

Stanford

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Ph.D. Student in Developmental Biology, admitted Autumn 2021

Publications

PUBLICATIONS

- **Champagne: Automated whole-genome phylogenomic character matrix method using large genomic indels for homoplasy-free inference.** *Genome biology and evolution*
Schull, J. K., Turakhia, Y., Hemker, J. A., Dally, W. J., Bejerano, G.
2022