

Figure S1. Viability of IPEC-J2 after mEVs exposure at 48h. The different concentrations of mEVs did not determine a significant difference in terms of cell viability after 48 h of treatment. Data are expressed as optical density (OD) \pm SD. Differences were evaluated through the Kruskal-Wallis test and applying the post-doc Dunn's Multiple Comparison Test. Statistical significance: $p < 0.05$.

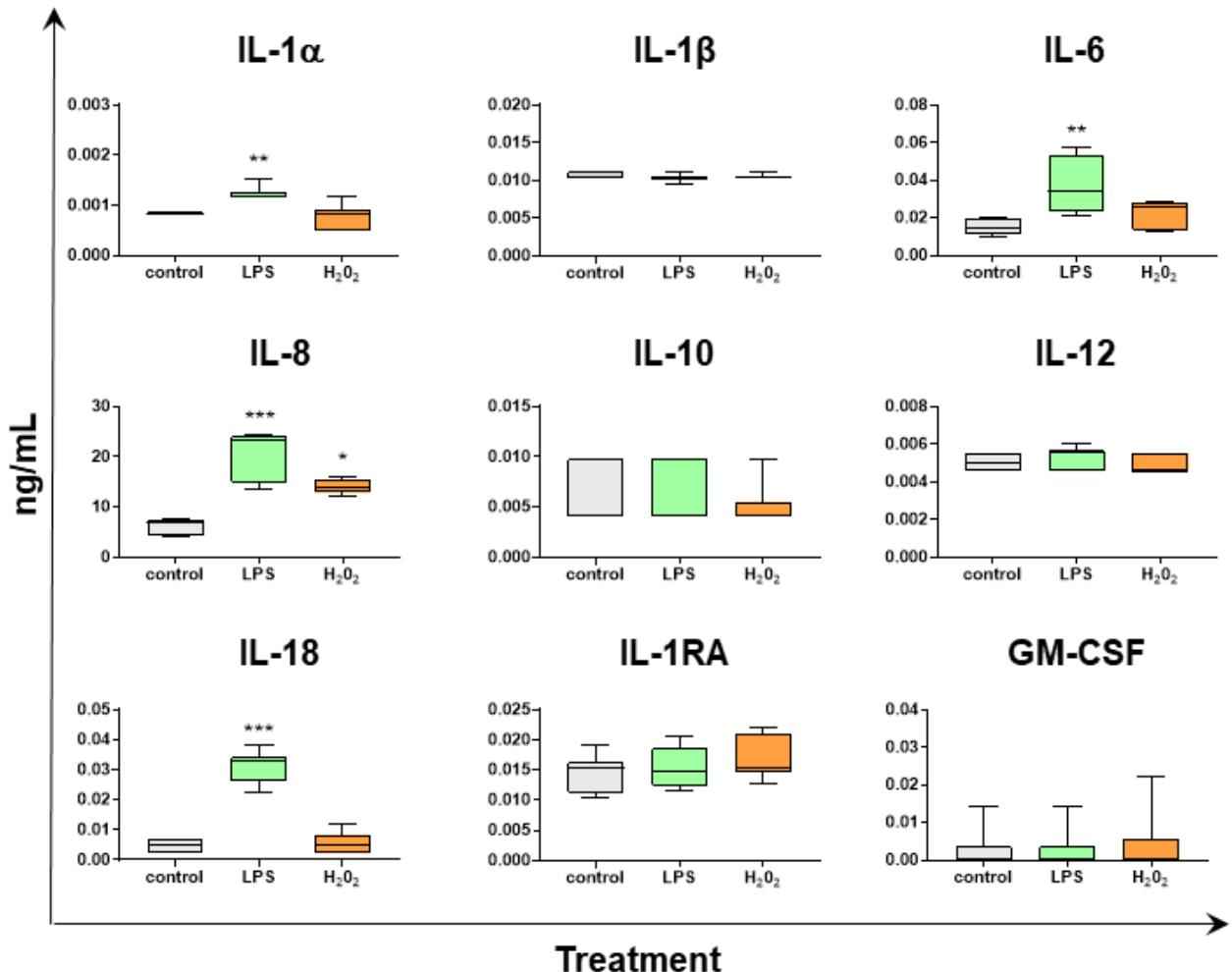


Figure S2. Cytokine release by IPEC-J2 after 24 h of treatment with LPS or H_2O_2 . IPEC-J2 were left untreated (control, grey), stimulated with LPS (green) or H_2O_2 (orange). After, culture supernatants were collected, and levels of several cytokines were determined through multiplex ELISA. Data are presented as box-and-whisker plots displaying median and interquartile range (boxes) and minimum and maximum values (whiskers). Differences (control vs LPS; control vs H_2O_2) were evaluated through ANOVA followed by Dunnett's multiple comparison test or a Kruskal-Wallis test followed by Dunn's multiple comparison test; * p<0.05, ** p<0.01 and *** p<0.001.

