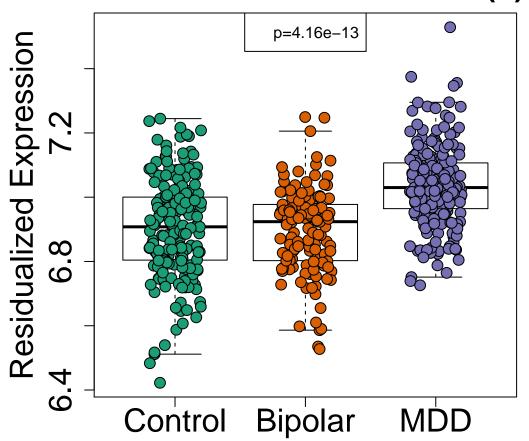
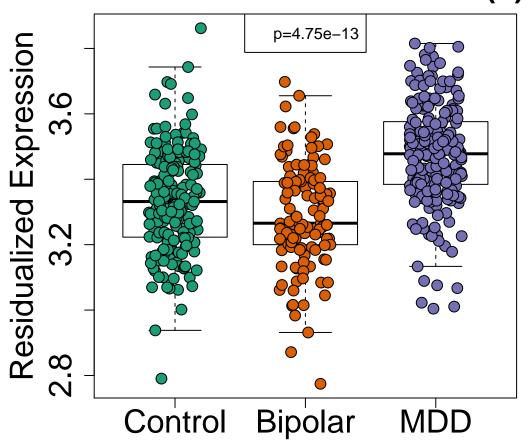
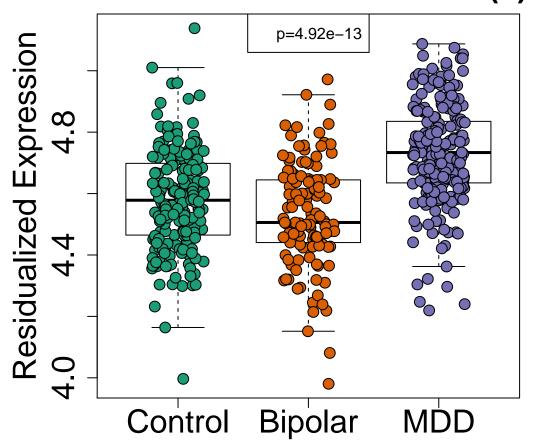
GNAS chr20:58891727-58891819(+)



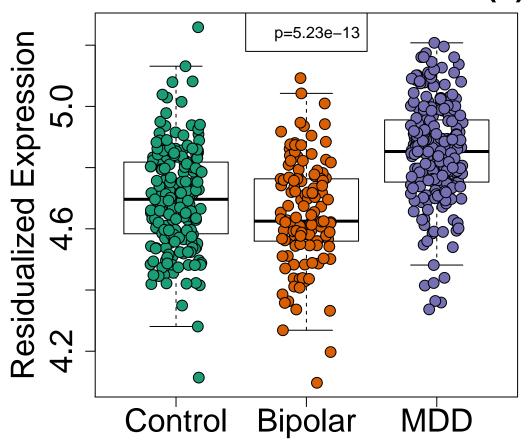
MAPT chr17:45894382-45894686(+)



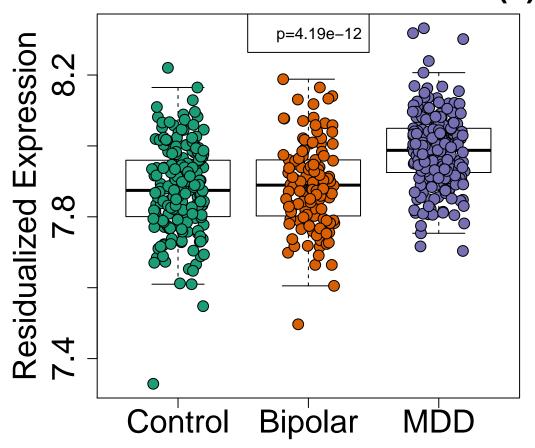
MAPT chr17:45894566-45894686(+)



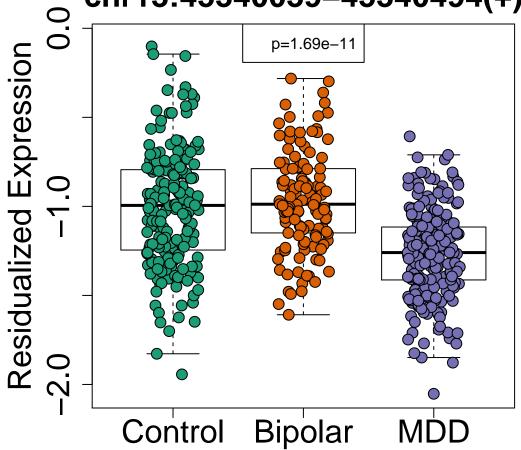
MAPT chr17:45894576-45894686(+)



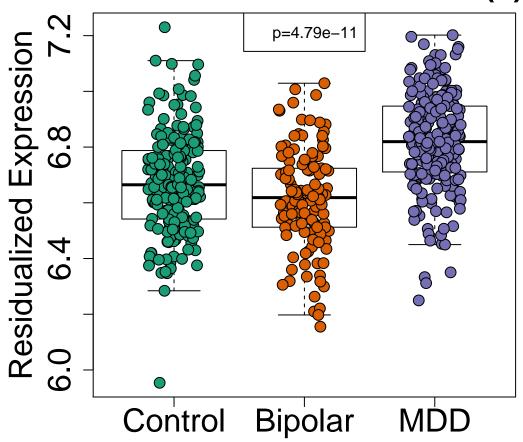
GPM6A chr4:175634546-175635057(-)



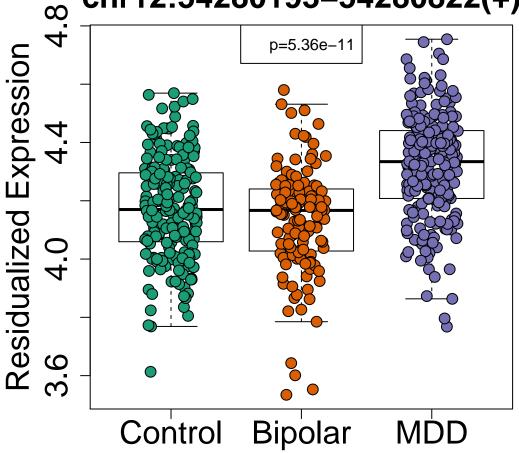
RP11-290D2.6 chr13:45340039-45340494(+)



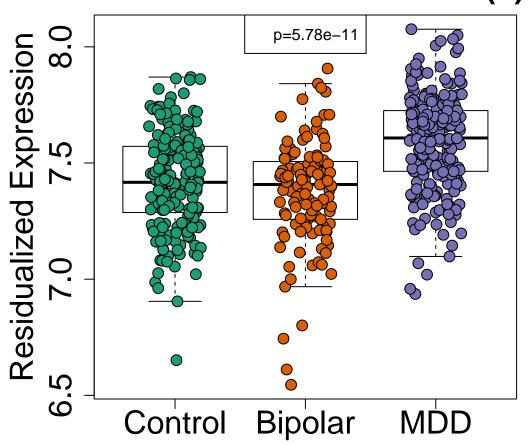
MAPT chr17:45894668-45894686(+)



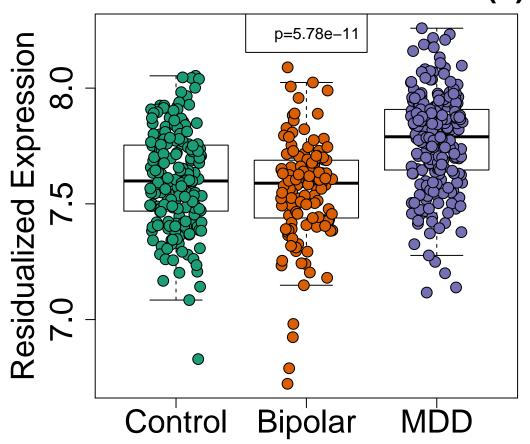
HNRNPA1 chr12:54280193-54280822(+)



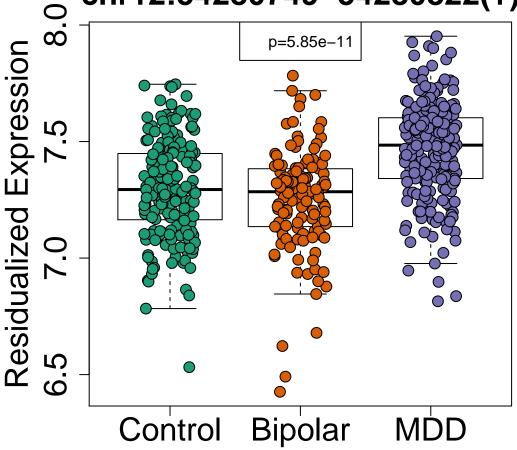
HNRNPA1 chr12:54280755-54280822(+)



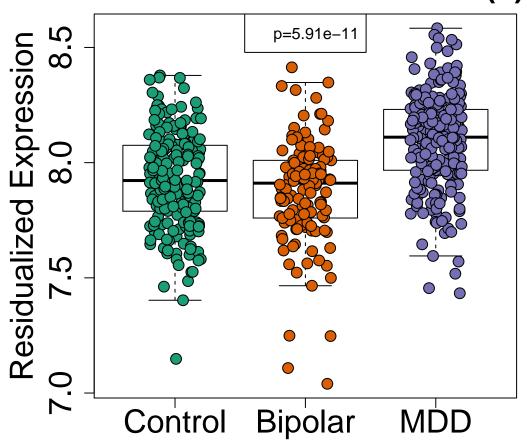
HNRNPA1 chr12:54280763-54280822(+)



