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Effects of toll like receptor 4 overexpression on blood parameters, body composition, and gastrointestinal microbial diversity in sheep

Xu- $Ting\ Song^1$, Duo- $Wei\ Zhao^1$, $Qi\ Lu^1$, $Jun\ Lu^1$, Mei- $Yu\ Qi^2$, Ming- $Hai\ Lu^3$, Xiu- $Qin\ Yang^1*$, Yu- $Chang\ Yao^1*$

Xu-Ting Song and Duo-Wei Zhao contributed equally to this work

The authors are fully responsible for both the content and the formal aspects of the electronic supplementary material. No editorial adjustments were made.

Electronic Supplementary Material (ESM)

Figure S1. Rarefaction curve analysis of all samples

¹College of Animal Science and Technology, Northeast Agricultural University, Harbin, P.R. China
²Institute of Animal Husbandry, Heilongjiang Academy of Agricultural Sciences, Harbin,
P.R. China

³Department of Animal Science, Heilongjiang State Farms Science Technology Vocational College, Harbin, P.R. China

^{*}Corresponding authors: xiuqin163@163.com; yaoyc@neau.edu.cn

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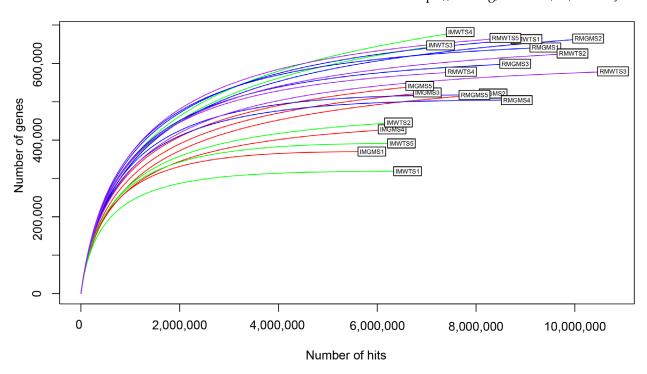


Figure S1. Rarefaction curve analysis of all samples
The *x*-axis shows the number of hits. The *y*-axis number of genes. RMGMS: rumen microbiota of genetically modified sheep. RMWTS: rumen microbiota of wild-type sheep. IMGMS: intestinal microbiota of genetically modified sheep. IMWTS: intestinal microbiota of wild-type sheep