLD FEMALE

Our next patient is a 38-year-old female with chief complaints of brain fog, low energy, joint pain, and stiffness with bloating, constipation, and distension. She's had a history of parasites and had self-treated with oregano oil but didn't retest after the self-treatment, and [she] had a tendency toward constipation. She did suspect issues with gluten but still consumed it occasionally.

			CO2 QC	Check	Pass				
	Gases	Expe	ected	Obse	erved	Norn	nal/Abnorm	al	
	$H_2^{\dagger}$	<25.58	ppm	6.	.54		Normal		
	CH₄	<10.00	ppm	53	.48		Abnormal		
	H <sub>2</sub> S	<5.00	ppm	1.	.52		Normal		
Indicative of In	estinal Meth	nanogenic	: Overgrow	Interp th	retation				
Indicative of In	estinal Meth	nanogenic	: Overgrow	Interp th	retation				
Indicative of In Samples	estinal Meth	nanogenic T2	: Overgrow	Interp th Re T4	retation esults T5	T6	17	T8	T9
ndicative of In Samples Interval (hr:mir	estinal Mett	nanogenic T2 16	: Overgrow T3 31	Interp th Re T4 46	esults T5 61	<b>T6</b> 76	<b>17</b> 91	<b>T8</b> 106	<b>T9</b> 121
ndicative of In Samples Interval (hr:mir Gases	estinal Mett	nanogenic T2 16	: Overgrow T3 31	Interp th Re T4 46	retation sults 15 61	<b>T6</b> 76	<b>17</b> 91	<b>T8</b> 106	<b>T9</b> 121
ndicative of In Samples Interval (hr:mir Gases H <sub>2</sub> (ppm	estinal Mett T1 0 5.58	nanogenic T2 16 IVR	<b>T3</b> 31 2.39	Interp th Re T4 46	retation sults 15 61 2.95	<b>T6</b> 76 6.54	<b>17</b> 91 6.95	<b>T8</b> 106 8.49	<b>19</b> 121 6.79
dicative of In amples nterval (hr:min ases H <sub>2</sub> (ppm CH4 (ppm	estinal Mett T1 0 5.58 53.48	<b>T2</b> 16 IVR IVR	<b>T3</b> 31 2.39 34.01	th Re T4 46 IVR IVR	retation results 15 61 2.95 34.56	<b>T6</b> 76 6.54 45.57	<b>17</b> 91 6.95 36.33	<b>T8</b> 106 8.49 36.27	<b>19</b> 121 6.79 35.69

So here you can see her trio-smart [small intestinal bacterial overgrowth] (SIBO) breath test is positive for intestinal methanogen overgrowth with a pretty high value, actually, up into the 50s. You'll also notice these IVR readings for two of the samples, which stands for insufficient volume received, meaning the tube didn't have enough air in it to sample. This happens from time to time for various reasons. If the lab thinks that the IVR results will impact [the] interpretation of the test, they'll generally reach out to you and ask if you want the patient to retest. I've only had to do that a few times in the course of using this test, and most of the time, you're still able to see the trend well enough with the results that you do have.



		and the stand	
Commensal Abundance	ommensal Microbio	ome Analysis	
	-30%	-10%	+10%
atient Total Commensal Abundance	-	Healthy	Cohort
Total Commenal Balance: The total commense healthy cohort. Low levels of commensal bacteria prebiotic-rich foods and may indicate the need for potential bacteria overgrowth or probiotic supplem Dysbiosis Patterns	I abundance is a sum-tota are often observed after a microbiome support. Com entation.	I of the reported comm Intimicrobial therapy, o versely, higher total co	vensal bacteria compared to a r in diets lacking fiber and/or mmensal abundance may indicate
Total Commenal Balance: The total commense healtry cohort. Low levels of commensal bacteria prebiolic-rich foods and may indicate the need for potential bacteria overgrowth or probiotic supplem Dysbiosis Patterns	al abundance is a sum-tota are often observed after a microbiome support. Conv lentation.	al of the reported comm untimicrobial therapy, c versely, higher total cc	ensal bacteria compared to a rin diets lacking fiber and/or mmensal abundance may indicate Dysbiosis Patterns: Genova's data ar
Total Commenal Balance: The total commenses healthy cohort. Low levels of commensal bacteria prebiotic-rich foods and may indicate the need for potential bacteria overgrowth or probiotic supplem Dysbiosis Patterns	al abundance is a sum-tota are often observed after a microbiome support. Com ientation.	al of the reported comm nntimicrobial therapy, c versely, higher total cc 110 High	ensal bacteria compared to a in diets lacking fiber and/or mmensal abundance may indicate Dysbiosis Patterns: Genova's data at has led to the development of unique dy patterns, related to key physiologic disr
Total Commenal Balance: The total commense healthy cohort. Low levels of commensal bacteria prebiotic-rich foods and may indicate the need for potential bacteria overgrowth or probiotic supplem Dysbiosis Patterns	al abundance is a sum-tota are often observed after a microbiome support. Com ientation.	I of the reported comm Intimicrobial therapy, c versely, higher total cc 110 High	ensal bacteria compared to a rin diets lacking fiber and/or mmensal abundance may indicate <u>Dyablosis Patterns</u> . Genova's data ar has led to the development of unique dy patterns, related to key physiologic dims such as immusuppression and inflam These patterns may represent dyabiotic changes that could pose cilicial signific

