

Supplemental Figure S4: FxrKO mice with HCC have a bloom of Verrucomicrobia, Tenericutes and Cyanobacteria. (A-C) Male WT (n=15) and FxrKO (n=16) mice were aged for 16 months, fecal samples were snap frozen and bacterial DNA was extracted for 16S microbiota analysis. (A) α -diversity, (B) β -diversity via Bray-Curtis, and (C) Linear discriminant analysis Effect Size (LEfSe) comparing WT and FxrKO. Results are expressed as means \pm SEM. N.S., non-significant.