Healthpath's Gut Health Tests

BIOMARKERS



Healthpath's **Essential, Advanced** and **Ultimate Gut Health Tests** show you what's going on in your gut. By looking at imbalances in bacteria, yeasts, parasites and other intestinal health biomarkers, you find out what's contributing to your symptoms. You also receive targeted diet, supplement and lifestyle recommendations to help you take back control.

The biomarkers provide clinical information on three key areas:



0

Digestion/Absorption

• pH

1

- Pancreatic elastase
- Zonulin
- Digestive Residues



2 | Immune activity/Inflammation

- Calprotectin
- Haemoglobin
- Secretory IgA
- H. Pylori
- Archaea/methanogens
- E. Coli, Lactobacillus species, Enterococcus species
- Akkermansia muciniphila, Faecalibacterium prausnitzii

3 Gut microbiome/Mycobiome

- Microbiome diversity
- Enterotype
- Dysbiosis index
- Actinobacteria
- Bacteroidetes
- Firmicutes
- Proteobacteria
- Fusobacteria
- Verrucomicrobia
- Hydrogen-sulphide production
- Oxalate-degrading bacteria
- Yeasts
- Parasites



Clinical advantages of the qPCR technology used in Healthpath's tests

This new method of analysis allows for a single sample. This makes the process easier for everyone, and it's particularly helpful for children and those struggling with diarrhoea or constipation.





Essential Advanced Gut Gut Health Health Test Test

Stool properties

Colour	\checkmark	\checkmark
Consistency	\checkmark	\checkmark
рН	\checkmark	\checkmark

Biodiversity

Diversity	\checkmark	\checkmark
Dysbiosis index	\checkmark	\checkmark

Bacterial distribution

Actinobacteria	\checkmark	\checkmark	
Bacteroidetes	\checkmark	\checkmark	
Firmicutes	\checkmark	\checkmark	
Fusobacteria	\checkmark	\checkmark	
Proteobacteria	\checkmark	\checkmark	
Verrucomicrobia	\checkmark	\checkmark	
Other	\checkmark	\checkmark	
Firmicutes/ Bacteroidetes Ratio	\checkmark	\checkmark	
Enterotype			
1, 2 or 3	\checkmark	\checkmark	



Actinobacteria

Bi	ifidobacteria	\checkmark	\checkmark
Equol-producing bacteria		\checkmark	\checkmark
	Adlercreutzia species	\checkmark	\checkmark
	Eggerthella lenta	\checkmark	\checkmark
	Slackia species	\checkmark	\checkmark

Bacteroidetes

Ba	acteroides	\checkmark	\checkmark
	Bacteroides uniformis	\checkmark	\checkmark
	Bacteroidesovatus	\checkmark	\checkmark
Pı	revotella	\checkmark	\checkmark
	Prevotella copri	\checkmark	\checkmark

Firmicutes

utyrate-producing acteria	\checkmark	\checkmark
Faecalibacterium prausnitzii	\checkmark	\checkmark
Eubacterium rectale	\checkmark	\checkmark
Eubacterium hallii	\checkmark	\checkmark
Roseburia species	\checkmark	\checkmark
Ruminococcus species	\checkmark	\checkmark
Coprococcus	\checkmark	\checkmark
Butyrivibrio species	\checkmark	\checkmark
Total bacterial count	\checkmark	\checkmark





Essential Gut Health Test Advanced Gut Health Test

Firmicutes

С	lostridia	\checkmark	\checkmark
	Clostridia total bacterial count	\checkmark	\checkmark
	Clostridia cluster 1	\checkmark	\checkmark

