



Fig. S1 MA plots of the differentially expressed genes. Differentially expressed genes with a p-value inferior to 0.05 are shown in red. The blue line indicates the FC 2 and -2 ($\log_2 \text{FC} > 1$ and < -1) threshold. The logCPM (count per million) is the average \log_2 count per million for a given gene taken over all the libraries in the dataset. Co-culture (CC) versus *P. aeruginosa* (PA) mono-culture (left panel) and co-culture (CC) versus *S. aureus* (SA) mono-culture (right panel).