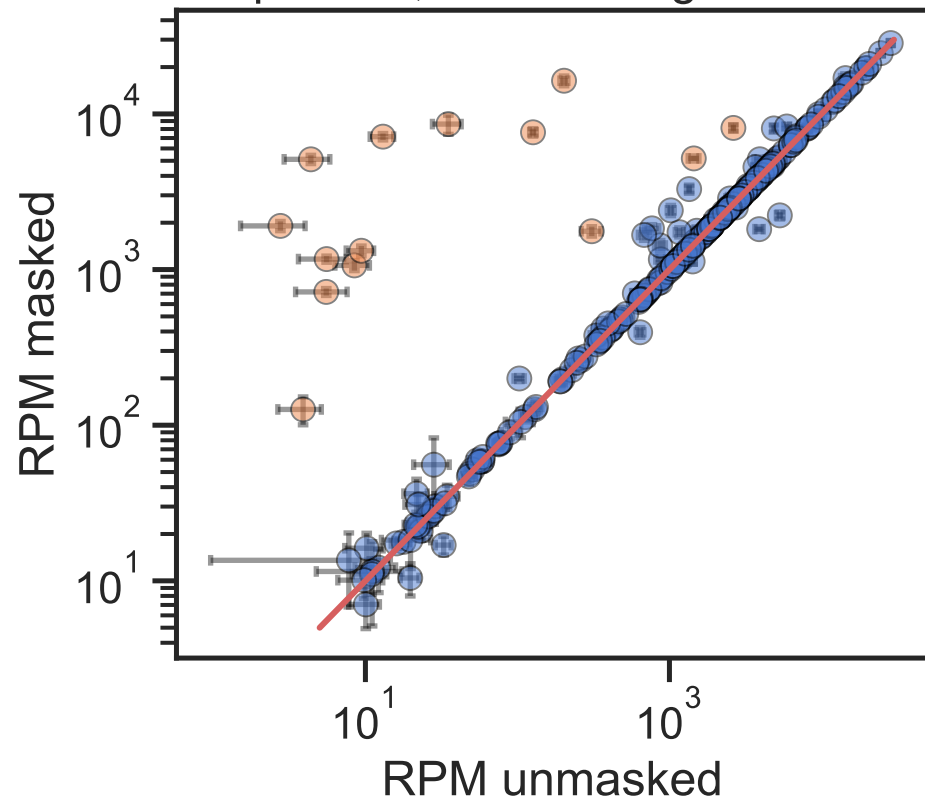
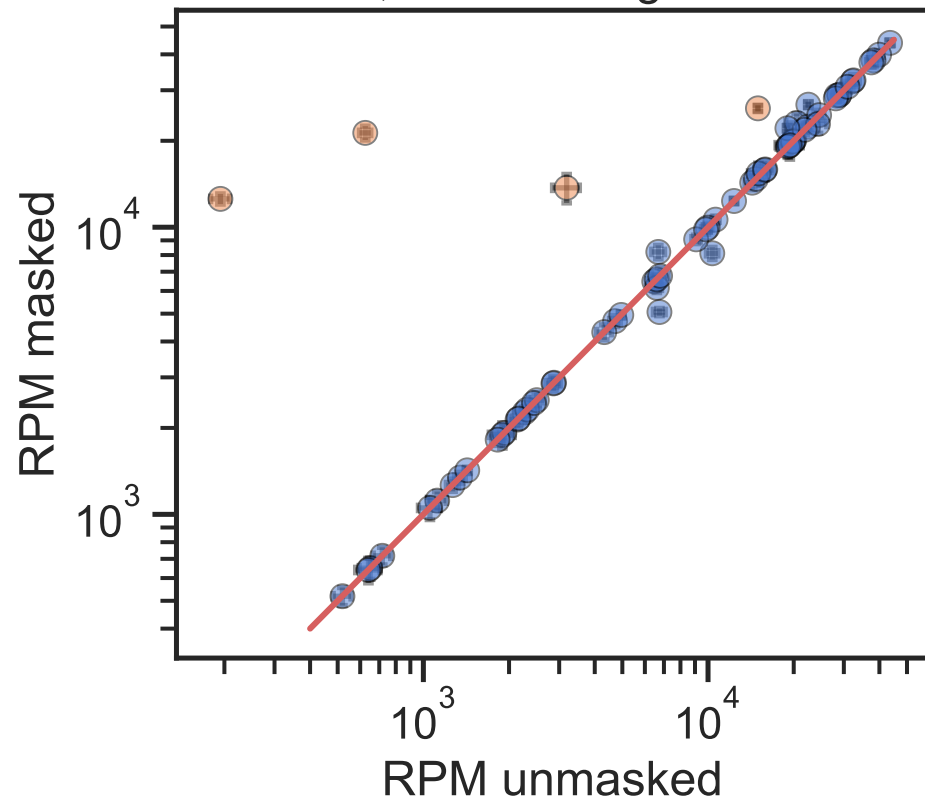


Transcript RPM, effect of alignment masking



Transcript	Fold change
Ser-AGA-2-1	1167.49
Ser-AGA-1-1	691.40
Pro-AGG-2-1	547.57
Val-AAC-1-1	244.99
Tyr-GTA-8-1	210.19
Leu-AAG-1-1	140.51
Ser-GCT-6-1	130.05
Cys-GCA-11-1	125.65
Asn-GTT-2-1	80.66
Tyr-GTA-1-1	60.18
Lys-TTT-7-1	32.46
Cys-GCA-6-1	5.72
Thr-TGT-4-1	3.58
Ala-CGC-2-1	3.09

Codon RPM, effect of alignment masking



Codon	: Fold change
Ser-UCU	: 64.82
Pro-CCU	: 34.09
Val-GUU	: 4.30
Ala-GCG	: 1.73