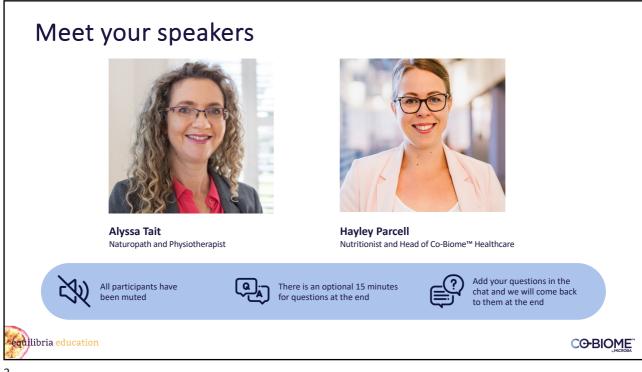
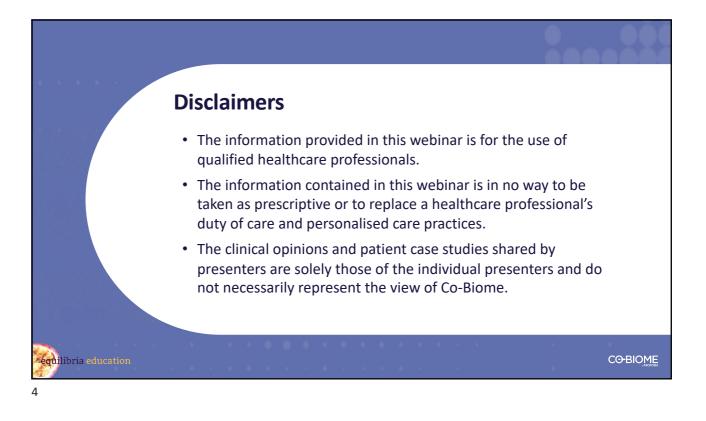
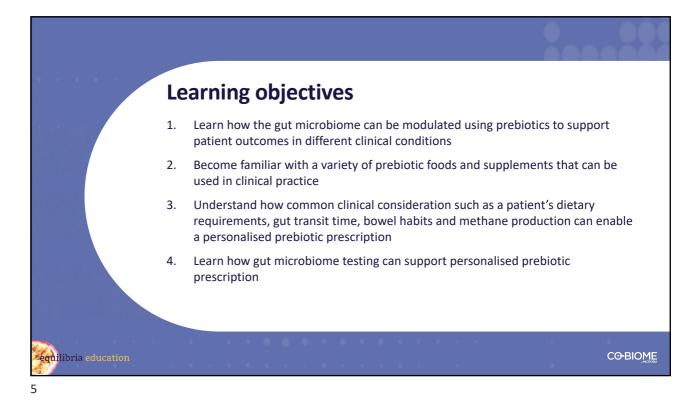
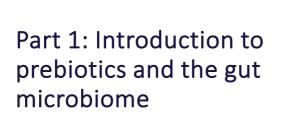
Personalised Prebiotic Prescription:				
Optimising Gut Microbiome Health with Prebiot	ics			
Tuesday 13 February 2024 7PM AEDT				

