

Papers for STAT 550 (DL) class, 2010

Papers available

This list is neither necessary nor sufficient.

That is, it is a somewhat random collection compiled over the years of teaching this class. It contains some papers that were a reasonable choice for a particular student, given their background or research direction, but bear little relationship to the details of the 550 course. Not all choices on the list are reasonable for all students.

Also, additions are welcome – if you have a particular area/paper you would like to choose, feel free to suggest.

The papers are listed first, somewhat sorted by category. The references follow on the last four pages (approx.)

Please note: in a given course offering, no two students may choose the same paper. To be fair, I operate on a first-come first-served basis, although I have occasionally negotiated switches.

So if you have a particular choice you want, let me know right away.

I Population-related issues/studies, including LD and LD mapping

Abecasis et al. (2005) – Using LD in association mapping.

Carlson et al. (2004) – selecting SNPs for Association Analysis.

Ceppellini et al. (1955) : the original “Gene counting” paper

Clayton (2000) – a review (as of that date) or fine-scale disequilibrium mapping

Graham and Thompson (1998) – disequilibrium mapping vi likelihoods

Falush et al. (2003) – another Stephens/Pritchard popn structure paper

Kaplan et al. (1995) – the first likelihood approach to disequilibrium mapping.

McPeck and Strahs (1999) – another disequilibrium mapping approach

Marth et al. (2004) or Pluzhnikov et al. (2002): Inference of human history and demography from genetic data.

Pritchard et al. (2000) – the first Stephens/Pritchard popn structure paper

Storey and Tibshirani (2003) – significance for genome-wide studies.

Thompson et al. (1988) – an interesting story about LD at three tightly linked loci

Thompson and Neel (1997) – coalescent ancestry of a mutant allele and implications for disequilibrium

Voight (2006) or Li and Stephens (2003) –selection and/or recombination in the human genome.

II Gene ibd and use in sib pair analyses etc.

Anderson and Weir (2007) – estimating relationships in structured populations.

Blackwelder and Elston (1985) – classic paper on sib pair data.

Dudoit and Speed (2000) – a score test for linkage based on ibd in sib pairs

Karigl (1981) recursive probabilities computations for more general IBD patterns

Spielman and Ewens (1996) – the original TDT test for linkage

Spielman and Ewens (1998) – the original extension TDT to S-TDT

Suarez et al. (1978) – formalization of sib pair linkage tests

Thompson (1974) – gene IBD for multiple individuals

III Linkage and Segregation Analysis

Abecasis et al. (2002) – the original Merlin paper.

Boehnke (1994) – paper on limits of resolution of linkage mapping

Elston and Stewart (1971): a fundamental paper in computing likelihoods on pedigrees.

George et al. (2005) – an application of new MCMC methods to estimate lod scores on some extended pedigrees.

Lander and Botstein (1987): the basic “Homozygosity mapping” paper

Lander and Green (1987); The original “Mapmaker” paper, which laid methodological foundation for “Genehunter” programs.

Luo and Lin (2003)– information from using marker data in fitting genetic models.

Morton (1955) – the classic introduction of the base-10 (!!!) lod score

Ploughman and Boehnke (1989) – the basis of SIMLINK

IV PHASE, Haplotyping, etc.

Browning (2006) – a newer and better model for haplotype clustering

Browning and Browning (2007) – use of this model in imputation

Browning (2008) - and in inferences for GWAS

Schouten et al. (2005) – combines haplotyping with ”pedigrees” – first solid study.

Stephens and Donnelly (2003)– a comparison of alternative computational algorithms/models for phasing

Stephens et al. (2001) – Mathew Stephens original PHASE paper

Scheet and Stephens (2006) The FastPHASE paper

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