Her GI Effects stool test showed [an] imbalance in commensal abundance with a high normal methane dysbiosis score, so we're starting to see some imbalance [in] the ecosystem and microbial balance already. There [are] also elevated markers of protein breakdown that you can see with exocrine pancreatic insufficiency, high-protein diet, SIBO, low [hydrochloric acid], and then certain levels of dysbiosis. So with this particular patient, SIBO and dysbiosis seem to be the top contenders that are impacting protein breakdown. Low fecal fats can sometimes be seen in low-fat diets, but for her, it's really only her triglycerides that are low. So I'm just not 100 percent sure what to make of it, but we'll keep an eye on this as we go.



0	Diges	tion and Absorption	
Pancreatic Elastase 1 †	382	100 200	>200 mcg/g
Products of Protein Breakdown (Total*) (Valerate, Isobutyrate, Isovalerate)	10.2 H		1.8-9.9 micromol/g
Fecal Fat (Total*)	25.7	+ + + +	3.2-38.6 mg/g
Triglycerides	<dl l<="" td=""><td><u>← I I I I I</u></td><td>0.3-2.8 mg/g</td></dl>	<u>← I I I I I</u>	0.3-2.8 mg/g
Long-Chain Fatty Acids	23.3	<u> </u>	1.2-29.1 mg/g
Cholesterol	1.4	+ + +	0.4-4.8 mg/g
Phospholipids	1.0		0.2-6.9 mg/g
1	Inflamm	ation and Immunology	
Calprotectin †	<16	50 120	<=50 mcg/g
Eosinophil Protein X (EPX)†	0.2	4.6	<=4.6 mcg/g
Fecal secretory IgA	1,272	680 2040	<=2,040 mcg/mL
	Gut Mic	crobiome Metabolites	
Metabolic Short-Chain Fatty Acids (SCFA) (Total*) (Acetate, n-Butyrate, Propionate)	57.9	<u>⊨ + + + + + </u> +	>=23.3 micromol/g
n-Butyrate Concentration	8.3	+ + + +	>=3.6 micromol/g
n-Butyrate %	14.3	<mark>⊢ ♦              </mark>	11.8-33.3 %
Acetate %	65.3	+ + + +	48.1-69.2 %
Propionate %	20.4		<=29.3 %
Beta-glucuronidase	3,152	<b>⊢</b> + + <b>→ →</b>	368-6,266 U/g

Overall, [the] metabolic activity of [the] microbiome is mostly good, with some ratio issues but not deficient overall.

	Parasitology
Microscopie OSB Results	
Microscopic O&F Results	with a destruction to a second and the second se
Microscopic O&P is capable of detecting all des	cribed gastrointestinal parasites. The organisms listed in the box represent those
commonly found in microscopic stool analysis. S	should an organism be detected that is not included in the list below, it will be report
in the Additional Results section. For an extensiv	ve reference of all potentially detectable organisms, please visit
www.gdx.net/product/gi-effects-comprehensive-	stool-test
Genus/species	Result
Nematodes - roundworms	
Ancylostoma/Necator (Hookworm)	Not Detected
Ascaris lumbricoides	Not Detected
Capillaria philippinensis	Not Detected
Enterobius vermicularis	Not Detected
Strongyloides stercoralis	Not Detected
Trichuris trichiura	Not Detected
Cestodes - tapeworms	
Diphyllobothrium Jatum	Not Detected
Dipulidium caninum	Not Detected
Hymonologia diminuta	Not Detected
Humonologia paga	Not Detected
Taonia eno	Not Detected
Tremetedee flukee	Not Detected
Trematodes - nukes	11 · 15 · · · ·
Clonorchis/Opisthorchis spp.	Not Detected
Fasciola spp./ Fasciolopsis buski	Not Detected
Heterophyes/Metagonimus	Not Detected
Paragonimus spp.	Not Detected
Schistosoma spp.	Not Detected
Protozoa	
Balantidium coli	Not Detected
Blastocystis spp.	Many Detected
Chilomastix mesnili	Not Detected
Cryptosporidium spp.	Not Detected
Cyclospora cayetanensis	Not Detected
Dientamoeba fragilis	Many Detected
Entamoeba coli	Not Detected
Entamoeba histolytica/dispar	Not Detected
Entamoeba hartmanii	Not Detected
Entamoeba polecki	Not Detected
Endolimax nana	Not Detected
Giardia	Not Detected
lodamoeba buetschlii	Not Detected
Cystoisospora spp.	Not Detected
Trichomonads (e.g. Pentatrichomonas)	Not Detected
Additional Findings	
White Blood Cells	Not Detected
Charcot-Levden Crystals	Not Detected
Other Infectious Findings	



She's been diagnosed with parasites before, and here we [see] *Blastocystis* and [*Dientamoeba*] *fragilis* on the GI Effects stool test.

		Parasitology	
PCR Parasitology - Proto	zoa	Ме	athodologies: DNA by PCR, Next Generation Sequencing
Organism	Result	Units	Expected Result
Blastocystis spp.	6.78e4	femtograms/microliter C&S stool	Detected Not Detected
Cryptosporidium parvum/hominis	<1.76e2	genome copies/microliter C&S stool	Not Detected Not Detected
Cyclospora cayetanensis	<2.65e2	genome copies/microliter C&S stool	Not Detected Not Detected
Dientamoeba fragilis	4.10e5	genome copies/microliter C&S stool	Detected Not Detected
Entamoeba histolytica	<9.64e1	genome copies/microliter C&S stool	Not Detected Not Detected
Giardia	<1.36e1	genome copies/microliter C&S stool	Not Detected Not Detected
Blastocystis spp. Reflex Subty	ping		
Type 1: Not Detected	Type 4:	Not Detected Type 7:	Not Detected
Type 2: Detected	Type 5:	Not Detected Type 8:	Not Detected
Type 3: Not Detected	Type 6:	Not Detected Type 9:	Not Detected
		Additional Results	
Methodology: Fecal Immunochemical Test	ing (FIT)		
Feed Oswitt Blood	Result	Expected Value	
Fecal Occuit Blood+	Negative	Negative	
Consistency <sup>††</sup>	Loose		

Here's the typing for [*Blastocystis*] and [polymerase chain reaction] (PCR) confirmation from the [ova and parasites] (O&P) we just saw. So based on the Genova interpretation guide, subtype two tends to be less pathogenic but can cause some bloating and diarrhea for patients.



IESI		н	SULI	
Array 3 – Wheat/Gluten Proteome Reactivity & Autoimmunity	IN RANGE (Normal)	EQUIVOCAL*	OUT OF RANGE	(ELISA Index)
Wheat IgG	0.38			0.3-1.5
Wheat IgA	0.54			0.1-1.2
Wheat Germ Agglutinin IgG	<0.40			0.4-1.3
Wheat Germ Agglutinin IgA		1.09		0.2-1.1
Native & Deamidated Gliadin 33 IgG	<0.20			0.2-1.2
Native & Deamidated Gliadin 33 IgA	0.34			0.1-1.1
Alpha Gliadin 17-mer IgG	0.53			0.1-1.5
Alpha Gliadin 17-mer IgA		0.94		0.1-1.1
Gamma Gliadin 15-mer IgG	<0.50			0.5-1.5
Gamma Gliadin 15-mer IgA	0.26			0.1-1.0
Omega Gliadin 17-mer IgG	< 0.30			0.3-1.2
Omega Gliadin 17-mer IgA	0.49			0.1-1.2
Glutenin 21-mer IgG		1.43		0.1-1.5
Glutenin 21-mer IgA	0.91			0.1-1.3
Gluteomorphin + Prodynorphin IgG	0.39			0.3-1.2
Gluteomorphin + Prodynorphin IgA	0.43			0.1-1.2
Gliadin-Transglutaminase Complex IgG	0.32			0.3-1.4
Gliadin-Transglutaminase Complex IgA	0.54			0.2-1.5
Transglutaminase-2 IgG	0.32			0.3-1.6
Transglutaminase-2 IgA	0.91			0.1-1.6
Transglutaminase-3 IgG	0.63			0.2-1.6
Transglutaminase-3 IgA	0.64			0.1-1.5
Transglutaminase-6 IgG	1.08			0.2-1.5
Transglutaminase-6 IgA	0.77			0.1-1.5

She was still consuming some small amounts of gluten, so we did a Cyrex Array 3, and she [had] three equivocal markers of gluten intolerance. But given her suspicion and her subjective reaction to gluten and everything else that's going on, I [advised] her to avoid gluten ongoing. So here are the diagnoses. We've got intestinal methanogen overgrowth based [on] the trio-smart breath test, *Blastocystis hominis* and *D. fragilis* on the GI Effects [test], some dysbiosis that we're seeing with high markers of protein breakdown, and gluten intolerance on the Cyrex [Array] 3X.