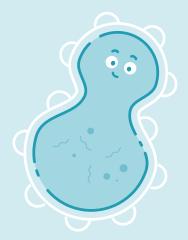
Fusobacteria

Fusobacterium species	\checkmark

Verrucomicrobia

Akkermansia muciniphila







Proteobacteria

	otentially athogenic bacteria	\checkmark	\checkmark
	Haemophilus	\checkmark	\checkmark
	Acinetobacter	\checkmark	\checkmark
	Proteus species	\checkmark	\checkmark
	Klebsiella species	\checkmark	\checkmark
	Enterobacter species	\checkmark	\checkmark
	Serratia species	\checkmark	\checkmark
	Hafnia species	\checkmark	\checkmark
	Morganella species	\checkmark	\checkmark
	Providencia species	\checkmark	\checkmark
	Citrobacter species	\checkmark	\checkmark
	Pseudomonas species	\checkmark	\checkmark
	istamine- roducing bacteria	\checkmark	\checkmark
Н	2S production	\checkmark	\checkmark
	Sulphate- reducing bacteria	\checkmark	\checkmark
	Desulfovibrio piger	\checkmark	\checkmark
	Desulfomonas pigra		\checkmark
	Bilophila wadsworthii		\checkmark



	Essential Gut Health Test	Advanced Gut Health Test	
Archaea			
Methanobrevibacter	\checkmark	\checkmark	
Immunogenically effective bacteria			
Escherichia coli	\checkmark	\checkmark	
Enterococcus species	\checkmark	\checkmark	
Lactobacillus species	\checkmark	\checkmark	
Mucin production/ mucosal barrier			
Akkermansia muciniphila	\checkmark	\checkmark	
Faecalibacterium prausnitzii	\checkmark	\checkmark	
Helicobacter pylori (H. pylori)			
Helicobacter AG		\checkmark	





Yeasts

Candida albicans	\checkmark	\checkmark
Candida krusei	\checkmark	\checkmark
Candida glabrata	\checkmark	\checkmark
Candida dubliniensis	\checkmark	\checkmark
Candida parapsilosis	\checkmark	\checkmark
Candida tropicalis		
Candida lusitaniae	\checkmark	\checkmark

Parasites

thobionts		\checkmark
Blastocystis hominis		\checkmark
Dientamoeba fragilis		\checkmark
Helicobacter AG		\checkmark
thogenic intestinal otozoa		\checkmark
Giardia lamblia		\checkmark
Entamoeba histolytica		\checkmark
Cryptosporidium species		\checkmark
Cyclospora cayetanensis		\checkmark
	Blastocystis hominis Dientamoeba fragilis Helicobacter AG thogenic intestinal otozoa Giardia lamblia Entamoeba histolytica Cryptosporidium species Cyclospora	Blastocystis hominisImage: ComparisDientamoeba fragilisImage: ComparisHelicobacter AGImage: Comparisthogenic intestinal otozoaImage: ComparisGiardia lambliaImage: ComparisEntamoeba histolyticaImage: ComparisCryptosporidium speciesImage: ComparisCyclosporaImage: Comparis



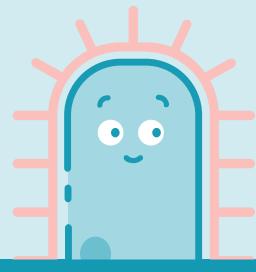


Functional markers

Calprotectin	\checkmark
Haemoglobin in faeces immunologically	\checkmark
Secretory IgA	\checkmark
Pancreatic elastase	\checkmark

Digestive residues

Determination of fat	\checkmark
Determination of nitrogen	\checkmark
Determination of sugar	\checkmark
Determination of water	\checkmark



Our Ultimate Gut Health test is the most comprehensive stool test. In addition to everything in the Advanced, you will also receive:

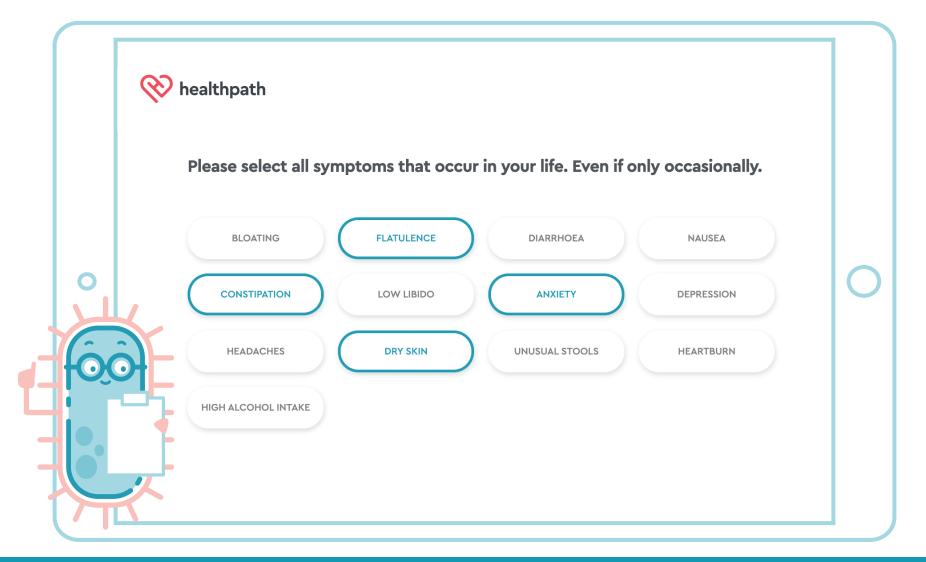
	Ultimate Gut Health Test
Functional markers	
Zonulin	\checkmark
Buytrate-producing bacteria	
Cl. buytricum	\checkmark
Clostridia	
Clostridia histolytium	\checkmark
Clostridia perfringens	\checkmark
Clostridia sporenges	\checkmark
Other Firmicutes	
Christensenellaceae	\checkmark
Dialister invisus	\checkmark
Proteobacteria	
Proteus mirabilis	\checkmark
Oxalate-degrading bacteria	\checkmark
Oxalobacter formigenes	\checkmark



www.healthpath.com

Gut Health Test results are delivered in your private Healthpath dashboard

Your practitioner considers your symptoms:





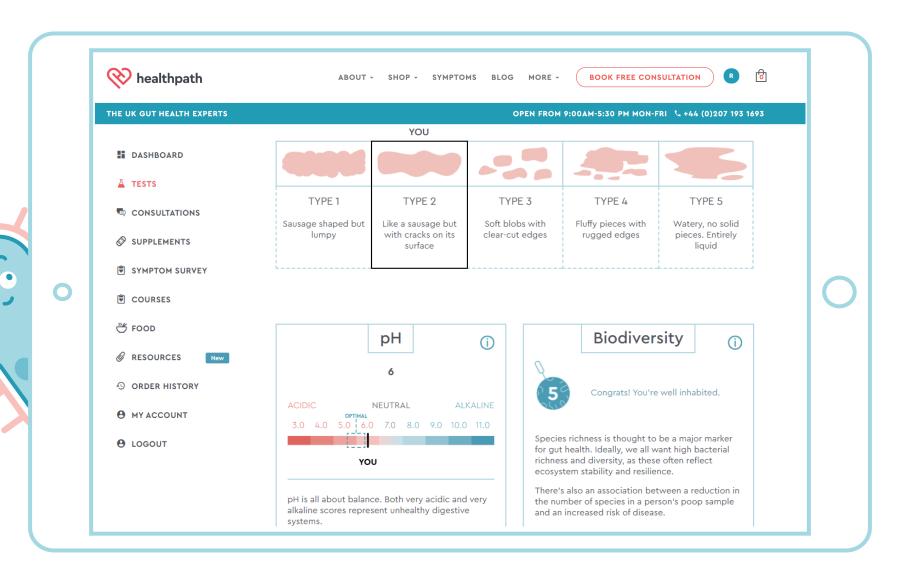


You get a personal note from your practitioner:

📎 healthpath	ABOUT - SHOP - SYMPTOMS BLOG MORE - BOOK FREE CONSULTATION
THE UK GUT HEALTH EXPERTS	OPEN FROM 9:00AM-5:30 PM MON-FRI 🕓 +44 (0)207 193 1693
	Dear Richard,
DASHBOARD	Congratulations on getting this far. I have reviewed your symptom questionnaire and test results to
TESTS	create your personalised program.
	Key Findings
	When viewing your test results, keep in mind that the microbiome is a new and complex area of science.
SUPPLEMENTS	You do not need to see 'out-of-range' markers as a diagnosis. Rather, see them as suggestions to focus on a particular aspect of health.
🗑 SYMPTOM SURVEY	Your key findings include:
🗑 COURSES	1. A good stool pH - improved since your last test
COURSES	 Good microbiome diversity – also improved since your last test. Dysbiosis
🐣 FOOD	4. Low levels of Bifidobacteria and key butyrate-producing (friendly) bacteria
	5. High levels of histamine-producing bacteria 6. High levels of candida species
	7. High secretory IgA (indicating immune activity) – this has increased since your last test.
S ORDER HISTORY	To learn more about your out-of-range markers, click the 'Show detailed description' button on the 'Out
	of range' page on your dashboard.
_	Your Recommendations
e logout	Your program has been created based on your results and your symptom questionnaire. It is designed to balance your microbiome and support your gut health, addressing the findings listed above.
	The primary aims of your program are to support the growth of the beneficial, gut microbiome regulating bacteria, lower the levels of the less beneficial organisms, and to support the immune system along the gut lining.



