

1. Verbiest M, Maksimov M, Jin Y, Anisimova M, Gymrek M, Bilgin Sonay T. Mutation and selection processes regulating short tandem repeats give rise to genetic and phenotypic diversity across species. *J of Evolutionary Biology*. 2023 Feb;36(2):321–36.
2. Bilgin Sonay T, Carvalho T, Robinson MD, Greminger MP, Krützen M, Comas D, et al. Tandem repeat variation in human and great ape populations and its impact on gene expression divergence. *Genome Res*. 2015 Nov;25(11):1591–9.
3. Sulovari A, Li R, Audano PA, Porubsky D, Vollger MR, Logsdon GA, et al. Human-specific tandem repeat expansion and differential gene expression during primate evolution. *Proc Natl Acad Sci USA*. 2019 Nov 12;116(46):23243–53.
4. Haller BC, Messer PW. SLiM 2: Flexible, Interactive Forward Genetic Simulations. *Mol Biol Evol*. 2017 Jan;34(1):230–40.
5. Rodrigues MF, Kern AD, Ralph PL. Shared evolutionary processes shape landscapes of genomic variation in the great apes [Internet]. *Evolutionary Biology*; 2023 Feb [cited 2024 Jan 18]. Available from: <http://biorxiv.org/lookup/doi/10.1101/2023.02.07.527547>
6. tskit [Internet]. Available from: https://tskit.dev/pyslim/docs/latest/vignette_parallel_phylo.html
7. Huang B, Durvasula A, Mousavi N, Ziaei-Jam H, Maksimov M, Lohmueller KE, et al. Genome-wide selection inference at short tandem repeats [Internet]. 2022 [cited 2024 May 21]. Available from: <http://biorxiv.org/lookup/doi/10.1101/2022.05.12.491726>