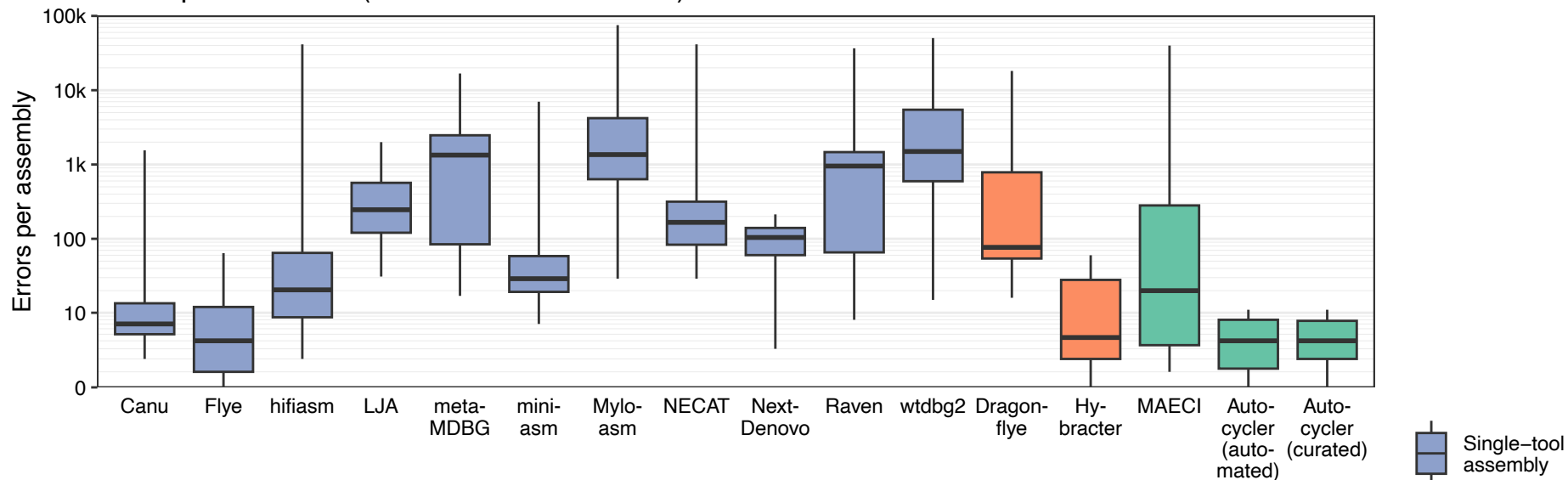


A. Sequence errors (substitutions and indels)



B. Total errors (sequence errors + structural errors)

