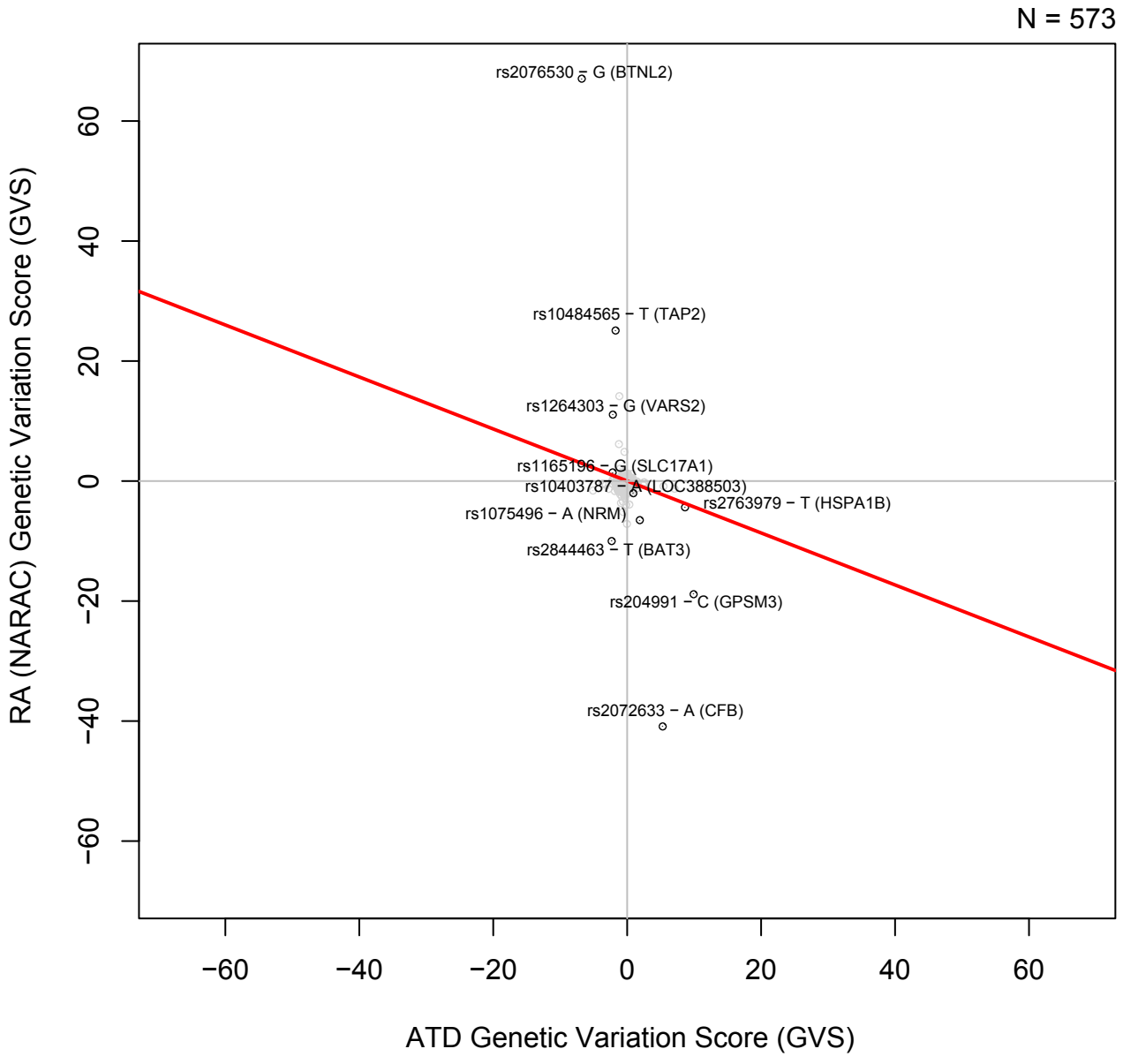
N = 573
rs2076530 − G (BTNL2)
rs10484565 − T (TAP2)
rs1264303 − G (VARS2)
rs1165196 − G (SLC17A1)
rs10403787 − A (LOC388503)
rs2763979 − T (HSPA1B)
rs1075496 − A (NRM)
rs2844463 − T (BAT3)
rs204991 − C (GPSM3)
rs2072633 − A (CFB)
RA (NARAC) Genetic Variation Score (GVS)
ATD Genetic Variation Score (GVS)
−60
−40
−20
0
20
40
60