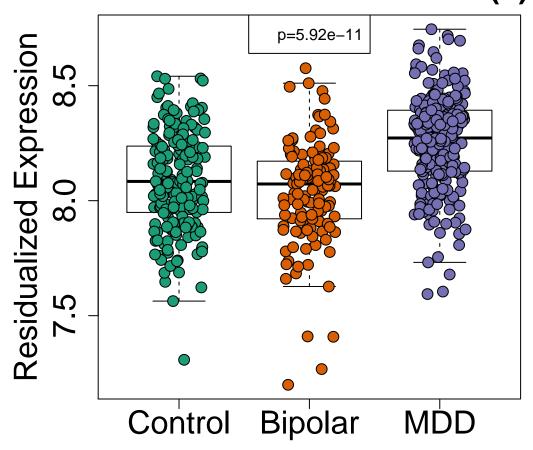
HNRNPA1 chr12:54280749-54280822(+)



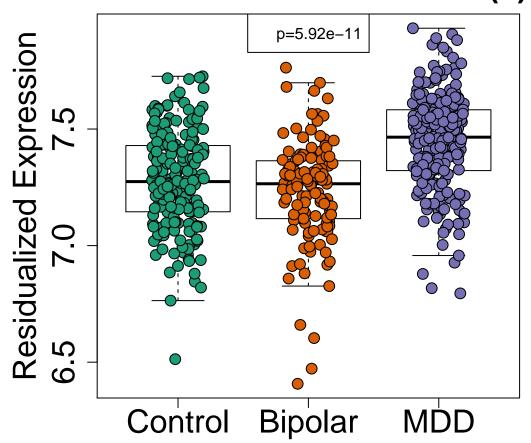
HNRNPA1 chr12:54280775-54280822(+)



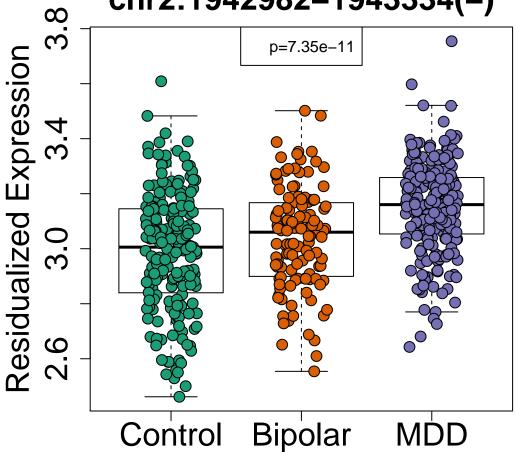
HNRNPA1 chr12:54280780-54280822(+)



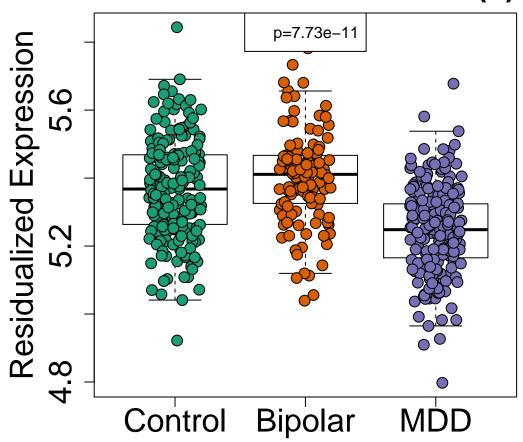
HNRNPA1 chr12:54280748-54280822(+)



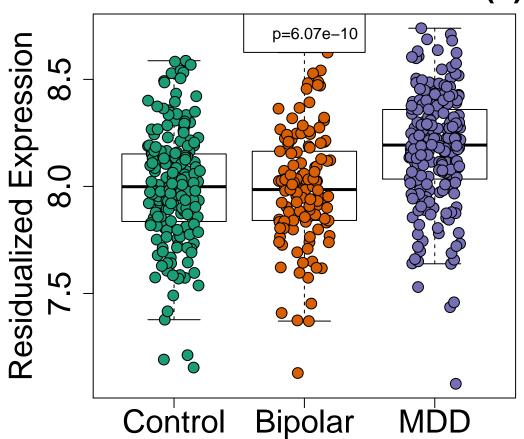
MYT1L chr2:1942982-1943334(-)



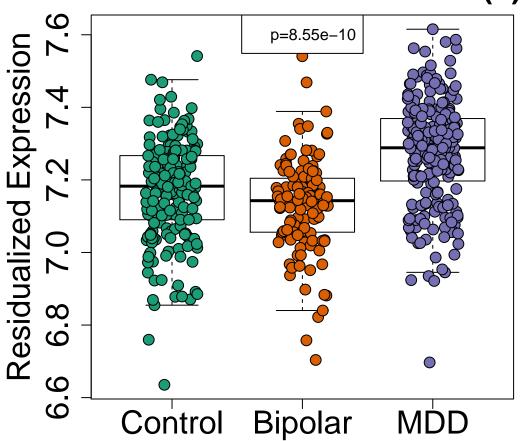
HNRNPD chr4:82359471-82359639(-)



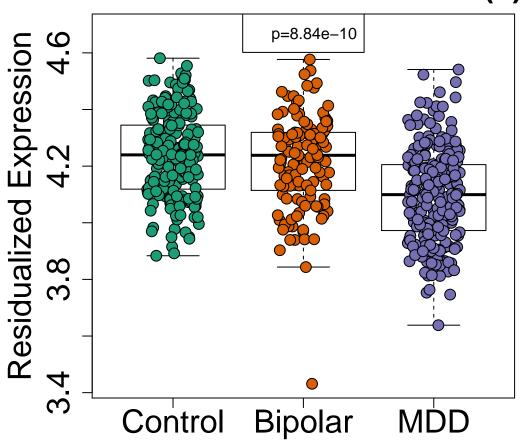
RPL17 chr18:49490454-49490468(-)



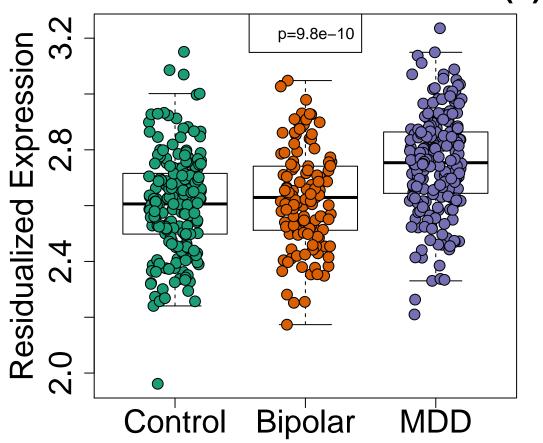
HNRNPA1 chr12:54281386-54281502(+)



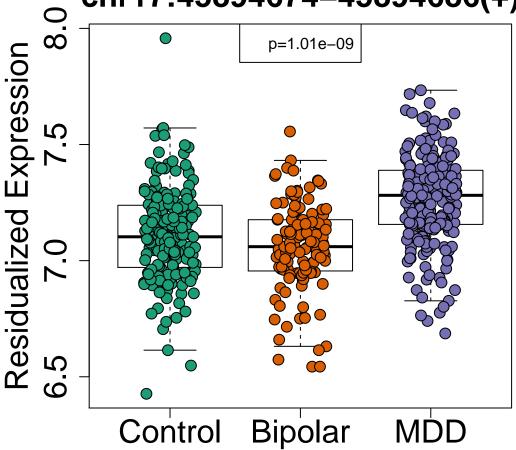
FUS chr16:31186802-31187245(+)



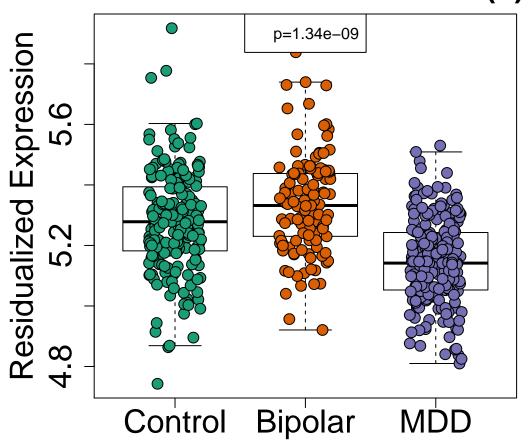
FGF14 chr13:102402127-102402457(-)



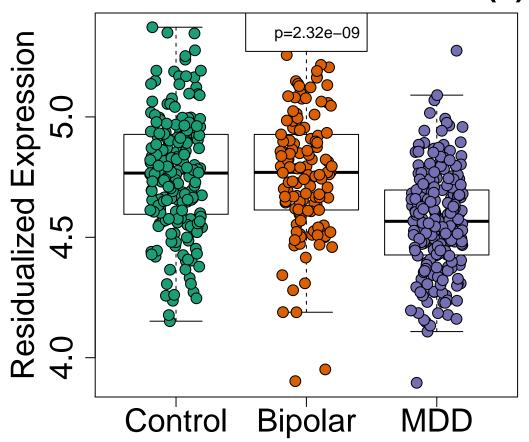
MAPT chr17:45894674-45894686(+)



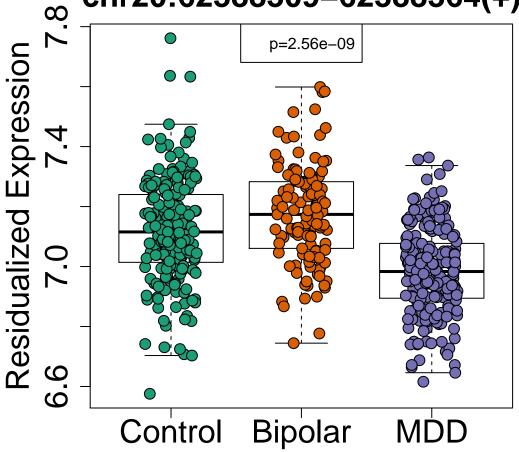
RPS21 chr20:62388309-62388519(+)



