

The underlying microbial mechanism of epizootic rabbit enteropathy triggered by a low fiber diet

DingXing Jin^{a*}, HuaWei Zou^{a*}, SiQiang Liu^a, LiZhi Wang^a, Bai Xue^a, De Wu^a, Gang Tian^a, Jingyi Cai^a, TianHai Yan^b, ZhiSheng Wang^a & QuanHui Peng^a

^a Institute of Animal Nutrition, Key Laboratory of Bovine Low-Carbon Farming and Safe Production, Sichuan Agricultural University, Ya'an, Sichuan, 625014, PR China

^b Agri-Food and Biosciences Institute, Hillsborough, Co. Down BT26 6DR, United Kingdom

*This two authors contributed equally to this paper.

Correspondence and requests for materials should be addressed to Q.P. and Z.W. (email: pengquanhui@126.com , wangzs67@163.com)

Address: Animal nutrition institute, Xinkang road 46#, Yucheng, Ya'an, Sichuan province. Post code: 625014

Tel: +86 15283511570

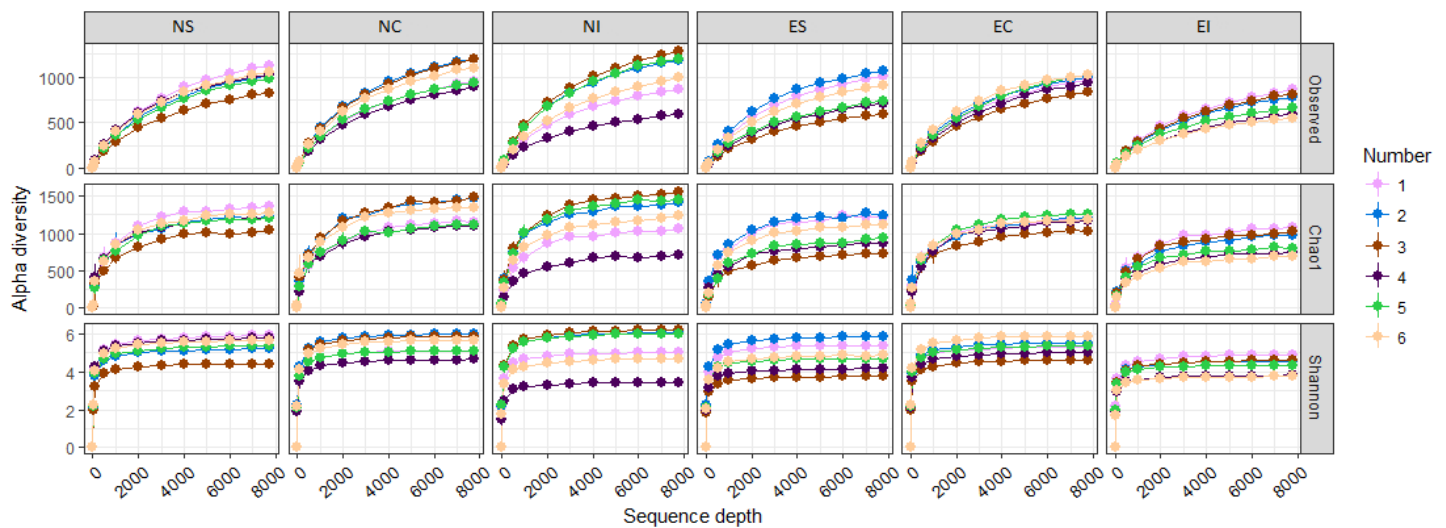


Figure S1. Diversity estimation of the 16S rRNA gene libraries in the stomach, small intestine and caecum of normal and ERE rabbits from the sequencing data.