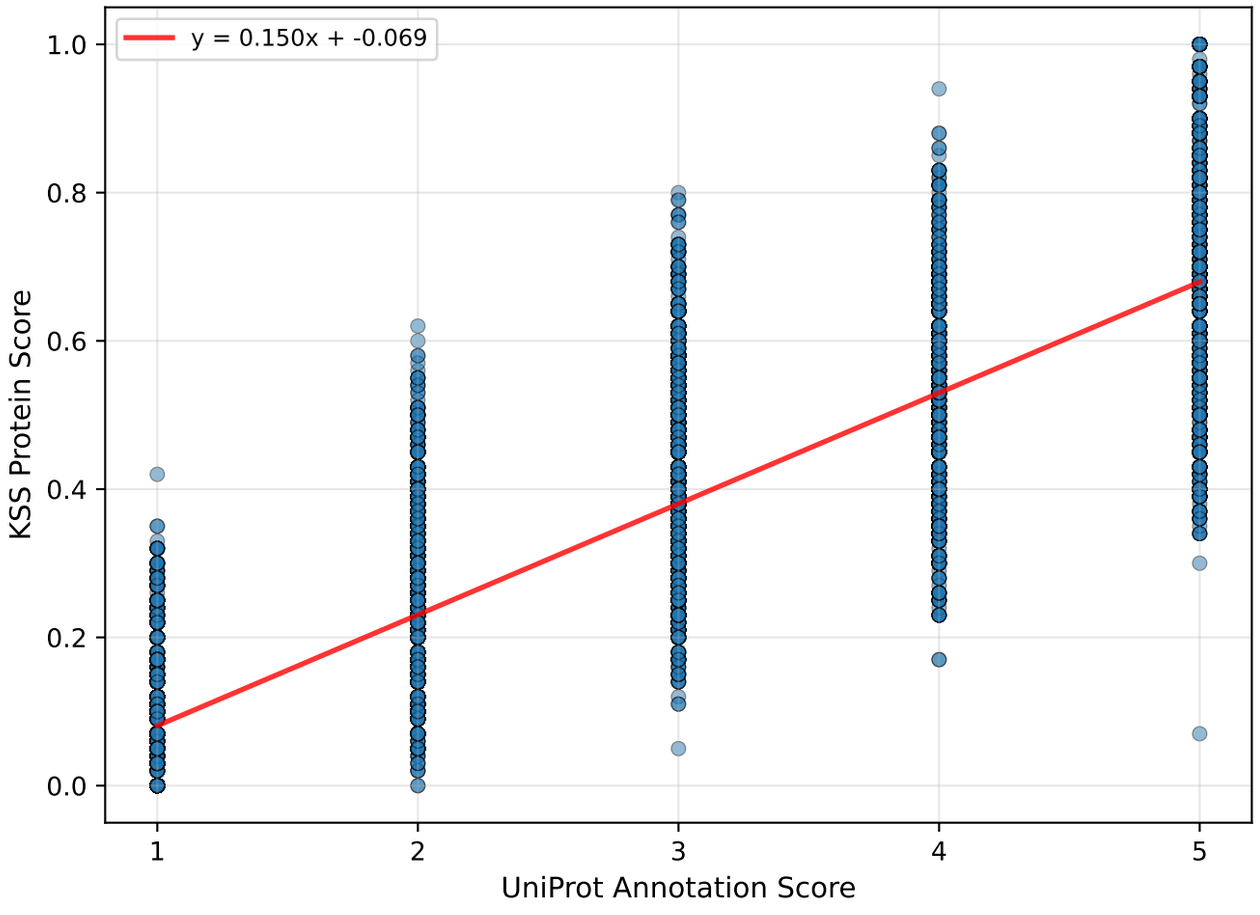
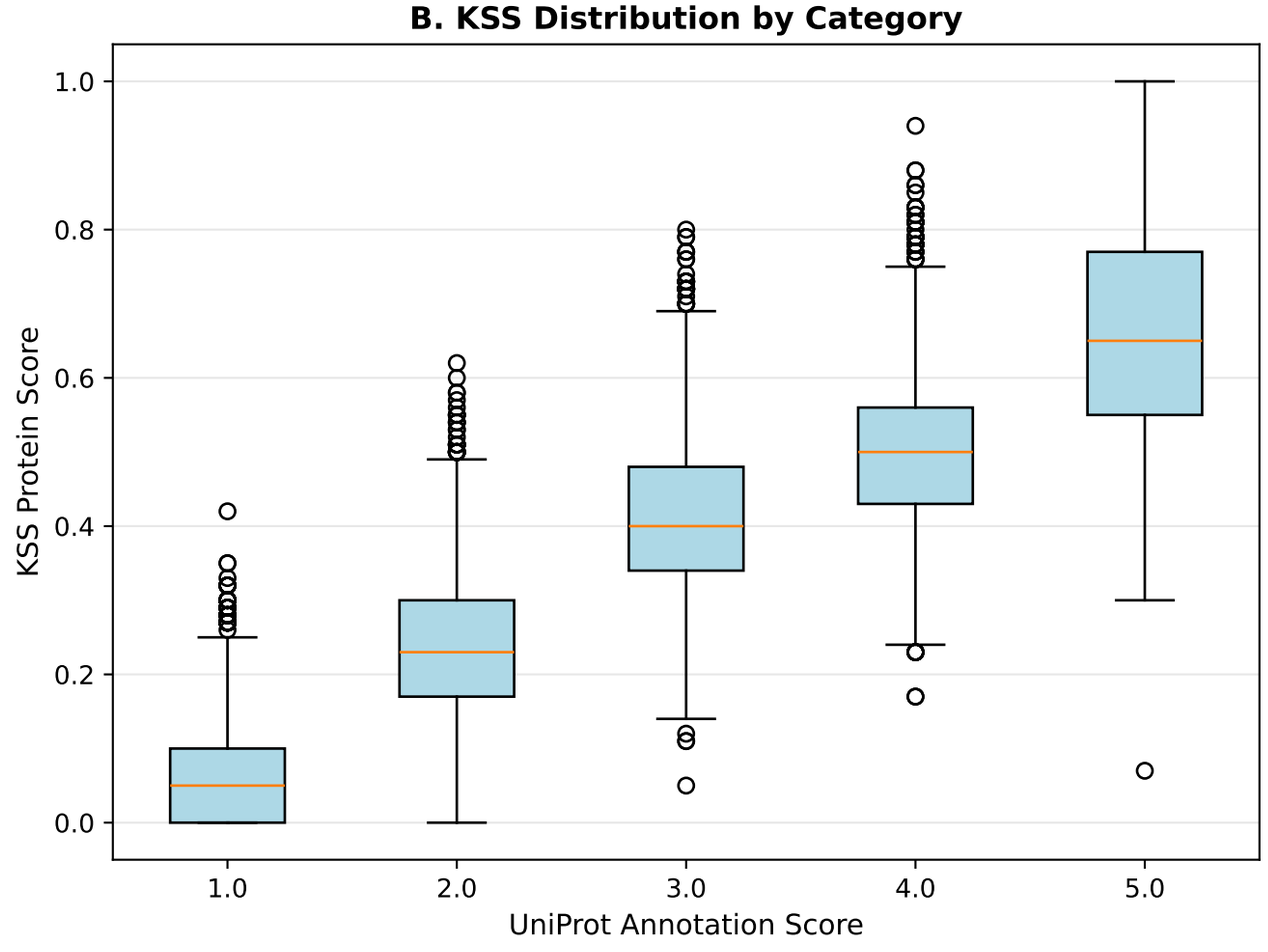
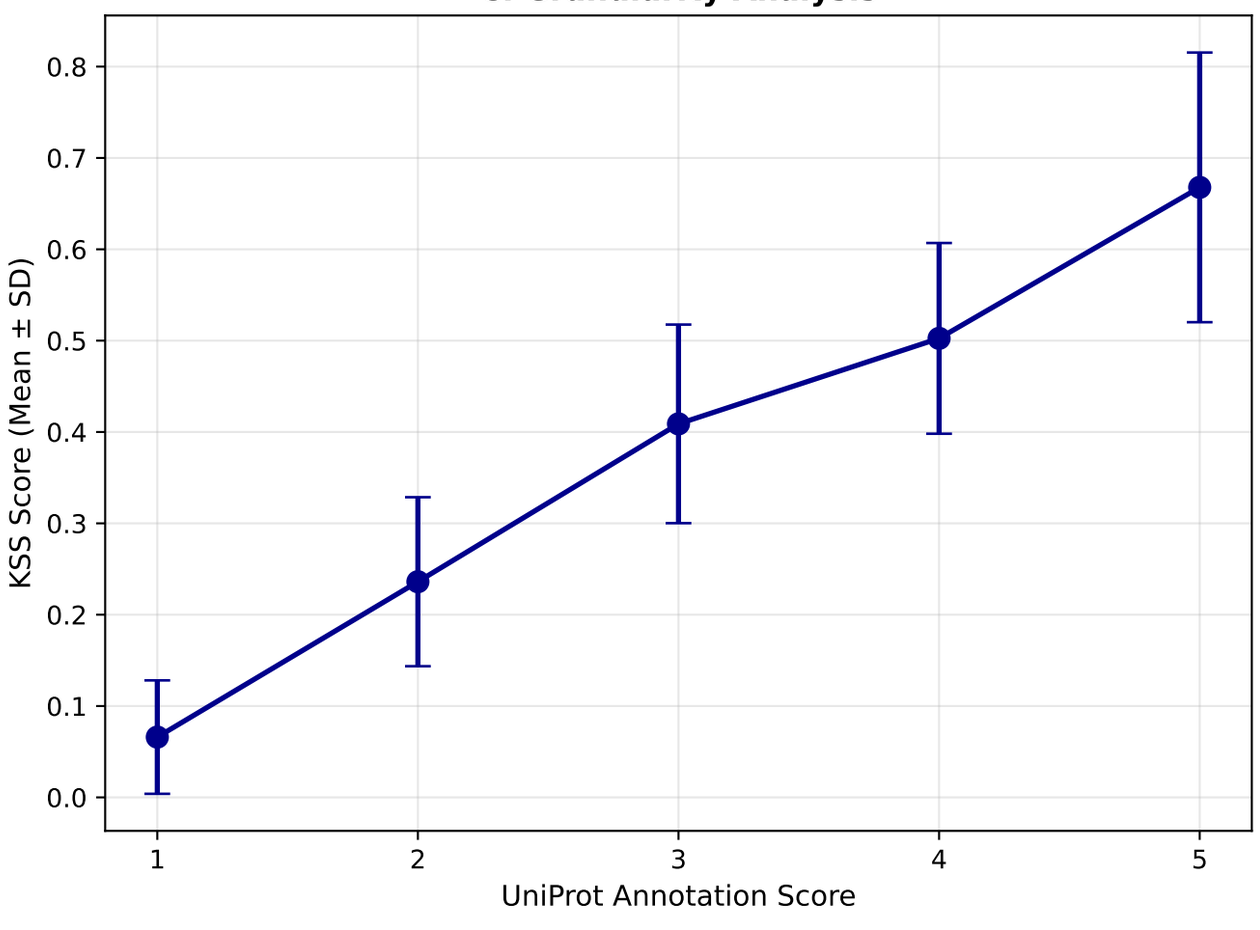
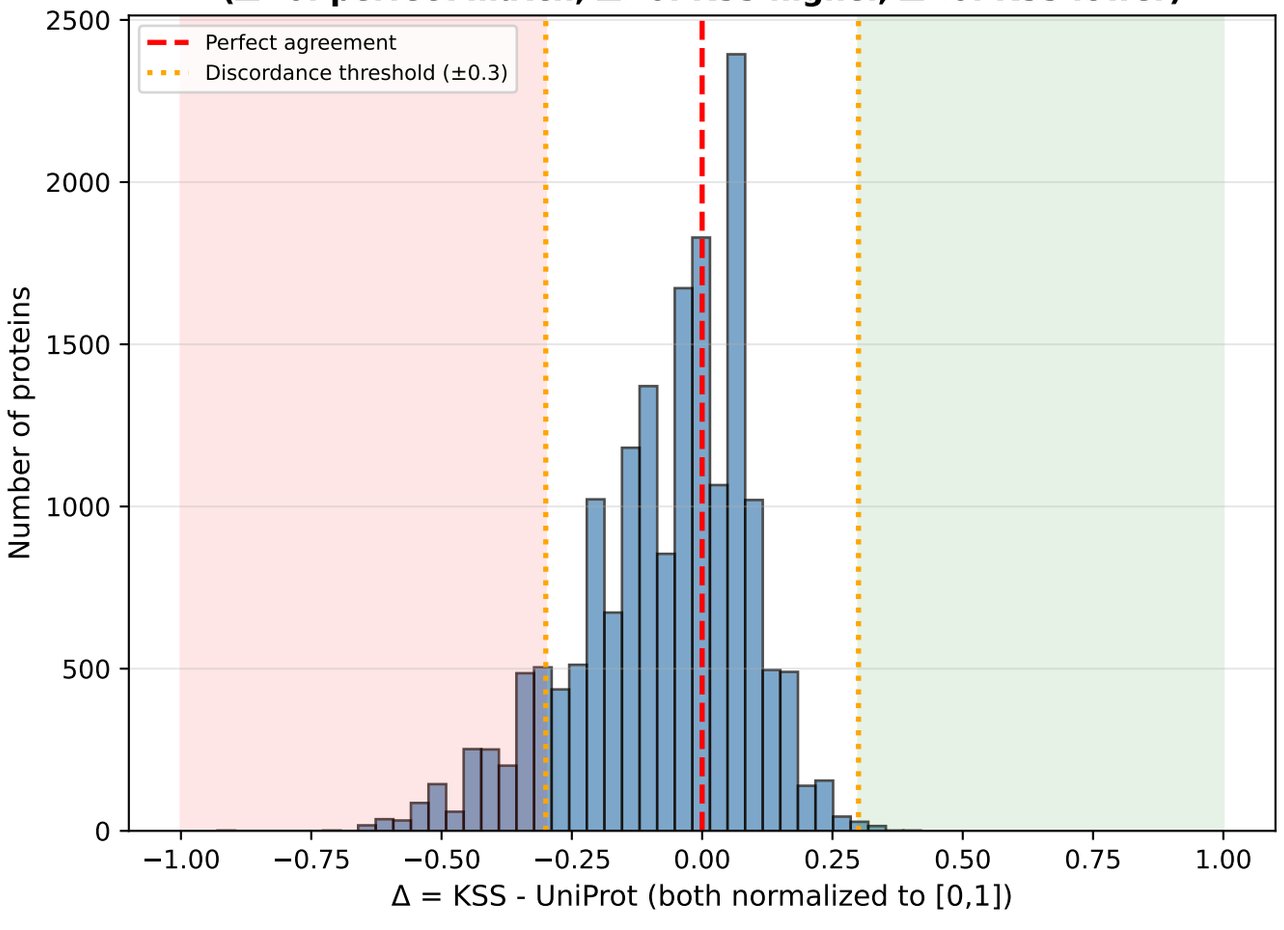

A. Correlation Analysis
ρ = 0.900, τ = 0.776, R² = 0.783
y = 0.150x + -0.069
KSS Protein Score
UniProt Annotation Score


B. KSS Distribution by Category
KSS Protein Score
UniProt Annotation Score


C. Granularity Analysis
KSS Score (Mean ± SD)
UniProt Annotation Score


D. Score Agreement Distribution
(Δ=0: perfect match, Δ>0: KSS higher, Δ<0: KSS lower)
Perfect agreement
Discordance threshold (±0.3)
Number of proteins
Δ = KSS - UniProt (both normalized to [0,1])