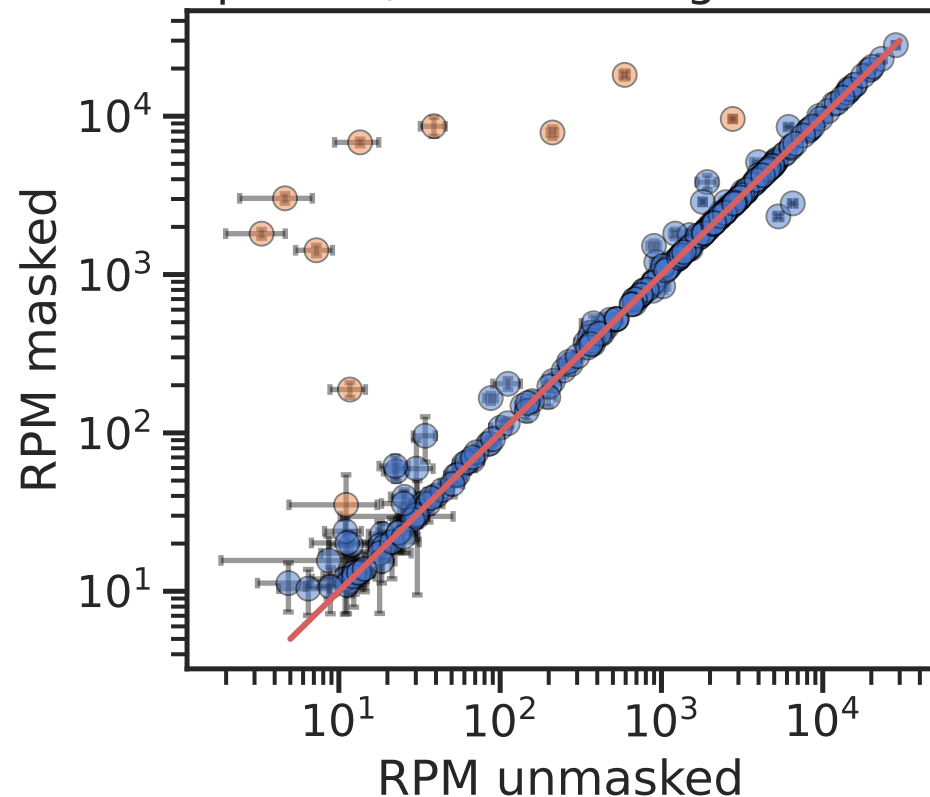
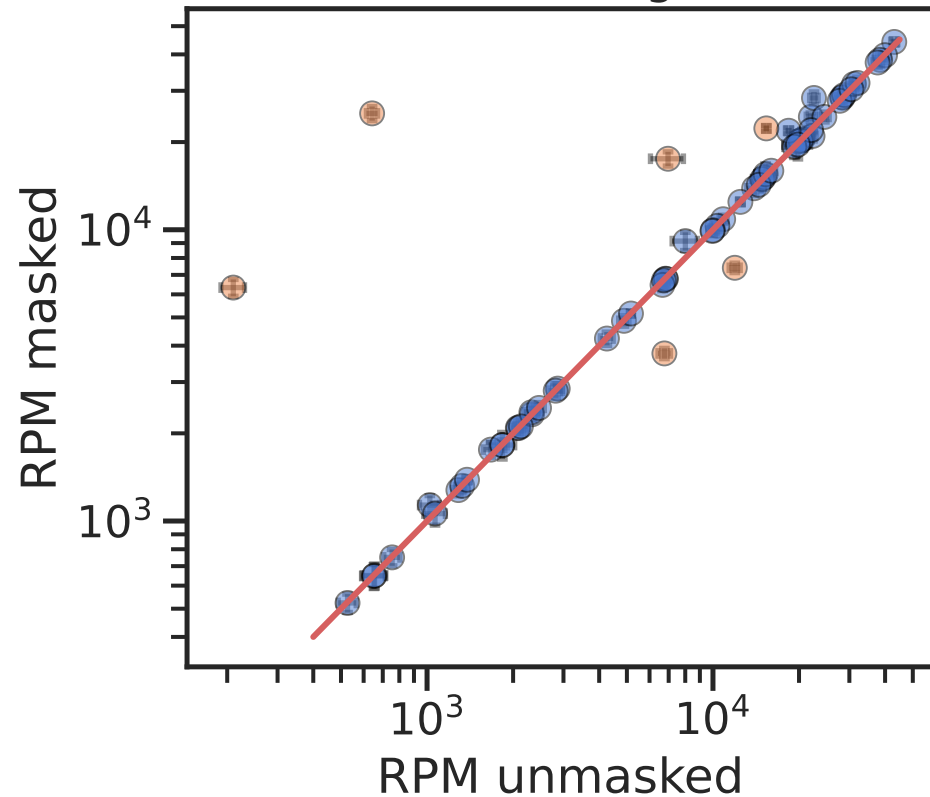


Transcript RPM, effect of alignment masking



Transcript	Fold change
Ser-AGA-2-6	653.63
Ser-AGA-1-1	545.79
Pro-AGG-2-1	504.82
Val-AAC-1-1	222.54
Leu-AAG-1-1	196.14
Asn-GTT-2-8	37.38
Pro-AGG-1-1	30.83
iMet-CAT-3-1	16.05
Ala-CGC-2-1	3.47
Asn-GTT-24-1	3.17

Codon RPM, effect of alignment masking



Codon	: Fold change
Pro-CCU	: 39.06
Ser-UCU	: 30.14
Val-GUU	: 2.52
Ala-GCG	: 1.45
Pro-CCA	: 0.62
Pro-CCG	: 0.56