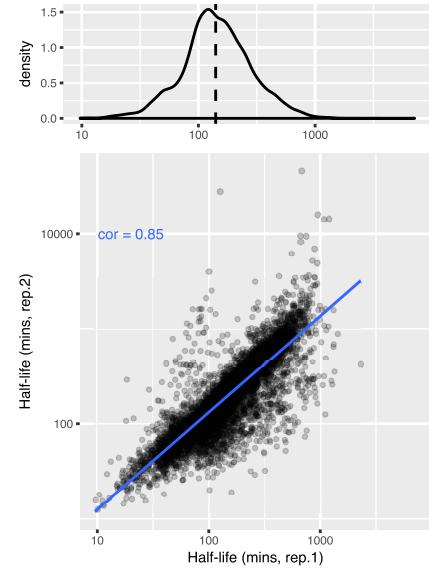
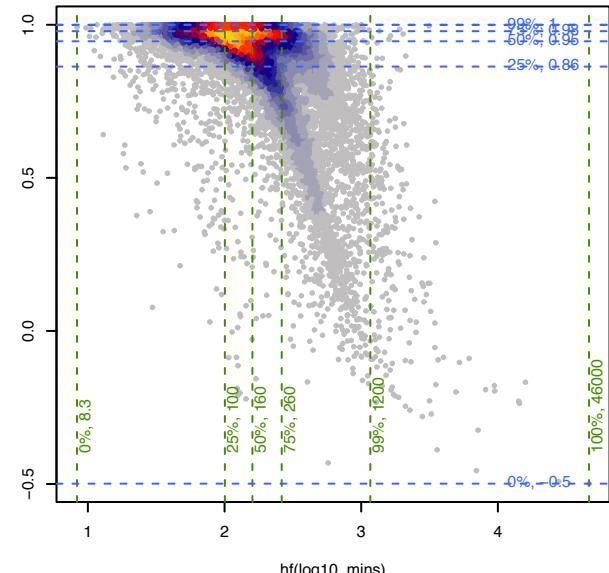


Whole genome scale half-life



Half-life with $\text{adjR2} \geq .86$



Error model (%5~95% genes)