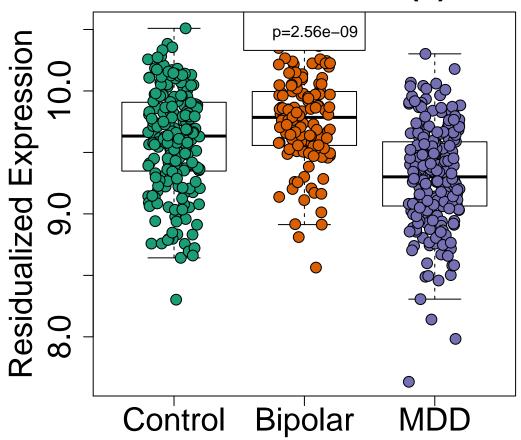
FUS chr16:31191399-31191558(+)



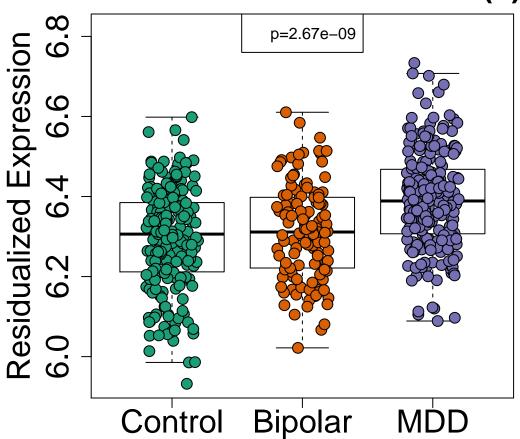
RPS21 chr20:62388309-62388364(+)



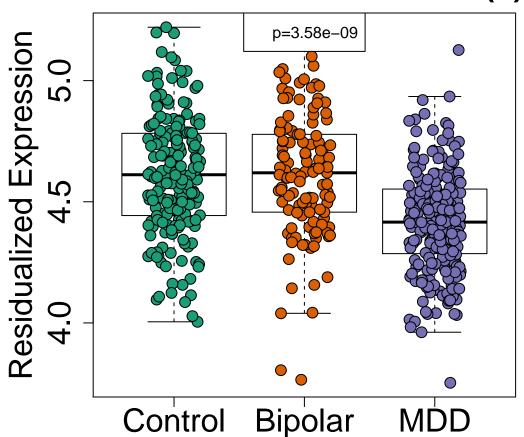
MT-RNR2 chrM:1671-3229(+)



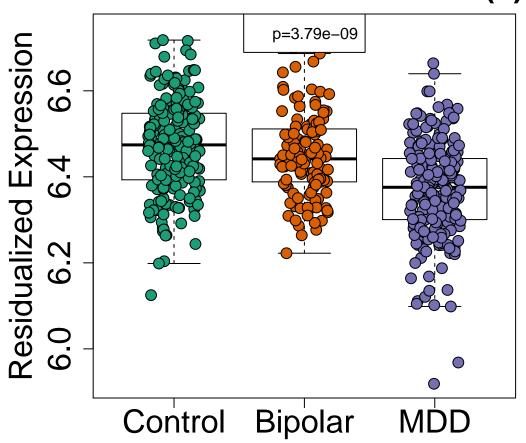
THRA chr17:40083835-40083982(+)



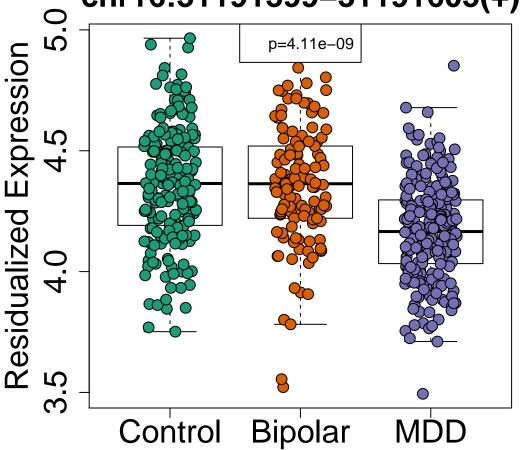
FUS chr16:31191399-31191575(+)



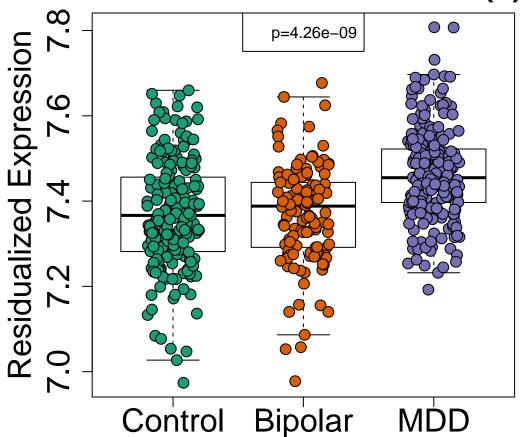
ATXN10 chr22:45700282-45700378(+)



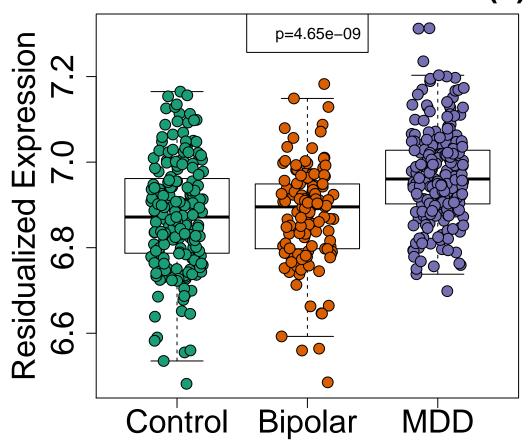
FUS chr16:31191399-31191605(+)



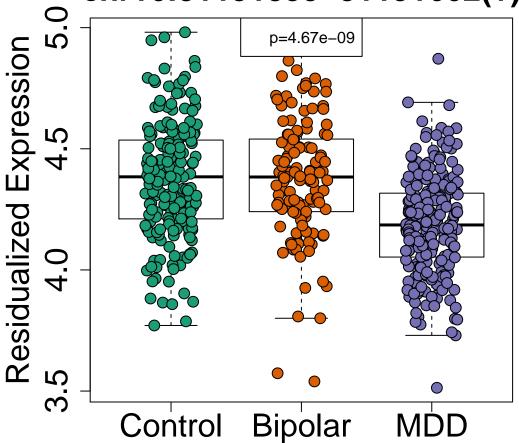
GNAS chr20:58891760-58891865(+)



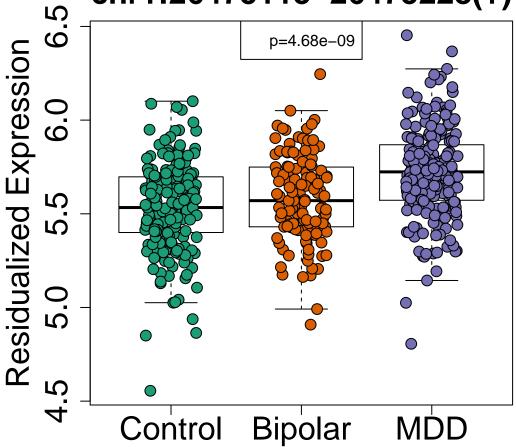
GNAS chr20:58891716-58891865(+)



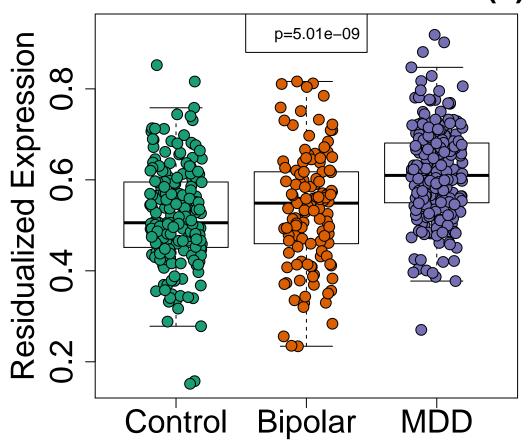
FUS chr16:31191399-31191602(+)



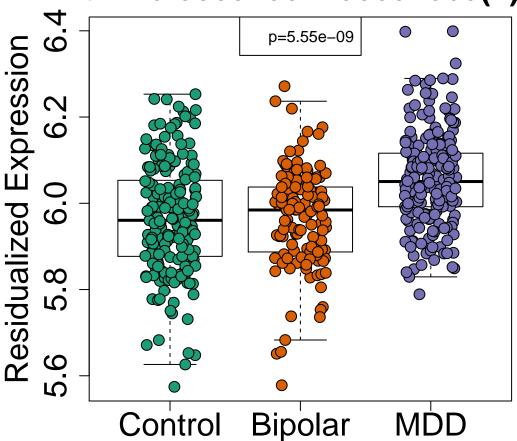
HMGN2 chr1:26475113-26475228(+)



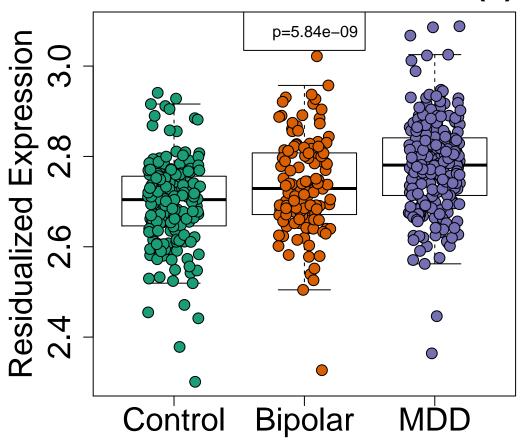
SLC25A42 chr19:19110569-19112888(+)



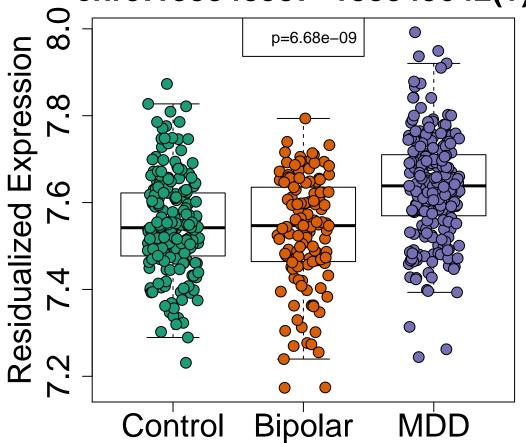
GNAS chr20:58891582-58891865(+)