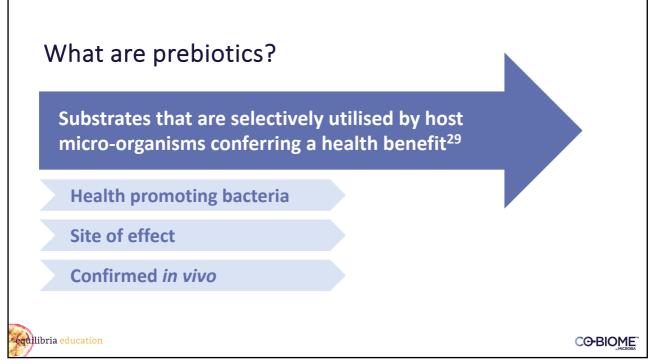




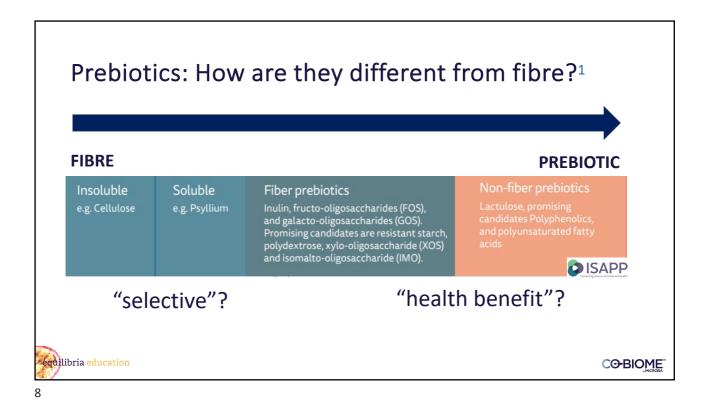


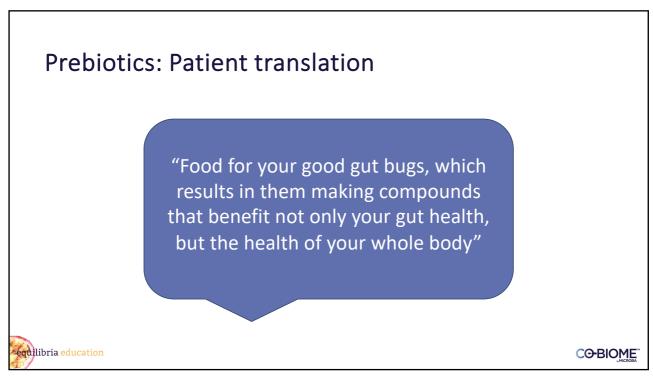


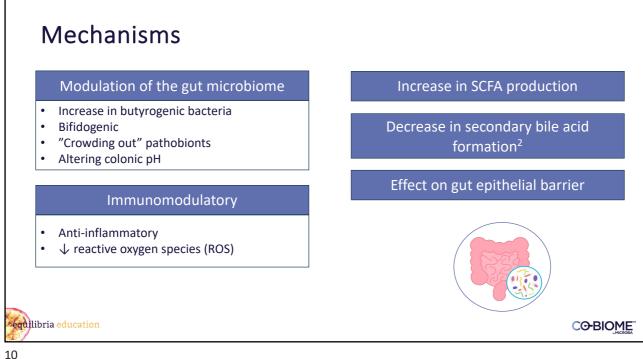




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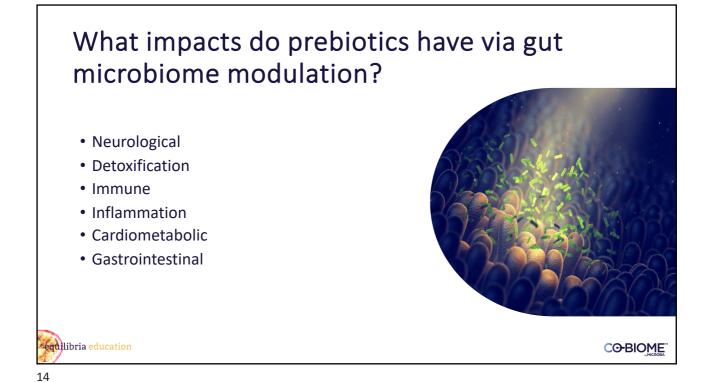
Prebiotics: Impact on gut microbiome 3,4,5,6,7,8,9

Prebiotic	Influence on microbial metabolites	Influence on microbial taxa
Inulin (medium/long- chain)	↑ methane production* ↓ faecal beta-glucuronidase activity*	 ↑ Bifidobacterium spp.^{3,4,5} ↑ Faecalibacterium spp.³ ↑ Akkermansia muciniphila⁶ ↑ Veillonella parvula/ atypica⁷ ↓ Bilophila spp^{4,5}
FOS and short-chain inulin	↓hydrogen sulphide production*	 ↑ Bifidobacterium spp.^{8,9} ↑ Faecalibacterium spp.^{8,9} ↓ Ruminococcus/ Faecalicatena spp.^{8,9}
ibria education		* Co-Biome Insight

Dat beta-glucan ↑ butyrate-production* ↑ Bifidobacterium spp. ^{16.17}	Prebiotic	Influence on metabolite production via microbial populations	Other microbiome impacts
\wedge Faecalibacterium spp. 15Dat beta-glucan \wedge butyrate-production* \wedge Bifidobacterium spp. 16.17	GOS	\downarrow faecal beta-glucuronidase activity*	↑ Anaerostipes spp ^{11,12}
	PHGG	\uparrow butyrate-producing microbes ¹⁰	
	Dat beta-glucan	↑ butyrate-production*	↑ Bifidobacterium spp. ^{16.17}
Resistant starch type 2		↑ acetate production*	↑ Ruminococcus_E bromii ¹⁸

Prebiotics: Impact on gut microbiome

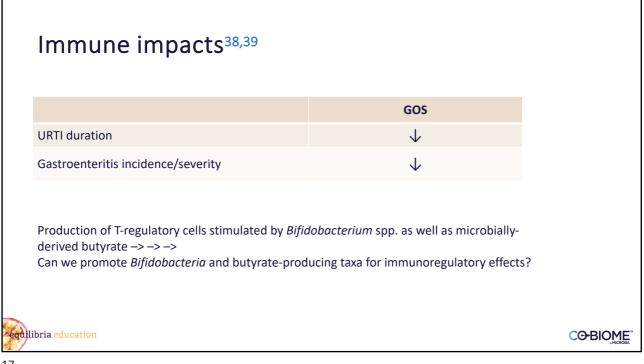
	Influence on metabolite production via microbial populations	Other microbiome impacts
Glucomannan	\downarrow faecal beta-glucuronidase activity*	↑ Bifidobacterium spp. ²⁴
Acacia gum	\uparrow butyrate production ¹⁹	↑ <i>Bifidobacterium</i> spp. ^{25,26}
Lactulose	 ↑ propionate production* ↓ colonic production and absorption of ammonia^{20,21,22} ↓ faecal beta-glucuronidase²³ 	 ↑ Bifidobacterium spp.²⁷ ↓ Faecalibacterium²⁸ ↓ Enterococcus²⁷ ↓ Escherichia-Shigella²⁹

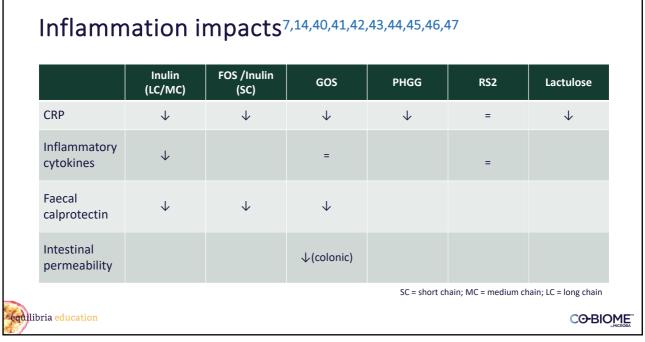


Prebiotic	Condition	Impact	Associated gut microbiome effects
Inulin (MC)	Obesity	Moderate ↑ mood, cognition	↑ Bifidobacterium spp,. ↑ response in those with个 baseline Coprococcus
Inulin (MC)	Alcohol use disorder	个serum BDNF (brain derived neurotrophic factor)	 ↑ Bifidobacterium spp., Faecalibacterium spp. ↓ Dorea, F. torques, E. ruminantium
GOS 6.9g/d	Anxious young females	No change in anxiety	↑ Bifidobacterium spp, Bacteroides spp ↓ Clostridium spp
	IBS	?↓depression	
Indirect res	earch / areas to v	watch	MC = medium chai

