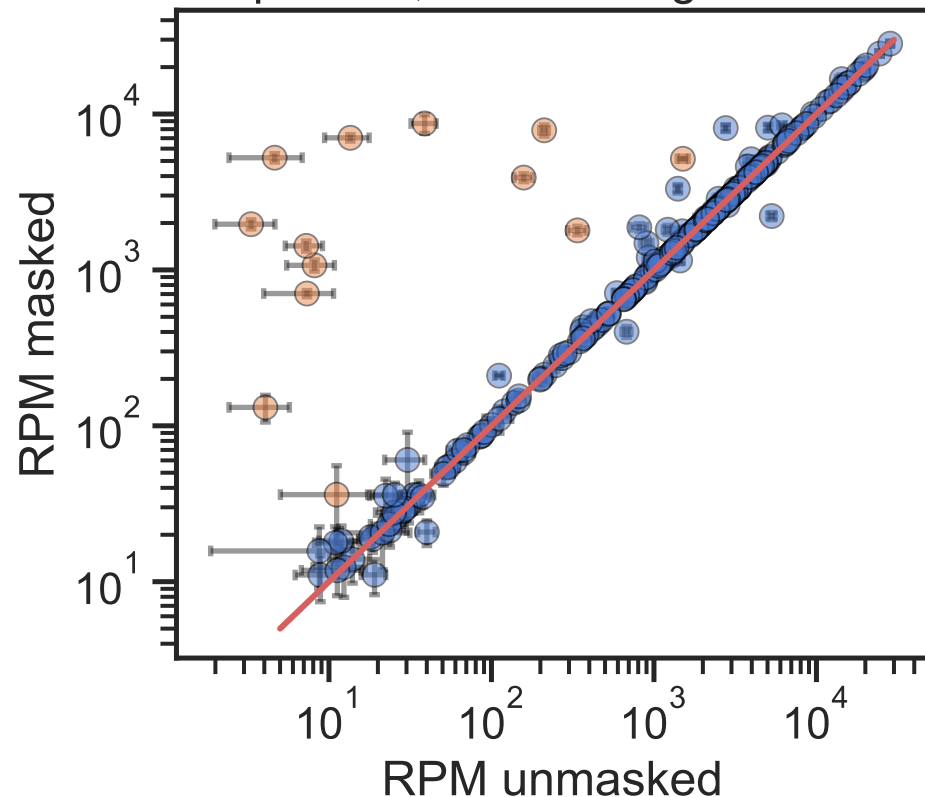
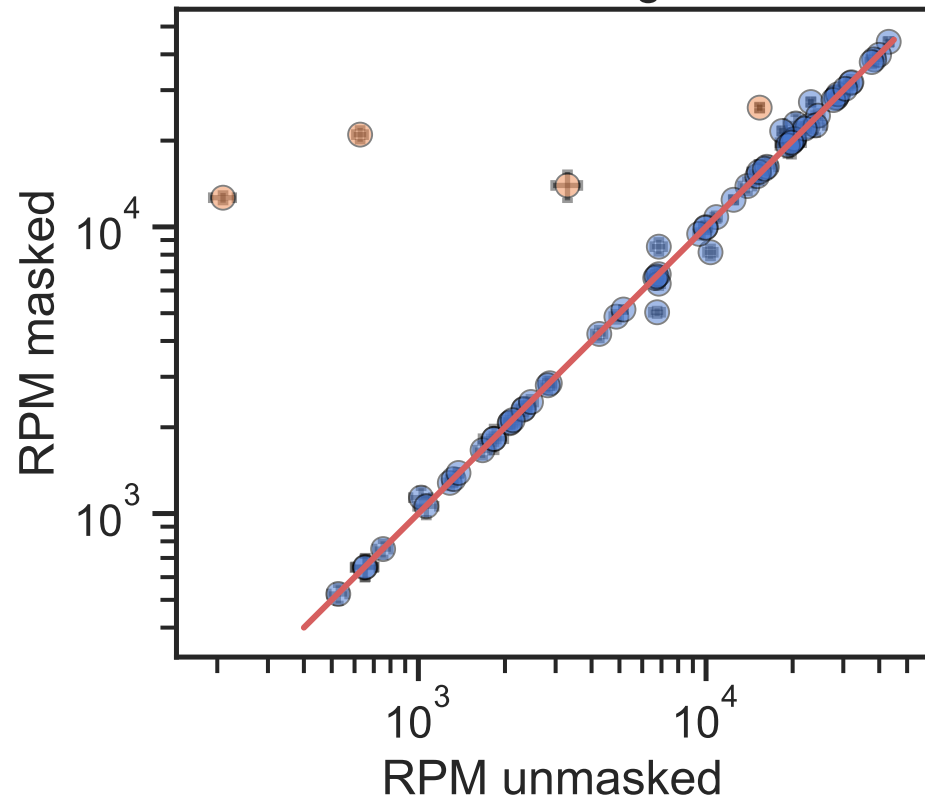


Transcript RPM, effect of alignment masking



Transcript	Fold change
Ser-AGA-2-1	1131.96
Ser-AGA-1-1	592.19
Pro-AGG-2-1	520.17
Val-AAC-1-1	224.08
Leu-AAG-1-1	196.65
Cys-GCA-11-1	131.43
Ser-GCT-6-1	96.38
Asn-GTT-2-1	37.26
Lys-TTT-7-1	32.29
Ser-GCT-4-1	24.78
Cys-GCA-6-1	5.30
Thr-TGT-4-1	3.43
Asn-GTT-24-1	3.24

Codon RPM, effect of alignment masking



Codon	: Fold change
Ser-UCU	: 60.37
Pro-CCU	: 33.45
Val-GUU	: 4.23
Ala-GCG	: 1.70