

## Host-probiotic interaction: new insight into the role of the endocannabinoid system by in vivo and ex vivo approaches.

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Histology	Control	VSL#3	P value
Enterocytes length	37.82 ± 1.97 µm	48.31 ± 1.93µm	<0.001
Villi length	0.839 ± 0.085 µm	1.200 ± 0.146 µm	<0.001
Crypt depth	0.088±0.081	0.100±0.022	0.202
Villi/cryptae ratio	12.8±3.73	14.2±5.51	0.016

**Table 1S:** Tabulation and statistical evaluation of enterocyte length, villus height, crypt depth and villus height/crypt depth ratio found in the front intestinal tracts. Values indicate mean ± SD, statistical differences were tested by variance analysis and confirmed by non-parametric methods (Wilcoxon).

<b>Gene</b>	<b>Forward</b>	<b>Reverse</b>
<i>cnr1</i>	TCTGTGGGAAGCCTGTTTC	ACCGAGTTGAGCCGTTTG
<i>cnr2</i>	TCAGAGAGACGATGCTAACC	CAAAAGGCTTCCCACAGAG
<i>trpv1</i>	TGATCGTCGCTGGTGCTT	GACTGGGCTCTCTCTGAACG
<i>faah</i>	CATTCTCGGACCAGCCTTCA	CCTTCACTGCCTGTTGGAAGA
<i>abhd4</i>	GAAGAGCAGTTTGTTCCTCCATAG	GACTCACTCTTTCTGGGTATTGGAT
<i>mgll</i>	CGAGAGGCCGCAGGATTTTA	TGAGTTTAGGAGCCAAGCG
<i>il1</i>	GCTGGGGATGTGGACTTC	GTGGATTGGGGTTTGATGTG
<i>tnfa</i>	TTGTGGTGGGTTTGATG	TTGGGGCATTATTTTGTAAAG
<i>myd88</i>	CAGCGACATCCAGTTTGTGC	GCGGCACCTCTTTTCGATGA
<i>il10</i>	GCCGTGGAGCAGGTGAAG	GAAGATGTCAAACCTCACTCATGGCT
<i>casp1</i>	AGAAAACGACTGGCCTTGCT	TGAGCAAAGTCCCTGATGGC
<i>nos2a</i>	CAAACCTCTACAGCGGCGG	GAGTACAGACGGGGCTTCAG
<i>tgfb1a</i>	CGTCTTCCAGCAAGCTCAGA	TGGAGACAAAGCGAGTTCCC
<i>nfkb</i>	GGCCAAAGACACTGTTCCGA	AGGACTTGCGGTTCTTCTCG
<i>tlr1</i>	CGGGAAAAACCATCGTGCAA	TAGTGGCACCATTGCTCTG
<i>tlr2</i>	CTGTCCGAGCACTTTGTGAG	CCACAACAGCTTAACACTCGTC
<i>tlr3</i>	GACTACGTGATAGTCCGCC	ACAAGCGTAGAACAAGGGCA
<i>tlr9</i>	GGATTCTGGAGTCTCGTGC	CTGCTGCACCAAAGGAAGG
<i>casp3</i>	CCGCTGCCCATCACTA	ATCCTTTCACGACCATCT
<i>baxa</i>	GGCTATTTCAACCAGGGTTCC	TGCGAATCACCAATGCTGT
<i>bcl2a</i>	CCTTCAATAAAGCAGTGGAGGAA	CGGGCTATCAGGCATTCAGA
<i>act1b</i>	GGTACCCATCTCCTGCTCAA	GAGCGTGGCTACTCCTTAC
<i>rplp</i>	CTGAACATCTCGCCCTTCTC	TAGCCGATCTGCAGACACAC

**Table 2S:** Lists of primers used in gene expression analyses by Real-Time qPCR.