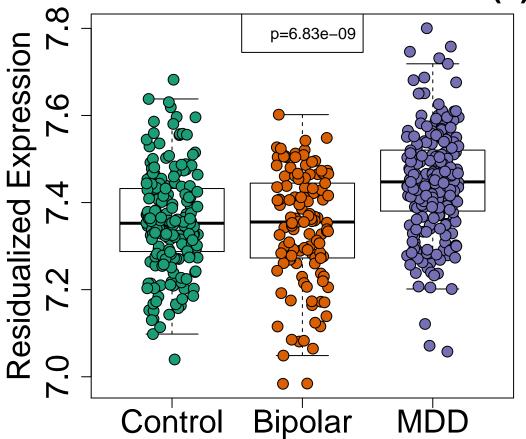
UBE2W chr8:73790605-73794115(-)



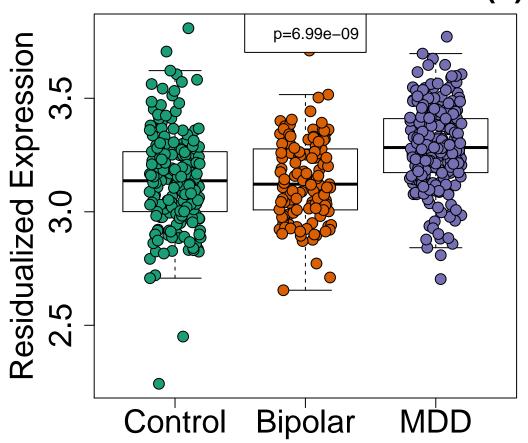
RPL7A chr9:133348937-133349042(+)



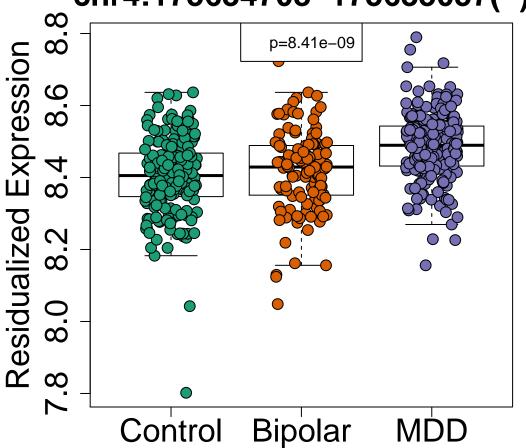
RPL7A chr9:133348922-133349042(+)



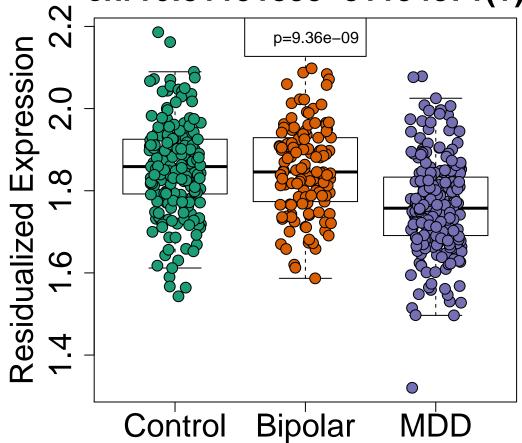
JPH4 chr14:23569415-23569717(-)



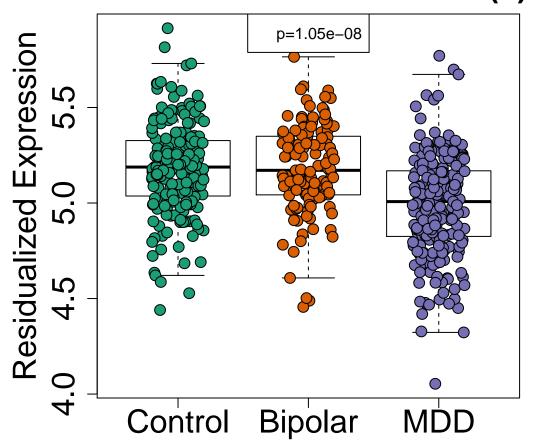
GPM6A chr4:175634708-175635057(-)



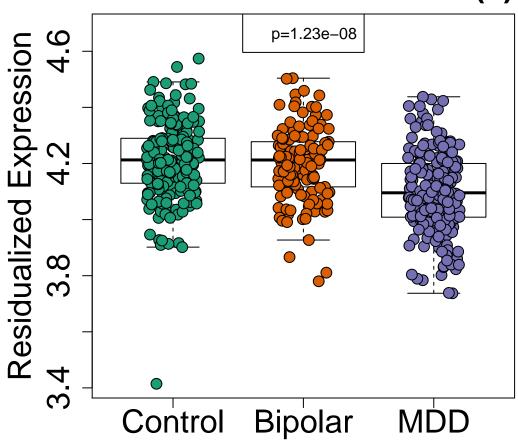
FUS chr16:31191399-31194871(+)



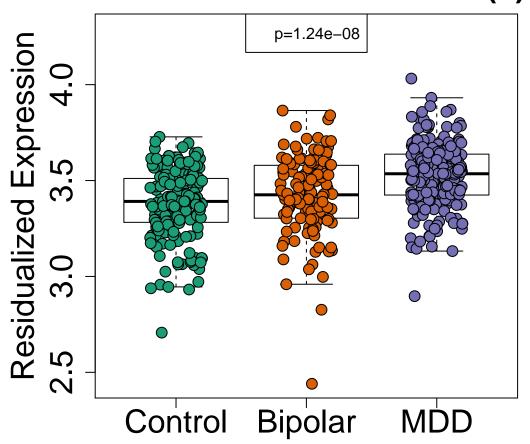
FUS chr16:31186802-31186836(+)



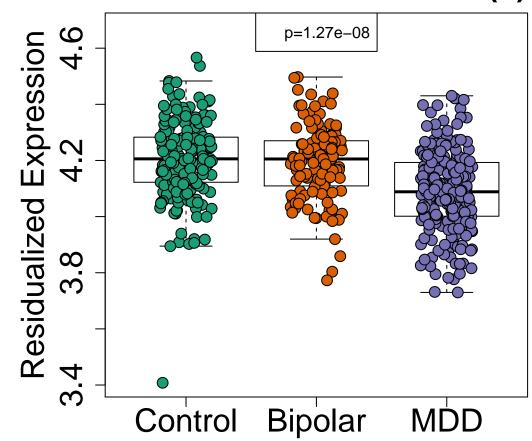
UQCRC1 chr3:48599003-48599192(-)



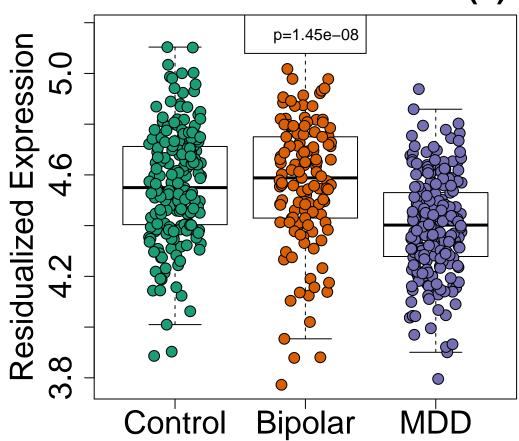
KCNQ2 chr20:63445238-63445364(-)



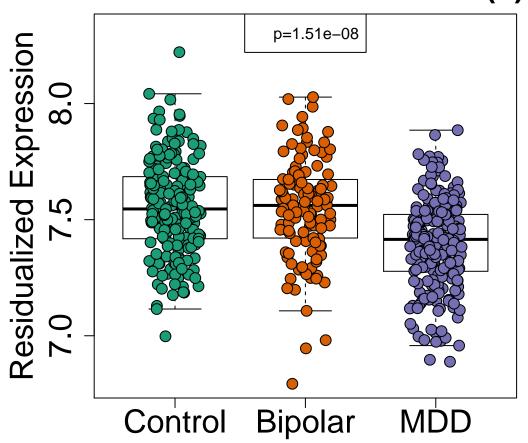
UQCRC1 chr3:48599002-48599192(-)



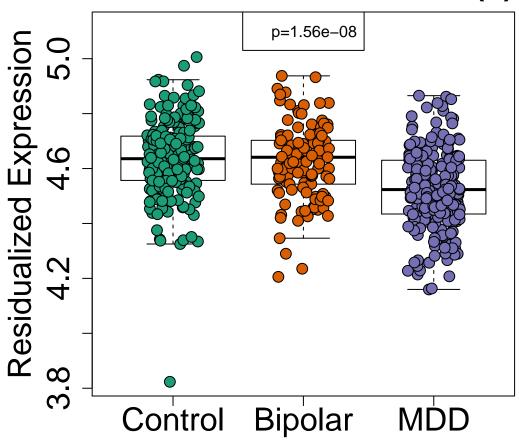
POLR2A chr17:7511811-7512014(+)



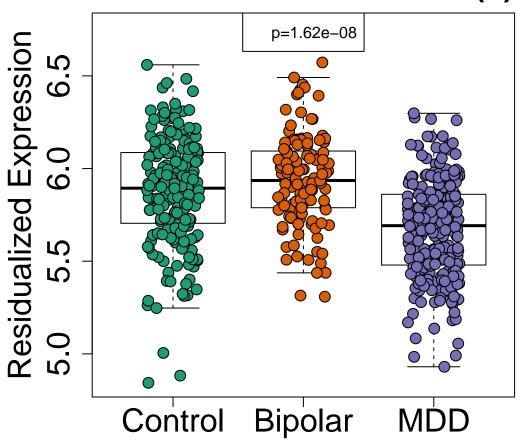
FUS chr16:31182398-31182422(+)