Detoxification	impacts7,23,24,32,33,34,35,36,37
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istant starch 2 (RS2) · · · · · · · · · · · · · · · · · · ·	esistant starch pe 2 (RS2) \checkmark RS2 \checkmark	rebiotic	↓Faecal beta- glucuronidase	Alters bile acid metabolism	↓Blood urea nitrogen	↓Colonic production/absorption ammonia
a 2 (RS2) Image: Company of the second sec	pe 2 (RS2)Image: Constraint of the second of th	ulin (MC/LC)	\checkmark	\checkmark	\checkmark	
sulose \checkmark \checkmark \checkmark	Account Image: Constraint of the second of	esistant starch /pe 2 (RS2)		\checkmark	\checkmark	
\sim	DS ✓ ✓ ✓ ✓ Instance of the second se	lucomannan	\checkmark			
	HGG ✓	actulose	\checkmark			\checkmark
GG ✓		OS	\checkmark	\checkmark		
	MC = medium chain; LC = long cha	HGG		\checkmark		
MC = medium chain; LC = long cha						MC = medium chain; LC = long cha

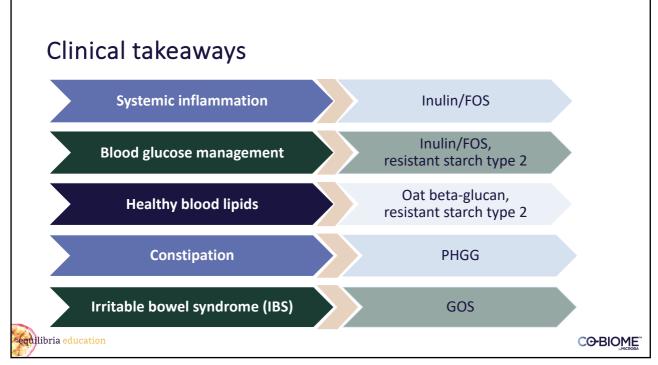


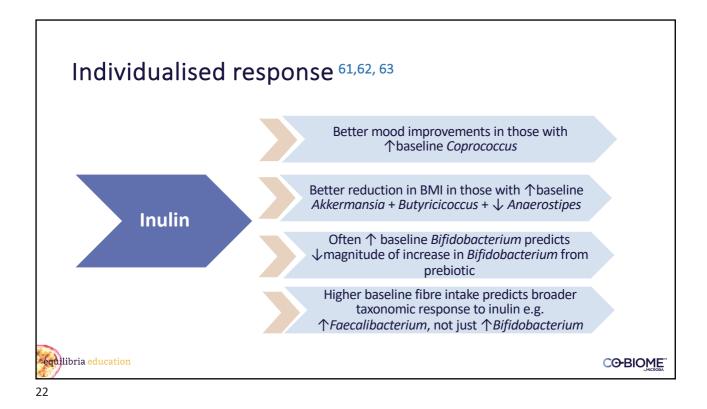


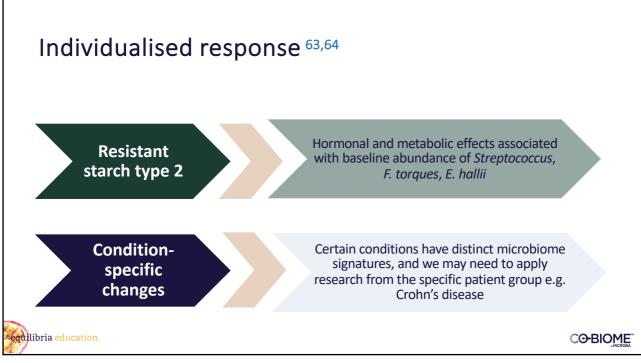
Cardiometabolic impacts^{41,43,48,49,50,51,52,53,54,55,56}

	Inulin (LC/MC)	FOS/Inulin (SC)	PHGG	Oat BG	RS2	Acacia gum	Lactulose
Fasting glucose	\checkmark	\checkmark			\checkmark	\downarrow	
Fasting insulin	\checkmark	\checkmark	\downarrow		\checkmark		
HbA1c	\checkmark	\checkmark			\checkmark		
HOMA-IR	\checkmark	\checkmark			\checkmark		
NASH/steatosis		\checkmark					
TG	=		\checkmark	\checkmark	\checkmark		\checkmark
тс	=			\checkmark	\checkmark		
LDL-C	\checkmark		\checkmark	\checkmark	\checkmark		
HDL-C	=		\uparrow	=			
АроВ				\checkmark			
BMI/BW					\checkmark	\checkmark	
BP						\checkmark	
Lactate threshold		\uparrow					
oria education					SC = short chain;	MC = medium chair	; LC = long chain CO-BIC

GI IMPACTS	Inulin (LC/MC)	FOS/Inulin (SC)	GOS	PHGG	RS2	Lactulose
Transit time				\checkmark	\uparrow	
Improved stool form	\uparrow		=	\uparrow	=/个	
Incomplete emptying	=		=	=		
Straining			\checkmark	=/↓		
Flatulence	\uparrow	\uparrow	\checkmark	√<6g	\uparrow	\uparrow
Bloating	\uparrow		\checkmark	^/=/↓	=	
Abdominal pain	\uparrow		\checkmark	=/↓	=	
Diarrhoea				\checkmark	\checkmark	
				SC = short chain;	MC = medium chain,	; LC = long chain
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*/MICROBA

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Prebiotics in the diet

Prebiotic	Food sources	
Fructooligosaccharides (FOS)	Jerusalem artichoke Broccoli stalks	Red onion Watermelon
Galactooligosaccharides (GOS)	Borlotti beans Chickpeas	Green lentils Soy flour
Inulin	Jerusalem artichoke Asparagus	Globe artichoke Leek
Pectin	Mandarins Kale	Blackberries Avocado
Arabinoxylan	Pumpernickel bread Wholegrain rye	Wheat bran Popcorn
Resistant starch (RS)	Green banana Green banana flour	Barley Lentils
Beta-glucan	Oats Wholegrain barley	Wheat bran Sorghum
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Why not just a high-fibre diet, rather than using prebiotics?

