

Supplementary Materials: Sub-chronic Difenoconazole Exposure Induced Gut Microbiota Dysbiosis in Mice

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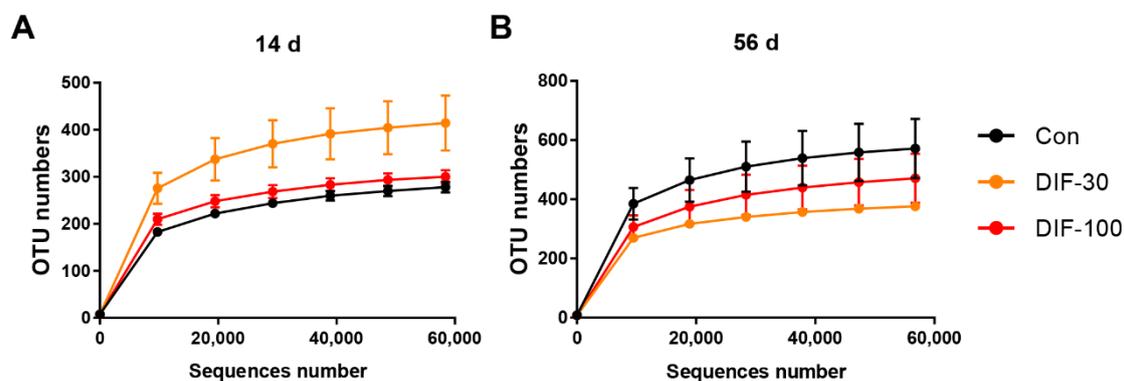


Figure S1. Rarefaction curves of 16S rRNA sequencing after (A) 14 days of DIF exposure and (B) 56 days of DIF exposure. The presented data are the Mean \pm SEM ($n = 6$).