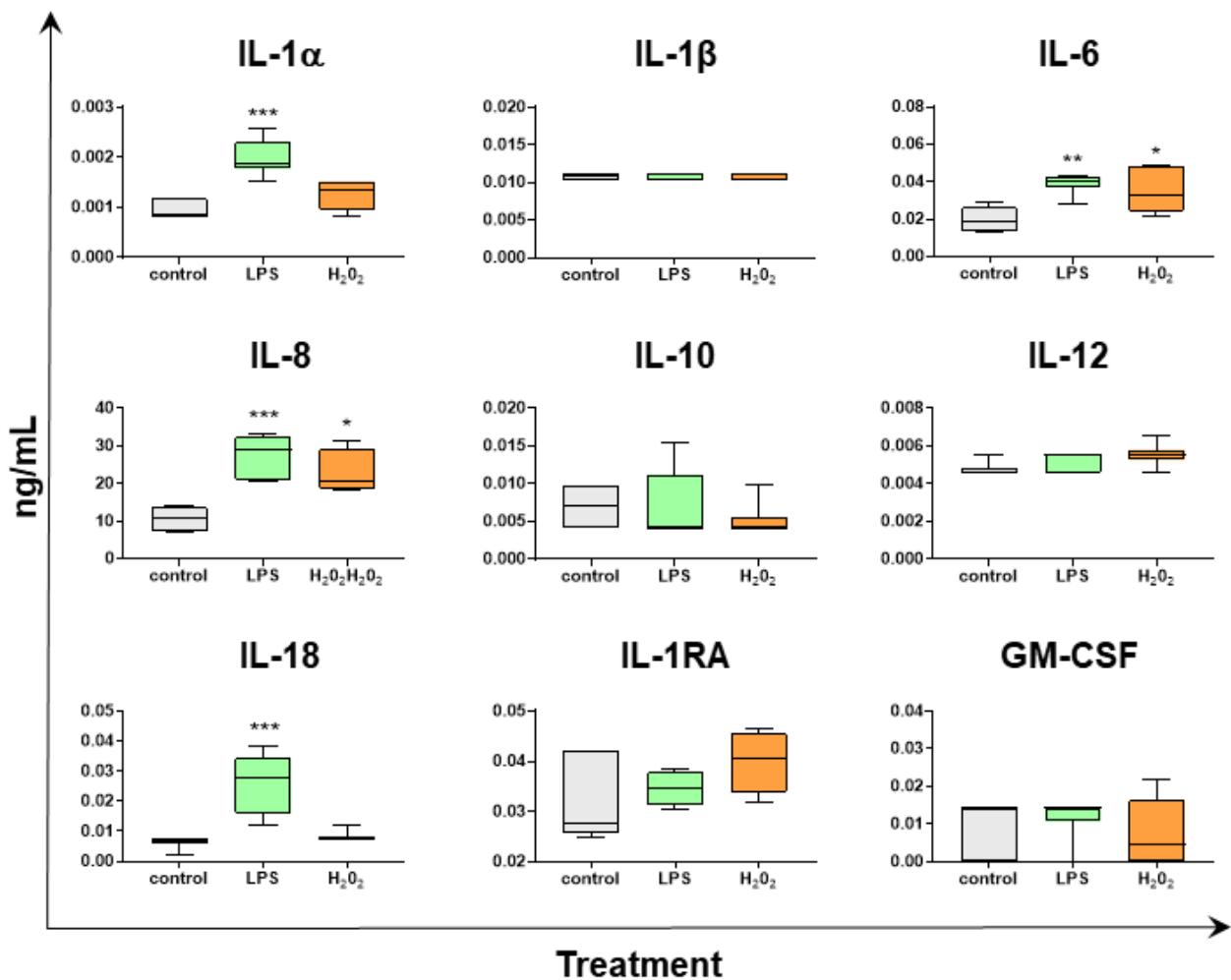


Figure S3. Cytokine release by IPEC-J2 after 48 h treatment with LPS or H_2O_2 . IPEC-J2 were left untreated (control, grey), stimulated with LPS (green) or H_2O_2 (orange). After 48 h, culture supernatants were collected, and levels of several cytokines were determined through multiplex ELISA. Data are presented as box-and -whisker plots displaying median and interquartile range (boxes) and minimum and maximum values (whiskers). Differences (control vs LPS; control vs H_2O_2) were evaluated through ANOVA followed by Dunnett's multiple comparison test or a Kruskal-Wallis test followed by Dunn's multiple comparison test; * p<0.05, ** p<0.01 and *** p<0.001.

Table S1. Primer set sequences for target and reference genes.