ffect on gut metabolites	Evidence for high-fibre diet/foods	Evidence for supplemental prebiotics
↑ butyrate microbes	Rye bran-enriched bread (6-10 pieces/d)	RS2 with high RS foods (combination) PHGG (6g/d)
↓ BCAAs	Rye bread (8 pieces/d)	
↑ IPA production	Very high fibre diet; wholegrain wheat, rye	
↓ TMAO	Brussel sprouts (300g/day)	
↓ beta-glucuronidase enzyme	Grain/legume fibres	Glucomannan with low-fat diet (4.5 g/d) Inulin (5-20g/d) Lactulose (3g/d)
↓ hydrogen sulphide nicrobes		FOS (12g/d)
↓ hexa-LPS microbes		GOS (5.5g/d)
↑ secretory IgA		GOS (5.5g/d)
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Summary of dietary fibre vs dietary prebiotics vs supplemental prebiotics

- Considerable overlap
- Some metabolites have fibre studies but not prebiotic studies to support
- Overall, effective dietary studies use (often unrealistically) high quantities of high-prebiotic food (e.g. 6, 8, 10 pieces of bread/day)
- Non-athletes and those on restricted diets may find it difficult to meet these dietary targets



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Questions I ask before prescribing prebiotics

Part 1: IF a prebiotic is indicated

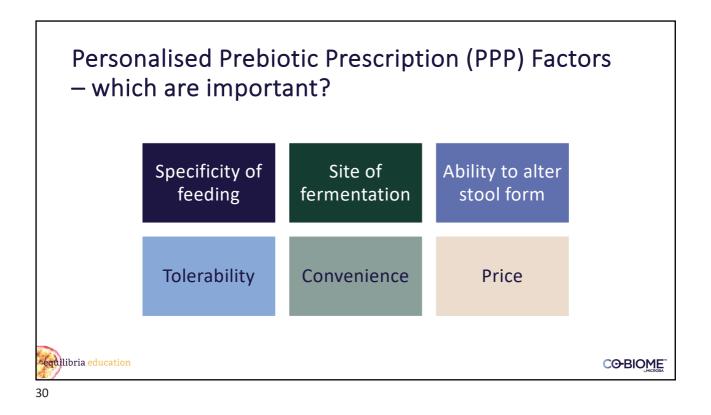
- How is a prebiotic likely to influence the clinical picture?
- Is now a good time to start a prebiotic?

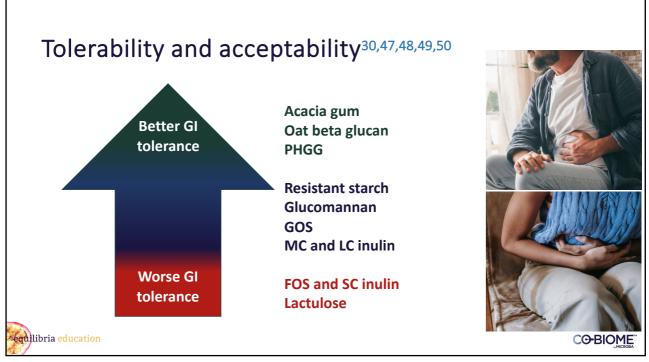
Part 2: WHICH prebiotic options we have

- Which of the six Personalised Prebiotic Prescription (PPP) factors are important here?
- In what order are they important?

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Prescribing practicalities: Flavour, texture, convenience

Prebiotic	Flavour	Texture	Convenience
Lactulose	✓ Sweet	Liquid	X Must be poured
FOS/Inulin	✓ Sweet	Soluble	✓ Mixes in food, drink
GOS	✓ Sweet	Soluble	✓ Mixes in food, drink
PHGG	✓ Neutral	Soluble	✓ Mixes in food, drink
Oat beta-glucan	Slight oat	Viscous	X Thickens, creates lumps
Resistant starch banana flour	Slight banana	Viscous	X Thickens, separates

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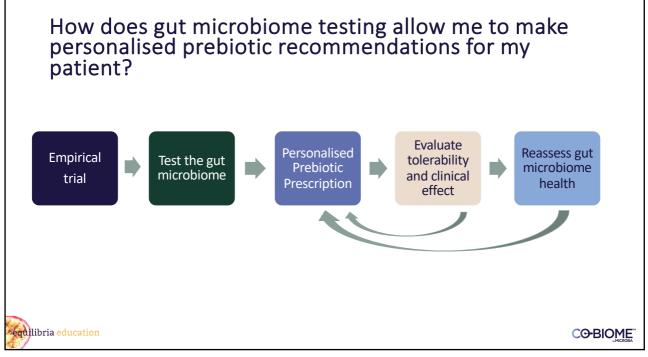
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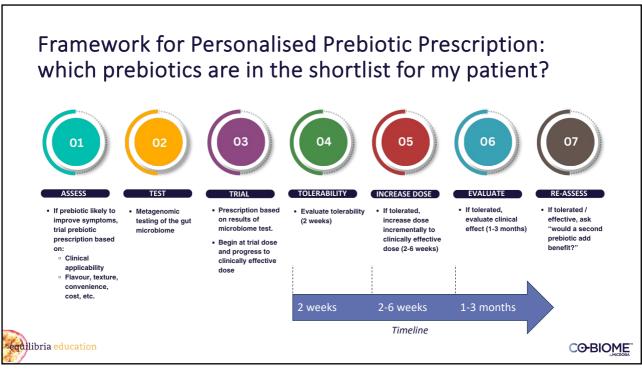
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Prescribing practicalities: Dosing guidelines

