

Interpreting and Applying GI360™ Test Data: The Three Stages of GI Dysfunction

Presented by
Dan Kalish, DC
and Julia Malkowski, ND, DC



Skill Acquisition and Overview

1. Show You...

- How to interpret stool test results and...
- assign patients to one of the three stages.

2. Your Turn...

Practice interpreting the results of a stool test (using the handouts provided).

3. Explain how...

To implement lifestyle and supplement modalities based on lab test results.

But first, a quick

Preface



Why perform GI testing in the first place?



- GI dysfunction occurs with or without GI symptoms
- If we are treating the “underlying cause” then why do we all only run GI tests on GI symptomatic patients?
- We should run GI test on every new patient as part of a basic functional medicine workup



3 Stages of GI Dysfunction

How it Happens:

The 3 Stages of GI Dysfunction

Stage 1

- Stress and poor diet cumulative effect, weakening of the adrenals and microbiome diversity

Stage 2

- GI organ function compromised, less HCl production, less enzyme and bile production, leaky gut starts with damage to gut lining and lowering of gut immunity (SIgA)

Stage 3

- Pathogens acquired, parasites, bacteria or yeast overgrowth



How to Identify Stage 1

A Disturbed Microbiome

- Low SCFA
- Commensal bacteria imbalance

How to Identify Stage 1

Low short chain fatty acids (esp. butyrate)

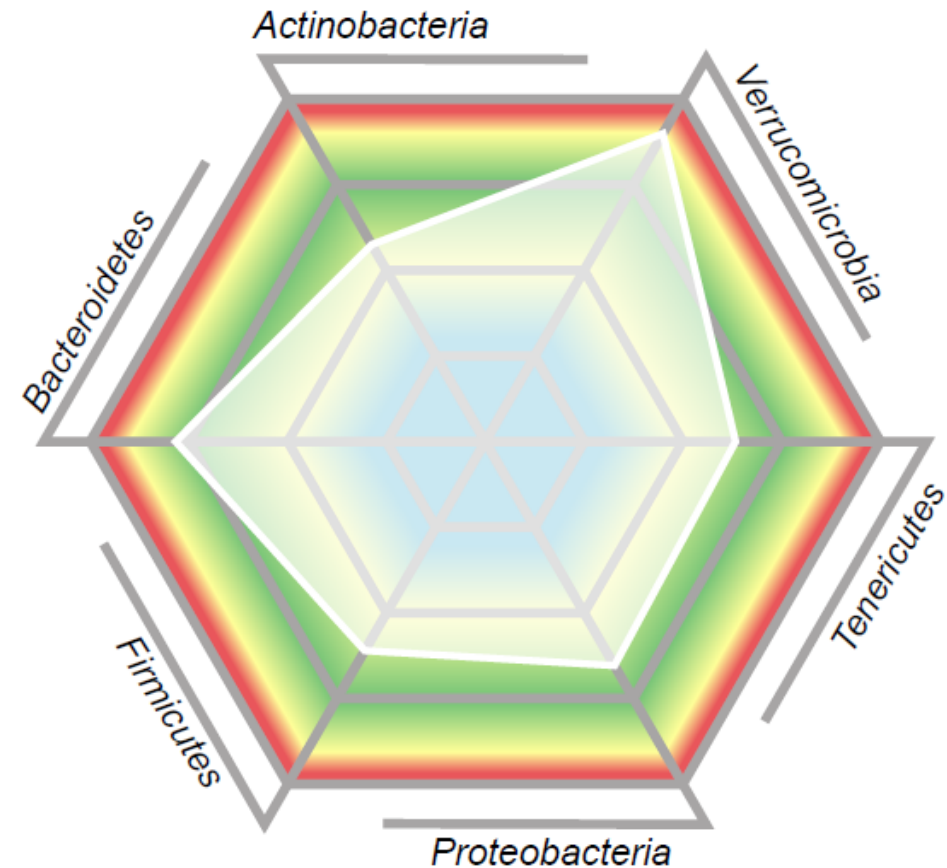
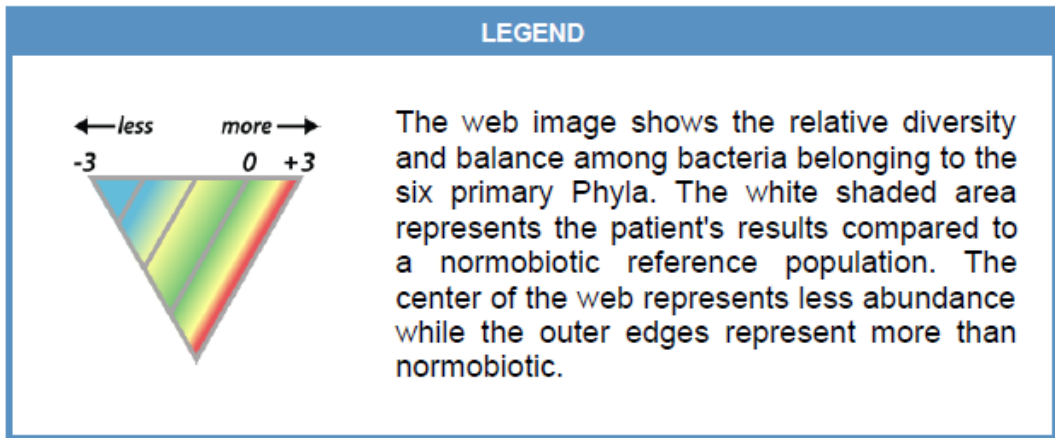
Short Chain Fatty Acids	Result	Unit	L	WRI	H	Reference Interval
% Acetate [‡]	66					50 – 72
% Propionate [‡]	16					11 – 25
% Butyrate [‡]	15					11 – 32
% Valerate [‡]	4.2					0.8 – 5.0
Butyrate [‡]	0.62	mg/mL				0.8 – 4.0
Total SCFA's [‡]	4.3	mg/mL				5.0 – 16.0

