

Combi Seminar

Wednesday, 1.13.21 | 1:30 | held remotely

<https://depts.washington.edu/gstrestrc/remote.htm>



Dr. Kelley Harris

University of Washington

“Whole genome sequencing of the BXD recombinant inbred mouse lines pinpoints a locus explaining a significant component of murine mutation spectrum differentiation”

My computational biology research group uses large datasets of genetic variation to study the evolutionary history of humans and other species, with particular interests in coalescent theory and the evolution of the mutation rate and spectrum.

<https://sites.google.com/site/harriskelley/home>

Questions? Contact Brian Giebel at bgiebel@uw.edu or visit the Combi website at <http://www.gs.washington.edu/news/combi.htm>

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