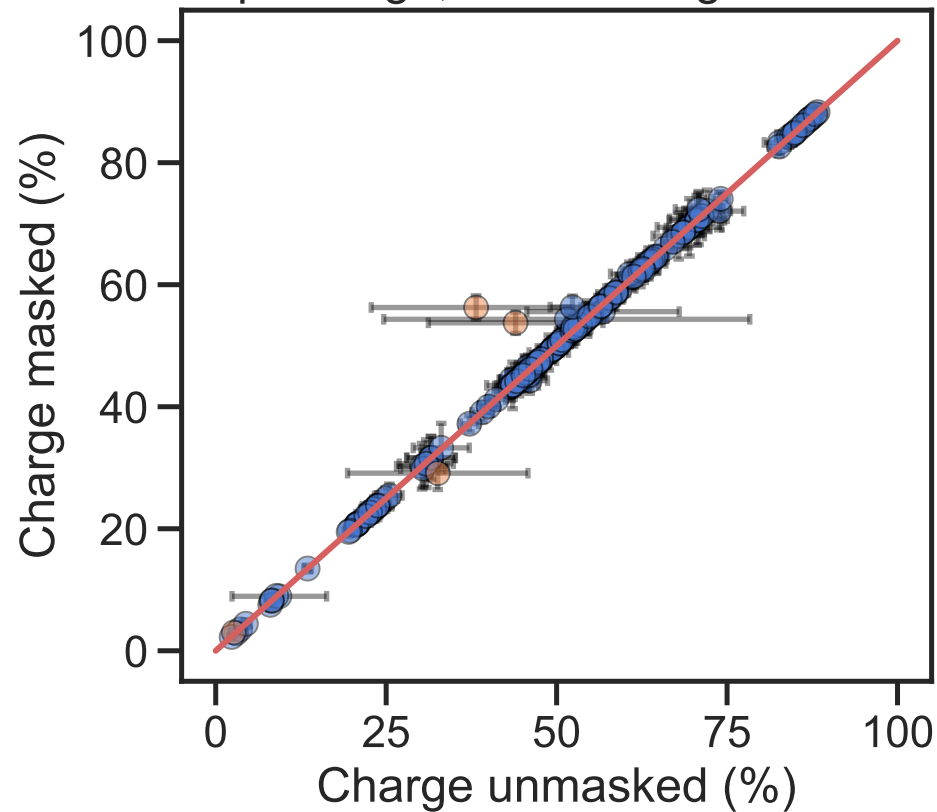
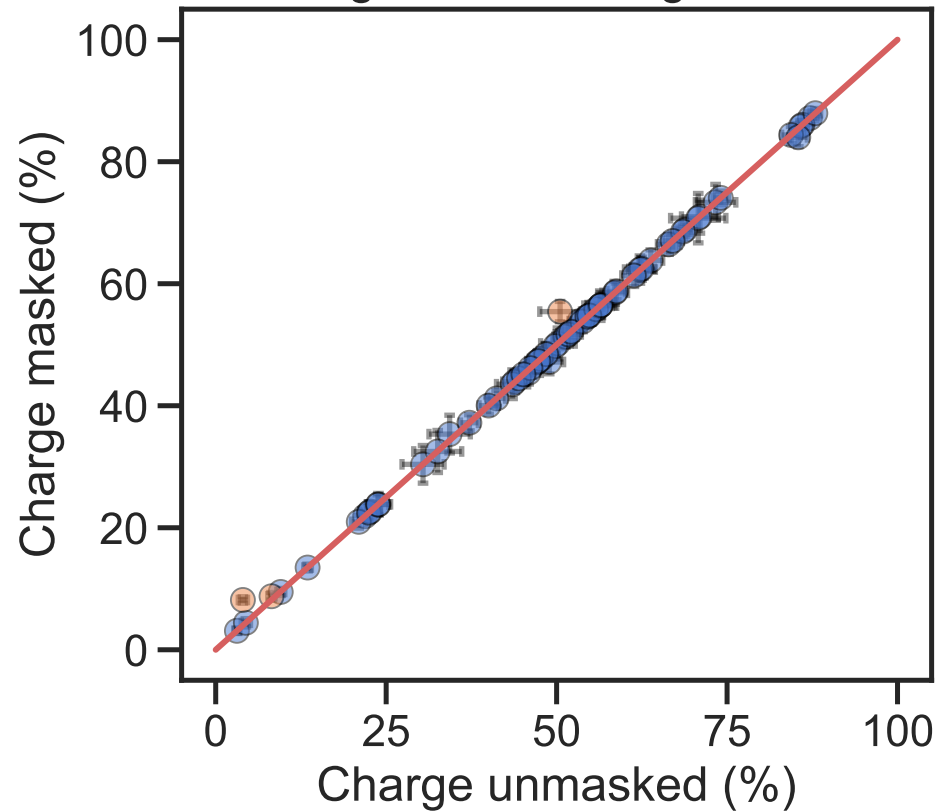


Transcript charge, effect of alignment masking



Transcript	:	Fold change
Tyr-GTA-8-1	:	1.47
Ser-AGA-2-1	:	1.22
Asn-GTT-3-1	:	1.13
Cys-GCA-11-1	:	0.89

Codon charge, effect of alignment masking



Codon	: Fold change
Pro-CCU	: 2.05
Ser-UCU	: 1.10
Pro-CCG	: 1.07