Microbiome Assessment Summary

How to Identify Stage 1

Microbiome Abundance and Diversity Summary

The abundance and diversity of gastrointestinal bacteria provide an indication of gastrointestinal health, and gut microbial imbalances can contribute to dysbiosis and other chronic disease states. The GI360™ Microbiome Profile is a gut microbiota DNA analysis tool that identifies and characterizes more than 45 targeted analytes across six Phyla using PCR and compares the patient results to a characterized normobiotic reference population. The web chart illustrates the degree to which an individual's microbiome profile deviates from normobiosis.



How to Identify Stage 2

GI Organs Impacted

- Evidence of poor digestion/absorption
- Degradation of immune response

How to Identify Stage 2

Markers of Digestion

Digestion Absorption	Result	Unit	L	WRI	H	Reference Interval
Elastase	83	µg/mL				> 200
Fat Stain	Few					None – Few
Carbohydrates [†]	Negative					Negative

How to Identify Stage 2

Markers of Immune Function

Inflammation	Result	Units	L	WRI	H	Reference Interval
Lactoferrin	8.1	µg/mL				< 7.3
Calprotectin*	59	µg/g				≤ 50
Lysozyme*	452	ng/mL				≤ 600
Immunology	Result	Units	L	WRI	H	Reference Interval
Secretory IgA*	48	mg/dL				51 – 204

How to Identify Stage 3

GI Pathogens

- Infection
 - Weakened microbiome and organ dysfunction
 - Pre-existing infection but flare up occurred due to stress
 - New infection acquired due to stress physiology and weakened immune response
- Bacterial, parasitic or yeast dominate, requires killing of bugs

Markers of Dysbiosis and Infection

How to Identify Stage 3

Parasites	Result	
<i>Cryptosporidium</i> (<i>C. parvum</i> and <i>C. hominis</i>)	Positive	<input checked="" type="checkbox"/>
<i>Entamoeba histolytica</i>	Negative	<input type="checkbox"/>
<i>Giardia duodenalis</i> (AKA <i>intestinalis</i> & <i>lamblia</i>)	Positive	<input checked="" type="checkbox"/>

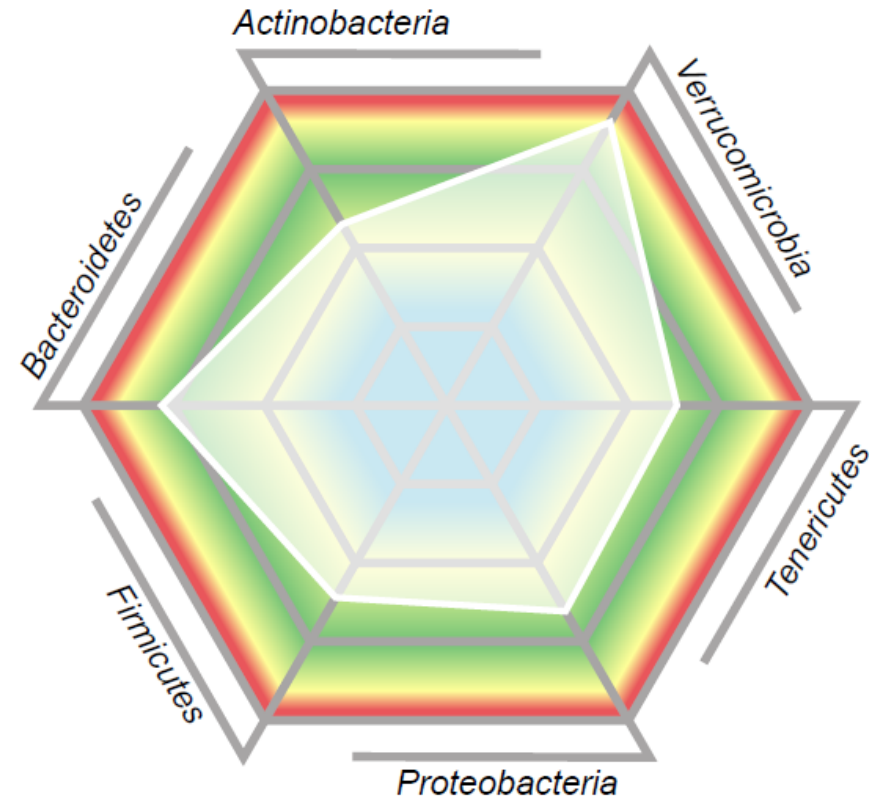
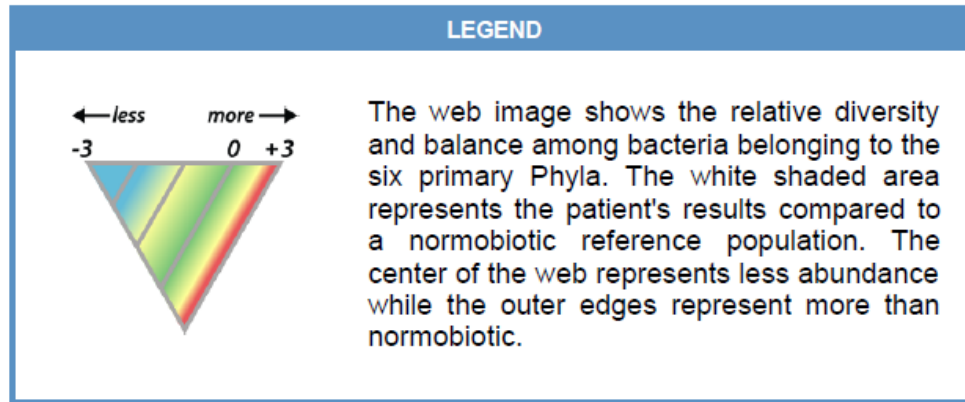
Viruses, pathogenic bacteria, parasites, protozoa, worms, yeast, etc.

