G_{eburtsdatum} Geschlecht

KBE/g Stuhl

1014

1019 - 10111

1015 - 1017

< 1013

KBE/g Stuhl

KBE/g Stuh

KBE/g Stuhl

KBE/g Stuhl

KBE/g Stuhl

KBE/g Stuh

KBE/g Stuhj

<10₄

<1014

<10₄

<101₄

<101₄

<1014

<1014

2x1016

3x10^9

1015

<10¹5

<1013

1045

negativ

<1013

Auftrag Nr.

Florastatus / q-PCR / Sequencing

bio vis

Escherichia coli Escherichia coli Biovare

Proteus species Klebsiella species

Pseudomonas species

Enterobacter species

Serratia species

Hafnia species

Bacterioides species

Lactobacillus species Clostridium species

Mykologische Stuhlunte

Candida species

Candida albicans

Eigenschaften des Stuhl

Schimmelpilze Geotrichum candidum

Konsistenz

Besides our Florastatus biovis now offers molecular genetic methods which allow you to obtain additional information on anaerobic bacteria, which can not be detected by conventional methods.

q-PCR methods allow indirect statements about the condition of the mucosal barrier, H2S formation or presence of toxin-forming cluster I-clostridia which, among other diseases, have been shown to be increased in autism related disorders. q-PCRs are offered in addition to our florastatus, improve significance and allow specific therapeutic measures.



- Consistency, colour and pH-value
- Cultivatable parts of the bacterial microbiome
- Yeasts / Mould fungi

q-PCR **Special Profiles**

- Mucin production / Butyrate production
- Mucin production / Butyrate production / H₂S formation
- A176 Autism / Clostridia
- A₁₆₅ Firmicutes / Bacteroidetes ratio
- **Parasites: Blastocystis hominis**
- A167 Toxine production / Butyrate production

biovis Innovation

Since the first of june 2015 *biovis* offers **complete microbiome sequencing** as a third pillar in fecal diagnostics. With our profiles Microbiome analysis Basic (A710) and Microbiome analysis Plus (A730) biovis is the first non-academic lab allowing information on the guts microbial diversity, being a major part of the colonization resistance and offering protection against endogenous infections. We also determine the enterotype with its resulting consequences on resorption of vitamins and minerals. The microbiome analysis includes over **200 parameters** of which the most relevant are

evaluated and used for the concluding diagnostic findings. Based upon recent research the results

Sequencing

Microbiome analysis Basic

are interpreted and appropriate therapeutic measures are compiled.

Diversity

Firmicutes / Bacteroidetes ratio

Enterotype

Bacterial genera with indicator species

Butyrate production

Detailed Clostridia diagnostics

Pathogenic /

Potentially pathogenic bacteria

Oxalate-degrading bacteria

Mucin production

H2S formation

Immunogenicity

Yeasts / Mould fungi

Microbiome analysis Plus

Digestion residues

Pancreatic elastase

Bile acids

Calprotectin

Alpha-1 antitrypsin

Secretory immunoglobulin A



Florastatus / q-PCR / Sequencing