	Diagnosis	
Pattern	Supporting Markers	Comments
SIBO (IMO)	TrioSmart	
Blastocystis hominis infection	GI Effects	+ D. Fragilis
Dysbiosis	GI Effects	High markers of protein breakdown
Gluten intolerance	Cyrex 3X	

So this patient has a lot going on, but we decided to focus primarily on the SIBO and [*Blastocystis*] first because of the high levels of methane on the test and her presenting symptoms. I don't know for sure if [*Blastocystis*] is a problem for her, but I figure we'll go ahead and address it in the protocol we're using for SIBO also.



## **Treatment protocol**

GI Synergy	1 seclect DID (with breakfast and disper)
	I packet BID (with breaklast and dinner)
Allimax Pro	1 capsules TID with food
Atrantil	2 capsules TID at beginning of meals
Interfase Plus	3-4 capsules BID (on an empty stomach)
Ideal Bowel Support	1 capsule BID; can be taken with food but away from antimicrobials or antibiotics
Mimosa Pudica	2 capsules twice daily on an empty stomach.
Saccharomyces boulardii	For Blastocystis

I have GI-Synergy and Interfase Plus from the core protocol, plus Allimax Pro, Atrantil, and Ideal Bowel Support for methane. I like using *Mimosa pudica* from Microbe Formulas for *Blastocystis* and *D. fragilis*, and then we also have some *Saccharomyces boulardii* in there.

				CHECK	Pass				
	Gases	Expe	ected	Obser	rved	Norn	nal/Abnorm	nal	
	H <sub>2</sub> <sup>†</sup>	<22.54	1 ppm	2.54	4		Normal		
	CH4	<10.00	) ppm	7.26	6		Normal		
	H <sub>2</sub> S	<5.00	) ppm	3.98	8		Normal		
Gases Detected	l at Normal	l Levels		Interpre	etation				
Gases Detected	l at Normal	l Levels		Interpre Resu	etation ults				
Gases Detected	l at Normal	l Levels T2	T3	Interpre Resu T4	etation ults T5	T6	17	T8	T9
Gases Detected Samples Interval (hr:min)	I at Normal	I Levels <u>T2</u> 17	<b>T3</b> 32	Interpre Resu T4 48	etation ults T5 65	<b>T6</b> 81	<b>17</b> 97	<b>T8</b> 113	<b>T9</b> 128
Gases Detected Samples Interval (hr:min) Gases	T1 0	<b>T2</b> 17	<b>T3</b> 32	Interpre Resu T4 48	etation ults 15 65	<b>T6</b> 81	<b>17</b> 97	<b>T8</b> 113	<b>T9</b> 128
Gases Detected amples Interval (hr:min) Gases H <sub>2</sub> (ppm)	T1 0 IVR	<b>T2</b> 17 2.54	<b>T3</b> 32 1.86	Interpre Resu T4 48 0.00 7.24	etation ults T5 65 2.07 4.11	<b>16</b> 81 0.28	<b>17</b> 97 0.00 3.45	<b>T8</b> 113 2.49	<b>T9</b> 128 0.77



So here's our follow-up SIBO breath test after doing that protocol for about 60 days, give or take a few extra weeks on there, and ramping up. The breath test showed resolution of intestinal methanogen overgrowth. This is really nice to see, considering how high her levels were. I don't always see those levels come down as nicely as this.

