Trimming of flow data to remove background and doublets: pMed\_None\_0\_10.fcs 6 log10(SSC-H) log10(SSC-A) all\_data bacteria single\_bacteria 2 log10(FSC-H) log10(SSC-H) all\_data count bacteria calibrated calibrated\_normalised normalised 7<sub>@\*07</sub> single\_bacteria BL1-H