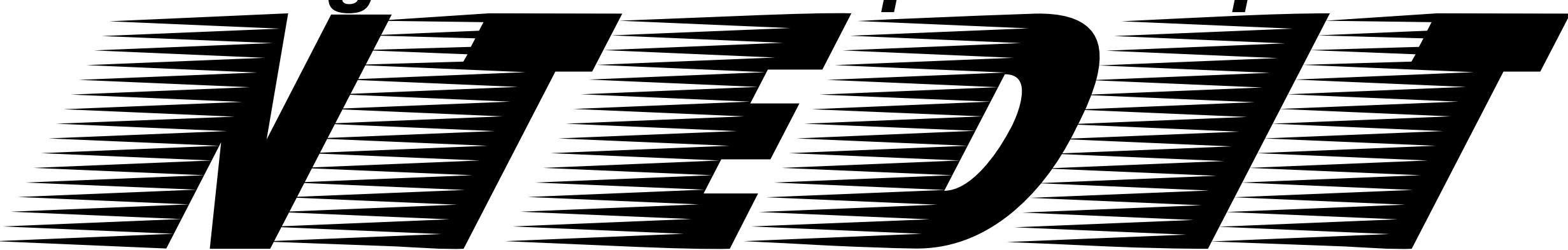
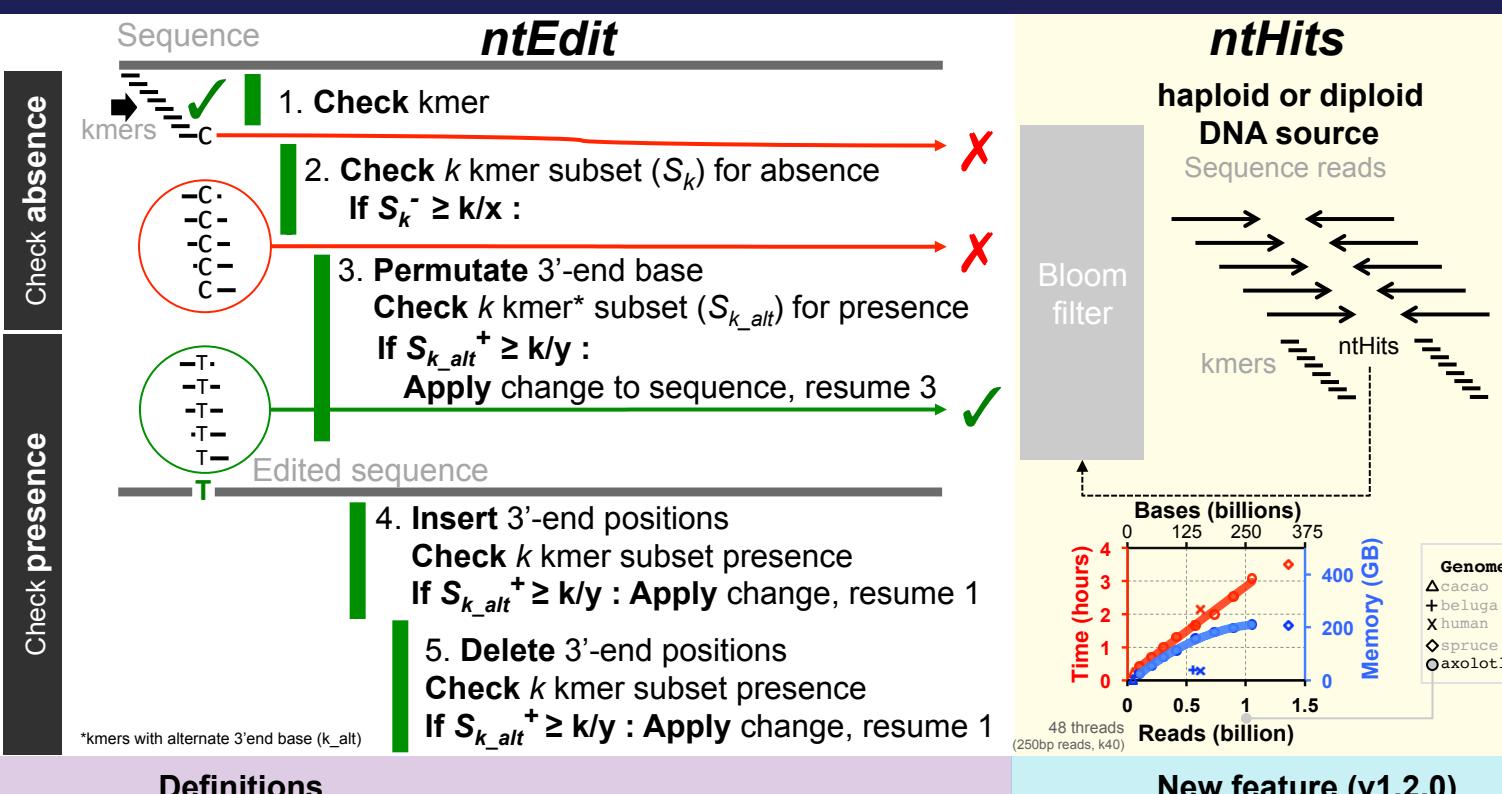