		Parasitology		
PCR Parasitology - Proto	zoa	Ме	thodologies: DNA by PCF	R, Next Generation Sequencing
Organism	Result	Units		Expected Result
Blastocystis spp.	6.78e4	femtograms/microliter C&S stool	Detected	Not Detected
Cryptosporidium parvum/hominis	<1.76e2	genome copies/microliter C&S stool	Not Detected	Not Detected
Cyclospora cayetanensis	<2.65e2	genome copies/microliter C&S stool	Not Detected	Not Detected
Dientamoeba fragilis	4.10e5	genome copies/microliter C&S stool	Detected	Not Detected
Entamoeba histolytica	<9.64e1	genome copies/microliter C&S stool	Not Detected	Not Detected
Giardia	<1.36e1	genome copies/microliter C&S stool	Not Detected	Not Detected
Blastocystis spp. Reflex Subty	ping			
Type 1: Not Detected	Type 4:	Not Detected Type 7:	Not Detected	
Type 2: Detected	Type 5:	Not Detected Type 8:	Not Detected	
Type 3: Not Detected	Type 6:	Not Detected Type 9:	Not Detected	
		Additional Results		
Methodology: Fecal Immunochemical Test	ing (FIT)			
Feed Oswitt Blood	Result	Expected Value		
	Negative	negative		
Consistency <sup>††</sup>	Loose			

*Blastocystis* and *D. fragilis* remained on follow-up testing, actually, and she did report some improvement in constipation and was now going daily, but found that she was actually leaning a little bit more toward loose stool after treating SIBO. She had also gone gluten-free at the time, so it's hard to know for sure what's the biggest contributor to her symptom improvement. But overall, she felt better, with bloating and distension still there most of the time, even after [the] first treatment.

So in this situation, we have a few options. We could continue with another botanical protocol and see if we have further progression for the parasites, or we could try pharmaceuticals. This patient wanted to try medications. She had already done a lot of botanical treatments with a history of unsuccessful treatments in the past.



Follow-up treatme	ent protocol (Blasto)
Intervention	Dosage
Secnidazole	400 mg TID (10 days)
Nitazoxanide (Alinia)	500 mg BID with meals (10 days)
Paramomycin sulfate	500 mg TID with meals (10 days)
SEED	2 capsules daily
Interfase Plus	3-4 capsules BID (on an empty stomach)
Saccharomyces boulardii	For Blastocystis and while on Rx

For [Blastocystis], the options are Alinia monotherapy of 500 milligrams twice daily for 30 days or triple-drug therapy from the Centre for Digestive [Diseases] (CDD). This drug therapy has changed over the years, and it used to be iodoquinol or Yodoxin, but that's no longer commercially available, at least based [on] what I can find and after talking to pharmacists. So at the time of this recording, the first-line [Blastocystis] treatment from the CDD is what this patient opted for, which was secnidazole 400 milligrams three times a day for 10 days, Alinia 500 milligrams twice daily, and paromomycin 500 milligrams three times daily. I did recommend continuing the probiotics, including the S. boulardii and a biofilm disruptor. [So] [she] continue[d] with the probiotics for an additional 30 days and then retested. Both [Blastocystis] and D. fragilis [were] resolved on the post-test, as did her joint pain and stiffness. The distension and bloating were continuing to improve, and her bowel patterns were normalizing, so that was good news. We're moving in the right direction. Considering she had so many infections and overgrowth present, I wasn't expecting her gut symptoms to just miraculously improve with resolution of the imbalances. So this is where we stick with it, continue with the restorative support, help with diet, [and] gut lining support with [inaudible] GI Revive and ION Gut Health and those type[s] of products.



## CASE #8: 32-YEAR-OLD MALE

[The] next patient is a 32-year-old male, [whose] chief complaint[s are] anxiety, panic attacks, irritability, [gastroesophageal reflux disease] (GERD), sinus congestion, and [minor?] joint pain. He was also recently diagnosed with immune thrombocytopenia (ITP). So also, that's the primary reason why he's here, to try to address or find underlying triggers of this disease. By the time he came to me, he was on Promacta already to address the low platelet values, and they did seem to be controlling the number some. If you look into this autoimmune disease more, you'll see that they just aren't sure what can trigger this process, but infections like HIV, hepatitis, or [*Helicobacter*] *pylori* have been associated with ITP. [It] also can be associated with viral illnesses like cytomegalovirus, varicella-zoster, and a few more. So the hunt is on to try to figure out, if you can, what triggered this process. We did start with gut testing, though. That's [the] baseline foundational start, even in this autoimmune process.



Here's the Genova SIBO three-hour breath test, positive for both intestinal methanogen overgrowth, at 12 part[s] per million at its highest value, and [a] hydrogen result of 35 at the 90-minute mark.



