

abi2link v1.0.0

2003/11/12

User documentation

Abi2link is designed to create linkage files out of ABI genotype and phenotype files. Please see example directory for detailed file description.

Usage: ./abi2link ARGUMENTS

Currently known arguments:

- map <haldane|kosambi>
 locus mapping function
- ped <file> pedigree file
- chr <file> chromosome description file
- trait <file> trait file (optional)

- estimate <all|founder>
 whether to estimate allele frequencies from all
 individuals or founders only (optional, default: all)
- prefix <name> output file prefix (optional, default: abi2link)

- v, --version print version information and exit
- h, --help print this text and exit

Copyright: Prof. Dr. Andreas Ziegler
For questions or comments please contact:
Andre Kleensang
Institute of Medical Biometry and Statistics
University at Lübeck, Germany
email: kleensang@imbs.uni-luebeck.de

Report errors to <daniel.franke@imbs.uni-luebeck.de>