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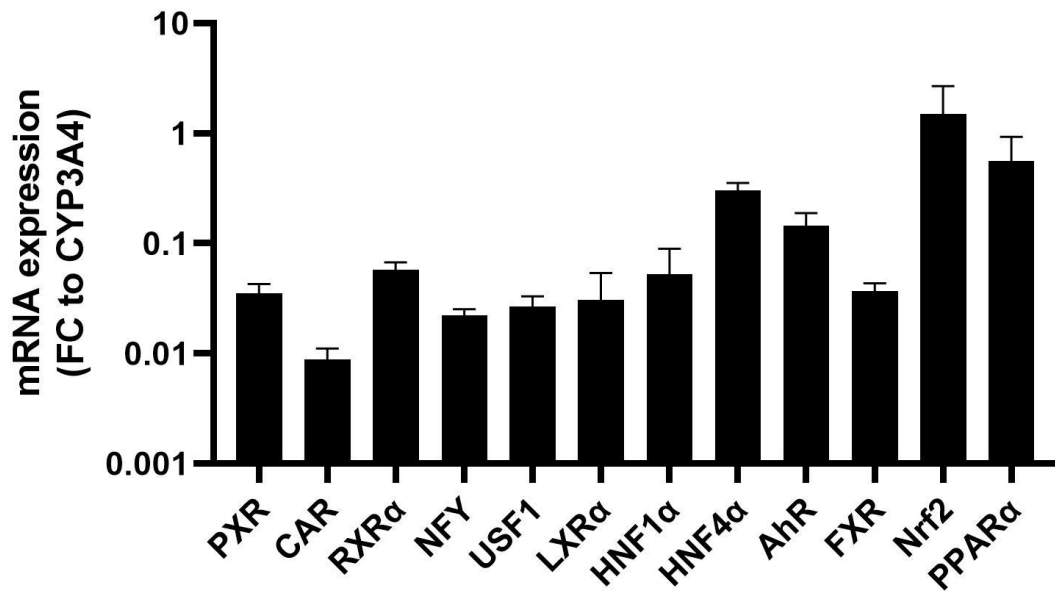
Supplemental Data

Drug Metabolism and Disposition

CYP and non-CYP drug metabolizing enzyme families exhibit differential sensitivities towards pro-inflammatory cytokine modulation

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Supplemental Fig. 2. Basal mRNA expression levels of DME regulating transcription factors and nuclear receptors in HepaRG cells. mRNA expression of the gene of interest was normalized to the housekeeping gene *RPLP0*, and presented as a fold change compared to basal *CYP3A4* expression in HepaRG cells. All values are means + SEM from 8 independent experiments.