

Additional file 12: Effect of gut microbiota-PXR interaction on hepatic fatty acid and xenobiotic metabolism in female mice

(A) RT-qPCR analysis of hepatic genes involved in fatty-acid homeostasis.

(B) Relative abundance of hepatic fatty acids.

(C) Hepatic neutral lipid quantification.

(D) RT-qPCR analysis of hepatic genes involved in xenobiotic metabolism.

