

**Changes in duodenal tissue-associated microbiota following hookworm infection  
and consecutive gluten challenges in humans with coeliac disease**

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**Supplementary Table S1** | Trial subject identification (ID) numbers or control subject (C) numbers displaying age, sex, baseline clinical diagnosis (T0 Marsh Score). Corresponding subjects IDs as indicated by Zaiss *et al.*<sup>20</sup> are also indicated where applicable.

Subject	Age	Sex	Marsh score	Subject ID according to Zaiss <i>et al.</i> <sup>20</sup>
ID1	44	F	M0	Pt2
ID2	40	F	M0	Pt3
ID3	48	F	M0	Pt4
ID6	40	M	M0	Pt8
ID7	64	M	M0	Pt1
ID10	52	M	M2	Pt7
C1	57	F	M3b	n/a
C2	22	M	M3a	n/a
C3	48	F	M3a	n/a
C4	52	F	M3b	n/a
C5	51	M	M3a	n/a
C6	75	M	M3b	n/a

**Supplementary Table S2** | Differentially abundant OTUs in the mucosally-associated microbiota of Trial subjects over the course of the study.

OTU	p-value	Adjusted p-value	FDR
Unassigned (kingdom)	0.0016	0.08	0.046
<i>Ralstonia</i> (genus)	0.0025	0.12	0.046
Chitinophagaceae (family)	0.0034	0.16	0.046
Chitinophagaceae (family)	0.041	0.19	0.046
Unassigned (kingdom)	0.0063	0.29	0.046
Unassigned (kingdom)	0.0066	0.3	0.046
<i>Phyllobacterium</i> (genus)	0.007	0.31	0.046
Unassigned (kingdom)	0.0074	0.32	0.046
Chitinophagaceae (family)	0.0092	0.39	0.05
Rhizobiales (order)	0.011	0.45	0.05
Rhizobiales (order)	0.011	0.45	0.05
Rhizobiales (order)	0.012	0.47	0.05

**Supplementary Figure S1 |** Microbial composition of duodenal biopsy samples, at the phylum (a) and genus (b) level, from Trial subjects (ID\*) prior to and following hookworm infection (T0), as well as following administration of escalating doses of gluten (10-50 mg/day – T24; 350 mg/day – T36) as predicted in the analysis of the V1-V3 16S rRNA gene. Bubble sizes reveal the relative abundance (%) of phylotypes (based on 97% sequence identity) in each sample. The composition of the microbial communities detected in duodenal biopsy samples from Control subjects with active coeliac disease (C\*) is also shown. OTUs listed were present in at least one sample at a relative abundance of  $\geq 7\%$ .



