

**Supplementary material**

**Supplementary Table S1** *Primer used for real-time quantitative PCR analysis of*

*ZO-1, occludin and β-actin*

Gene <sup>1</sup>	primer sequence (5'-3')	accession number
<i>ZO-1</i>	Forward: GAGGATGGTCACACCGTGGT Reverse: GGAGGATGCTGTTGTCTCGG	XM_021098896.1
<i>occludin</i>	Forward: ATGCTTCTCAGCCAGCGTA Reverse: AAGGTTCCATAGCCTCGGTC	NM_001163647.2
<i>β-actin</i>	Forward: ATGCTTCTAGACGGACTGCG Reverse: GTTCAGGAGGCTGGCATGA	XM_003357928.4

<sup>1</sup> ZO-1, zonula occludens-1

**Supplementary Table S2** Effects of the early lactoferrin intervention on the diversity and richness of jejunal and ileal bacterial community of suckling piglets at the 0.03 dissimilarity level<sup>1</sup>

Items <sup>2</sup>	Day 8				Day 21			
	CON	LF	SEM <sup>3</sup>	P-value	CON	LF	SEM	P-value
<b>Jejunum</b>								
Ace	200.88	176.69	21.45	0.297	385.55	416.59	41.25	0.714
Chao 1	188.61	180.86	23.66	0.743	372.91	393.82	45.33	0.374
Shannon	1.39	0.911	0.28	0.086	2.11	2.26	0.35	0.787
Simpson	0.54	0.71	0.09	0.121	0.35	0.32	0.15	0.846
<b>Ileum</b>								
Ace	217.95	188.58	15.41	0.265	339.84	289.98	18.66	0.214
Chao 1	218.32	178.89	14.22	0.068	215.06	296.47	18.56	0.364
Shannon	2.17	1.59	0.45	0.202	2.86	2.64	0.18	0.364
Simpson	0.32	0.51	0.11	0.127	0.14	0.17	0.06	0.514

<sup>1</sup> Values are means of 6 replicates per treatment.

<sup>2</sup> CON, Control; LF, Lactoferrin; ACE and Chao 1 represented the species richness; Simpson index and Shannon index represented species diversity. Higher values (Ace, Chao 1 and Shannon index) represented the higher richness and diversity, while higher values (Simpson index) represented the lower diversity.

<sup>3</sup> SEM, standard error of mean.