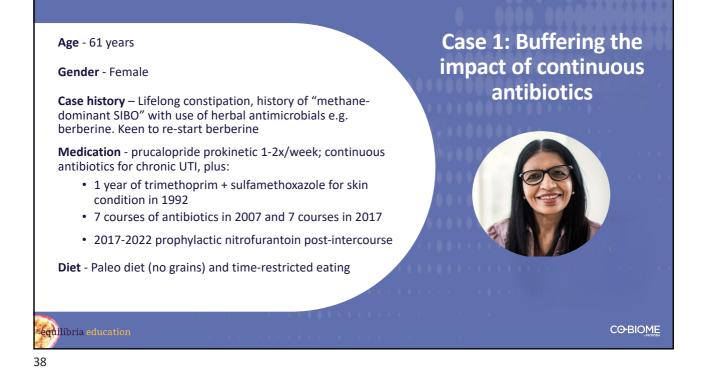
Prebiotic	Research - supported daily dosage range	Approx. equivalent	Common clinically effective dose	Approx. equivalent	References
Inulin	10g – 15g	8 tsp – 12 tsp	10g	8 tsp	7,8,31,32,33,40,42, 47,49,51,56, 58,59
FOS	7.5g – 30g	3.75 tsp – 15 tsp	10g	5 tsp	7,8,31,32,33,40,42, 47,49,51,56,58,59
GOS	3g – 11g	½ tsp – 2 tsp	5.5g	1 tsp	13,14,30,33,39,43,4 4,58,59
PHGG	5g – 20g	2.5 tsp – 10 tsp	8g	4 tsp	34,58
Oat beta-glucan	3g – 6.6g	1 ½ tsp – 3 ¼ tsp	3.5g (lipids) 10g (stool form)	1 ¾ tsp 5 tsp	50,55
RS2	10g – 40g	3 ⅓ tsp	15g	5 tsp	35,37,41,45,46,53,5 4,57,60
Glucomannan	4.5g	7 x 600mg capsules	4.5g	7x 600mg capsules	24
Lactulose	3g – 25g	5-40 mL	10g	15mL	23,48,52
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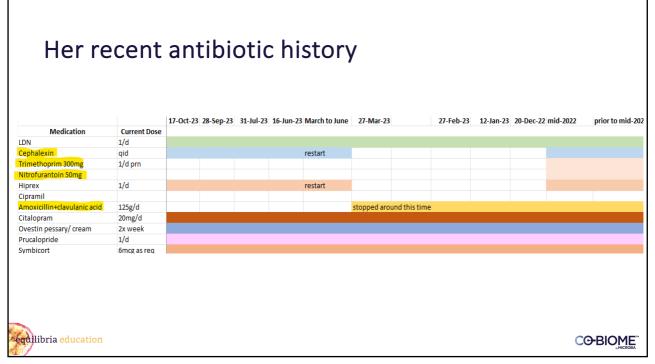
	A	A	August and former think
Prebiotic	Approx. cost/serve (low end dosage)	Approx. cost for common clinical serve	Approx. cost/serve (high end dosage)
nulin (long-chain)	\$1.20 for 10g	\$1.20 for 10g	\$1.80 for 15g
FOS/Inulin	\$0.70 for 5g	\$1.40 for 10g	\$2.80 for 20g
GOS	\$0.70 for 3g	\$1.20 for 5.5g	\$2.60 for 11g
PHGG	\$0.50 for 5g	\$0.80 for 8g	\$2.00 for 20g
Oat beta-glucan	\$0.80 for 3.5g	\$0.80 for 3.5g	\$2.30 for 10g
Resistant starch banana flour	\$1.00 for 10g	\$1.50 for 15g	\$4.00 for 40g
actulose	\$0.10 for 3g	\$0.33 for 10g	\$0.80 for 25g (\$1.60 if bd)
Glucomannan	\$2.20 for 4.5g in capsules	\$2.20 for 4.5g in capsules	\$2.20 for 4.5g in capsules



