

BGS under non-equilibrium demography (paper 1)

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Creating folders and files

This builds a look-up table on a dense parameter grid that will allow for fast assembly of whole chromosomes
Creates files that will be used as input for moments++

Running moments++

Computes summary statistics for each parameter combination in lookup_tbl
Collects lookup table with Hr values on the parameter grid

TODO

Add Python chunk with fwdpy11 simulations