A Summary of the 3 Stages – What to look for

1 Stage	<ul style="list-style-type: none">• Evidence of disturbed commensal microbiota• Low short chain fatty acids
2 Stage	<ul style="list-style-type: none">• Evidence of poor digestion/absorption<ul style="list-style-type: none">- Elastase, Fat stain, Carbohydrates• Degradation of immune response<ul style="list-style-type: none">- Lactoferrin, Calprotectin, Lysozyme, sIgA
3 Stage	<ul style="list-style-type: none">• Evidence of parasites, dysbiotic bacteria and/or yeast

Microbiome Abundance and Diversity Summary

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Dysbiosis Index

The Dysbiosis Index (DI) is a calculation with scores from 1 to 5 based on the overall bacterial abundance and profile within the patient's sample as compared to a reference population. Values above 2 indicate a microbiota profile that differs from the defined normobiotic reference population (i.e., dysbiosis). The higher the DI above 2, the more the sample is considered to deviate from normobiosis.

DI Score

4





Key Findings

<i>Bifidobacterium</i> spp.,	↑	Cryptosporidium (<i>C. parvum</i> and <i>C. hominis</i>), Detected	←	Butyrate, Low	←
Clostridia Class, Very Low	↓	Giardia duodenalis (AKA <i>intestinalis</i> & <i>lamblia</i>), Detected	←	Total SCFA's, Very Low	←
<i>Lactobacillus</i> spp.,	↑	WBC, Abnormal			
		Color, Abnormal			
		Consistency, Abnormal			
		Lactoferrin, High	←		
		Calprotectin, High	←		
		% Propionate, High			
		% Butyrate, Low	←		
		% Valerate, High			

Key Findings // Practice Test // Stage 1

Short Chain Fatty Acids [‡]	Result	Units	L	WRI	H	Reference Interval
% Acetate	59.7	%				40 – 75
% Propionate	26.5	%				9 – 29
% Butyrate	10.2	%				9 – 37
% Valerate	6.5	%				0.5 – 7
Butyrate	0.74	mg/mL				0.8 – 4.8
Total SCFA's	3	mg/mL				4 – 18

Key Findings // Practice Test // Stage 2

Inflammation	Result	Units	L	WRI	H	Reference Interval
Lactoferrin	8.1	µg/mL				< 7.3
Calprotectin*	59	µg/g				≤ 50
Lysozyme*	452	ng/mL				≤ 600
Immunology	Result	Units	L	WRI	H	Reference Interval
Secretory IgA*	48	mg/dL				51 – 204

*This test was developed and its performance characteristics determined by Doctor's Data Laboratories in a manner consistent with CLIA requirements. The U. S. Food and Drug Administration (FDA) has not approve or cleared this test; however, FDA clearance is not currently required for clinical use. The results are not intended to be used as a sole means for clinical diagnosis or patient management decisions.



Key Findings // Practice Test // Stage 3

Parasites	Result	
<i>Cryptosporidium</i> (<i>C. parvum</i> and <i>C. hominis</i>)	Positive	<input checked="" type="checkbox"/>
<i>Entamoeba histolytica</i>	Negative	<input type="checkbox"/>
<i>Giardia duodenalis</i> (AKA <i>intestinalis</i> & <i>lamblia</i>)	Positive	<input checked="" type="checkbox"/>

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Summary of Key Findings

<p>1 Stage</p>	<ul style="list-style-type: none">• Low butyrate• Low short chain fatty acids
<p>2 Stage</p>	<ul style="list-style-type: none">• High Lactoferrin• High Calprotectin• Low sIgA
<p>3 Stage</p>	<ul style="list-style-type: none">• <i>Giardia duodenalis</i> (AKA <i>intestinalis</i> & <i>lamblia</i>)• <i>Cryptosporidium</i> (<i>C. parvum</i> and <i>C. hominis</i>)

Stage-Specific Recommendations

Stage 1 – Microbiome

Eat to encourage commensal bacterial growth and SCFA production (fiber, polyphenols, fermented foods).

- Prebiotics capsules or powders
 - Polyphenol and prebiotics on their own or in combinations to boost bifidobacteria and microbiome diversity‡
- Probiotics
 - To promote GI and immune health across all ages‡
- Butyrate
 - Short chain fatty acid liquid that promotes gut barrier integrity, cytokine balance, bowel motility, and abdominal comfort‡



Stage 2 – Organ Function

Eat to stimulate normal digestive processes: chew food properly, avoid liquid with meals and relax before and after meals.

- Betaine HCl Pepsin
 - Promotes healthy digestive function and may help stimulate digestive function and absorb nutrients[‡]
- Digestive Enzymes
 - Digestive enzymes to support protein, carbohydrate, fat, fiber digestion[‡]
- GI Repair powders
 - Glutamine based powders to enhance immune cell function and support GI barrier integrity[‡]
- GB Support
 - Promote healthy gallbladder function and lipid utilization in addition to general digestive function[‡]



Stage 3 – Microbial balance

Consider introducing a low FODMAP or gluten-free diet.