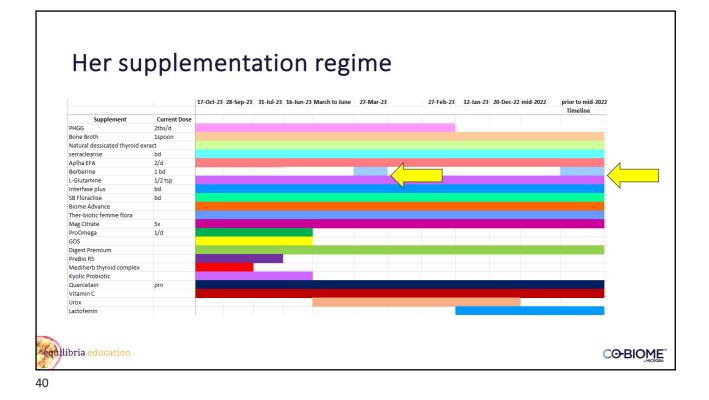


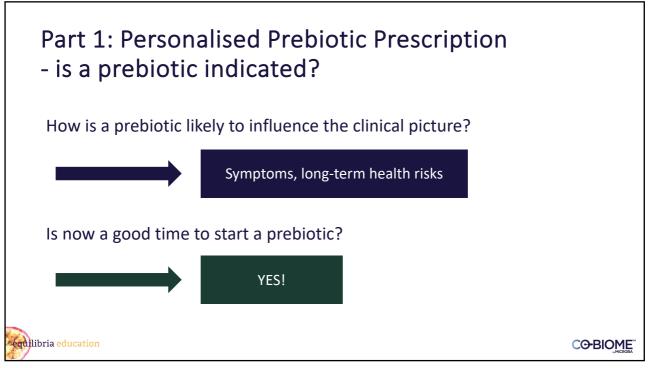


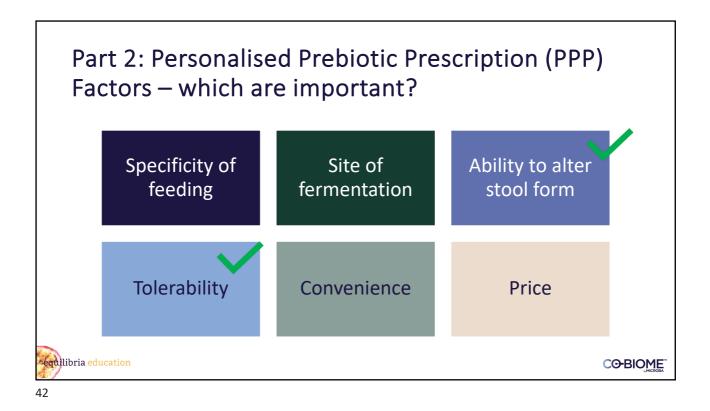






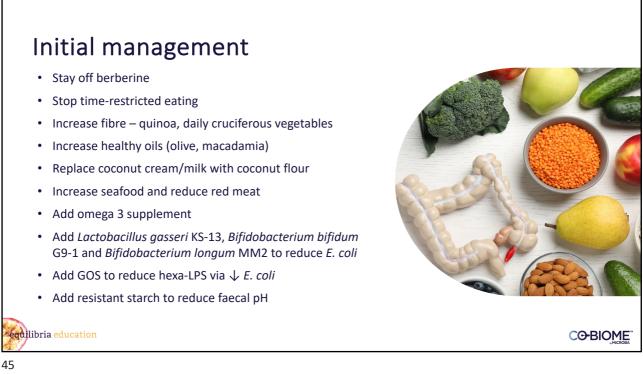




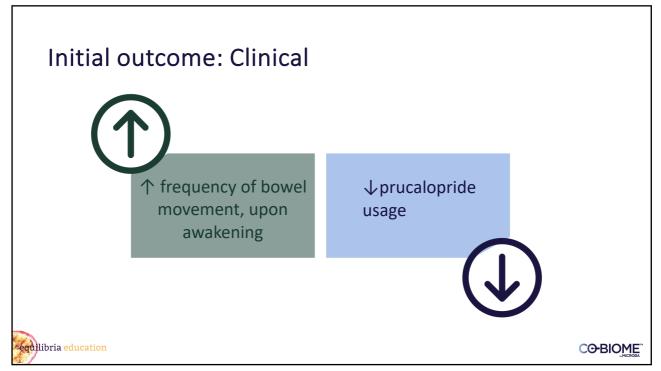


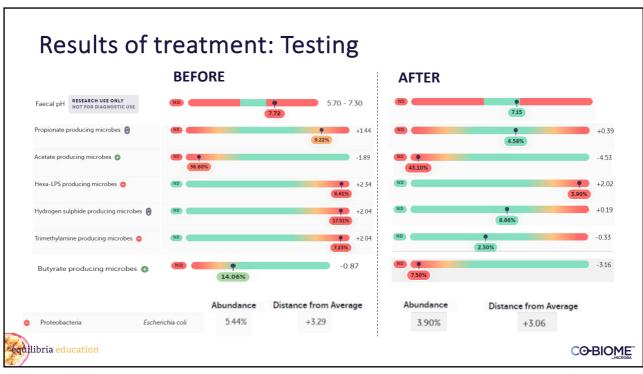


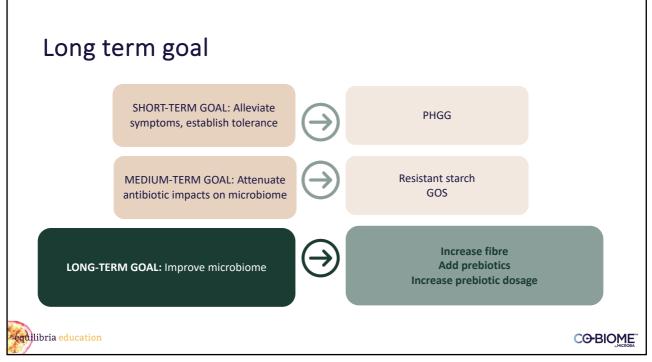
Faecal pH RESEARCH USE ONLY NOT FOR DIAGNOSTIC USE	7.72	5.70 - 7.30		
Methane producing archaea 😑	0.00%			INTESTINAL INFLAMMATION
Propionate producing microbes 🏮	ND		9.22%	+1.44
Acetate producing microbes ④	ND •	%		systemic INFLAMMATION
Hexa-LPS producing microbes 🤤	(ND)		9 - 6.61%	
Hydrogen sulphide producing microbes 🏮	ND		• 17.51%	+2.04 INTESTINAL BARRIER
Trimethylamine producing microbes 🤤	ND		7.23%	FUNCTION
e Proteobacteria Escheric	chia coli	5.44%	Less common	+3.29



