Sex dependent effects of silver nanoparticles on the zebrafish gut microbiota

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(14 pages)

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## Table S1 MiSeq sequences quality information

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SD: 5718 ± 74

Average: 28281 ± 5718
SD: 6090 ± 49
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Figure S8 Boxplots (percentage of sequences) of the six most represented genera in male zebrafish groups.
Figure S9 Boxplots (percentage of sequences) of the six most represented genera in female zebrafish groups.
Figure S10 The relative abundance of *Cetobacterium* and *Aeromonas* in male and female zebrafish. Asterisks indicates significant differences (* p < 0.05, ** p \leq 0.01).