- Anti-Microbial Combination Products
 - Combination of phytonutrients designed to support healthy gastrointestinal tract function and microbial balance
- Anti-Microbial Single Ingredient products
 - Options include oregano, artemisia, berberine, grapefruit seed extract, morinda, black walnut



So what Protocol should you use?

Protocol 1 // Protocol 2 // Protocol 3

It depends...

- Practitioners experience
- Severity of patient symptoms
- Cost
- Patient preference
- Time



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Thank You



Daniel Kalish, DC

Founder of The Kalish Institute



Cases

Case #1

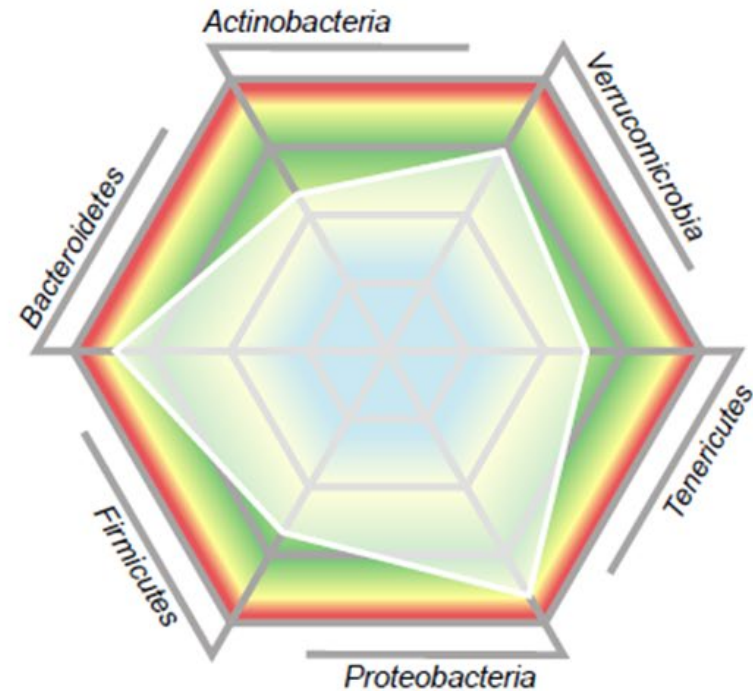
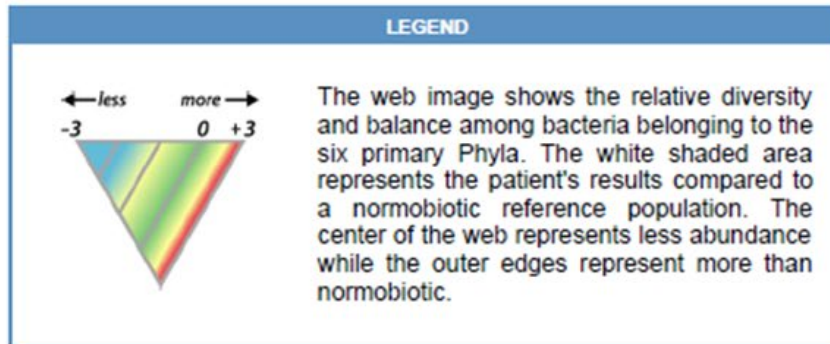
- Previously overweight depressed in the care of a psychiatrist
- Within last 3 months switched to less processed diet and lost 20lbs.
- Treated with three rounds of antibiotics for H. pylori



Case #1

Microbiome Abundance and Diversity Summary

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DI Score

5



Case #1



Key Findings

<i>Dialister invisus</i> & <i>Megasphaera micronuciformis</i> , Very High	↑	Fat Stain, High
<i>Parabacteroides</i> spp., Very High	↑	% Propionate, High
<i>Eubacterium siraeum</i> , Very Low	↓	% Valerate, Very Low
<i>Faecalibacterium prausnitzii</i> , Very Low	↓	Secretory IgA, High
Actinobacteria, Low	↓	
<i>Alistipes</i> spp., Low	↓	
Proteobacteria, High	↑	
<i>Alistipes onderdonkii</i> , Low	↓	
<i>Bacteroides fragilis</i> , High	↑	
<i>Escherichia</i> spp., High	↑	



Case #1

<i>Firmicutes</i>	Result	-3	-2	-1	0	+1	+2	+3	Reference Interval
<i>Dialister invisus</i>	+2						▲		0 to +1
<i>Dialister invisus</i> & <i>Megasphaera micronuciformis</i>	+3							▲	0 to +1
<i>Dorea</i> spp.	0				▲				0 to +1
<i>Eubacterium bifforme</i>	0				▲				0 to +1
<i>Eubacterium hallii</i>	-1			▲					-1 to +1
<i>Eubacterium rectale</i>	0				▲				0 to +1
<i>Eubacterium siraeum</i>	-3	▲							-1 to +1
<i>Faecalibacterium prausnitzii</i>	-3	▲							-1 to +1
Lachnospiraceae	-1			▲					-1 to +1
<i>Lactobacillus ruminis</i> & <i>Pediococcus acidilactici</i>	0				▲				0 to +1
<i>Lactobacillus</i> spp.	0				▲				0 to +1
<i>Phascolarctobacterium</i> spp.	0				▲				0 to +1
<i>Ruminococcus albus</i> & <i>R. bromii</i>	0				▲				0 to +1
<i>Ruminococcus gnavus</i>	+2						▲		0 to +1
<i>Streptococcus agalactiae</i> & <i>Eubacterium rectale</i>	0				▲				0 to +1
<i>Streptococcus salivarius</i> ssp. <i>thermophilus</i> & <i>S. sanguinis</i>	0				▲				0 to +1
<i>Streptococcus salivarius</i> ssp. <i>thermophilus</i>	-1			▲					-1 to +1
<i>Streptococcus</i> spp.	0				▲				0 to +1
<i>Veillonella</i> spp.	+2						▲		-1 to +1



