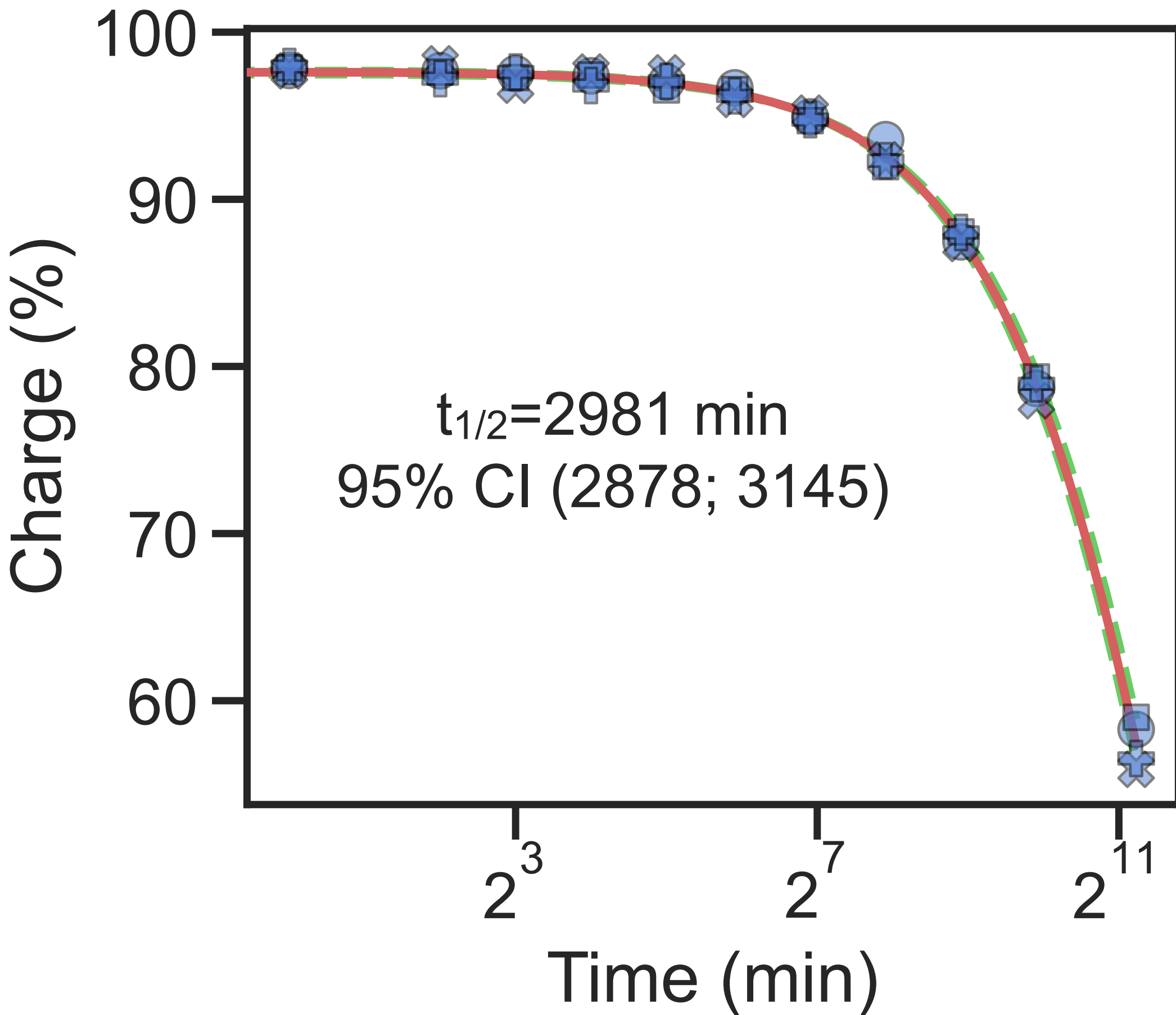


Best fitting aminoacylation decay
mito-Ile-CAT



Worst fitting aminoacylation decay
Gln-TTG-1-1