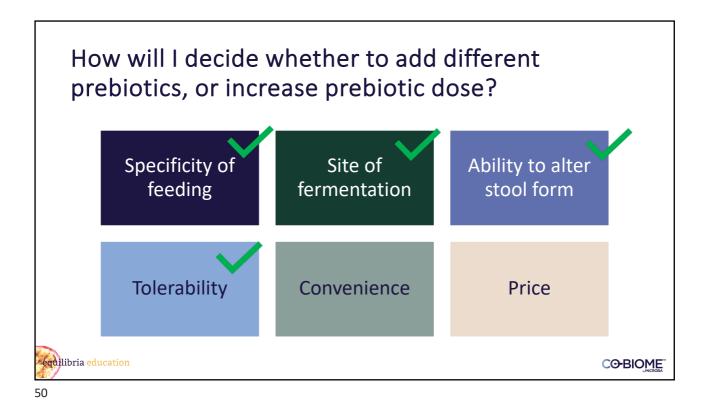






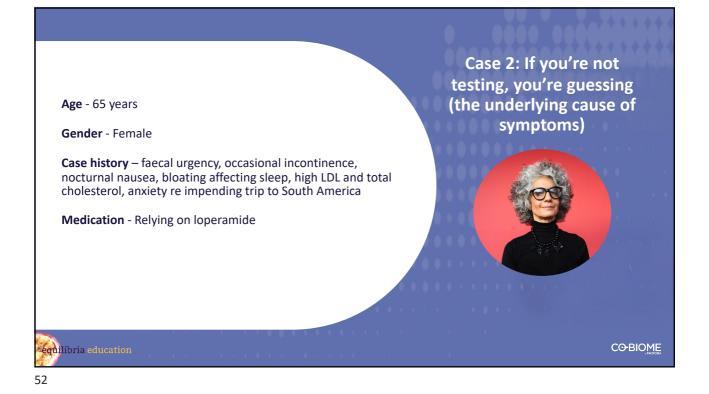


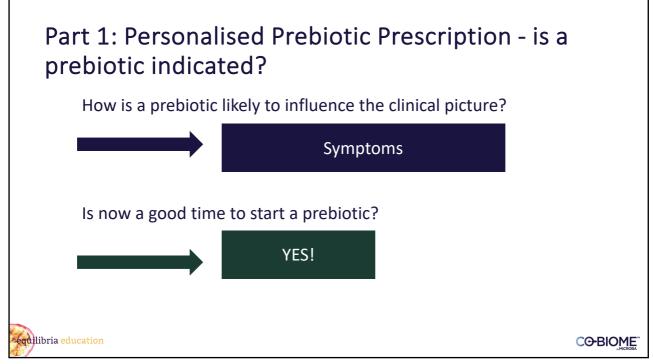




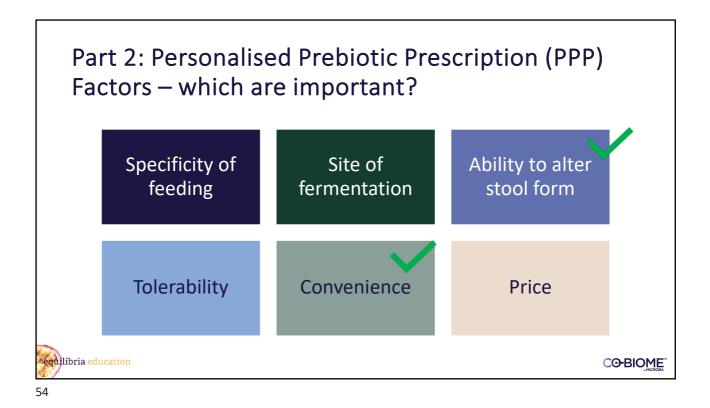
Current issu	e	Possible action		Potential problen	n	Alternative	
pH remains	high	Add lactulose		Tolerability Site of fermentati (too proximal)	on	Increase RS2 dose	
Hexa-LPS mi remain high		Add inulin or lact	ulose	Tolerability		Increase GOS dose	
↓ Acetate a microbes	nd butyrate	Add inulin		Tolerability		Increase RS2/GOS Counsel re trial off ant	ibiotics
Persistent "t viruses" aro fibre	hought und SIBO and	Gradual increase	in fibre				
	Prebiotic		Current	: dose	Dose t	to trial	
GOS Resistant stard			1 tsp (5	g)	2 tsp (10g)	
		ch banana flour	2 scoop	os (30g)	2.5 sco	oops (37.5g)	

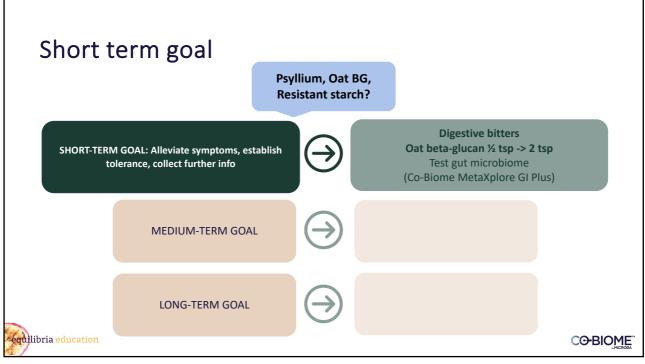


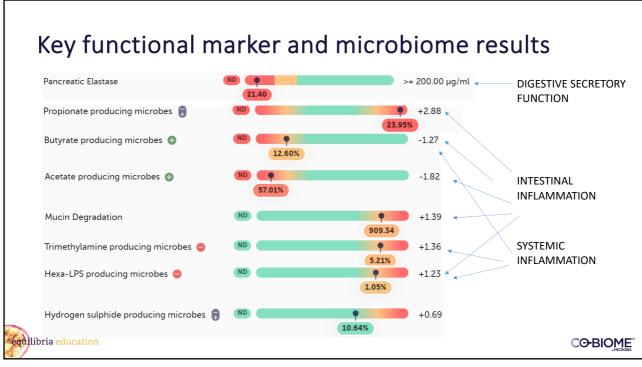




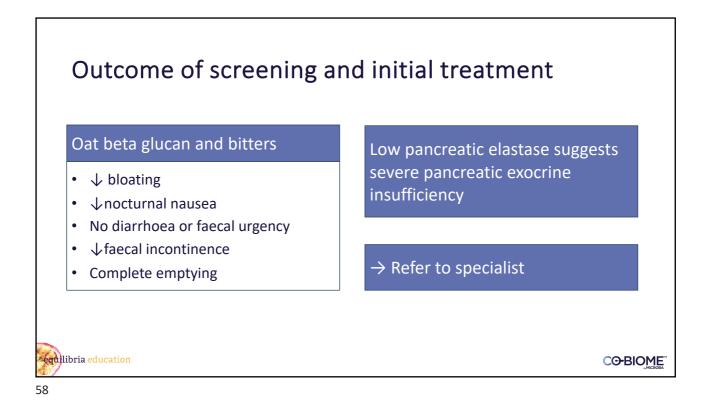


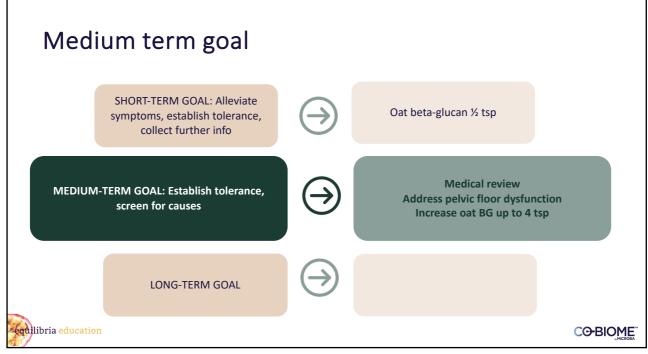




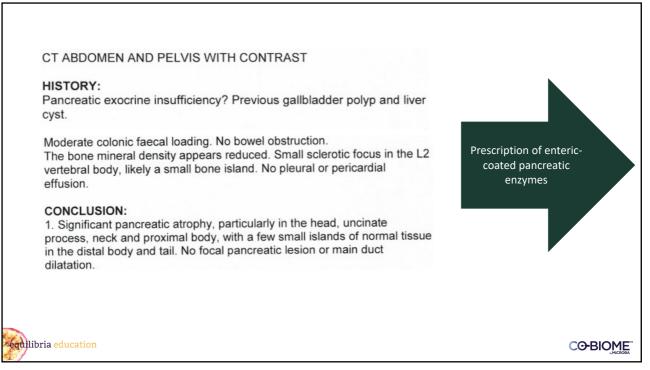


Species results			
Oral Species			
Streptococcus salivarius	0.11%	Common	
Streptococcus anginosus	0.06%	Less common	
Streptococcus parasanguinis	0.04%	Rare	
Fusobacterium vincentii	0.02%	Rare	
Peptostreptococcus anaerobius	0.02%	Less common	
allibria <mark>education</mark>			

