Normalized regression coefficients: mixed vs fixed effects model (1) Age (7) DxAFF (13) RNA_batchF (19) Ancestry.3 (14) RNA_batchG (15) RNA_batchH (16) RNA_batch0 (2) InstitutionPenn (8) RIN (20) Ancestry.4 (3) InstitutionPitt (9) RNA_batchB (21) Ancestry.5 (10) RNA_batchD (11) RNA_batchD (12) RNA_batchE (5) PMI (17) Ancestry.1 signif.level 0.01 (6) DxSCZ (18) Ancestry.2 -5 5 5 0 -5 0 MAGEL2 **TMEM261P1** AL132709.5 RP11-909M7.3 SNHG14 17 ⁸ 5 .5¹**9**0 178928 20 0 8¹789 -5 17 ZIM2 NAP1L5 MEG3 PEG3 PWAR6 5 18 1120 29 29 .17 17⁵9 516 **29**⁵⁸ 0 -5 8 **SNURF** SNRPN FAM50B NDN PEG10 19° 18²⁰ 5 tpg mixed effects (mixed.1) 19¹81²07⁸ **39** 1800 1898 18 0 20 -5 KCNQ10T1 ZDBF2 GRB10 SNORD116-20 KCNK9 5 8³²⁸ 21 **49** 0 -5 INPP5F RP13-487P22. **MEST ZNF331** hsa-mir-335 8 5 17 1**9**08 0 8 -5 IGF2 UBE3A **DIRAS3** PWRN1 NLRP2 5 18 73²89⁵ 19 10 1133 1188 1800 0 17 -5 0 5 -5 -5 0 5 -5 0 5 tpafixed effects (unlm.Q)