Supplementary Material

GPR35-mediated kynurenic acid sensing contributes to maintenance

of gut microbiota homeostasis in ulcerative colitis

Di Wang^{1,2,†}, Wenbao Wang^{4,†}, Xue Bing², Chenguang Xu², Jiahua Qiu², Jiangang Shen², Jinwen Huang², Junda Li², Pi Liu², Biao Xie^{2,3}*

¹ School of Life and Health Technology, Dongguan University of Technology, Dongguan 523808, P.R. China

² Department of Gastroenterology, Affiliated Longhua People's Hospital, Southern Medical University (People's Hospital of Longhua), Shenzhen 518109, P.R. China
³ Department of Gastroenterology, Guangzhou Eighth People's Hospital, Guangzhou Medical University, Guangzhou 510440, P.R. China

⁴ College of Pharmacy, Qiqihar Medical University, Qiqihar 161006, P.R. China

*Correspondence should be addressed to Biao Xie, Chief physician, Department of Gastroenterology, People's Hospital of Longhua, NO.38 Jinglong Construction Road, Longhua District, Shenzhen 518109, China. Tel/Fax: +0755-27741585; E-mail: xiebiaohxq@163.com

[†]These authors contributed equally to this work



Figure S1. Schematic representation of rat disposal process in each group.



Figure S2. α-diversity analysis of gut microbiota by Observed_species, Shannon, Simpson, Faith_pd, Goods_coverage and Pielou_e indexes.



Figure S3. The LEfSe analysis of the differential gut microbiome in Cont_T and DSS_T groups (A), Cont_T and KA_T groups (B), Cont_T and C_KA_T groups (C), KA_T and C_KA_T groups (D). The size of the circles was based on the relative abundance, the solid circle represented LDA > 3.0, p < 0.05, and the hollow circle represented p > 0.05.



Figure S4. (A, B) Rarefaction curve (A) and Rank abundance curve (B) of each group.(C) α-diversity analysis of gut microbiota by Chao1, Observed_species, Shannon, Simpson, Faith_pd, Goods_coverage and Pielou_e indexes.

The raw unedited western blot images presented in Figure 1E

p-ERK1/2 ERK1/2 β -actin

The raw unedited western blot images presented in Figure 4E

p-ERK1/2 ERK1/2 β -actin
