Supplementary Figure 1. Meta analysis of the association between the rs11200638 (HTRA1) SNP and risk of AMD across different studies. The boxes indicate the estimated ORs for the homozygous (left panel) and heterozygous (right panel) risk alleles. Each box plot represents an individual study and the size of the box is proportional to the respective sample sizes. The OR corresponds to the centre point of the box and the horizontal line indicates the 95% CI for each study. The diamond represents the pooled estimate of the OR under the fixed-effect model. The vertical dotted line marks the point estimate of the pooled OR.