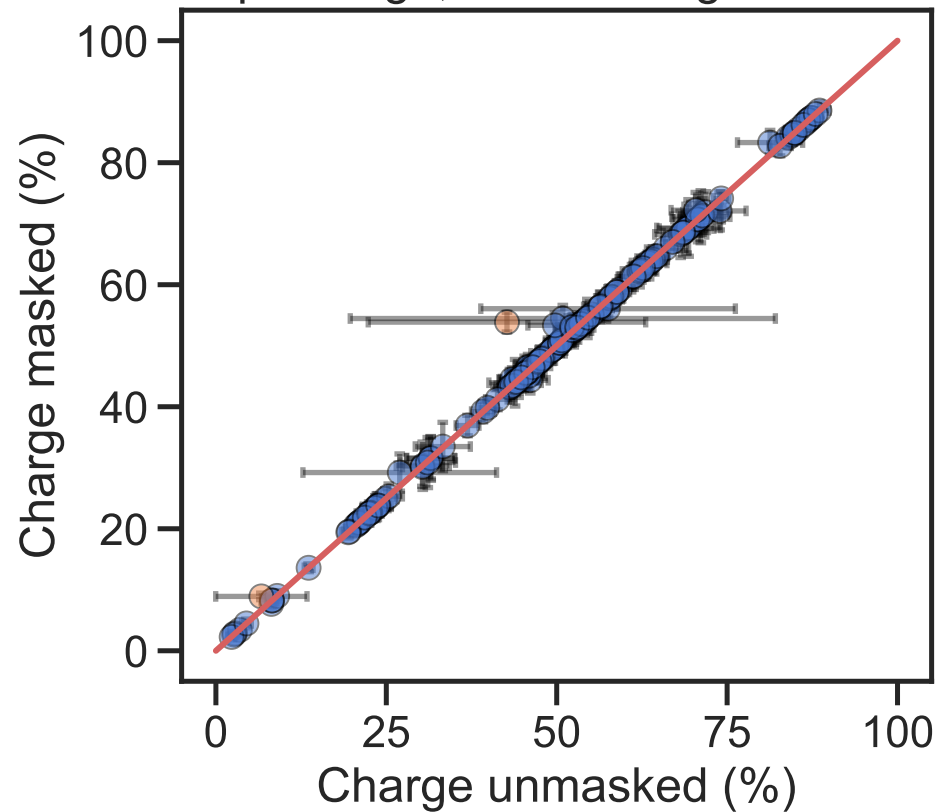
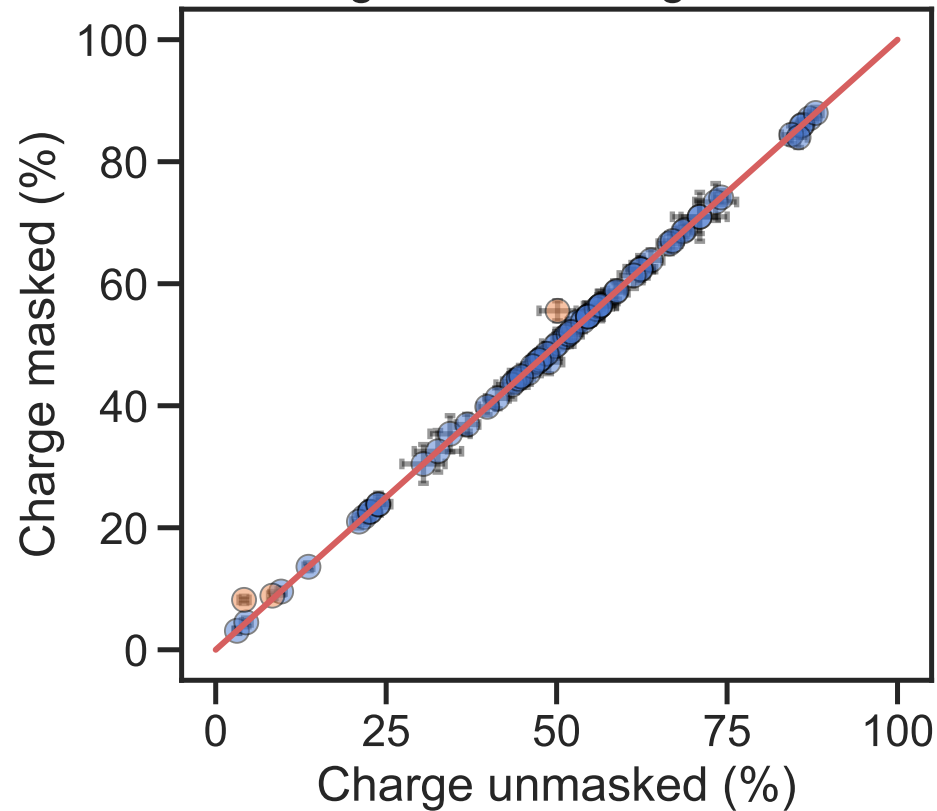


Transcript charge, effect of alignment masking



Transcript	:	Fold change
Pro-AGG-2-1	:	1.34
Ser-AGA-2-1	:	1.26

Codon charge, effect of alignment masking



Codon	: Fold change
Pro-CCU	: 1.97
Ser-UCU	: 1.11
Pro-CCG	: 1.07