Manual for MultiWaver 2.1

MultiWaver 2.1 is a modified version of MultiWaver 2.0 and it is specially used to estimate the introgression wave utilizing the information output by ArchaicSeeker 2.0.

Compile

\$ cd getAS2Seg

\$ g++ getAS2Seg.cpp -o getAS2Seg

\$ cd ../MultiWaver2.1

\$ make

Step1: To estimate the introgression waves, we need to first use the executable file 'getAS2Seg' to convert the output of *ArchaicSeeker 2.0* into the format of *MultiWaver 2.1*. getAS2Seg filepath_to_genetic_map seg_output_by_ArchaicSeeker2.0 output_file Archaic Here, the 'filepath_to_genetic_map' need to have genetic map of chromosomes 1-22, and named as genetic map chr* combined b37.txt. Archaic can be Altai or Denisova.

Step2: Use the executable file 'MultiWaver2.1' to estimate the introgression waves.

MultiWaver2.1 -i output_by_getAS2Seg -l threshold -o output_filename

The detailed arguments of *MultiWaver* 2.1 are list as follows:

Arguments	Types	Descriptions
-i/input	<string></string>	Input of the ancestral tracks [required]
-l/lower	[double]	Lower bound to discard short tracks [optional, default 0]
-b/bootstrap	[integer]	Number of bootstrapping [optional, default 100]
-a/alpha	[double]	Significance level to reject null hypothesis in LRT [optional, default 0.001]

-e/epsilon	[double]	Epsilon to check whether a parameter converge or not [optional, default 1.0e-6]
-p/minProp	[double]	Minimum survival proportion for a wave at the final generation [optional, default 0.05]
-m/maxIter	[integer]	Maximum number of iterations to scan for waves of admixture events [optional, default 10000]
-t/thread	[integer]	Number of threads [optional, default 1]
-o/output	<string></string>	Prefix of output [required]

Options	Descriptions
-h/help	Print help message, default is OFF

Examples

Step1:

getAS2Seg/getAS2Seg MWexamples/genetic_map_b37/ ../examples/Han.seg MWexamples/Han
Denisova

Step2:

MultiWaver2.1/MultiWaver2.1 -i MWexamples/Han.seg -l 0.00015 -o MWexamples/Han. The introgression model can be found in MWexamples/Han.sum.

Questions

Questions and suggestions are welcome, feel free to contact Kai Yuan (yorkklause@gmail.com) or Rui Zhang (zhangrui2018@picb.ac.cn).