Case #1

Digestion Absorption	Result	Unit	L	WRI	H	Reference Interval
Elastase	>500	µg/mL				> 200
Fat Stain	Moderate					None – Few
Carbohydrates [†]	Negative					Negative

Inflammation	Result	Unit	L	WRI	H	Reference Interval
Lactoferrin	1.1	µg/mL				< 7.3
Lysozyme*	248	ng/mL				< 500
Calprotectin*	<10	µg/g				< 50

Immunology	Result	Unit	L	WRI	H	Reference Interval
Secretory IgA*	329	mg/dL				30 – 275

Short Chain Fatty Acids [†]	Result	Unit	L	WRI	H	Reference Interval
% Acetate	61	%				50 – 72
% Propionate	26	%				11 – 25
% Butyrate	13	%				11 – 32
% Valerate	0.2	%				0.8 – 5.0
Butyrate	1.6	mg/mL				0.8 – 4.0
Total SCFA's	12	mg/mL				5.0 – 16.0

Intestinal Health Markers	Result	Unit	L	WRI	H	Reference Interval
pH	5.8					5.8 – 7.0
Occult Blood	Negative					Negative



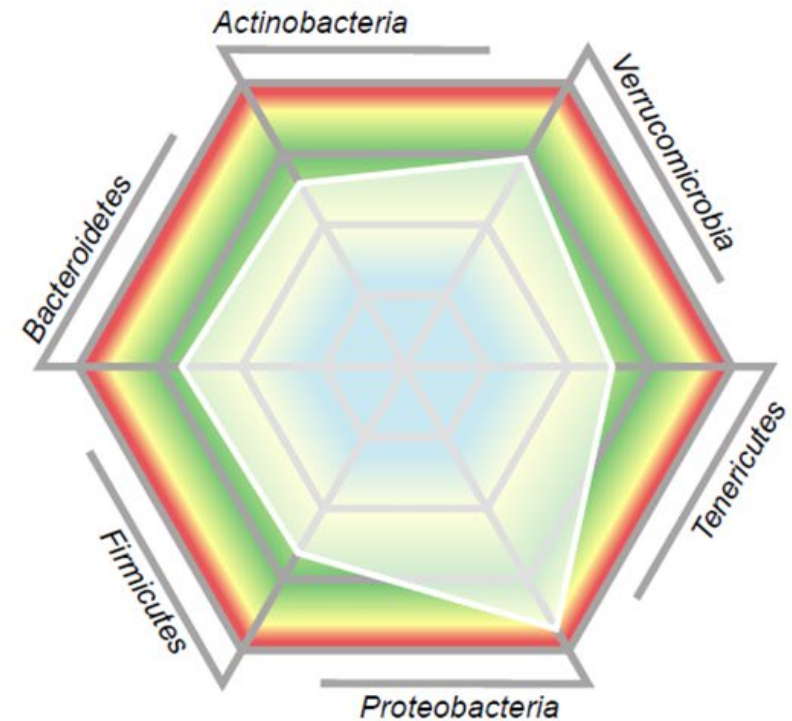
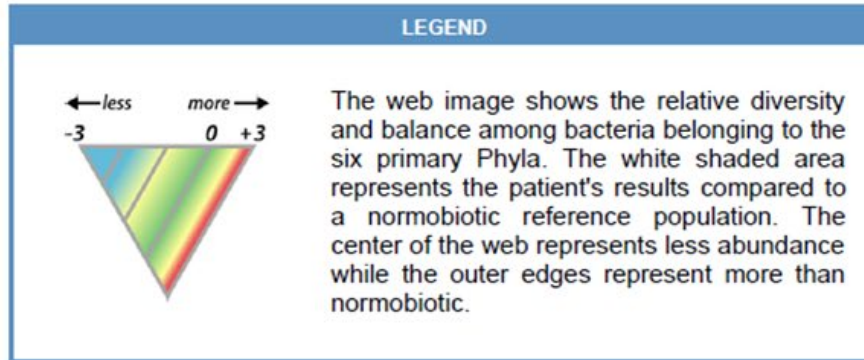
Case #2

- Client presents with ongoing arthralgia, underactive thyroid and alternating IBS symptoms, which has worsened considerably since a trip to Hong Kong in Oct 2019.
- She's taking pain medications (morphine x 2, gabapentin, NSAIDs, paracetamol) having had a bike accident June 2019

