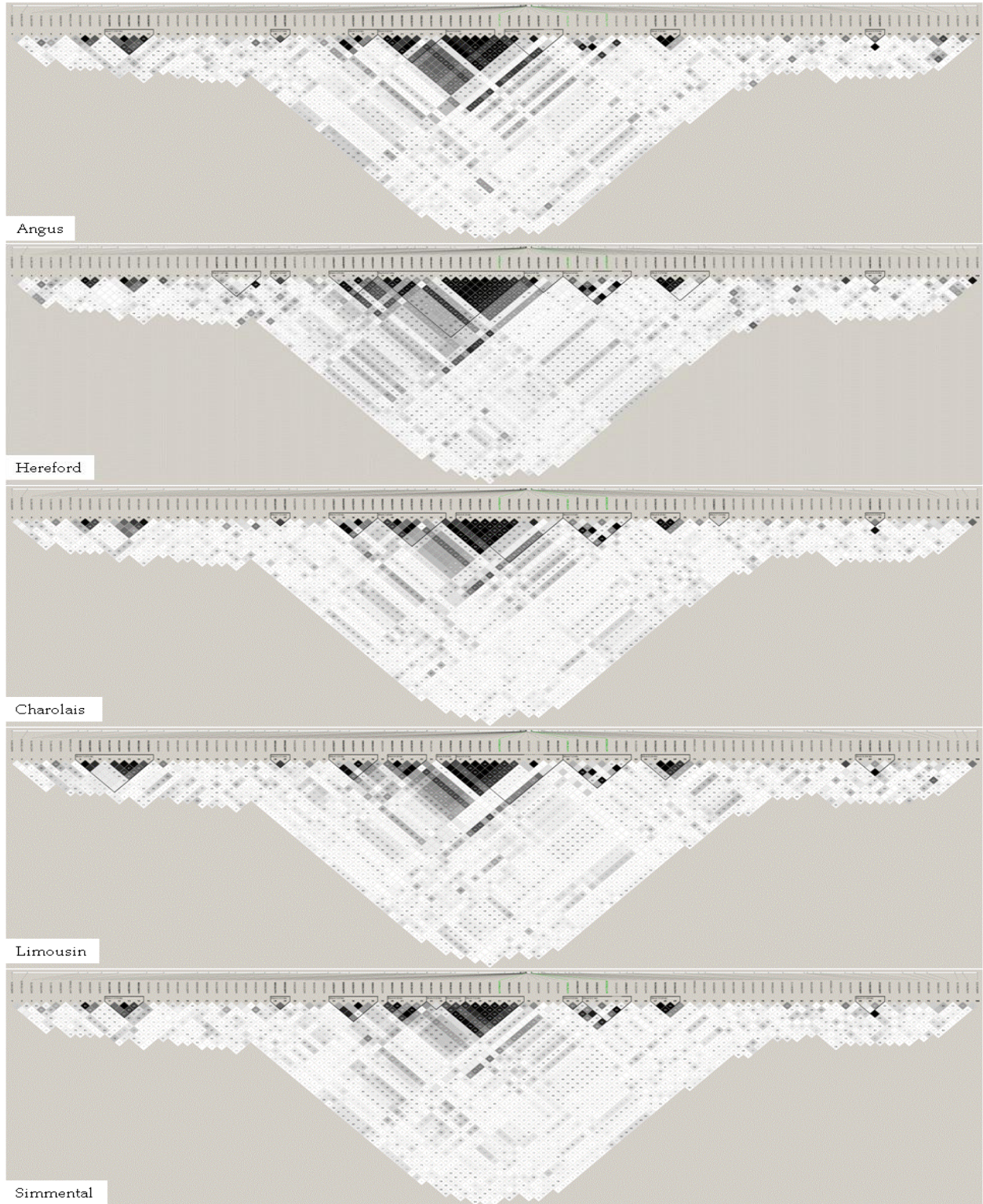**Figure S3** Linkage disequilibrium (LD) plots (r^2) created in Haploview v4.1 (Barrett *et al*. 2005) for the 100 SNP spanning 3.12 Mb centered on *CAPN1* on BTA29. Darker shading indicates regions of higher LD. The commercially tested SNP, *rs17812000* (#51), *rs17871051* (#58), and *rs17872050* (#62) are 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.