An overview of your gut health:



www.healthpath.com

A detailed breakdown of in-range and out-of-range markers:

ᅇ healthpath	ABOUT - SHOP - SYMPTOMS	BLOG MORE - BOOK FREE CONSULTATION	
THE UK GUT HEALTH EXPERTS		OPEN FROM 9:00AM-5:30 PM MON-FRI し+44 (0)207 193 1693	
DASHBOARD	Tests > Ultimate Gut Health Test Results - out of range	Notes Overview In range Out of range Next steps	
TESTS	Show detailed description 🚾 🧻	DOWNLOAD LAB REPORT 🗈	
CONSULTATIONS Image: Consultations<	< PREV	NEXT>	
⑦ COURSES	Bacterial distribution	Out of range	
RESOURCES New	- Actinobacteria	Borderline low	
 ORDER HISTORY MY ACCOUNT 	which reduce the pH within the intestines. A lower pl bad bacteria.	hey produce special substances called short-chain fatty acids, H is a good thing because it helps to prevent the growth of n to intestinal inflammation. Low levels of Bifidobacteria (a	
e logout	type of Actinobacteria) are seen in IBS, IBD and color		
	- Verrucomicrobia	Borderline low	



A personalised program:

🛞 healthpath	
THE UK GUT HEALTH EXPERTS	OPEN FROM 9:00AM-5:30 PM MON-FRI & +44 (0)207 193 1693
 DASHBOARD TESTS CONSULTATIONS 	Tests > Ultimate Gut Health Test Notes Overview In range Out of range Next steps It's time to view your Healthpath dashboard and explore your personal recommendations: Notes Overview In range Out of range Next steps
SUPPLEMENTS	01 Food program recommendations GET STARTED
♥ COURSES	These recipes and meal planning guidelines have been handpicked to aid the repair of your digestive health and optimise your wellbeing, based on your individual test results and personal information.
RESOURCES New	02 Supplement recommendations Get STARTED
• ORDER HISTORY • MY ACCOUNT	Your practitioner has made these highly targeted supplement recommendations to fulfil your individual needs, based on your personal health assessment, alongside your food and lifestyle plans.
e Logout	03 Resource recommendations Get STARTED
	These articles, videos and comprehensive guides will educate and guide you on your journey to better health, having been carefully selected based on your personal details and individual results.

