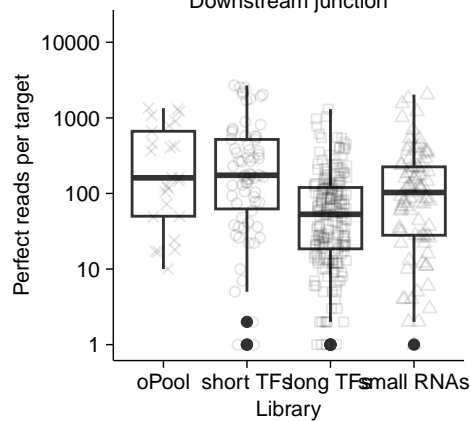
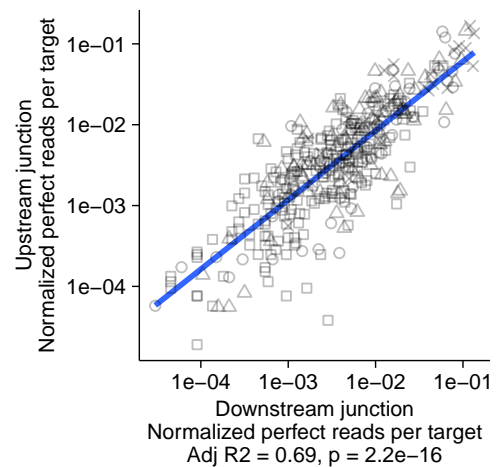
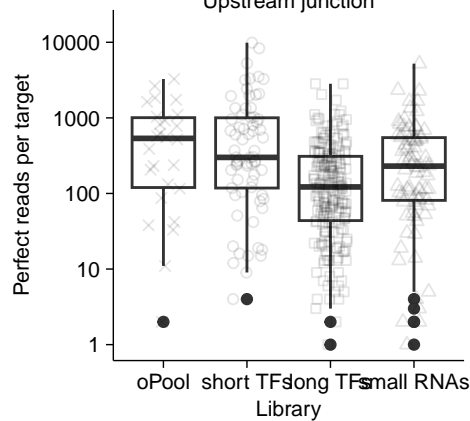


Downstream junction

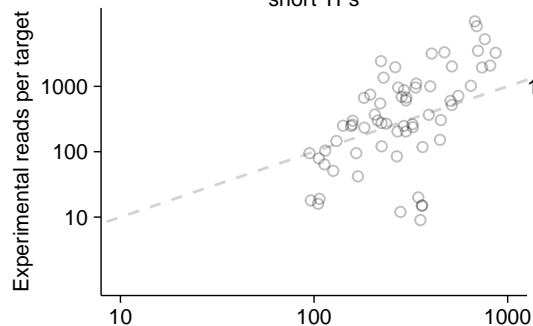


Upstream junction

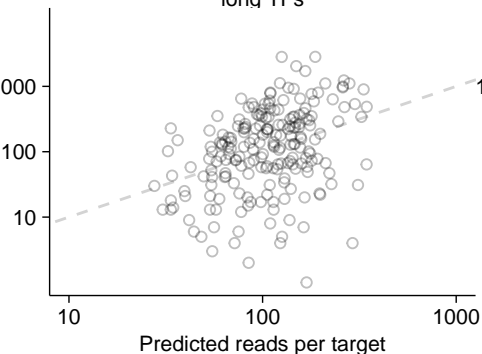


3 var model
Reads per target ~ oligo abundance + homology arm free energy + distance to ori

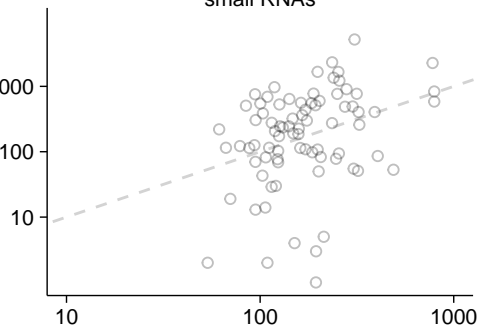
short TFs



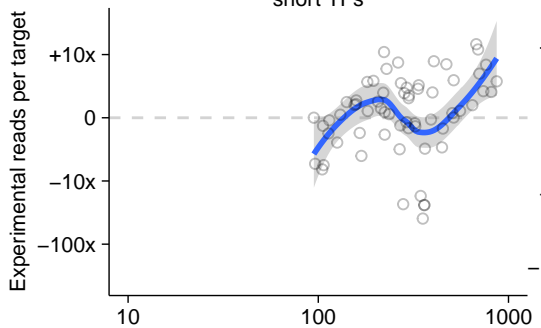
long TFs



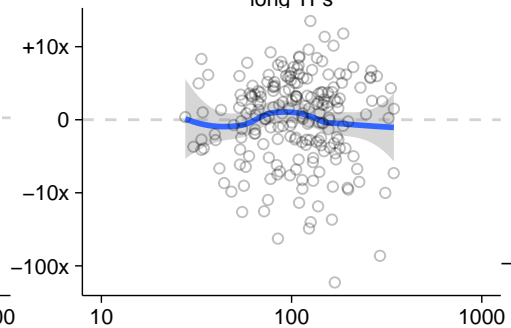
small RNAs



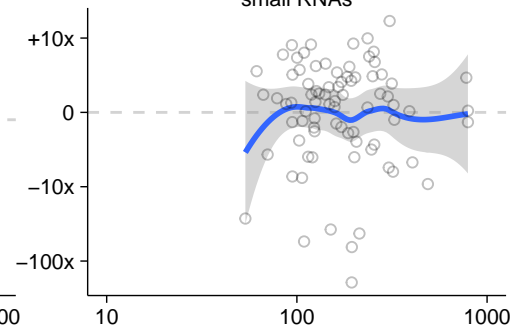
short TFs



long TFs



small RNAs



Predicted reads per target
 short TFs: adj R2 = 0.13 p = 0.0071
 long TFs: adj R2 = 0.115 p = 4.1e-06
 small RNAs: adj R2 = 0.072 p = 0.035