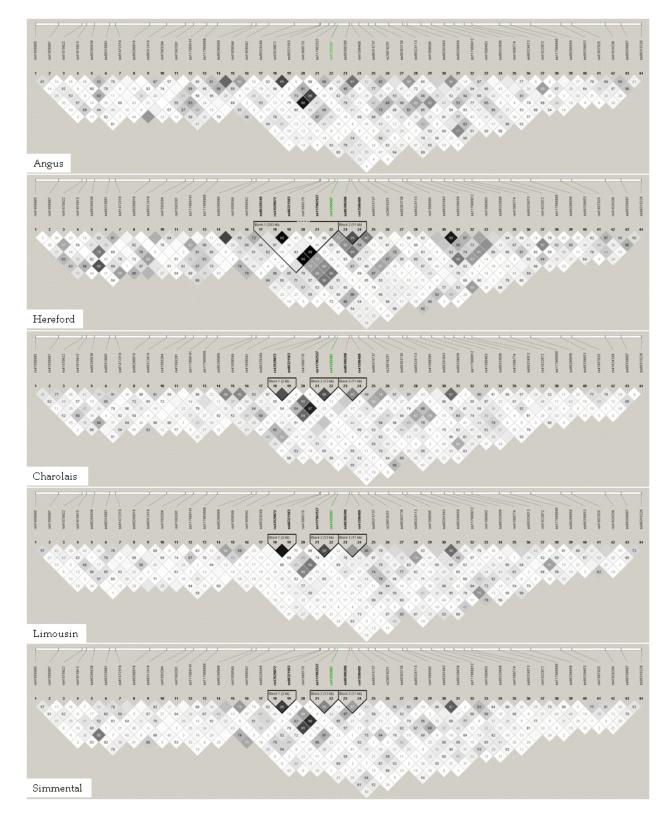
Figure S2 Linkage disequilibrium (LD) plots (r^2) created in Haploview v4.1 (Barrett *et al.* 2005) for 44 SNP spanning 2.86 Mb centered on *CAST* on BTA7. Darker shading indicates regions of higher LD. The commercially tested SNP, *rs41255587*, (#22) is highlighted in green.



References

Barrett J.C., Fry B., Maller J. & Daly M.J. (2005) Haploview: analysis and visualization of LD and haplotype maps. *Bioinformatics* **21**, 263-5.