

Leicester Cardiovascular BRU Publications, April 2009 - March 2010

1. Codd V, Mangino M, van der Harst P, Braund PS, Kaiser M, Beveridge AJ et al. Common variants near TERC are associated with mean telomere length. *Nat Genet.* 2010;42:197-199
2. Elliott P, Chambers JC, Zhang W, Clarke R, Hopewell JC, Peden JF et al. Genetic Loci associated with C-reactive protein levels and risk of coronary heart disease. *JAMA.* 2009;302:37-48
3. Wellcome Trust Case-Control Consortium. Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. *Nature.* 2010;464:713-720
4. Soranzo N, Spector TD, Mangino M, Kühnel B, Rendon A, Teumer A et al. A genome-wide meta-analysis identifies 22 loci associated with eight hematological parameters in the HaemGen consortium. *Nat Genet.* 2009;41:1182-1190
5. Coronary Artery Disease Consortium, Samani NJ, Deloukas P, Erdmann J, Hengstenberg C, Kuulasmaa K et al. Large scale association analysis of novel genetic loci for coronary artery disease. *Arterioscler Thromb Vasc Biol.* 2009;29:774-780
6. Williams B, Lacy PS; CAFE and the ASCOT (Anglo-Scandinavian Cardiac Outcomes Trial) Investigators. Impact of heart rate on central aortic pressures and hemodynamics: analysis from the CAFE (Conduit Artery Function Evaluation) Study: CAFE-Heart Rate. *J Am Coll Cardiol.* 2009;54:705-713
7. Sleight P, Redon J, Verdecchia P, Mancia G, Gao P, Fagard R et al. ONTARGET investigators. Prognostic value of blood pressure in patients with high vascular risk in the Ongoing Telmisartan Alone and in combination with Ramipril Global Endpoint Trial study. *J Hypertens.* 2009;27:1360-1369.
8. Narayan H, Mohammed N, Quinn PA, Squire IB, Davies JE, Ng LL. Activation of a novel natriuretic endocrine system in humans with heart failure. *Clin Sci.* 2010;118:367-374