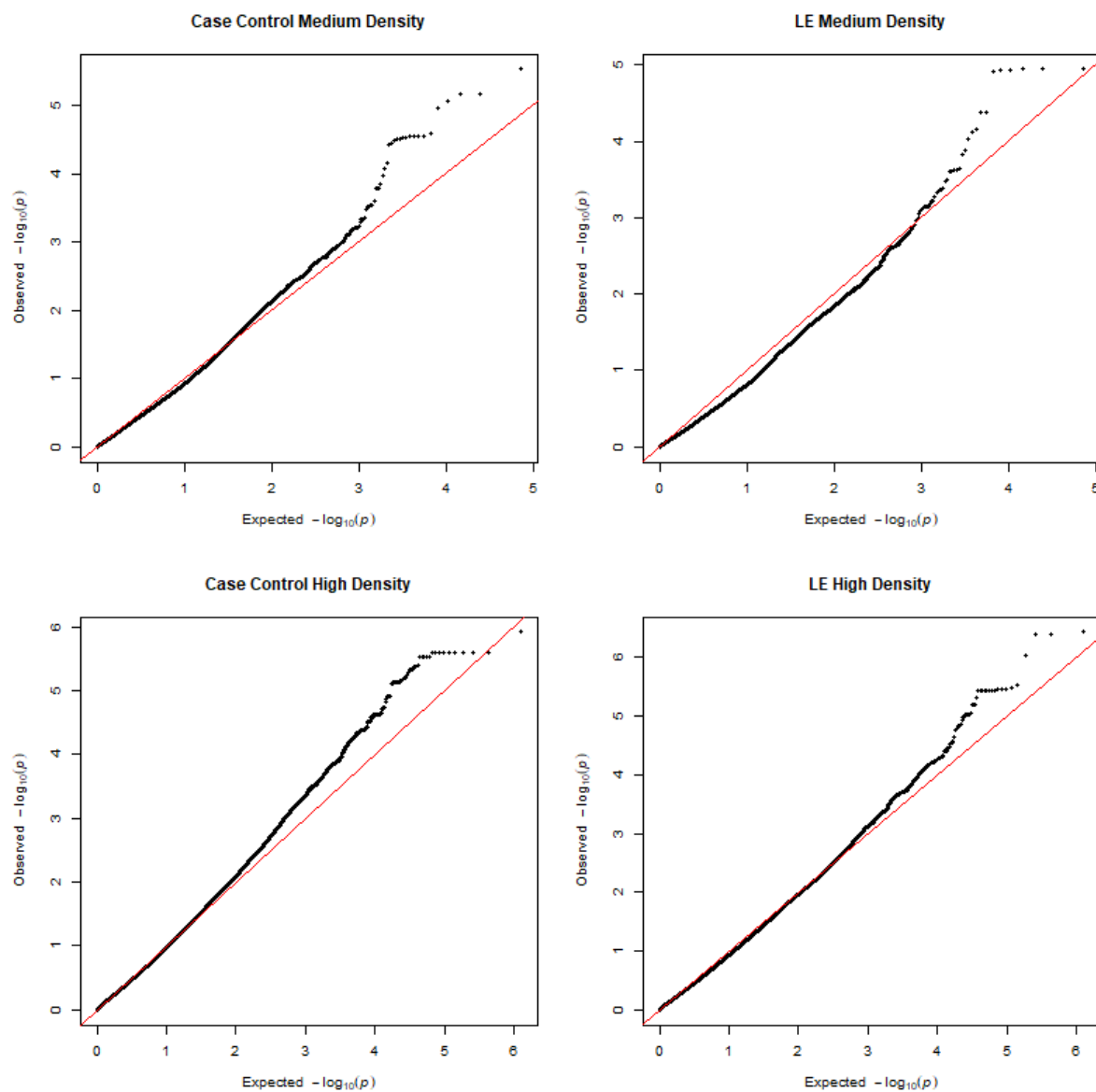


Additional file 5 Figure S4



QQ-plots of the four haplotype-based GWAS for mange susceptibility in Belgian Blue. Top left shows the case-control approach on the medium density dataset (29,010 SNPs), top right shows the quantitative (lesion extent, LE) approach on medium density, bottom left the case-control approach on the high density dataset (633,512 SNPs) and bottom right the quantitative analysis on high density.