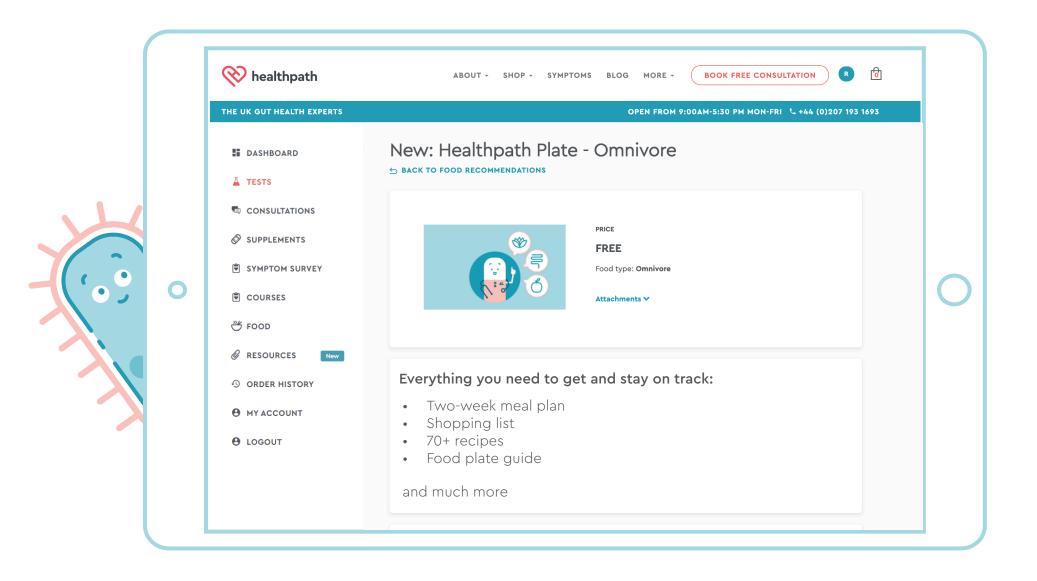


Advanced supplement protocol:

🛞 healthpath	ABOUT - SHOP -	SYMPTOMS BLOG MORE -	BOOK FREE CONSULTATION	<u>ह</u> ो
THE UK GUT HEALTH EXPERTS		OPEN FROM	2:00AM-5:30 PM MON-FRI 🥾 +44 (0)207 193 169	3
DASHBOARD TESTS	Your supplements schedul Remember, this recommendation is best required.		SEND TO MY EMAIL	
 CONSULTATIONS SUPPLEMENTS 	Upon rising			
 SYMPTOM SURVEY COURSES 	SUPPLEMENT Bio.Me Prebio GOS 90g I suggest starting with a ¼ of the suggested the fibre.	DOSAGE 0.5 tsp in 200ml water dosage and build this up slowly over a couple of	DURATION 8-12 weeks weeks. This allows the microbiome time to adjust to	0
FOOD	Bio.Me Barrier 90's	3 capsules	8-12 weeks	2
• ORDER HISTORY	Breakfast			5
O LOGOUT	SUPPLEMENT Oil of Oregano 120's	DOSAGE 1 capsule	DURATION 4 weeks	
	I suggest starting with just 1 capsule a day, a	nd if tolerated, increasing to the suggested dosa	ge slowly over a few weeks.	



An evidence-based food plan, hand-picked by our practitioners:



Plus a copy of the original lab report

bio vis ' diagnos ⁻	ТІК					Biovis Diagnostik MVZ. Pot Or, med. Burkhard Sch Wossendichte Lettrag Tomas Gaprel 3 Artthe Lettrag De. med. Herbert Schnäft 1 Pot De. med. Herbert Schnäft 2 Pot De. med. Herbert Schnäft 2 Pot De. De. Medical Kram Jours-Bouch Scrube 2 65555 Limberg-Offnein Vishard für Likentrisersechtin	9tt	
xternal ID						Tel.: 0.64.31 / 21.248-0 Fax: 0.64.31 / 21.248-6	0 E-mail: 66 Web:	nfo@biovis.de www.biovis.de
lame irst Name	Demo A713AE1	Date Sex	of Birth		.1964 emale	Order ID Order Date		1263093 10.12.202
ampling Date ample Material	10.12.2021 00:00 FE		ation Date ation on	10.1	2.2021	Findings Status Findings Date		Final Repo 10.12.202
Test		Result	Unit	Standard Range			Pr	evious Resul
Microbiome Healthpath								
Molecular genetic micro	blome analysis 3.0							
Stool Properties								
Colour		brown						NUT VE
Consistency		mushy						NPQ VE
pН		6,5		5,8 - 6,5				NA) TES
Biodiversity								
Diversity		6,42		> 5				NA) MOS
			nsiderably icreasing age, ersity.				6	
Enterotype Bacteroides Human intestinal micr Enterotypes are defini properties.	obiomes can be different ed by dominant bacterial	iated into clusters v	three Enterot	VDES.	Enterot	ур	1	NA) MGS
Bacteroides Human intestinal micr Enterotypes are define	obiornes can be different ed by dominant bacterial	iated into clusters v	three Enterot	VDES.	Enterot	ур	1	NU, MGO
Bacteroides Human intestinal micr Enterotypes are defini properties. Dysbiosis Index The dysbiosis index re	obiomes can be different ed by dominant bacterial presents a measure of o presents a measure of the evance, all detected phy	leviations	three Enterot vith distinct m	ypes. etabolic	Enterot	yp 1	1	NQ NGS
Bacteroides Human intestinal micr Enterotypes are define properties. Dysbloata Index Dysbloata Index The dysbloats index re Depending on their re	ed by dominant bacterial	leviations	three Enterot vith distinct m	ypes. etabolic		yp 1	1	
Bacteroldes Human intestinal micr Enterolypes are defin properties. Dysbloais Index The dysbloais index re Depending on their re considered.	ed by dominant bacterial	leviations	three Enterot vith distinct m	ypes. etabolic		yp 1	1	Nų REO
Bacterolides Human intestinal micr Enterotypes are defin properties. Dyabiosis index Dyabiosis index Depending on heir re considered. Ratio	ed by dominant bacterial	leviations la, genera	three Enterot with distinct m within the mic	ypes. etabolic zrobiome. are		1	1	Nų REO
Bacteroides Human intestinal mar Enterolytes are defin properties Dysbioals Index The dysbiols index Despring on her re considered. Ratio Firmicules/Bacteroidetes	ed by dominant bacterial	leviations la, genera	three Enterot with distinct m within the mic	ypes. etabolic zrobiome. are	Index	1	1	MA REC
Bacterolides Human intestinal mice Erriterolytes are defin properties. Dyabioals Index The dyabioals Index re Depending on their re consistence. Ratio Firmicules/Bacterolidetes Phylia	ed by dominant bacterial	leviations la, genera 1,73	three Enterot with distinct m within the min a and species Quotient	ypes etabolic problome. are <1.5	Index	1	1	
Bacterolides Human intestimal mice Friendorges are defin properties, see defin properties, see defin properties, see defin properties, see defin The dysbiosis index rc Depending on heir re considered, Firmica.des/Bacterolidetes Phyla Actinobacteria	ed by dominant bacterial	leviations la, genera 1,73	three Enterot with distinct m within the min a and species	yres. eabolic arobiome. are <1.5	Index	1	0	NU KG
Bacterolides Human intestinal micr Enterotypes are defin properties. are defin properties. are defin properties. are defin properties. and the Depending on their re considered. Fatio Ratio Califorabeteria Bacterolidetes	ed by dominant bacterial	leviations la, genera 1,73 1,7 28,6	three Enterot with distinct m within the min a and species Quotient	yres etabolic zrobiome. are <1.5 1.0-5.0 30-60	Index	1	1	NU NCO NU NCO NU NCO NU NCO
Bacteroides Human intestinal more Enterolytes are defin properties. Dysbloals Index The dysbloals Index The dysbloals Index The dysbloals Index Ratio Ratio Bacteroidetes Firmicules Firmic	ed by dominant bacterial	leviations la, genera 1,73 1,7 28,6 49,4	three Enterot with distinct m within the min a and species Quotient	ypes etabolic xobiome. are <1.5 	Index	1	1	ing pool book pool book pool book pool book pool book pool
Bacterolides Human intestinal mice Enterolytes are defin properties. Dyabioals Index The dyabioals Index The dyabioals Index re Depending on their re consistence. Ratio Firmicates/Bacterolidetes Firmicates Fuedocateria Bacterolidetes Fuedocateria	ed by dominant bacterial	leviations la, genera 1,73 1,7 28,6 49,4 0,0	three Enterot with distinct m within the mix and species Quotient	ypes etabolic arobiome. < 1.5 1.0-5.0 30-60 30-60 0.0-1.0	i i i i i i i i i i i i i i i i i i i	1	1	ing they well well well
Bacterolides Human intestimal micr Enterolytes are defin properties. Dyabiosis index r Depablosis index r Departing on heir re considered. Firmica.das/Bacterolidetes Plyia Actinobacteria Bacterolidetes Firmica.das/Bacterolidetes	ed by dominant bacterial	1,73 1,7 28,6 49,4 0,0 8,9	three Enterot tilt distinct m within the min and species Quotient	ypes etabolic crobiome. 	i i i i i i i i i i i i i i i i i i i	1	1	COM PAR COM PAR Lical Par
Bacteroldes Human intestinal micr Enterolytes are defin properties, ere defin properties, ere defin properties, ere defin Depending on their re considered. The dysbiols index r Depending on their re considered. Firmicutes/Bacteroldetes Phyla Actinobacteria Bacteroldetes Firmicutes Fusobacteria Vertucomicrobia	ed by dominant bacterial	Leviations 1,73 1,7 28,8 49,4 0,0 8,9 1,9	three Enterot tilt distinct m within the min a and species Quotient Cuotient	ypes etabolic crobiome. 	i i i i i i i i i i i i i i i i i i i	1		NU 400