Case #2

Microbiome Abundance and Diversity Summary

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DI Score

5



Case #2



Key Findings

Proteobacteria, Very High



Salmonella spp., Detected

Clostridia Class, Very Low



Giardia duodenalis (AKA *intestinalis* & *lamblia*), Detected

Streptococcus spp., Very High



Citrobacter freundii complex, Cultured

Bacilli Class, High



Klebsiella pneumoniae, Cultured

Eubacterium siraeum, Low



Geotrichum spp., Cultured

Lachnospiraceae, Low



Lactoferrin, Very High

Streptococcus salivarius ssp. *thermophilus*, High



Secretory IgA, High

Veillonella spp., Low



Case #2

Viruses	Result	
Adenovirus F40/41	Negative	<input type="checkbox"/>
Norovirus GI/GII	Negative	<input type="checkbox"/>
Rotavirus A	Negative	<input type="checkbox"/>
Pathogenic Bacteria	Result	
<i>Campylobacter</i> (<i>C. jejuni</i> , <i>C. coli</i> and <i>C. lari</i>)	Negative	<input type="checkbox"/>
<i>Clostridium difficile</i> (Toxin A/B)	Negative	<input type="checkbox"/>
<i>Escherichia coli</i> O157	Negative	<input type="checkbox"/>
Enterotoxigenic <i>Escherichia coli</i> (ETEC) lt/st	Negative	<input type="checkbox"/>
<i>Salmonella</i> spp.	Positive	<input checked="" type="checkbox"/>
Shiga-like toxin-producing <i>Escherichia coli</i> (STEC) stx1/stx2	Negative	<input type="checkbox"/>
<i>Shigella</i> (<i>S. boydii</i> , <i>S. sonnei</i> , <i>S. flexneri</i> & <i>S. dysenteriae</i>)	Negative	<input type="checkbox"/>
<i>Vibrio cholerae</i>	Negative	<input type="checkbox"/>
Parasites	Result	
<i>Cryptosporidium</i> (<i>C. parvum</i> and <i>C. hominis</i>)	Negative	<input type="checkbox"/>
<i>Entamoeba histolytica</i>	Negative	<input type="checkbox"/>
<i>Giardia duodenalis</i> (AKA <i>intestinalis</i> & <i>lamblia</i>)	Positive	<input checked="" type="checkbox"/>

Case #2

Pathogenic Bacteria	Result	NG	1+	2+	3+	4+	Reference Interval
<i>Aeromonas</i> spp.	NG	▲					No Growth
<i>Edwardsiella tarda</i>	NG	▲					No Growth
<i>Plesiomonas shigelloides</i>	NG	▲					No Growth
<i>Salmonella</i> group	NG	▲					No Growth
<i>Shigella</i> group	NG	▲					No Growth
<i>Vibrio cholerae</i>	NG	▲					No Growth
<i>Vibrio</i> spp.	NG	▲					No Growth
<i>Yersinia</i> spp.	NG	▲					No Growth
Imbalance Bacteria	Result	NG	1+	2+	3+	4+	Reference Interval
<i>Hafnia alvei</i>	4+					▲	No Growth
<i>Proteus vulgaris</i> group	2+			▲			No Growth
Dysbiotic Bacteria	Result	NG	1+	2+	3+	4+	Reference Interval
<i>Citrobacter freundii</i> complex	4+					▲	No Growth
<i>Klebsiella pneumoniae</i>	3+				▲		No Growth
Yeast	Result	NG	1+	2+	3+	4+	Reference Interval
<i>Geotrichum</i> spp.	4+					▲	0+ – 1+



