

# Error Analysis of PSMC'

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## 1 Miscellaneous

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GitHub:

## 2 Objective

Determine if there's a relationship between the lengths of contigs being analysed and the error given by PSMC' (MSMC [1] with two haplotypes).

## References

- [1] Stephan Schiffels and Richard Durbin. Inferring human population size and separation history from multiple genome sequences. *Nature genetics*, 46(8):919–925, 2014.