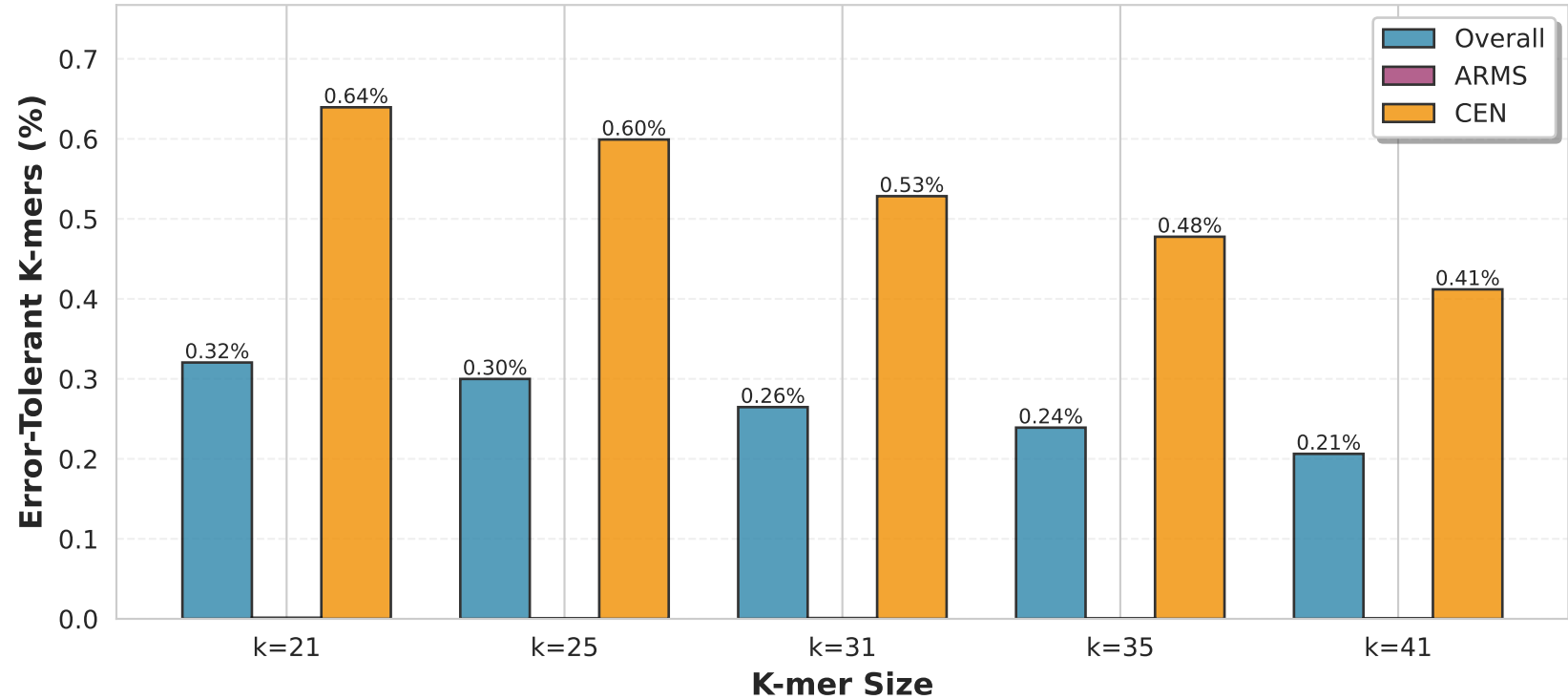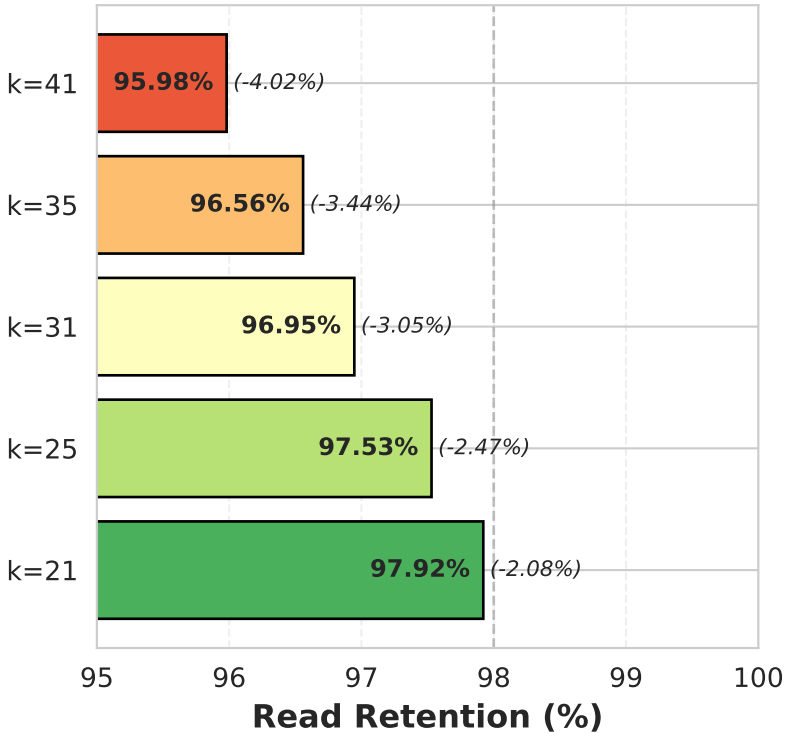Sequencing Error Resilience Analysis Across K-mer Sizes
Impact of Single-Base Sequencing Errors on Marker Specificity

A. Error Tolerance Comparison Across K-mer Sizes

B. Expected Read Retention
(0.1% sequencing error)

C. Error Tolerance Distribution by Region