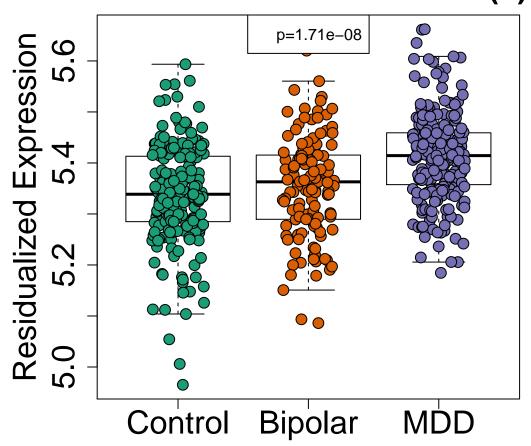
UQCRC1 chr3:48599054-48599192(-)



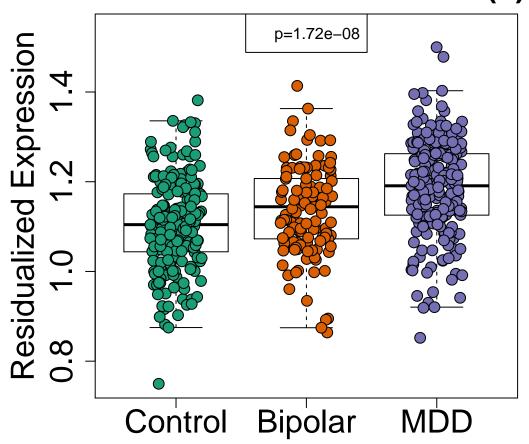
QDPR chr4:17504379–17504475(-)



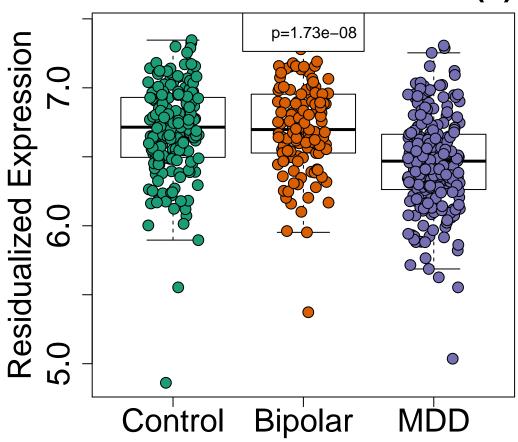
SUMO2 chr17:75167868-75168401(-)



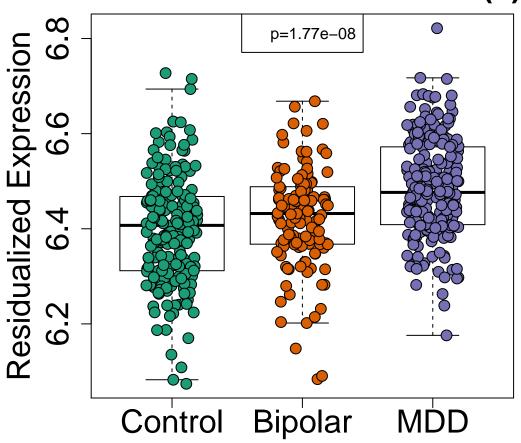
MKL1 chr22:40411144-40412892(-)



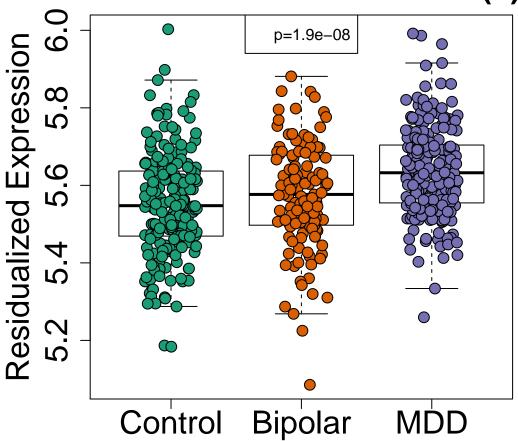
QDPR chr4:17511950-17511981(-)



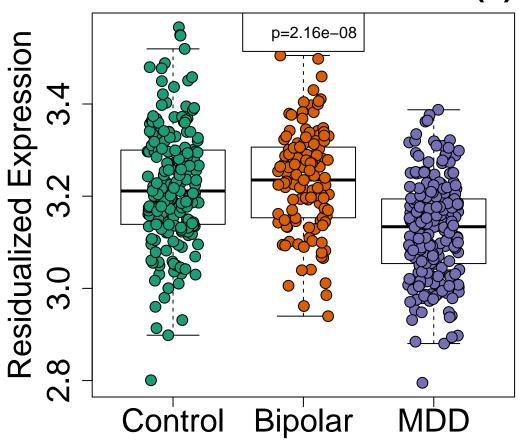
TMBIM6 chr12:49762873-49763094(+)



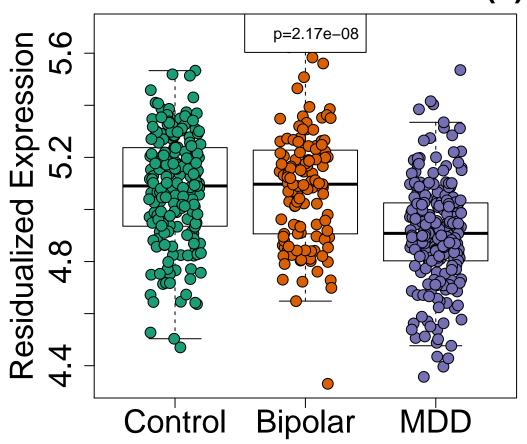
FKBP8 chr19:18532664-18532795(-)



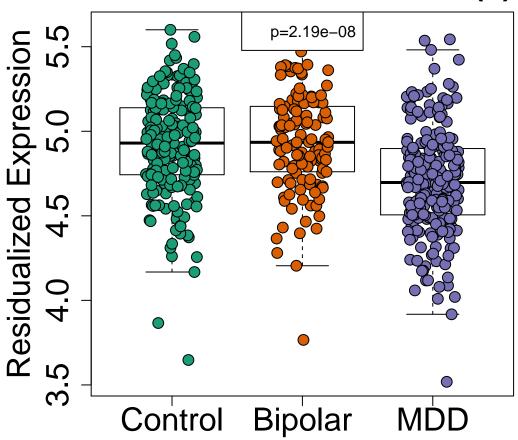
HNRNPD chr4:82357313-82358842(-)



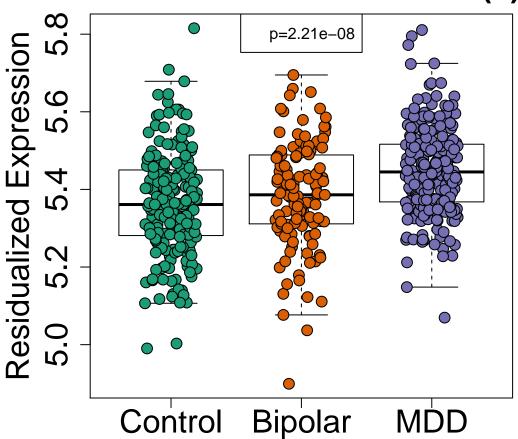
PITPNB chr22:27853579-27853663(-)



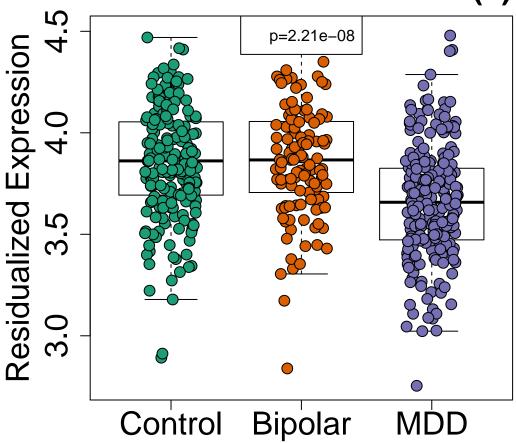
QDPR chr4:17511950-17512069(-)



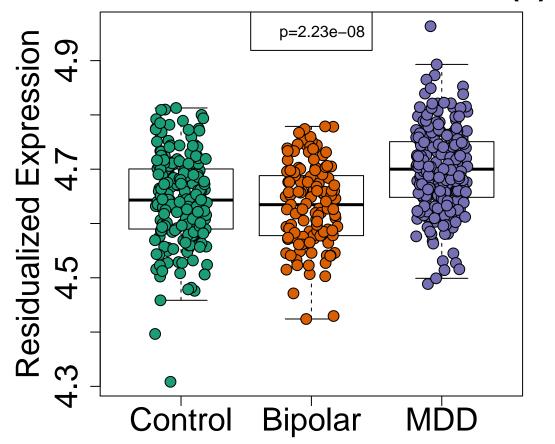
FKBP8 chr19:18532664-18532814(-)



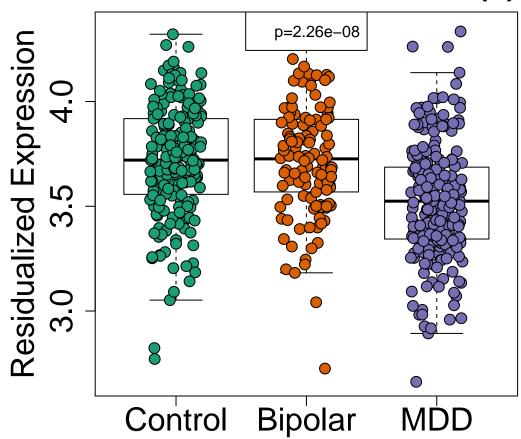
QDPR chr4:17511950-17512206(-)



