A species-specific Nanopore basecalling model improves sequencing accuracy for *Mycobacterium tuberculosis*

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Authors

- **Michael B. Hall**
  0000-0003-3683-6208 · mbhall88 · mbhall88
  European Bioinformatics Institute, Hinxton, Cambridge, CB10 1SD, UK

- **Zamin Iqbal**
  0000-0001-8466-7547 · iqbal-lab · ZaminIqbal
  European Bioinformatics Institute, Hinxton, Cambridge, CB10 1SD, UK · Funded by Grant XXXXXXXX

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