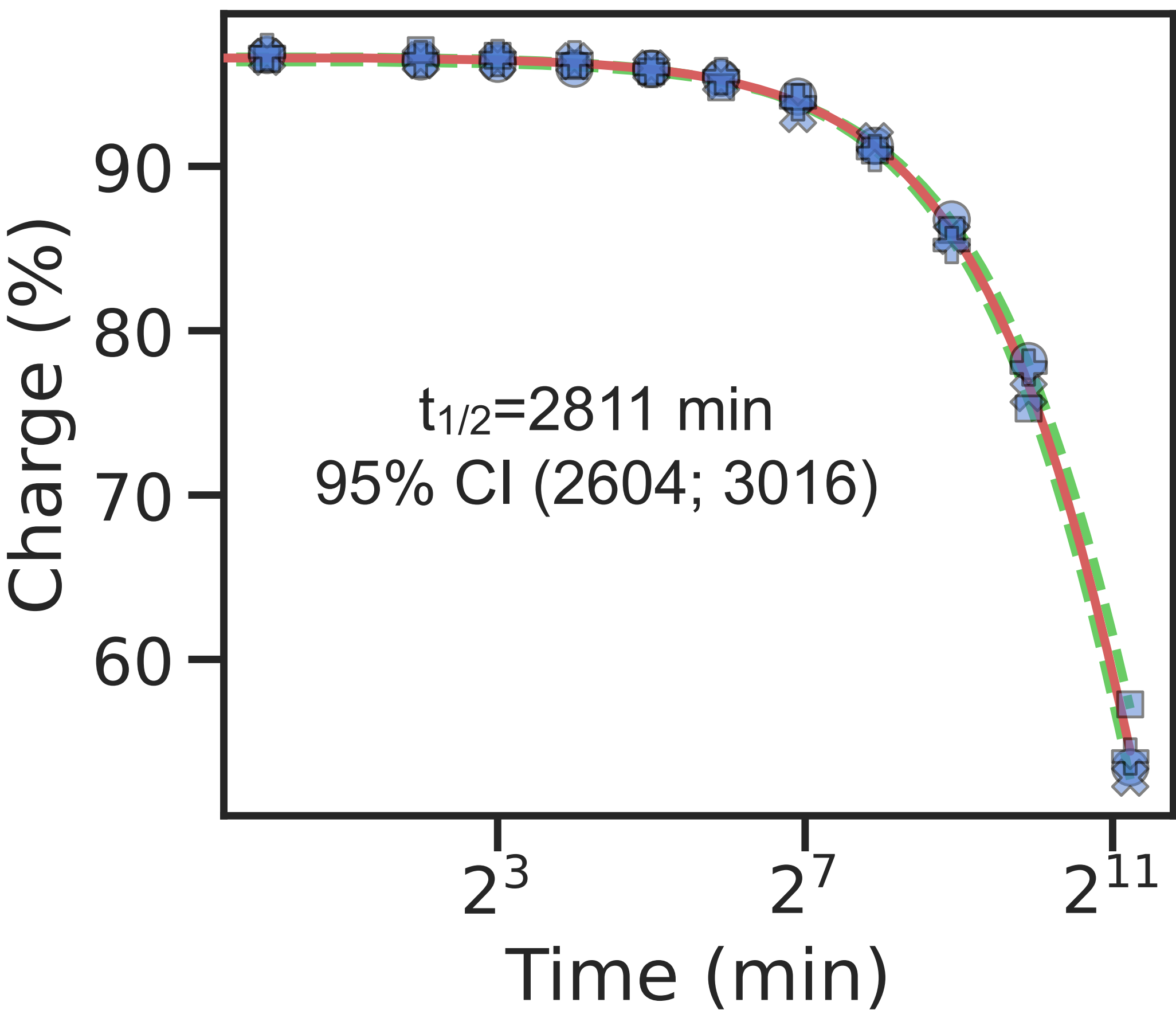


Best fitting aminoacylation decay  
Ile-AAT-5-4



Worst fitting aminoacylation decay  
Gln-TTG-2-1

