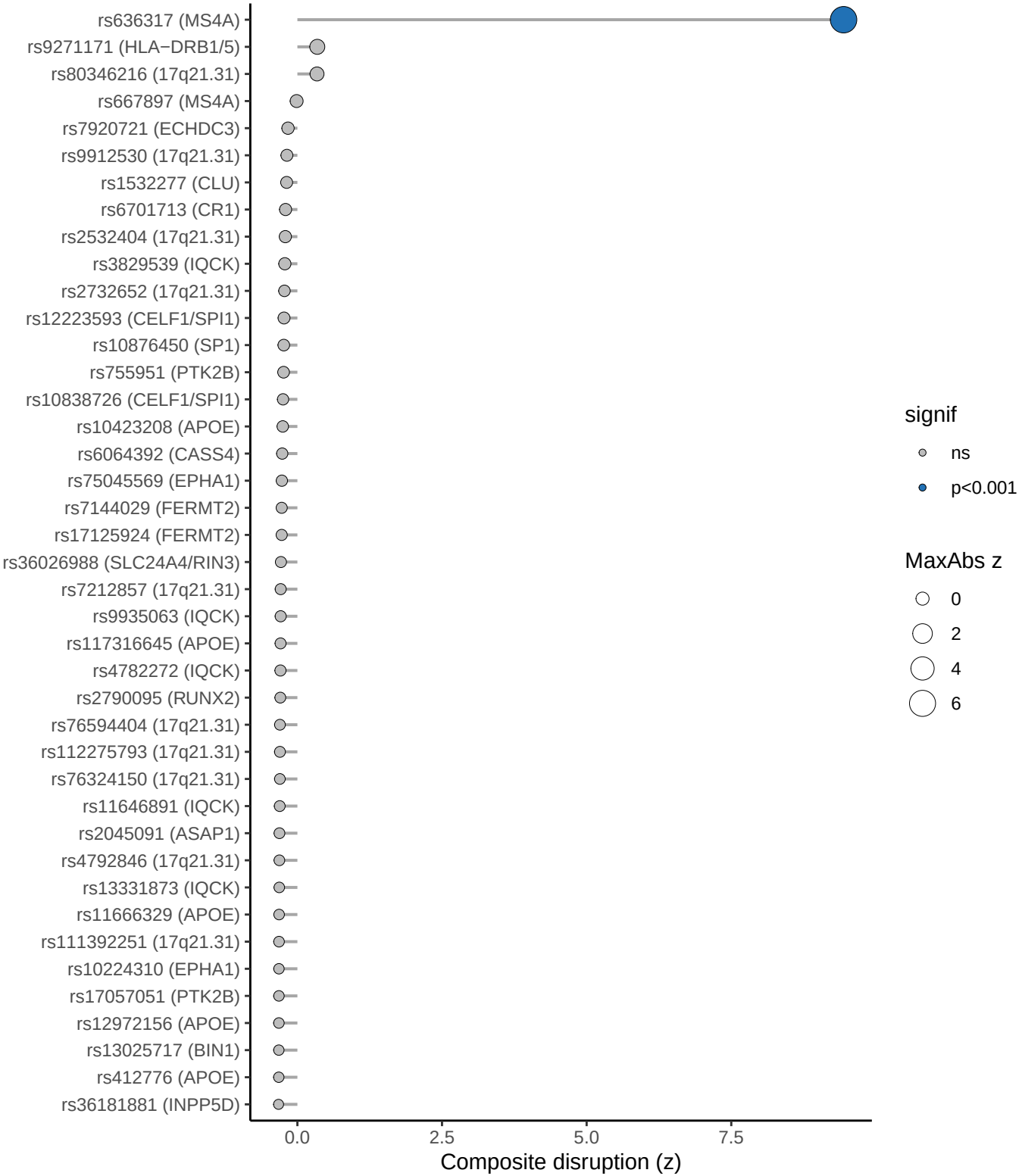
rs636317 (MS4A)
rs9271171 (HLA−DRB1/5)
rs80346216 (17q21.31)
rs667897 (MS4A)
rs7920721 (ECHDC3)
rs9912530 (17q21.31)
rs1532277 (CLU)
rs6701713 (CR1)
rs2532404 (17q21.31)
rs3829539 (IQCK)
rs2732652 (17q21.31)
rs12223593 (CELF1/SPI1)
rs10876450 (SP1)
rs755951 (PTK2B)
rs10838726 (CELF1/SPI1)
rs10423208 (APOE)
rs6064392 (CASS4)
rs75045569 (EPHA1)
rs7144029 (FERMT2)
rs17125924 (FERMT2)
rs36026988 (SLC24A4/RIN3)
rs7212857 (17q21.31)
rs9935063 (IQCK)
rs117316645 (APOE)
rs4782272 (IQCK)
rs2790095 (RUNX2)
rs76594404 (17q21.31)
rs112275793 (17q21.31)
rs76324150 (17q21.31)
rs11646891 (IQCK)
rs2045091 (ASAP1)
rs4792846 (17q21.31)
rs13331873 (IQCK)
rs11666329 (APOE)
rs111392251 (17q21.31)
rs10224310 (EPHA1)
rs17057051 (PTK2B)
rs12972156 (APOE)
rs13025717 (BIN1)
rs412776 (APOE)
rs36181881 (INPP5D)
0.0
2.5
5.0
7.5
Composite disruption (z)
signif
ns
p<0.001
MaxAbs z
0
2
4
6