

Figure S4. Calibration of the posterior inclusion probabilities from B-LORE. The SNPs were put into 10 bins of width 0.1 according to their posterior inclusion probabilities (PIPs). Each point on the plot represents a single bin, with the center of the PIP within that bin on the *x*-axis and the proportion of SNPs which were true positives in that bin on the *y*-axis. Vertical bars show ± 2 standard errors of the proportions, assuming a binomial distribution. Panel (a) is the result of B-LORE using maximum of 2 causal SNPs and panel (b) using maximum of 3 causal SNPs.