H. pylori				Devention		
	Result		Normal	Protozoa	Besult	Normal
Helicobacter pylori	1.2e3	High	<1.0e3	Blastocystis hominis	<di< td=""><td>&lt;2.00e3</td></di<>	<2.00e3
Virulence Factor, babA	Negative		Negative	Chilomastix mesnili	<dl< td=""><td>&lt;1.00e5</td></dl<>	<1.00e5
virulence Factor, cagA	Negative		Negative	Cyclospora spp.	<dl< td=""><td>&lt;5.00e4</td></dl<>	<5.00e4
firulence Factor, dupA	Negative		Negative	Dientamoeba fragilis	<dl< td=""><td>&lt;1.00e5</td></dl<>	<1.00e5
Arulence Factor, iceA	Negative		Negative	Endolimax nana	<dl< td=""><td>&lt;1.00e4</td></dl<>	<1.00e4
/irulence Factor, oipA	Negative		Negative	Entamoeba coli	<di< td=""><td>&lt;5.00e6</td></di<>	<5.00e6
/irulence Factor, vacA	Negative		Negative	Pentatrichomonas hominis	<di< td=""><td>&lt;1.00e2</td></di<>	<1.00e2
firulence Factor, virB	Negative		Negative	Worms	Result	Normal
virulence Factor, virD	Negative		Negative	Ancylostoma duodenale	Not Detected	Not Detected
Normal Bacterial Flora				Ascaris lumbricoides	Not Detected	Not Detected
	Result		Normal	Necator americanus	Not Detected	Not Detected
Bacteroides fragilis	5.11e10		1.60e9 - 2.50e11	Trichuris trichiura	Not Detected	Not Detected
Bifidobacterium spp.	2.89e11		>6.70e7	Taenia spp.	Not Detected	Not Detected
Enterococcus spp.	2.51e6		1.9e5 - 2.00e8	Intestinal Health		
Escherichia spp.	4.14e8		3.70e6 - 3.80e9	Digestion	Result	Normal
actobacillus spp.	3.86e7		8.6e5 - 6.20e8	Steatocrit	<dl< td=""><td>&lt;15 %</td></dl<>	<15 %
Clostridia (class)	2.93e6	Low	5.00e6 - 5.00e7	Elastase-1	312	>200 ug/g
Enterobacter spp.	2.42e6		1.00e6 - 5.00e7	GI Markers	Result	Normal
%kermansia muciniphila	3.95e5	High	1.00e1 - 5.00e4	b-Glucuronidase	1215	<2486 U/mL
Faecalibacterium prausnitzii	5.62e4		1.00e3 - 5.00e8	Occult Blood - FIT	0	<10 ug/g
Phyla Microbiota	Result		Normal	Immune Response	Result	Normal
Bacteroidetes	1.30e12		8.61e11 - 3.31e12	Secretory IgA	549	510 - 2010 ug/g
Firmicutes	7.80e10		5.70e10 - 3.04e11	Anti-gliadin IgA	108	0 - 157 U/L
Firmicutes:Bacteroidetes Ratio	0.06		<1.00	Inflammation	Result	Normal
				Calprotectin	11	<173 ug/g

We did a [Diagnostic Solutions Laboratory] (DSL) GI-MAP stool test that showed high levels of *H. pylori* without any virulence factors, but mostly normal bacterial flora and optimal digestion markers. I would like to see that elastase up just a touch higher to 500, but nowhere near pancreatic insufficiency.



	Diagnosis	
Pattern	Supporting Markers	Comments
SIBO	Genova breath	IMO and H2
H.pylori	DSL GI MAP	No virulence factors

Initial testing results: [the] diagnosis was SIBO, intestinal methanogen overgrowth, and hydrogen, *H. pylori* on the DSL GI-MAP without any virulence factors.

Treatmen	t protocol	
Nutraceutical	Dosage	
GI Synergy	1 packet BID (with breakfast and dinner)	
Lauricidin	1 scoop TID with each meal	
Interfase Plus	3-4 capsules BID on empty stomach	
TerraFlora	One capsule with lunch	
Bio HPF	2-3 capsules before meals, TID	
Atrantil	Slowly build to 20-30 drops BID with meals	
Saccharomyces boulardii	3 billion CFU BID at lunch and before bed	



For treatment, considering he had both SIBO and *H. pylori*, I used portions of the core protocol with some SIBO-specific botanicals and *H. pylori*-specific treatments. This was a little tricky because he wasn't terribly symptomatic from a gut perspective other than mild GERD, but with the association between *H. pylori* and ITP, I felt that I had to at least focus on the *H. pylori* a little bit more. So I added Bio-HPF for *H. pylori* along with *S. boulardii*, Atrantil for SIBO, and I honestly can't remember why we went with Terraflora for this patient over Seed, but I think at the time, Seed was probably on backorder. He did this protocol for about 10 weeks or so, including ramp-up, took a two- to four-week break, and then did follow-up testing. Toward the end of the protocol, he wasn't feeling much different other than his nasal congestion was gone and joint pain seemed to have [improved] insignificantly. He also changed his diet and removed inflammatory foods. So [it's] always a little tricky to know when we're doing multiple interventions at a time, but the goal is to improve symptoms and function and health, so we don't always have time or space to do every intervention by itself in order to tease [out] all that information.