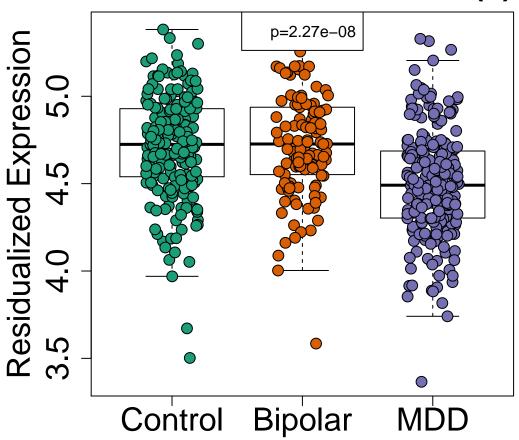
RPL6 chr12:112406747-112408656(-)



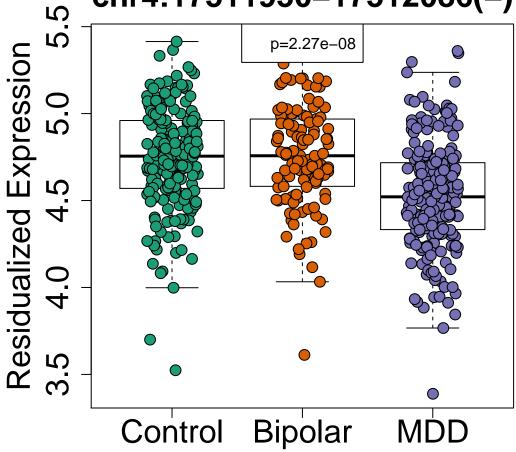
QDPR chr4:17511950-17512234(-)



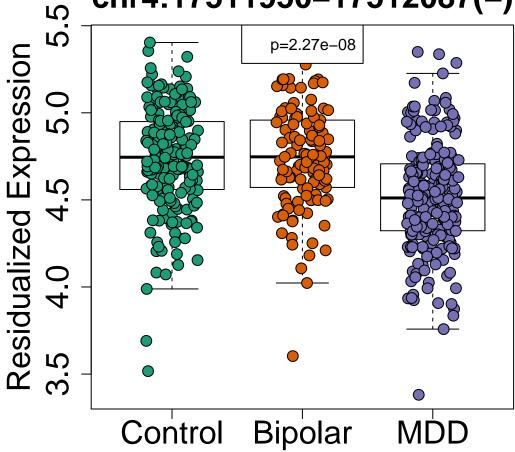
QDPR chr4:17511950-17512089(-)



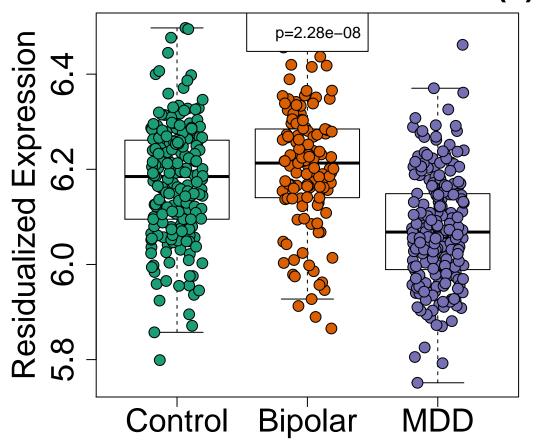
QDPR chr4:17511950-17512086(-)



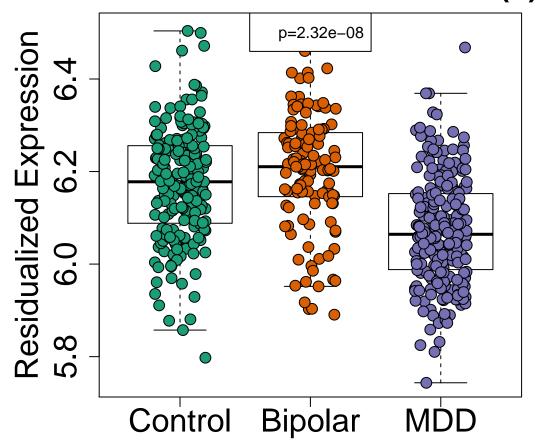
QDPR chr4:17511950-17512087(-)



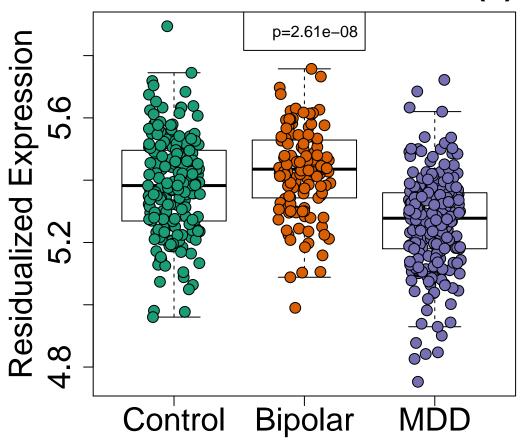
RAD21 chr8:116857279-116857473(-)



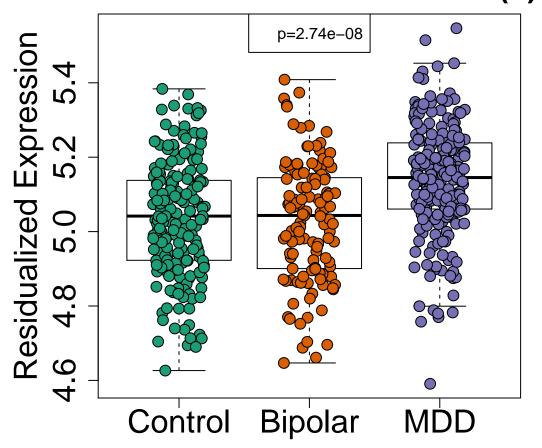
RAD21 chr8:116857271-116857473(-)



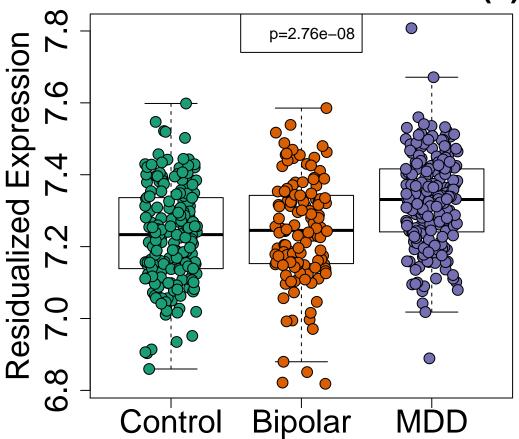
HNRNPD chr4:82359516-82359639(-)



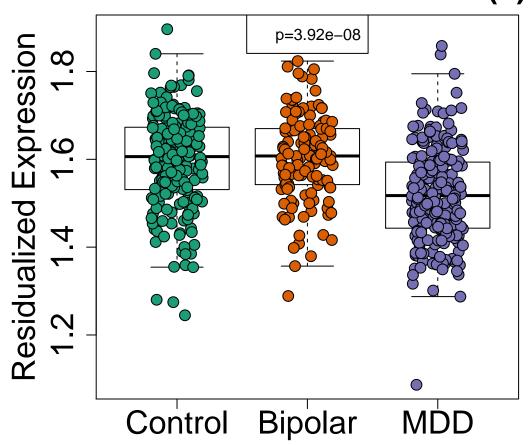
PCNX2 chr1:233198939-233199030(-)



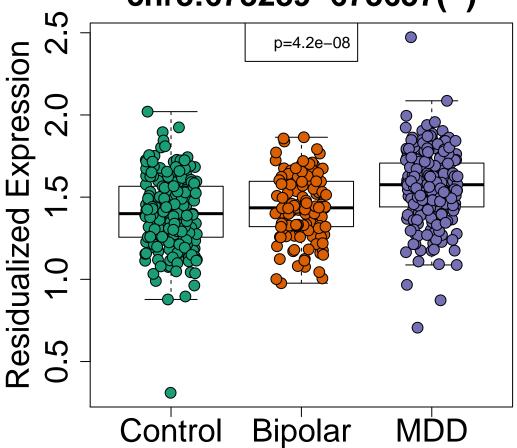
RPL4 chr15:66499575-66499652(-)



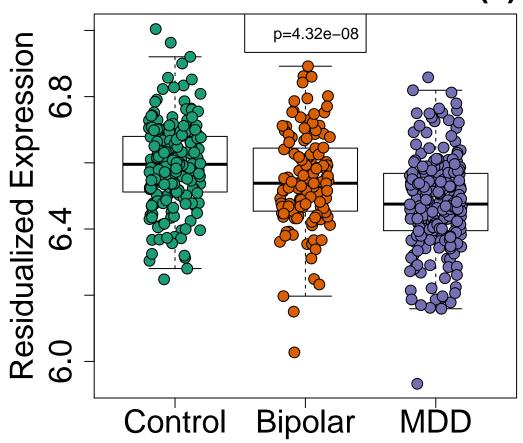
SEPHS2 chr16:30443631-30446181(-)



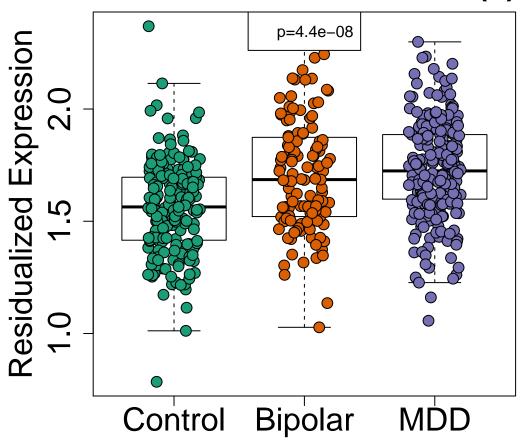
ERICH1 chr8:673289-673657(-)



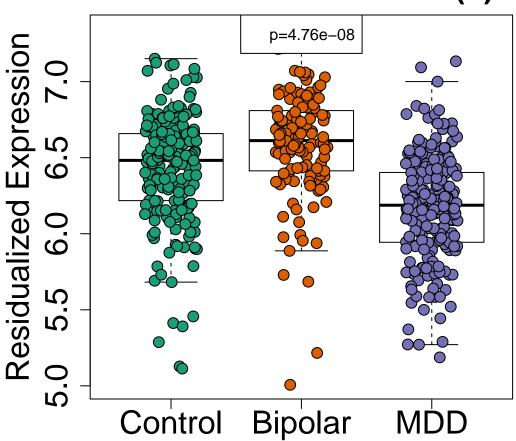
DDX3X chrX:41341484-41341616(+)