Gene	Primer sequences	Amplicon length	Source
<i>IL12A</i>	For - 5' - ATGCCTCAACCACCTCCAAA - 3' Rev - 5' - TGTGCTGGTTTATCTTGGTGA - 3'	135	[12]
<i>IL12B</i>	For - 5' - TCAGGGACATCATCAAACCA - 3' Rev - 5' - GAACACCAAACATCAGGGAAA - 3'	140	[58]
<i>EBI3</i>	For - 5' - CAACGTCACAGCCATCCAC - 3' Rev - 5' - CGTTTCCACTGCACCCAA - 3'	140	[12]
<i>IL6</i>	For - 5' - TGGCTACTGCCTCCCTACC - 3' Rev - 5' - CAGAGATTTGCCGAGGATG - 3'	131	[59]
<i>CXCL8</i>	For - 5' - TTGATGCCAGTGCATAAATA - 3' Rev - 5' - CTGTACAACCTCTGCACCCA - 3'	175	[59]
<i>IL18</i>	For - 5' - CGTGTGAGGATATGCTGATT - 3' Rev - 5' - TGTTACTGCCAGACCTCTAGTGA - 3'	106	[59]
<i>TNFA</i>	For - 5' - TGCTACTGCACCTCGAGGTTATC - 3' Rev - 5' - GTGGCGACGGCTTATCTG - 3'	125	[60]
<i>TGFB1</i>	For - 5' - CGC GTG CTA ATG GTG GAA AG - 3' Rev - 5' - CCG ACG TGT TGA ACA GCA TA - 3'	87	[12]
<i>NOS2</i>	For - 5' - CGTTATGCCACCAACAATGG - 3' Rev - 5' - AGACCCGGAAGTCGTGCTT - 3'	84	[61]
<i>MUC2</i>	For - 5' - GCAAAGACCTCTAACATGGCC - 3' Rev - 5' - AGACCAACTCAGCCTCAG - 3'	132	[12]
<i>MMP9</i>	For - 5' - TGT GGA CCA GAT GTT CCC C - 3' Rev - 5' - AGT CCA CCT GAT TCA CCT CG - 3'	130	[12]
<i>TLR1</i>	For - 5' - AGA TTT CGT GCC ACC CTA TG - 3' Rev - 5' - CCT GGG GGA TAA ACA ATG TG - 3'	276	[61]
<i>TLR2</i>	For - 5' - CGG CTT CCA AGG ATG GAG AAA - 3' Rev - 5' - TCC AGA GAG TTG ACC TTG CAG - 3'	71	[61]
<i>TLR3</i>	For - 5' - TGAAGAACTTGATTCCTGGCA - 3' Rev - 5' - GGCATGAAAACACCCCTGGAG - 3'	93	[61]
<i>TLR4</i>	For - 5' - TGGCAGTTCTGAGGAGTCATG - 3' Rev - 5' - CCGCAGCAGGGACTTCTC - 3'	71	[59]
<i>TLR5</i>	For - 5' - TCAAAGATCCTGACCATCACA - 3' Rev - 5' - CCAGCTGTATCAGGGAGCTT - 3'	59	[59]
<i>TLR7</i>	For - 5' - GTGAAATTGCCCTCGTTGT - 3' Rev - 5' - GATGGATCTGTAGGGGAGCA - 3'	77	[61]
<i>TLR8</i>	For - 5' - AAGACAACCAGTTACGTGAAATACC - 3' Rev - 5' - GGGTGTAAAAGATAATGACAGCAC - 3'	236	[62]
<i>TLR9</i>	For - 5' - AGGACTTCATGCCAAACTGC - 3' Rev - 5' - CGAGCAAACATCTCCGACTG - 3'	90	[61]
<i>DEFB1</i>	For - 5' - TGCCACAGGTGCCGATCT - 3' Rev - 5' - CTGTTAGCTGCTTAAGGAATAAAGGC - 3'	80	[59]
<i>DEFB4A</i>	For - 5' - CCAGAGGTCCGACCACTA - 3' Rev - 5' - GGTCCCTTCAATCCTGTT - 3'	87	[59]
<i>MYD88</i>	For - 5' - GCA GCT GGA ACA GAC CAA CT - 3' Rev - 5' - GTG CCA GGC AGG ACA TCT - 3'	62	[63]
<i>NFKB1</i>	For - 5' - CCCATGTAGACAGCACCCACCTATGAT - 3' Rev - 5' - ACAGAGGCTCAAAGTTCTCCACCA - 3'	131	[63]
<i>RELA</i>	For - 5' - CGAGAGGAGCACGGATACCA - 3' Rev - 5' - GCCCGTGTAGCCATTGA - 3'	61	[63]
<i>IFNA1</i>	For - 5' - GGCTCTGGTGCATGAGATGC - 3' Rev - 5' - GCCTTCTCCTGAATCTGTCTCA - 3'	546	[64]

<i>IFNB</i>	For - 5' - AGTTGCCTGGGACTCCTCAA - 3' Rev - 5' - CCTCAGGGACCTCGAAGTTCAT - 3'	59	[65]
<i>IRF3</i>	For - 5' - GGGAAAGGAGGCCTGTTGAC - 3' Rev - 5' - ACCAGAGGGTGTACCGTGTT - 3'	92	[63]
<i>GAPDH</i>	For - 5' - ATGGTGAAGGTGGAGTGAA - 3' Rev - 5' - AGTGGAGGTCAATGAAGGGG - 3'	61	[12]