# scalable genome sequence polishing

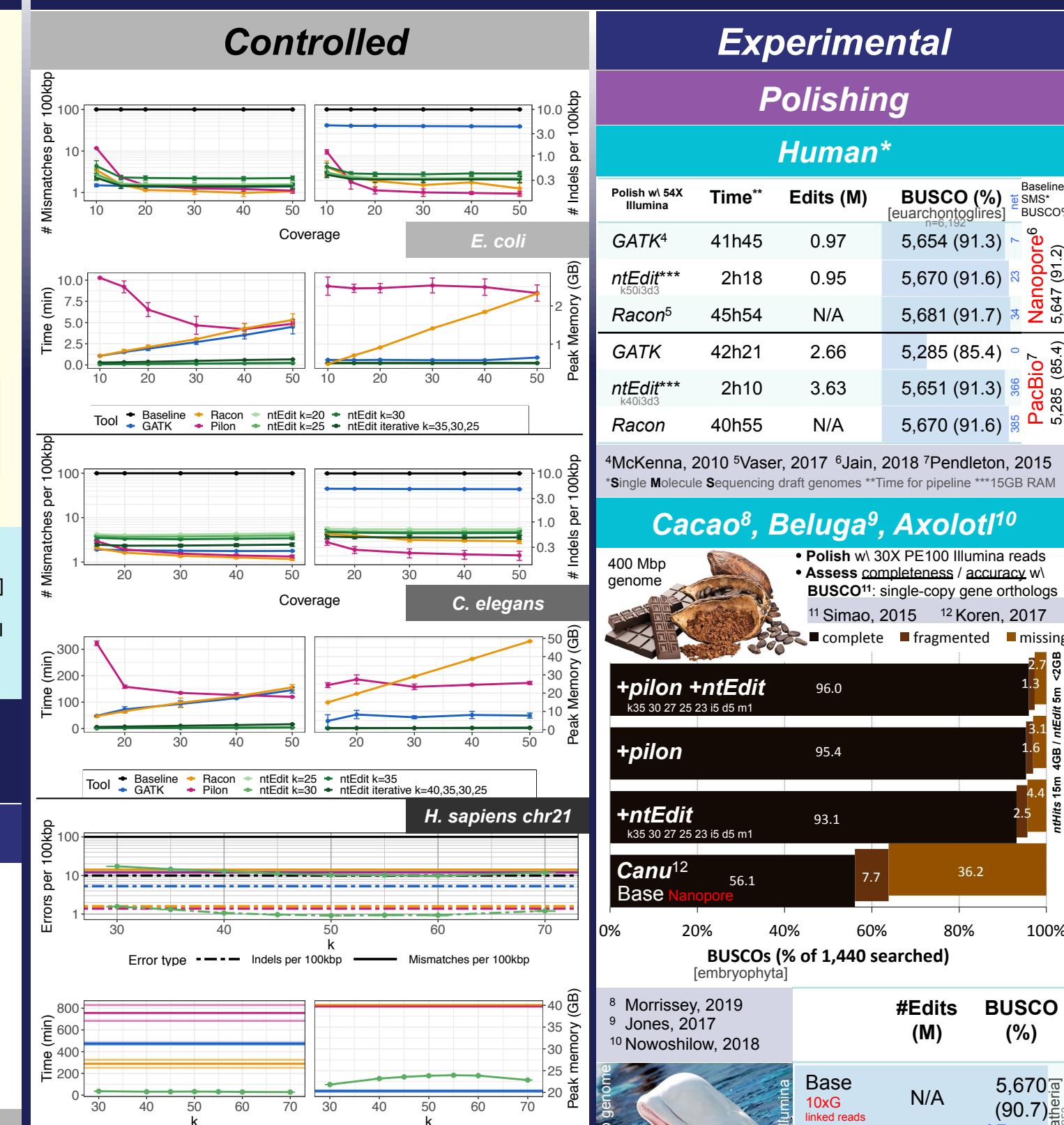


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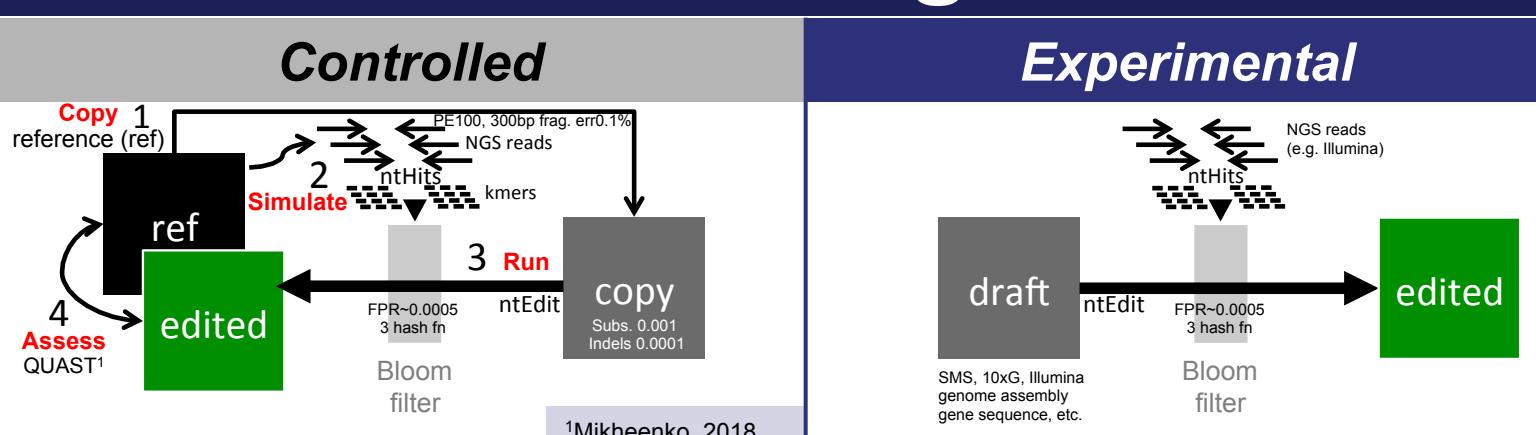
## Method