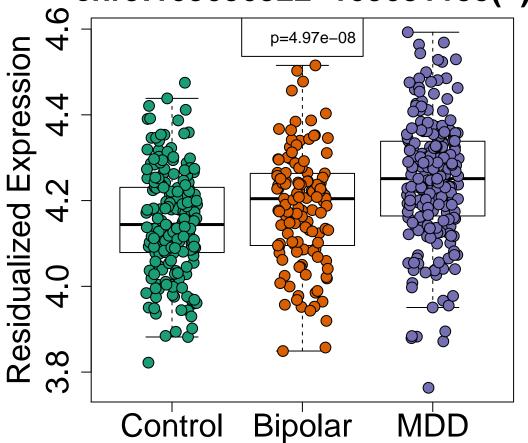
ELK1 chrX:47636167-47636927(-)



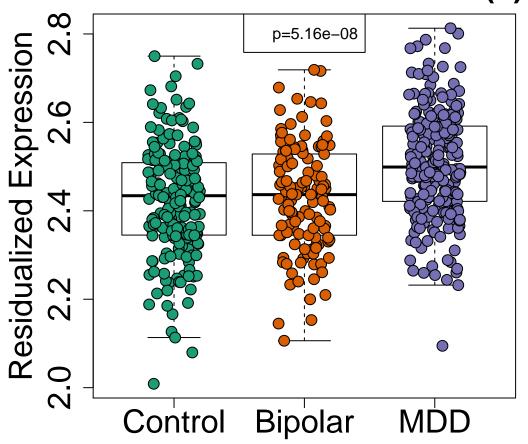
RNA5SP202 chr6:4427963-4428081(+)



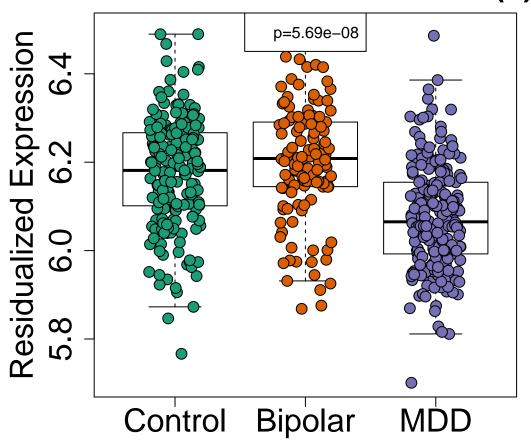
TMEM245 chr9:109090922-109091155(-)



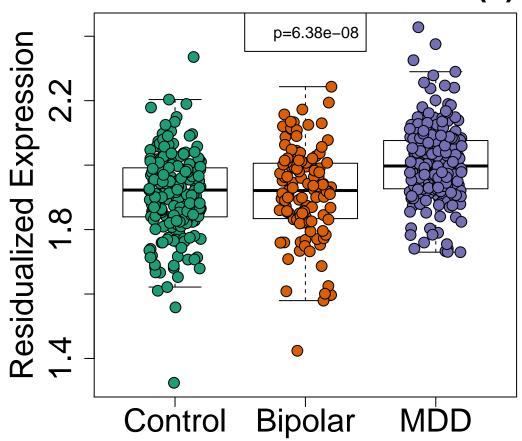
JOSD1 chr22:38699803-38700618(-)



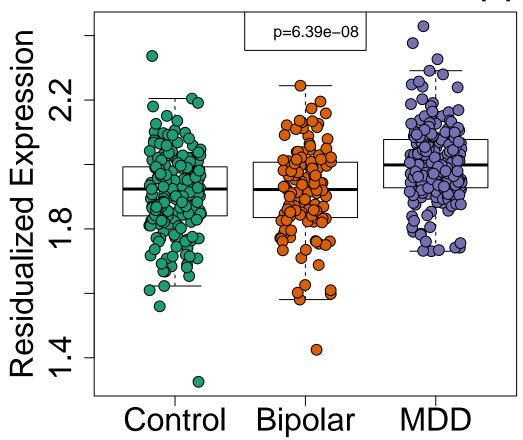
RAD21 chr8:116857289-116857473(-)



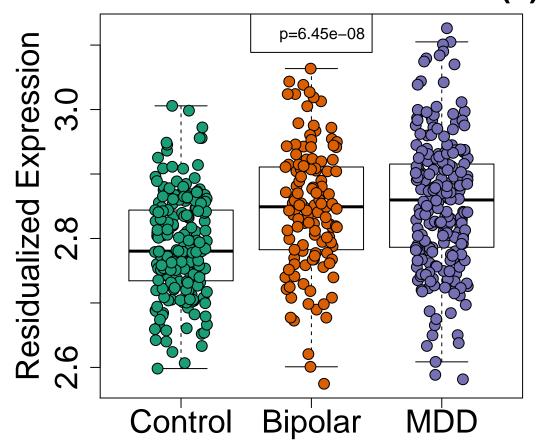
HMGA1 chr6:34244831-34246231(+)



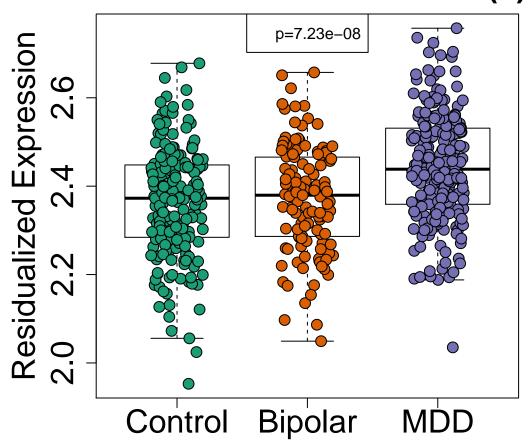
HMGA1 chr6:34244831-34246230(+)



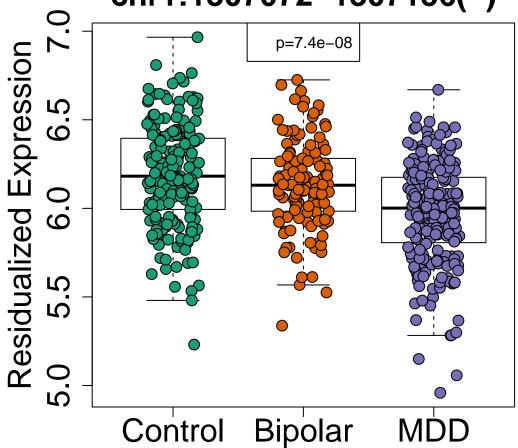
TMEM167B chr1:109094417-109096934(+)



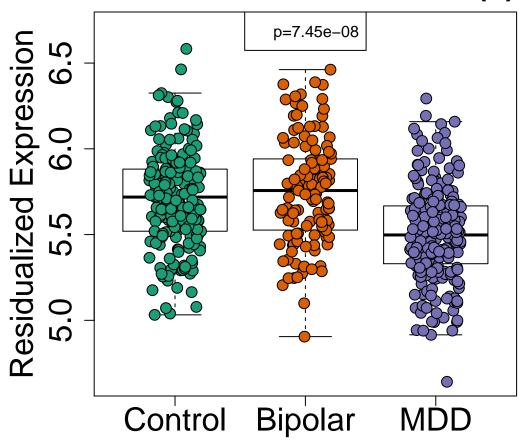
JOSD1 chr22:38699803-38700667(-)



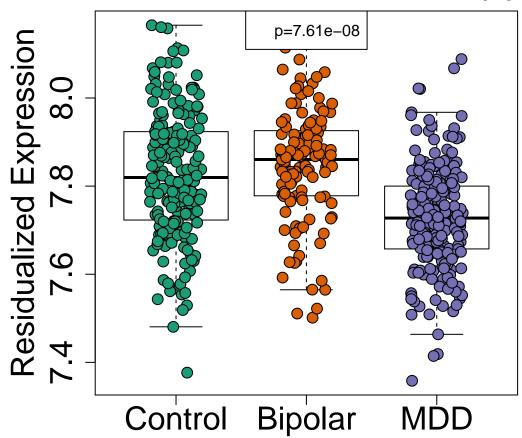
ACAP3 chr1:1307072-1307156(-)



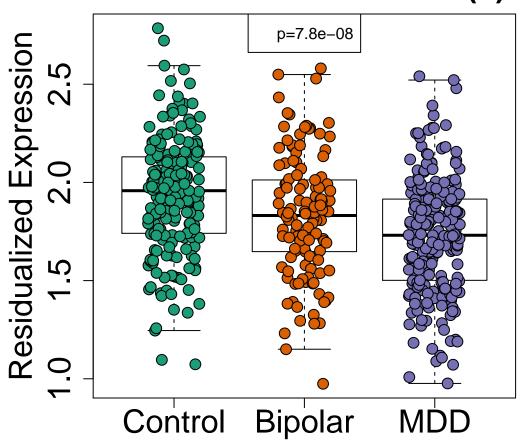
SNORA47 chr5:77080434-77080571(-)



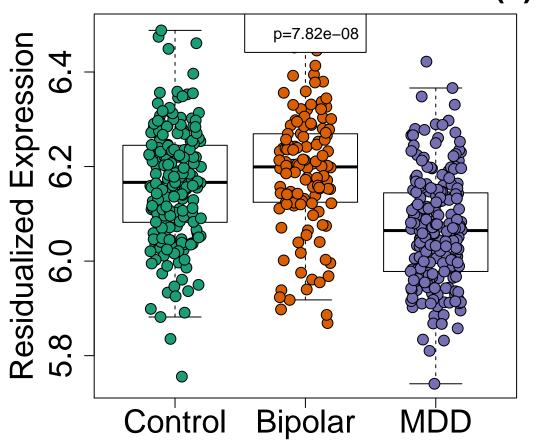
RPL37 chr5:40832442-40832555(-)



