

S1 Fig: Relative abundance ofbacterial genera and number oftotal readsincontrol samples

Negative control samples repeatedly showed high abundances of ASVs belonging to Anaerobacillus, Alkalibacterium, Bacteroides, Shewanella and Serratia.

<u>blank PCR</u>: negative control from extraction buffer subjected to PCR and sequencing; <u>E. coli PCR</u>: positive control with E. coli DNA in extraction buffer subjected to PCR and sequencing; <u>extracted blank</u>: negative control from extraction buffer subjected to DNA extraction, PCR and sequencing; <u>extracted blank tube</u>: negative control from sterile saline sampled through urinary catheters and tubes imitating preanalytics of catheter urine samples