

Patient
Birthdate
Analysis #

Average

Unusual

PHYLUM	FAMILY	GENDER	% of total	Ref.
Firmicutes (gram +)	Lachnospiraceae	Anaerostipes	8,24	<1
		Coprococcus	0,06	<10
		Dorea	25,84	<15
		Moryella	0	<1
		Roseburia	5,14	<50
	Ruminococcaceae	Sporobacterium	0	<1
		Syntrophococcus	0	<1
		Acetanaerobacterium	0	<5
		Acetivibrio	1,24	<1
		Ethanoligenens	0	<1
		Faecalibacterium	0	<25
		Papillibacter	0	<1
		Ruminococcus	0	>2
		Sporobacter	0	<1
		Subdoligranulum	0	<25
	Clostridiaceae	Butyricicoccus	0	<5
		Clostridium	20,57	<5
		Lactonifactor	0,12	0
	Eubacteriaceae	Anaerofustis	0	<0,5
		Eubacterium	0	0
	Blautia	Blautia	25,34	<50
	Howardella	Howardella	0	<1
	Lactobacillaceae	Lactobacillus	0,12	<1
	Enterococcaceae	Enterococcus	2,66	0
	Streptococcaceae	Lactococcus	0	<1
		Streptococcus	2,42	<5
	Leuconostoc	Leuconostoc	0	<0,3
	Erysipelotrichaceae	Catenibacterium	0	<0,3
		Coprobacillus	5,7	<1
		Holdemania	0,19	<1
		Turicibacter	0	>0,5
		Dialister	0	0-1
	Veillonellaceae	Megamonas	0	0
		Megasphaera	0	0
		Oscillibacter	0	<4
	Oscillospiraceae	Oscillibacter	0	<4
	Staphylococcus	Staphylococcus	0	<0,05
Bacteroidetes (gram-)	Bacteroidaceae	Bacteroides	0	<10
	Rikenellaceae	Alistipes	0	<3
	Porphyromonadaceae	Barnesiella	0	<2
		Odoribacter	0	<0,5
		Parabacteroides	0	<3
		Prevotella	0	<5
		Xylanibacter	0	<1
Actinobacteria (gram+)	Bifidobacteriaceae	Bifidobacterium	0,81	>5
	Actinomycineae	Actinomyces	0,06	<1
	Micrococcineae	Rothia	0	<0,2
	Coriobacterineae	Asaccharobacter	0	>0,1
		Collinsella	0	<25
		Olsenella	0	0
		Slackia	0	<1
Proteobacteria (gram-)	Enterobacteriaceae	Escherichia/Shigella	1,12	<0,5
		Klebsiella	0	<0,5
	Sutterellaceae	Sutterella	0	<1
	Desulfovibrionaceae	Lawsonia	0	<0,5

	Value	Ref.
Total Lachnospiraceae	39,28	>5
Total Ruminococcaceae	1,24	>5
Total Clostridiaceae	20,69	<5
Enterococcus	2,66	0
Streptococcus	2,42	<5
Ruminococcus	0	>2
Lactonifactor	0,12	0
Turicibacter	0	>0,5
Bacteroides	0	<10
Prevotella	0	<5
Bifidobacterium	0,81	>5
Asaccharobacter	0	>0,1

Firmicutes	97,64 %
Bacteroidetes	0 %
Actinobacteria	0,87 %
Proteobacteria	1,12 %
Other	0,37 %

Range of Firmicutes % in European population: 50-85%

Firmicutes/Bacteroidetes ratio	
High	#DIV/0!
Average	#DIV/0!
Low	#DIV/0!

Low ratio may be associated with gut inflammation

Gram+ / Gram- ratio	
High	87,96
Average	
Low	

Diversity Index	2,72
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Low <4, Average 4-5, High >5

Dysbiosis associated with low diversity

Electronically validated on:14/05/2013

Requesting physician: Himmunitas

Observations: