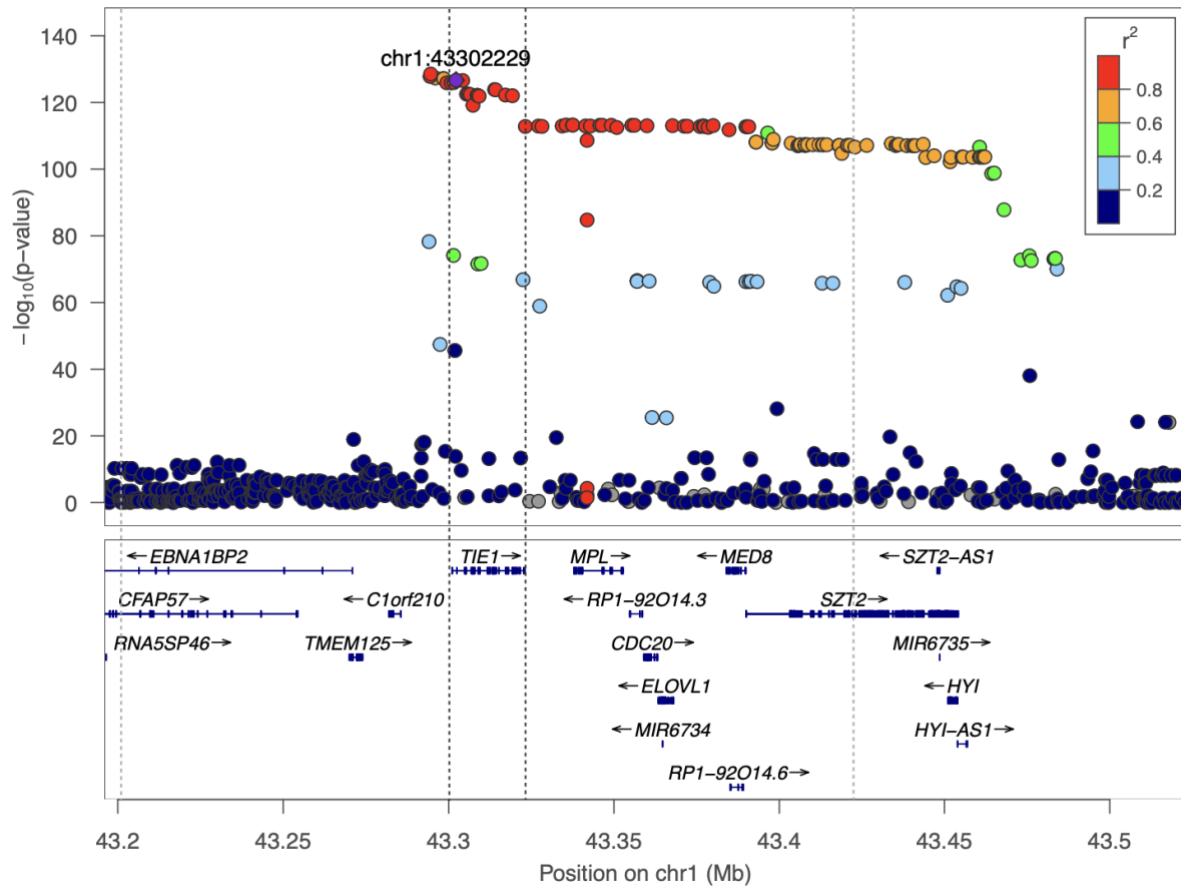


Figure S7 – LocusZoom plot of genetic associations with sTIE1 protein levels in the *TIE1* gene region with 100kB window



LocusZoom plot of genetic associations in *TIE1* gene region with circulating sTIE1 protein levels. The purple diamond is the genetic variant with the highest posterior probability of being the shared causal variant underlying the two traits, as determined by colocalization. Dotted grey lines indicate boundaries of *TIE1* gene body and 100kB window either side.