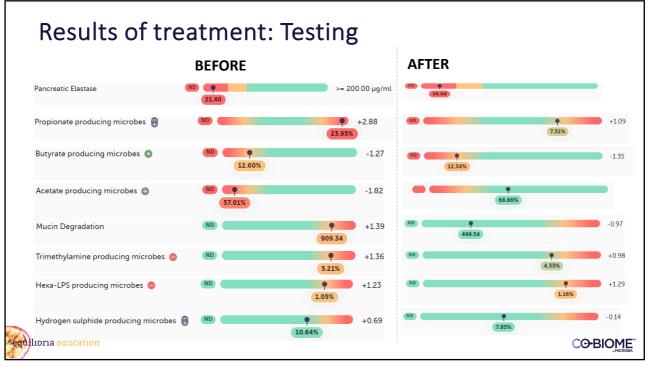














Meal	Day 1	Fibre (g)	Day 2	Fibre (g)
Breakfast	1 x low-FO bread Cheese 2 tsp protein powder	1	1x low-FO bread Goat's cheese 2 tsp protein powder	1
Morning tea	1 tb muesli 3 tb low-fat Greek yoghurt Strawberries	1	-	
Lunch	2 x low-FO bread Goat's cheese 5 cherry tomatoes	2 1	Quinoa Leek Celery Tomato Chicken	2.5 1 1 1
Afternoon tea	Ice cream with oat beta-glucan	4	Ice cream with oat beta-glucan	4
Dinner	Quinoa Chicken Leek, celery, tomato Watermelon	2.5 1 0.3	Broccolini 2g ½ cup brown rice 1.8g Chicken Celery 1g Mandarin 1.8g	2 1.8 1 1.8
TOTAL FIBRE		<mark>~13.8g</mark>		~17.1g
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Outcome and plan

Outcome	Plan
Nocturnal nausea if has richer food like pizza for dinner	Discuss Creon dosage with specialist
Moderate compliance with oat beta-glucan ("if I put it in juice or yoghurt it gets gluggy")	Try mixing with protein powder smoothie If all else fails -> almond milk with a little soy ice cream mixed in -> Trial GOS
BMs occasionally too firm	Kiwifruit 2/day 2 nd daily legumes Track fibre intake on My Fitness Pal
Total cholesterol reduced from 6.3 -> 5.6 mmol/L LDL-C reduced from 4.5 -> 4.0 mmol/L (no change in HDL-C)	
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Key takeaways

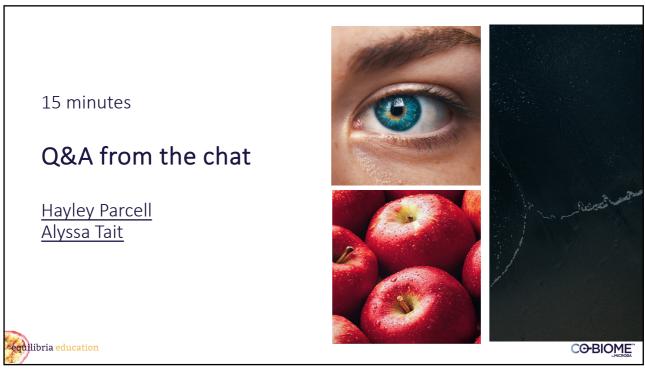
- Personalise your prebiotic to the patient, taking into account the clinical picture, goals and preferences, while being guided by the gut microbiome results.
- Dosage varies considerably according to condition be aware of the research.
- Tolerance and acceptability can be a stumbling block

 when in doubt, start conservatively and proceed with caution.
- Dosing via the instructions on the bottle is a minefield be specific about dosing.
- Even if the clinical outcome improves, re-test the gut microbiome.





	The M	etaXplore™ range	
	Hashing and the second se	MetaXplore [™] MetaXplore [™] provides a metagenomic driven gut microbiome profile, together with the latest research insights for healthcare professionals. Technology: metagenomics	\$369
	Headport G	MetaXplore [™] GI MetaXplore [™] GI provides the same comprehensive microbiome profile as MetaXplore [™] as well as reporting on seven gastrointestinal health markers and science backed clinical insights to assist clinical decision-making and intervent Technology: metagenomics + diagnostic GI health markers + faecal pH	\$489 ion.
	Heatpear or Deal Monte and the second	MetaXplore [™] GI Plus MetaXplore [™] GI Plus is Co-Biome's most comprehensive functional gut microbiome profile. It provides all the features found in MetaXplore [™] and MetaXplore [™] GI, plus targeted pathogen panels. Technology: metagenomics + diagnostic GI health markers + faecal pH + RT-PCR	\$529
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- Prebiotic Guide
- Low FOMAP Prebiotic Guide
 - Patient Handouts Ellagic acid; Arabinoxylan; Inulin; FOS; GOS; Pectin
- Dietary Impacts on the Gut Microbiome Guide
- Interpretation Guide
 - Pathogen and Pathobiont Management Guide



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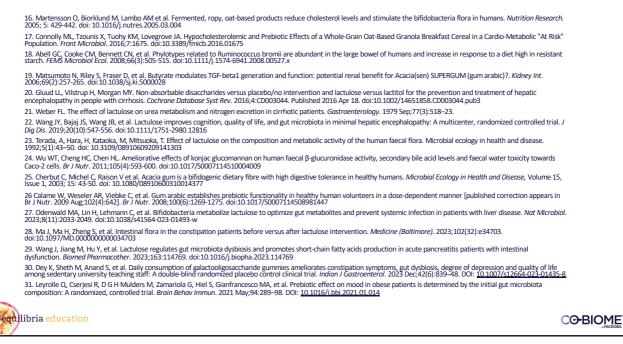
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