|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Lead SNP** | **Locus** | **DNAse I Hypersensitivity Site** | **Promotor Histone Marks** | **Enhancer Histone Marks** | **eQTL** | **Conservation** |
| rs532436 | ABO | Y | Y | N | N | N |
| rs2107595 | HDAC9 | Y | N | Y | N | Y |
| rs2723334 | PITX2 | N | N | N | N | N |
| rs12932445 | ZFHX3 | N | Y | Y | N | N |

**Table e-4**: Functional information on lead SNPs of genome-wide significant stroke loci. Data is extracted from HaploReg v2. Shown are presence of functional features “DNAse 1 Hypersensitivity Site”, “Promotor Histone Marks”, “Enhancer Histone Marks”, “eQTL” and “Conservation in other species”. “Y” signifies presence of this feature in at least one cell type/line. “N” signifies absence of this feature.