lame irst Name		Demo A713AE1	Date of I Sex	Sirut	23.01.1964 Female	Order ID Order Date		1263093 10.12.202
Test			Result	Unit	Standard Range	Order Date		revious Result
Bacteria Phyla - most i	mportant ger							
Actinobacteria								
Bifidobacterium		1,2 X	10^10 CFU/g	faeces	>5,0 x 10*9			NA) MOST
Bifidobacterium	adolescentis		81	%				NAU MOSE
Bifidobacterium	longum		16	%				NAU MOSE
Equol producing bacteri	a	4,3 x	10^9 CFU/g	faeces	> 5,0 x 10/9			NAU MOSE
Adlercreutzia spp.			•					NA) MGS
Eggerthella lenta			•					NA) MOS
Slackia spp.			•					NA) MOS
Bacteroldetes			-					NAV MODE
Bacteroides		1,5 X	10^11 CFU/g	faeces	> 1,5 x 10^11			NAU MOSE
Bacteroides	uniformis		15	%				NAU MOSE
Bacteroides	ovatus		12	%				NAU MOSE
Prevotella		4,0 X	10^10 CFU/g	faeces	> 1,0 x 10^10			NA) MGSE
Firmicutes								NA) MUSE
Butyrate producing ba	cterla							
Total bacteria count		3,0 X	10^11 CFU/g	faeces	> 1,2 x 10^11			NA) MGS
Faecalibacterium prausr	nitzii	8,4 X	10^10 CFU/g	faeces	> 5,0 x 10^10			NA) MOS
Eubacterium rectale		3,7 X	10^10 CFU/g	faeces	> 1,0 x 10^10			NA) MOST
Eubacterium hallii		3,3 X	10^10 CFU/g	faeces	> 5,0 x 10*9			NAU MOSE
Roseburia spp.		5,0 X	10^10 CFU/g	faeces	> 2,0 x 10^10			NA) MGSE
Ruminococcus spp.		4,1 X	10^10 CFU/g	faeces	> 3,0 x 10^10			NAL MERS
Coprococcus spp.		3,0 X	10^10 CFU/g	faeces	> 2,0 x 10^10			NA) MOSE
Butyrivibrio spp.		2,1 X	10^10 CFU/g	faeces	> 5,0 x 10^9			NA) MUSE
Clostridia								NA) MODE
Clostridia total bacteria	xount	2,7)	(10^9 CFU/g	faeces	< 4,0 x 10^9			NA) MOST
Clostridia Cluster I		1,00	10^5 CFU/g	faeces	< 2,0 x 10*9			NAU MOSE
Fusobacteria								They include
Fusobacterium		< 1,0)	(10^5 CFU/g	faeces	< 1,0 x 10^7			NAU MOSE
Verrucomicrobia								
Akkermansia muciniphili	1	1,9 X	10^9 CFU/g	faeces	> 5,0 x 10^9			NA) MOSE
Proteobacteria								
Pathogenic or potentia	lly pathogen							
Haemophilus spp.		2,2 X	10^9 CFU/g	faeces	< 1,0 x 10^9			NA) MOSE
Acinetobacter spp.		< 1,0)	(10^5 CFU/g	faeces	< 1,0 x 10^6			NA) MOST
Proteus spp.			< 10^5 CFU/g		< 1,0 x 10^6			NA) MGS
Klebsiella spp.			10^5 CFU/g		< 1,0 x 10^6			NA) MGSE
Enterobacter spp.		< 1,0)	10^5 CFU/g	faeces	< 1,0 x 10^6			NA) MGSE
Serratia spp.		< 1,0)	10^5 CFU/g	faeces	< 1,0 x 10^6			NA) MGSE
Hafnia spp.		< 1,0)	10^5 CFU/g	faeces	< 1,0 x 10^6			NA) MSSE
Morganella spp.		< 1,0)	10^5 CFU/g	faeces	< 1,0 x 10^6			NA) MOST
Citrobacter spp.		< 1,0)	10^5 CFU/g	faeces	< 5,0 x 10^8			NA) MOSE
Pseudomonas spp.		< 1,0)	10^5 CFU/g	faeces	< 5,0 x 10^7			NAU MOSE
Providencia spp.		< 1,0)	10^5 CFU/g	faeces	< 5,0 x 10^7			NA) MOSE
H2S production								
-stool						1 00000011	dice (P) A) according	ed. NA) not accredite
		W 7LT London				cooperate ana	field (iii), rij accredi	ou, reij not accrouit

