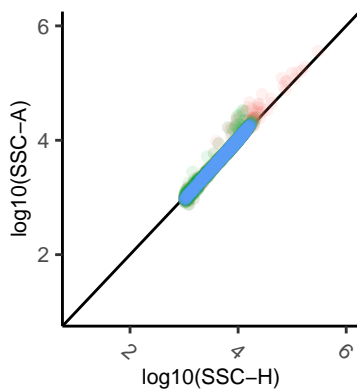
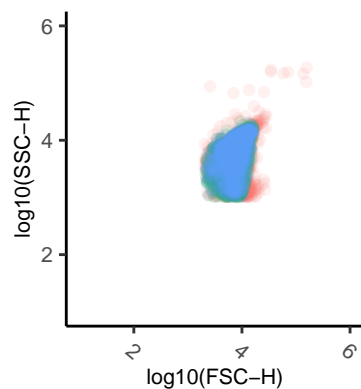
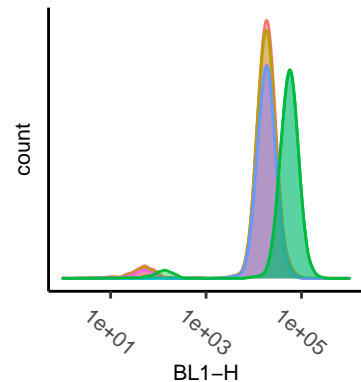


# Trimming of flow data to remove background and doublets:

pMed\_None\_0\_1.fcs



- all\_data
- bacteria
- single\_bacteria



- all\_data
- bacteria
- calibrated
- calibrated\_normalised
- normalised
- single\_bacteria