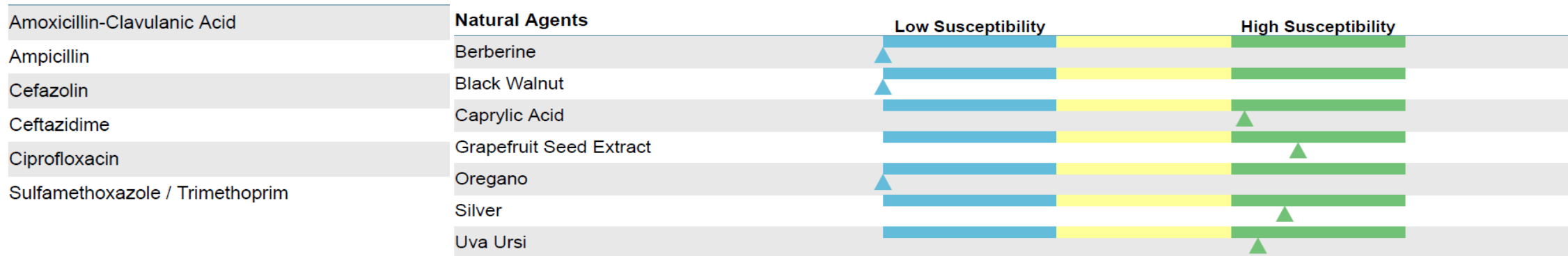
Citrobacter freundii complex

Natural Agents

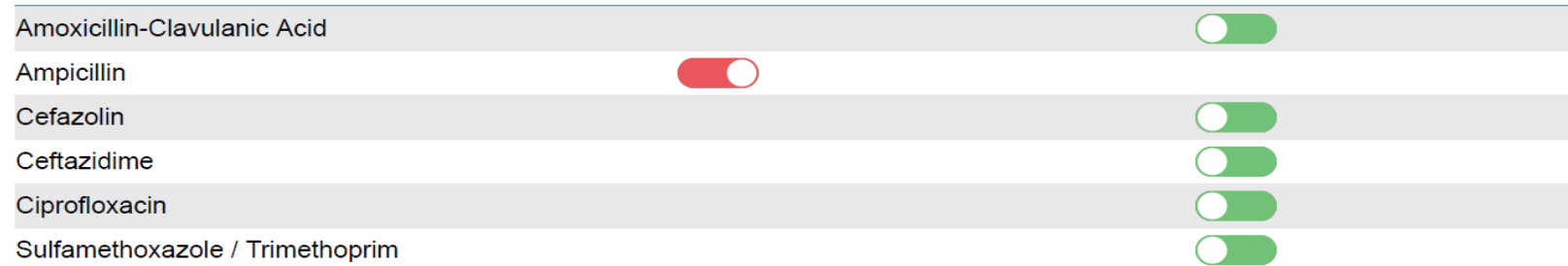


Case #2

Prescriptive Agents



Prescriptive Agents



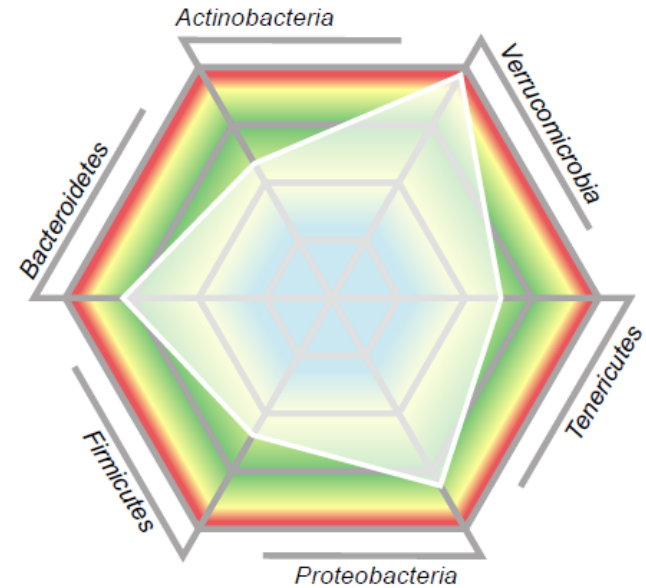
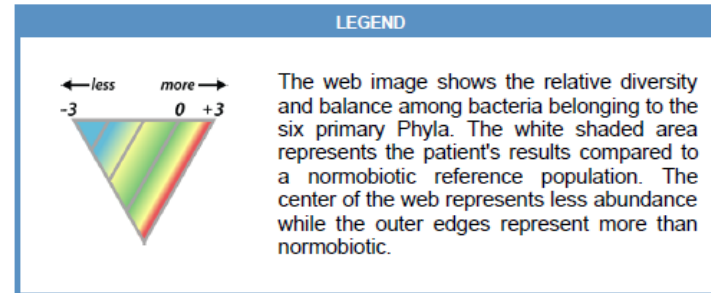
Case #3

- Pt on Ketogenic diet
 - Pt is WNL weight prior to gestation
 - No GI history
 - Develops gestational Type II DM
-
- Avoiding all carbs in an effort to destroy dyslglycemia

Case #3

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DI Score

5



Key Findings

<i>Akkermansia muciniphila</i> , Very High	↑	<i>Candida albicans</i> , Cultured
<i>Faecalibacterium prausnitzii</i> , Very Low	↓	pH, High
<i>Actinobacteria</i> , Low	↓	% Propionate, High
Bacilli Class, Low	↓	Butyrate, Low
Lachnospiraceae, Low	↓	Total SCFA's, Very Low
<i>Veillonella</i> spp., Low	↓	Secretory IgA, Low



Case #3

Digestion Absorption	Result	Unit	L	WRI	H	Reference Interval
Elastase	286	µg/mL		▲		> 200
Fat Stain	None		▲			None – Few
Carbohydrates†	Negative			▲		Negative
Inflammation	Result	Unit	L	WRI	H	Reference Interval
Lactoferrin	2.3	µg/mL		▲		< 7.3
Lysozyme*	113	ng/mL		▲		≤ 500
Calprotectin	<10	µg/g	▲			≤ 50
Immunology	Result	Unit	L	WRI	H	Reference Interval
Secretory IgA*	28.2	mg/dL		▲		30 – 275
Short Chain Fatty Acids†	Result	Unit	L	WRI	H	Reference Interval
% Acetate	53	%		▲		50 – 72
% Propionate	26	%			▲	11 – 25
% Butyrate	17	%		▲		11 – 32
% Valerate	4.1	%			▲	0.8 – 5.0
Butyrate	0.59	mg/mL		▲		0.8 – 4.0
Total SCFA's	3.4	mg/mL	▲			5.0 – 16.0
Intestinal Health Markers	Result	Unit	L	WRI	H	Reference Interval
pH	7.1				▲	5.8 – 7.0
β-glucuronidase	298	U/L		▲		100 – 1200
Occult Blood	Negative			▲		Negative

Case #4

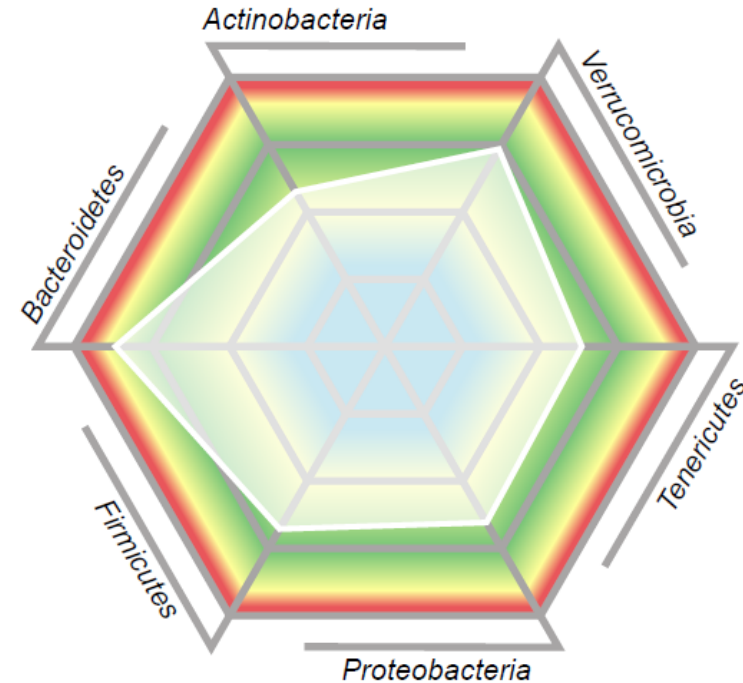
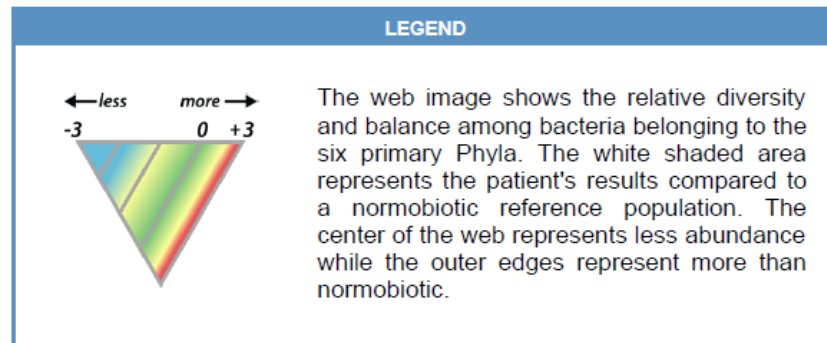
- Client presents with GI distress
- Pt is on self prescribed carnivore diet to avoid plant reactions