Name	Demo	Date of E	Birth	23.01.	1964	Order ID		12630932	
First Name	A713AE1	A713AE1 Sex		Female		Order Date		10.12.2021	
Test		Result	Unit	Standard Range				Previous Result	
Sulphate reducing bacteria	5,0)	K 10^9 CFU/g	faeces	< 2,0 x 10^9				P NU MOSE	
Desulfovibrio piger	< 1,0	x 10^5 CFU/g	faeces	< 1,0 x 10^9				P NU MOSE	
Desulfomonas pigra	< 1,0	x 10^5 CFU/g	faeces	< 1,0 x 10^9				P NA MGSE	
Bilophila wadsworthii	< 1,0	x 10^5 CFU/g	faeces	< 2,0 x 10^9				P. NAJ MGSEI	
Histamine producing bacteria									
Histamine producing bacteria	< 1,0	x 10^5 CFU/g	faeces	< 5,0 x 10^8				P NU MOSE	
Immunogenicity / Mucus produc	ction								
Immunogenically effective bact	erla								
Escherichia coli	5,2)	K 10^5 CFU/g	laeces	10^6 - 10^7				P NU MOSE	
Enterococcus spp.	< 1,0)	K 10^5 CFU/g	faeces	10^6 - 10^7				P NO MOSE	
Lactobacillus spp.	2,6	x 10^5 CFU/g	faeces	10^5 - 10^7				NA) MODE	
Mucin production / Mucosal bar	rler								
Akkermansia muciniphila	1,90	K 10^9 CFU/g	faeces	> 5,0 x 10^9				F NALMOSE	
Faecalibacterium prausnitzii	8,4 X	10^10 CFU/g	faeces	> 5,0 x 10^10				P NAU MISSE	
Archaea									
Methanogens									
Methanobrevibacter spp.	2,6	x 10^7 CFU/g	faeces	< 1,0 x 10^8				ni) MGSE	
Mycoblome: relevant yeasts									
Candida albicans (CA)	4,1)	K 10^3 KBE /	g Stuhl	<1,0 x 10^3				FI NU OFC	
Candida krusei (CK)	<1,0	x 10^3 KBE /	g Stuhl	< 1,0 x 10^3				NA OPO	
Candida glabrata (CG)	<1,0	x 10^3 KBE /	g Stuhl	< 1,0 x 10^3				NAL OPC	
Candida dubliniensis (CD)	<1,0	x 10^3 KBE /	stuhi	< 1,0 x 10^3				NAL OPC	
Candida parapsilosis (CP)	<1,0	x 10^3 KBE /	stuhi	< 1,0 x 10^3				RA OFC	
Candida tropicalis (CTp)	<1,0	x 10^3 KBE /	g Stuhl	< 1,0 x 10^3				NAJ OPC	
Candida lusitaniae (CL)	<1,0			< 1.0 x 10^3				NA) OPC	

	Healthpath 85 Great Portland Street GB-W1W 7LT London	Seite 3 von 4
dited	FE-stool	* cooperate analytics (R), A) accredited, NA) not accredited

