

SUPPLEMENTAL TABLES

Table S1

Confirmatory q-PCR for several key genes from microarray results

Gene	Symbol	Wt CEHPOBA-treated vs. <i>Fah</i>^{-/-} on NTBC	<i>Fah</i>^{-/-} off NTBC vs. <i>Fah</i>^{-/-} on NTBC
Tyrosine aminotransferase	<i>Tat</i>	-26.18	-17.40
Cytochrome P450, family 2, subfamily e, polypeptide 1	<i>Cyp2e1</i>	-8.12	-6.55
Apolipoprotein A2	<i>Apoa2</i>	-2.34	-6.10
Alpha-fetoprotein	<i>Afp</i>	1042.3	1001.2
Cyclin-dependent kinase inhibitor 1A	<i>Cdkn1a</i>	250.7	200.1
Regulator of calcineurin 1	<i>Rcan1</i>	102.3	76.8

*Footnotes: Genes that are similarly regulated in wild-type mice treated with CEHPOBA and *Fah*^{-/-} mice off NTBC compared to *Fah*^{-/-} on NTBC. Values represent fold change.*