Case #4

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Dysbiosis Index

The Dysbiosis Index (DI) is a calculation with scores from 1 to 5 based on the overall bacterial abundance and profile within the patient's sample as compared to a reference population. Values above 2 indicate a microbiota profile that differs from the defined normobiotic reference population (i.e., dysbiosis). The higher the DI above 2, the more the sample is considered to deviate from normobiosis.

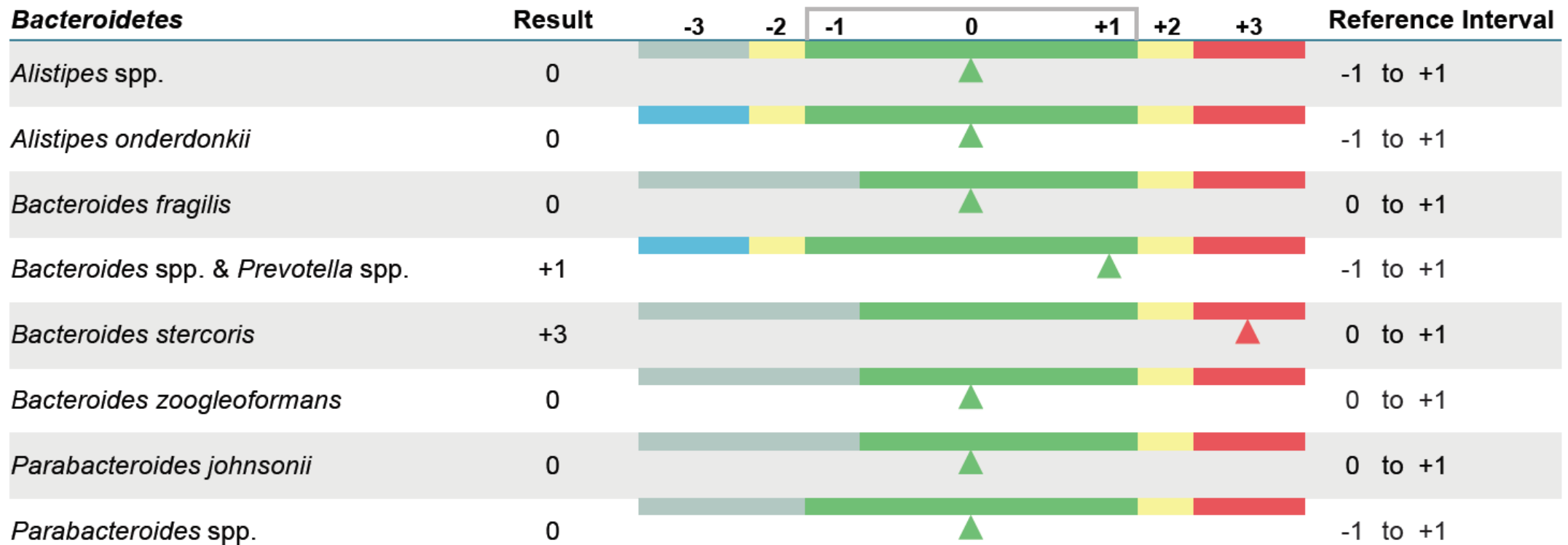
DI Score

4



Case #4

Bacteroides stercoris



Case #4

Digestion Absorption	Result	Unit	L	WRI	H	Reference Interval
Elastase	239	µg/mL				> 200
Fat Stain	None					None – Few
Carbohydrates [†]	Negative					Negative
Inflammation	Result	Unit	L	WRI	H	Reference Interval
Lactoferrin	5.3	µg/mL				< 7.3
Lysozyme*	368	ng/mL				≤ 500
Calprotectin	<10	µg/g				≤ 50
Immunology	Result	Unit	L	WRI	H	Reference Interval
Secretory IgA*	80.0	mg/dL				30 – 275
Short Chain Fatty Acids	Result	Unit	L	WRI	H	Reference Interval
% Acetate [‡]	66	%				50 – 72
% Propionate [‡]	16	%				11 – 25
% Butyrate [‡]	15	%				11 – 32
% Valerate [‡]	4.2	%				0.8 – 5.0
Butyrate [‡]	0.62	mg/mL				0.8 – 4.0
Total SCFA's [‡]	4.3	mg/mL				5.0 – 16.0
Intestinal Health Markers	Result	Unit	L	WRI	H	Reference Interval
pH	7.0					5.8 – 7.0
β-glucuronidase*	424	U/L				100 – 1200
Occult Blood	Positive					Negative



Thank you for watching



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For more information:

doctorsdata.com

GI360.com

800.323.2784

info@doctorsdata.com