Figure S2 (A-C): Negative log(10) statistical significance plots of the each local region with 500 kbp on either side of the top SNP significantly associated with mean platelet volume

Chromosomal position is on x-axis and –log(10) of p-value is on the y-axis. The color of a circle is based on the r^2 from HapMap II data. A gray circle represents a SNP for which r^2 data is not available from HapMap II. These plots were generated using LocusZoom (http://csg.sph.umich.edu/locuszoom/).