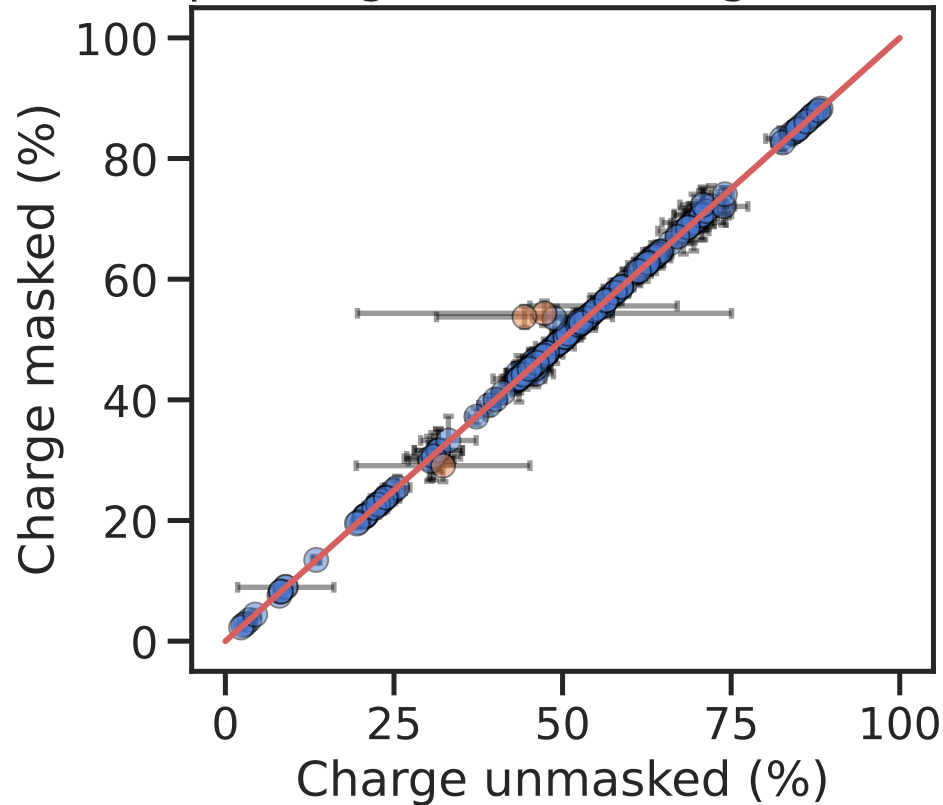
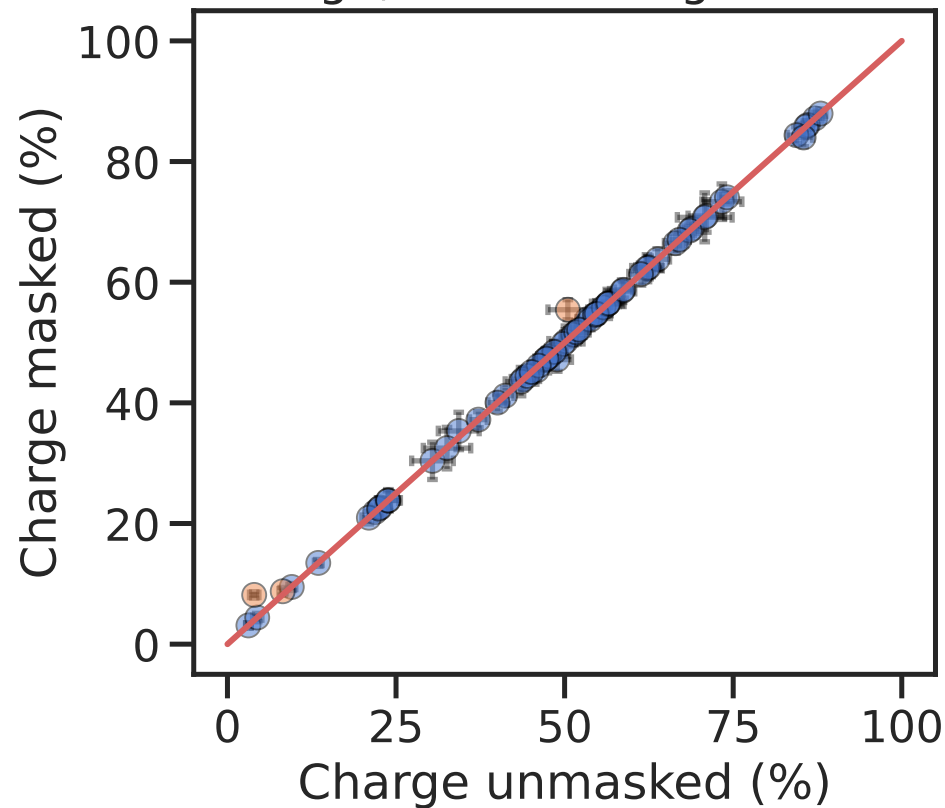


Transcript charge, effect of alignment masking



| | | |
|--------------|---|-------------|
| Transcript | : | Fold change |
| Ser-AGA-2-1 | : | 1.21 |
| Ser-AGA-1-1 | : | 1.15 |
| Cys-GCA-11-1 | : | 0.90 |

Codon charge, effect of alignment masking



| | | |
|---------|---|-------------|
| Codon | : | Fold change |
| Pro-CCU | : | 2.06 |
| Ser-UCU | : | 1.10 |
| Pro-CCG | : | 1.07 |