

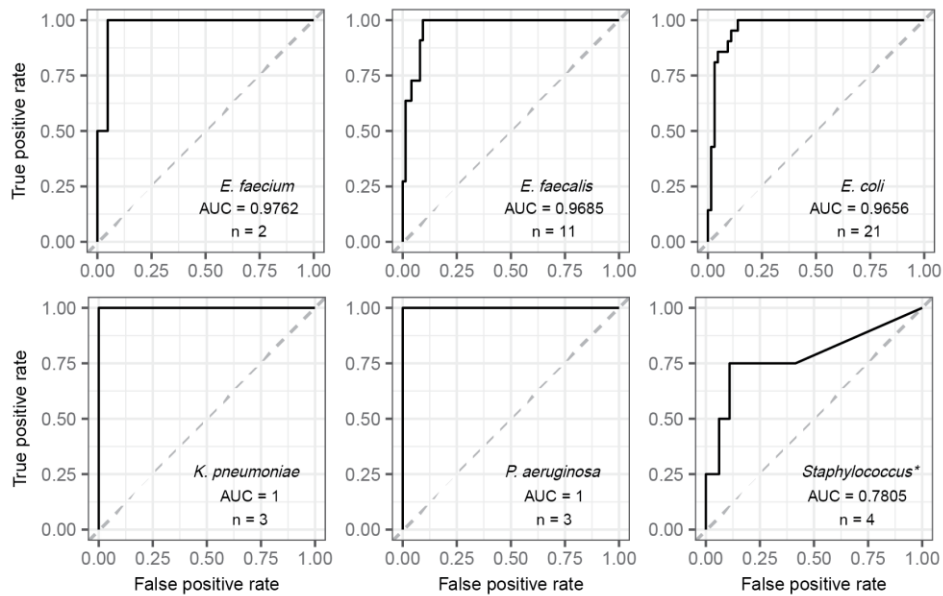
Supplementary Information for:

Urinary cell-free DNA is a versatile analyte for monitoring infections of the urinary tract

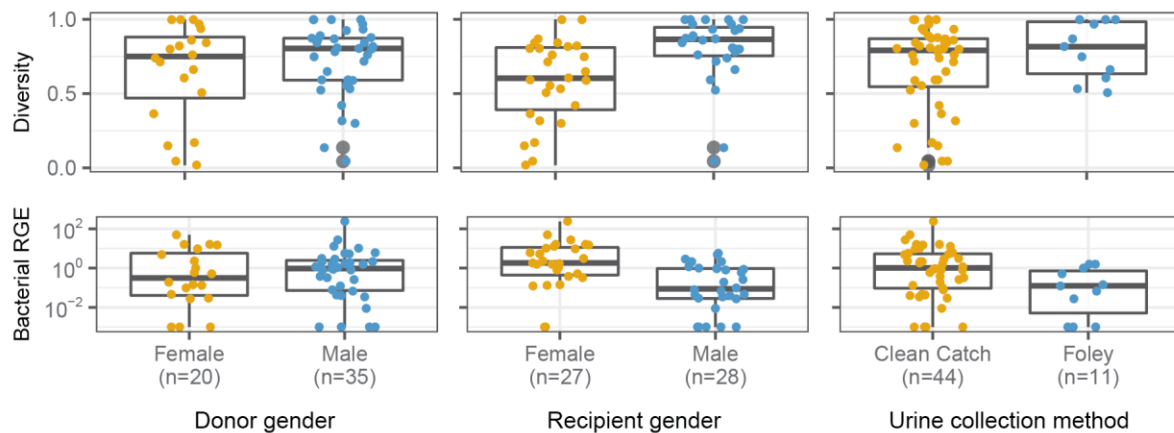
Burnham et al.

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Supplementary Figures 1-2



Supplementary Figure 1: Analysis of performance of urinary cfDNA assay to identify bacterial UTI against conventional bacterial culture. Receiver operating characteristic analyses of the performance of urinary cfDNA in identifying UTIs with the five suspected causative bacterial species and one suspected causative bacterial genus (marked with *). Area under curve = AUC, n = number of positive cultures. Total number of samples consisted of those clinically determined to have UTI based on positive urine culture, and those negative for UTI based on a bacterial count of < 10,000 CFU/mL in bacterial culture (total $n = 86$).



Supplementary Figure 2: Species-level diversity (Simpson index) and total relative genome abundance (RGE) as function of gender and urine collection method. The significance of clinical parameters on the bacteriome is tested using two-tailed Wilcoxon rank sum test for samples that were either UTI negative or taken within three days of transplantation ($n = 55$): donor gender ($p_{\text{diversity}} = 0.59$ and $p_{\text{RGE}} = 0.75$), urine collection method ($p_{\text{diversity}} = 0.26$ and $p_{\text{RGE}} = 0.02$), and recipient gender ($p_{\text{diversity}} = 1.7 \times 10^{-3}$ and $p_{\text{RGE}} = 3.2 \times 10^{-4}$).