This repeat SIBO breath test showed improvement in hydrogen levels but pretty similar methane levels. At this point, I decided to wait and see what the repeat stool test showed before deciding how to move forward. I wasn't really planning on aggressively treating the methane of 11 at this point for this particular patient, but I was open to it depending on what we found.



H pylori				Basasilas		
	Result		Normal	Protozoa	Result	Normal
Helicobacter pylori	1.1e3	High	<1.0e3	Blastocystis hominis	<dl< td=""><td>&lt;2.00e3</td></dl<>	<2.00e3
Virulence Factor, babA	Negative		Negative	Chilomastix mesnili	<dl< td=""><td>&lt;1.00e5</td></dl<>	<1.00e5
Virulence Factor, cagA	Negative		Negative	Cyclospora spp.	<dl< td=""><td>&lt;5.00e4</td></dl<>	<5.00e4
Virulence Factor, dupA	Negative		Negative	Dientamoeba fragilis	5.70e4	<1.00e5
Virulence Factor, iceA	Negative		Negative	Endolimax nana	<di< td=""><td>&lt;1.00e4</td></di<>	<1.00e4
Virulence Factor, oipA	Negative		Negative	Entamoeba coli	<dl< td=""><td>&lt;5.00e6</td></dl<>	<5.00e6
Virulence Factor, vacA	Negative		Negative	Pentatrichomonas hominis	<dl< td=""><td>&lt;1.00e2</td></dl<>	<1.00e2
Virulence Factor, virB	Negative		Negative	Worms	Result	Normal
Virulence Factor, virD	Negative		Negative	Ancylostoma duodenale	Not Detected	Not Detected
Normal Bactorial Flora				Ascaris lumbricoides	Not Detected	Not Detected
Normal Bacterial Flora	Result		Normal	Necator americanus	Not Detected	Not Detected
Bacteroides fragilis	2.75e11	High	1.60e9 - 2.50e11	Trichuris trichiura	Not Detected	Not Detected
Bifidobacterium spp.	8.69e11		>6.70e7	Taenia spp.	Not Detected	Not Detected
Enterococcus spp.	1.12e5	Low	1.9e5 - 2.00e8	Intestinal Health		
Escherichia spp.	7.95e8		3.70e6 - 3.80e9	Digestion	Result	Normal
Lactobacillus spp.	1.11e7		8.6e5 - 6.20e8	Elastase-1	397	>200 ug/g
Clostridia (class)	1.1266	Low	5.00e6 - 5.00e7	Gi Markers	Popult	Normal
Enterobacter spp.	6.56e6		1.00e6 - 5.00e7	b-Glucuronidase	531	<2486 U/mL
Akkermansia muciniphila	7.45e4	High	1.00e1 - 5.00e4	Occult Blood - FIT	0	<10 ug/g
Faecalibacterium prausnitzii	1.36e6		1.00e3 - 5.00e8	Immune Response	Result	Normal
Phyla Microbiota	Recult		Normal	Secretory IgA	250 Low	510 - 2010 ug/g
Bacteroidetes	2.45e12		8.61e11 - 3.31e12	Anti-gliadin IgA	94	0 - 157 U/L
Firmicutes	1.80e11		5.70e10 - 3.04e11	Inflammation	Result	Normal
Firmicutes:Bacteroidetes Ratio	0.07		<1.00	Calprotectin	0	<173 ug/g
	/					

Here's the repeat DSL follow-up test. *H. pylori* is still present in pretty similar levels. Normal flora was impacted a bit. An appearance of *D. fragilis* showed up. [There was] some slight improvement in elastase, but secretory [immunoglobulin A] also went down a bit, so it's a little hard to know exactly what is the result of the protocol here. He did the retest four weeks after treatment but hadn't really continued any of the supportive portions of the protocol that I had recommended. I see small changes in the stool test after protocols, like these little tiny fluctuations of changes in normal bacterial flora, a little bit of uptick in other—potential pathogens. So I generally like to wait it out and make a plan to treat the primary target, especially in this case, when the patient isn't terribly symptomatic and GERD symptoms did improve by about 50 percent.

