

NCSU Summer Institute 2004

QTL II

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- Model: selection for multiple QTL
- Pheno: extensions beyond normal data
- Bayes: interval mapping with prior info
- Traits: multiple phenotypes & microarrays

contact information & resources

- email: byandell@wisc.edu
- web: www.stat.wisc.edu/~yandell/statgen
 - QTL & microarray resources
 - references, software, people
- thanks:
 - students: Chunfang “Amy” Jin, Fei Zou, Pat Gaffney, Jaya Satagopan
 - faculty/staff: Alan Attie, Hong Lan, Michael Newton, Christina Kendziorski, Tom Osborn, Jason Fine