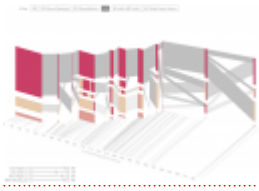
GENETICS, SOFTWARE

3D LD

15.12.2006 1 COMMENT

While waiting for genomewide SNP data to be re-partioned into LD blocks I found this page with some neat progamming tricks. It is part of the dissertation of Ben Fry / MIT about computational information design. Page 74 ff has a history of redesigning the widely used haploview pogram.

The design of these diagrams was first developed manually to work out the details, but in interest of seeing them implemented, it was clear that HaploView needed to be modified directly in order to demonstrate the improvements in practice. Images of the redesigned version are seen on this page and the page following. The redesigned version was eventually used as the base for a subsequence â€˜version 2.0â€™ of the program,
which has since been released to the public and is distributed as one of the analysis tools for the HapMap [www.hapmap.org] project.



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23.04.2008 AT 07:10

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COMMENTS ARE CLOSED.