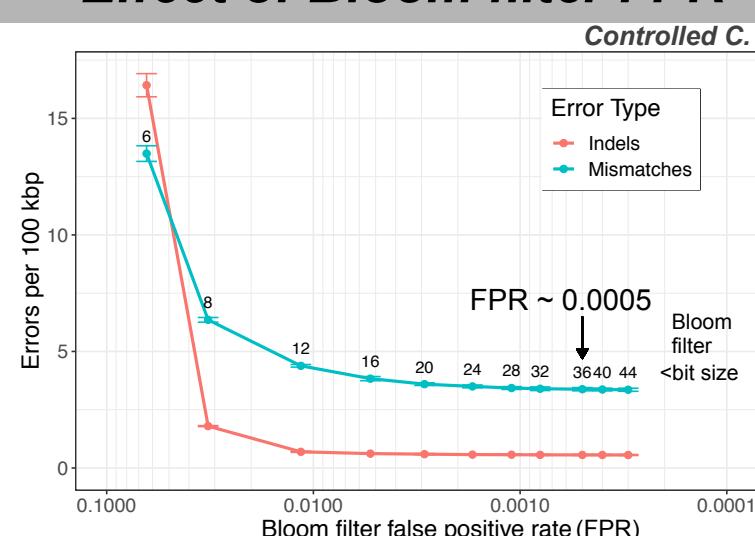
## Results



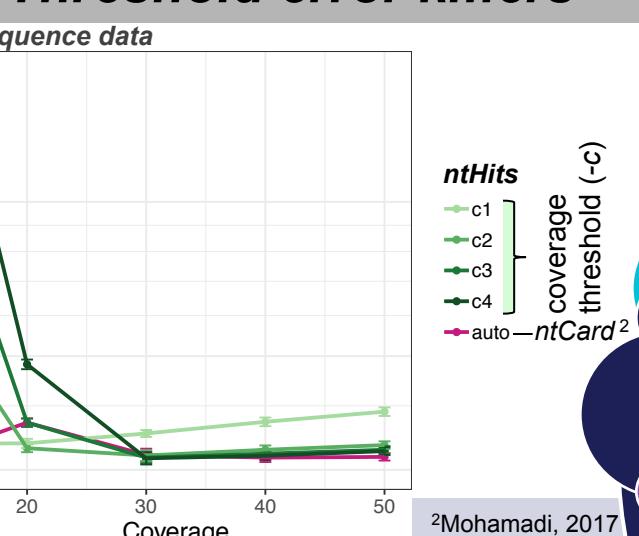
## Testing



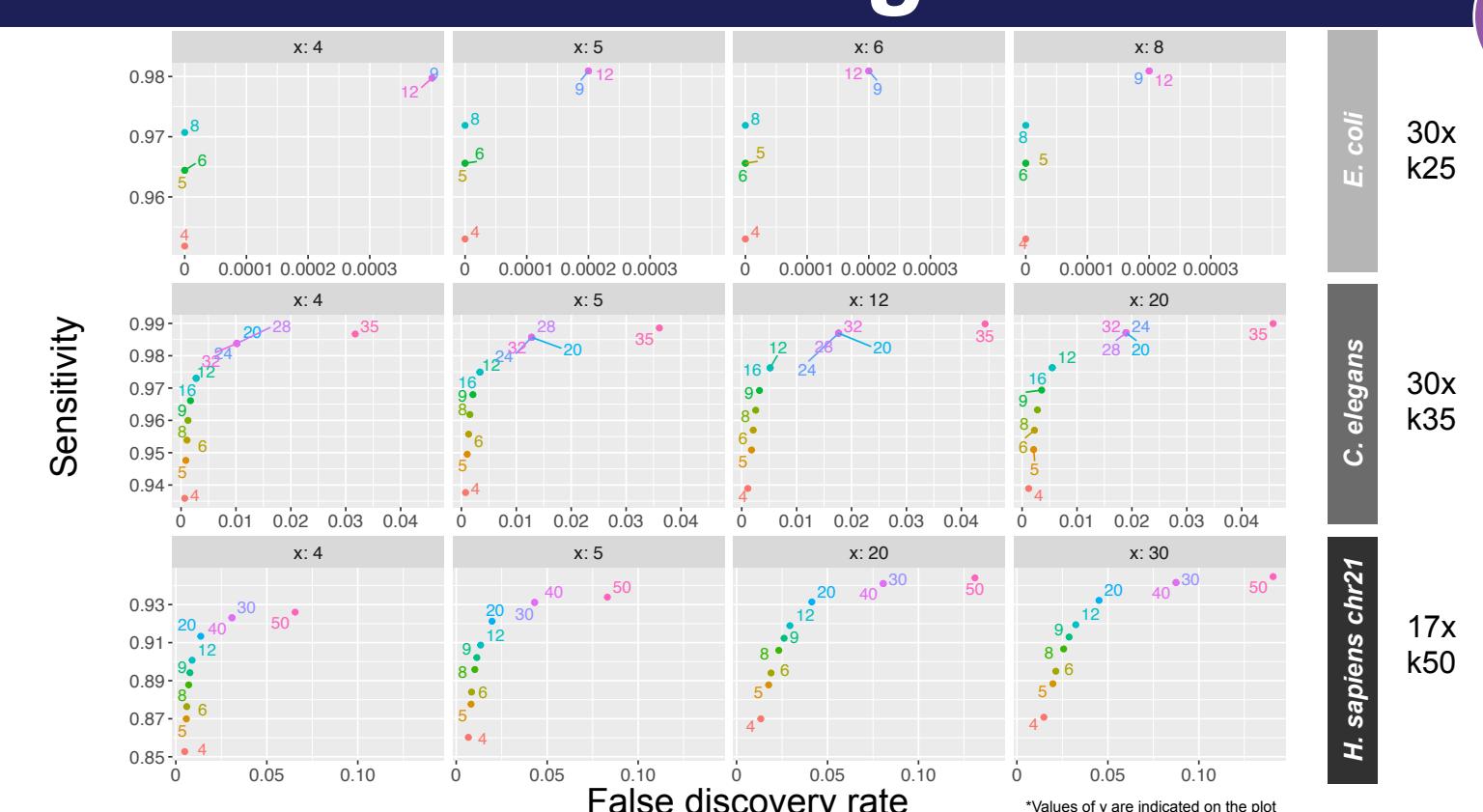
## Effect of Bloom filter FPR



## Threshold error kmers



## Tuning



## Summary

Scalable solutions for genome assembly

Assembly Correction, Scaffolding, Gap-filling, Polishing

Tintinni, ABSS, OROS, RAILS, Sealer, MEDIT

Read Technology: Short, Linked, Long

Illumina, SMS drafts (Nanopore/PacBio)

<https://github.com/bcgsc/bioinformatics-lab>

<https://github.com/bcgsc/ntedit>

<https://github.com/bcgsc/nthits>

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