



Supplementary Figure 1. BMMs from wild-type mice were treated with indicated concentrations of BaP and I3C, respectively, or DMSO as control for 6 h. Changes in gene transcription were assessed by real-time PCR after activation with hk-S.E. for 20 h. Relative *Il12rb1* (A), *Slpi* (B), *Ido2* (C), *Gsta3a* (D), *Spn* (E) and *Igf1r* (F) expression for BaP and I3C treatment. Data represent the mean of the relative expression \pm SEM after normalization with housekeeping genes *Alas1* and *Hprt* (n=4). * $p \leq 0.05$ indicates significant differences between treated and untreated cells.