Supplementary information:

Fusimonas intestini gen. nov., sp. nov., a novel intestinal bacterium of the family *Lachnospiraceae* associated with diabetes in mice

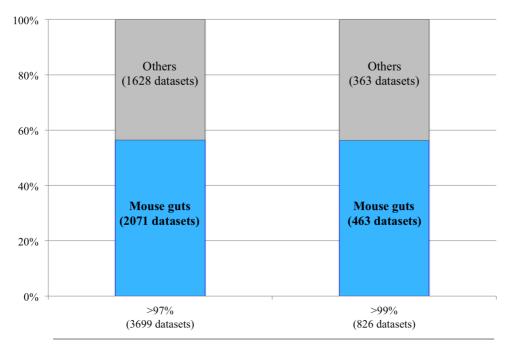
Hiroyuki Kusada¹[§], Keishi Kameyama²[§], Xian-Ying Meng¹, Yoichi Kamagata¹ & Hideyuki Tamaki^{1*}

^{1.} Bioproduction Research Institute, National Institute of Advanced Industrial Science and Technology (AIST), Central 6, 1-1-1 Higashi, Tsukuba, Ibaraki 305-8566, Japan ² Frontier Research Laboratories, Institute for Innovation, Ajinomoto Co., Inc., 1-1 Suzuki-cho, Kawasaki, Kanagawa 210-8681, Japan

[§]These authors contributed equally to this work.

*Correspondence: Hideyuki Tamaki, Bioproduction Research Institute, National Institute of Advanced Industrial Science and Technology (AIST), Central 6-10, Higashi 1-1-1, Tsukuba, Ibaraki 305-8566, Japan

Tel: +81-29-861-6591. Fax: +81-29-861-6587. E-mail: tamaki-hideyuki@aist.go.jp



Total 88,579 of 16S rRNA gene amplicon datasets from 96 environments

Supplementary Figure S1. Potential habitability of *F. intestini* strain AJ110941^P based on IMNGS search using the 16S rRNA gene sequence (AB861470). Prevalence of *F. intestini*-related sequences (>97% and >99% sequence similarity) against all 16S rRNA sequence datasets. The relative abundances of *F. intestini*-related 16S rRNA gene sequences in mouse guts and non-mouse guts are shown in blue and gray, respectively.