10 Mar



Follow-up treatment protocol (H.pylori)			
Intervention	Dosage		
PPI (lansoprazole, omeprazole, pantoprazole, etc.)	40 mg twice daily (used Omeprazole)		
Amoxicillin	1 g BID		
Clarithromycin	500 mg BID		
SEED	2 capsules daily, continue ongoing		
Saccharomyces boulardii	3 billion CFU BID at lunch and before bed, continue for 30 days		

For this patient, we discussed the options for additional treatment of *H. pylori*. I might not have traditionally pursued this as much, given he didn't have peptic ulcer disease that I know of, and he didn't have a strong family history of gastric cancer, although his parents passed away pretty early on in life, so that's also a little difficult to know for sure. But for me, it was hard to ignore the connection between ITP and *H. pylori*, so I gave him the option of repeating [an] herbal protocol or moving to prescription therapy. I really try to avoid prescription therapy and use [it] only in certain cases. He really wanted prescription therapy because of the timeline it presented and the other stressors and responsibilities that he had going on in life. If you aren't able to prescribe, I still think a second herbal protocol could be helpful and useful if you really suspect *H. pylori* to be playing a role.



						Ç
H. pylori						
					Range CFU/g	
Helicobacter pylori		3.6e2			<1.0e3	
Virulence Factor, babA		N/A			Negative	
Virulence Factor, cagA		N/A			Negative	
Virulence Factor, dupA		N/A			Negative	
Virulence Factor, iceA		N/A			Negative	
Virulence Factor, oipA		N/A			Negative	
Virulence Factor, vacA		N/A			Negative	
Virulence Factor, virB		N/A			Negative	
Virulence Factor, virD		N/A			Negative	
Antibiotic Resistance	e Genes, phenotype	\$				
Helicobacter						
Clarithromycin		N/A			Negative	
A2142C	N/A	A2142G	N/A	A2143G	N/A	
Fluoroquinolones		N/A			Negative	
gyrA N87K	N/A	gyrA D91N	N/A	gyrA D91G	N/A	
gyrB S479N	N/A	gyrB R484K	N/A			
Tetracycline		N/A			Negative	
PBP1A S414R	N/A	PBP1A T556S	N/A	PBP1A N562Y	N/A	
Amoxicillin		N/A			Negative	
A926G	N/A	AGA926-928TTC	N/A			

I took a look at the antibiotic-resistant gene phenotype section of the DSL, and it showed no resistance for his *H. pylori*. So we went with the first line of treatment with a [proton pump inhibitor], amoxicillin, and clarithromycin for 14 days. I also had him continue Seed and *Saccharomyces boulardii*, and then we planned to retest four weeks after cessation of treatment.



Test Name	In Range Out Of Range	Lab
		NL1
HELICOBACTER PYLOR	I AG, EIA, STOOL	
Micro Number: Test Status: Specimen Source: Specimen Quality: H.pylori Ag.	11019640 Final Stool Adequate	
	Antimicrobials, proton pump inhibitors, and bismuth preparations inhibit H. pylori and ingestion up to two weeks prior to testing may cause false negative results. If clinically indicated the test should be repeated on a new specimen obtained two weeks after discontinuing treatment.	
Reference Range:	Not Detected	
Test Name HELICOBACTER PYLORI, UREA BREATH TEST	In Range Out Of Range Reference Range	Lab NL1
Antimicrobials, pr preparations are k ingestion of these may lead to false indicated, the tes obtained two weeks	oton pump inhibitors, and bismuth nown to suppress H. pylori, and prior to H. pylori diagnostic testing negative results. If clinically t may be repeated on a new specimen after discontinuing treatment.	

For follow-up testing, I decided to do the *H. pylori* portion of the DSL test this time only, but I also added on a Quest [Diagnostics] H. pylori stool antigen and the urea breath test just to have more information to base my next treatment plan on. I'll often do this when I only have the DSL [and] PSR to go off of, just because of what I mentioned in the past of having this huge portion of patients test positive for *H. pylori* on the DSL. And it's really not [the] gold standard for testing at this time, so I just want to be diligent and make sure that before deciding to do more aggressive treatments, that we're really being thorough. You can see here that the value of H. pylori on DSL did go down a bit and is no longer considered elevated by the lab. So that's good news, I guess. [It's] something to look at, some improvement in that number. His stool antigen and urea breath test results were both negative. So, all in all, with the DSL H. pylori test being just slightly elevated, these two tests being negative, and him no longer having GERD-type symptoms, we decided not to continue treatment. And he had already done a prescription treatment for this once, and I'm not convinced it was worth pursuing given these lab results. I gave him the option of consulting with a gastroenterologist for a biopsy, but we both agreed that was extremely invasive, and he opted not to do it at that time. So we continued working our way through looking for triggers for ITP.