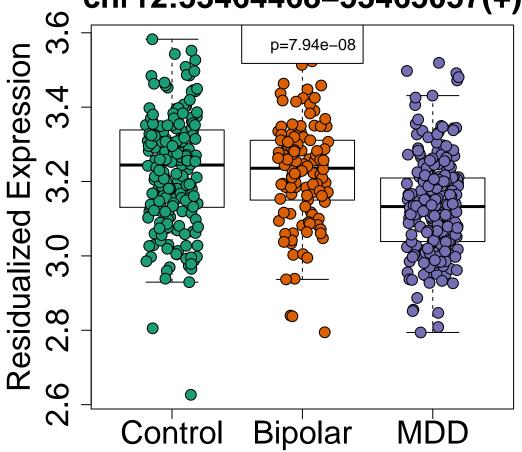
COL5A3 chr19:9966536-9966746(-)



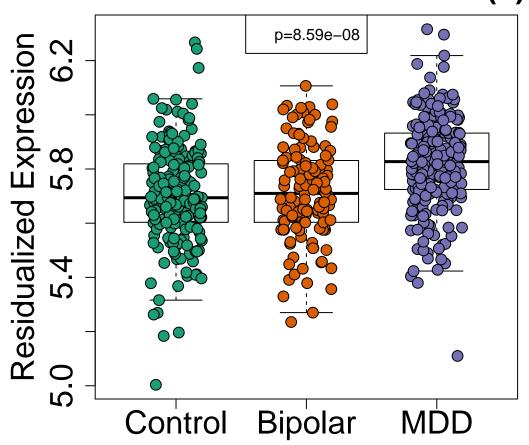
RAD21 chr8:116857267-116857473(-)



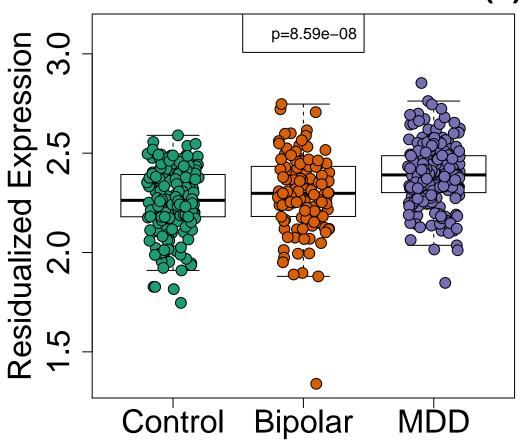
PCBP2-OT1 chr12:53464468-53465057(+)



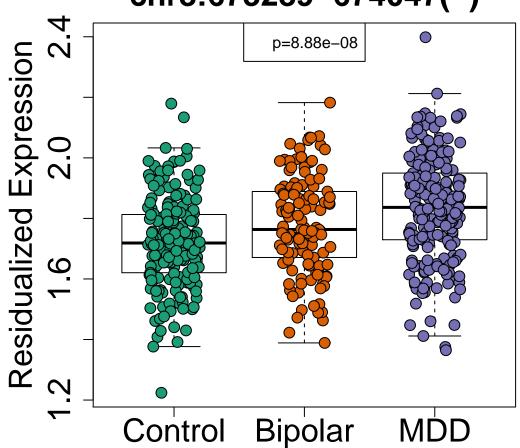
RPL17 chr18:49489471-49489550(-)



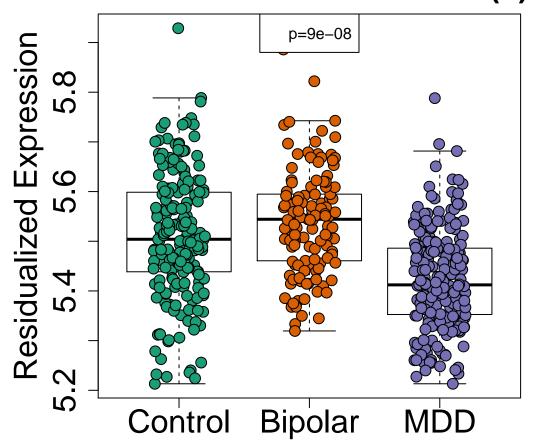
KCNQ2 chr20:63445238-63445576(-)



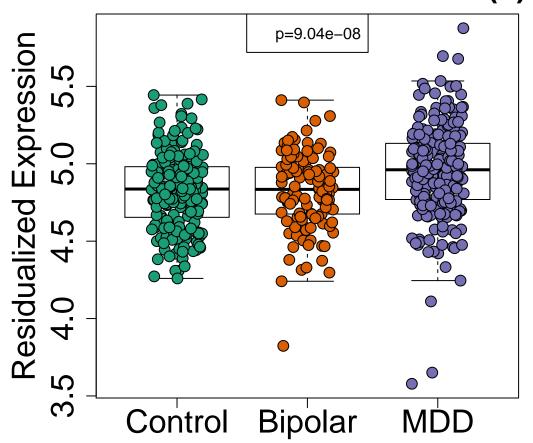
ERICH1 chr8:673289-674047(-)



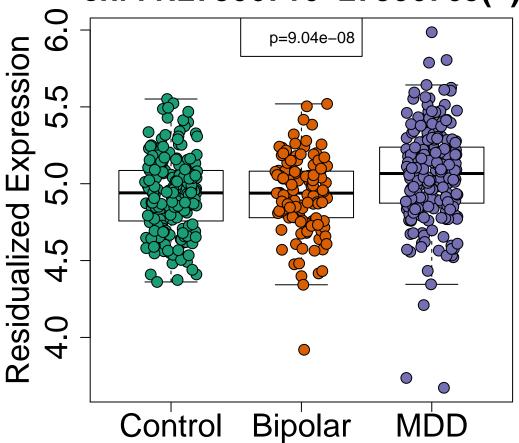
RPS21 chr20:62387611-62388520(+)



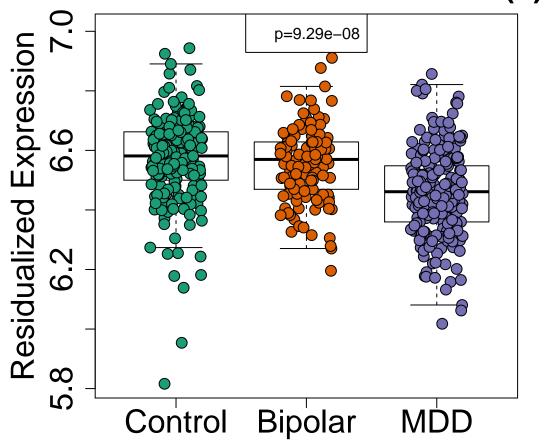
LIN7C chr11:27506716-27506773(-)



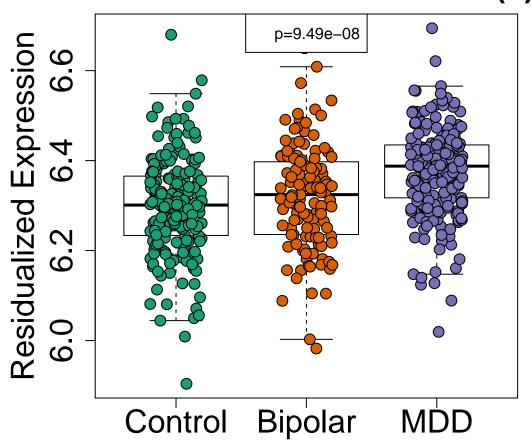
LIN7C chr11:27506716-27506769(-)



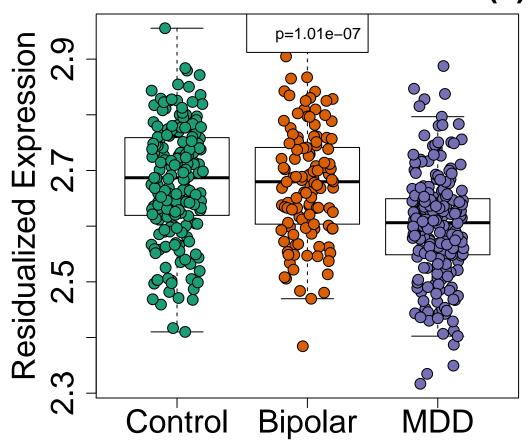
FEZ1 chr11:125455952-125456106(-)



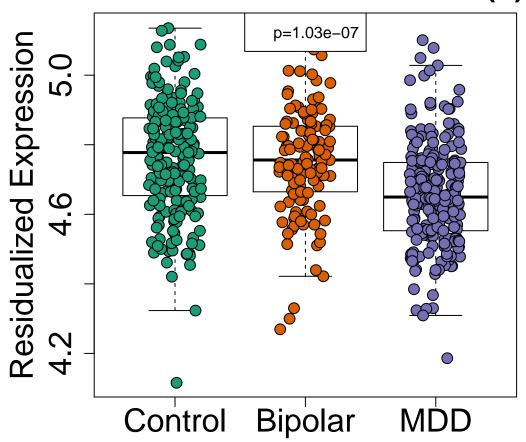
GPM6A chr4:175640753-175641745(-)



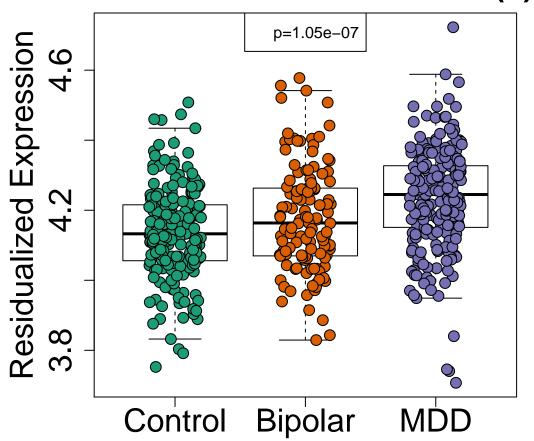
PSMC5 chr17:63828858-63830501(+)



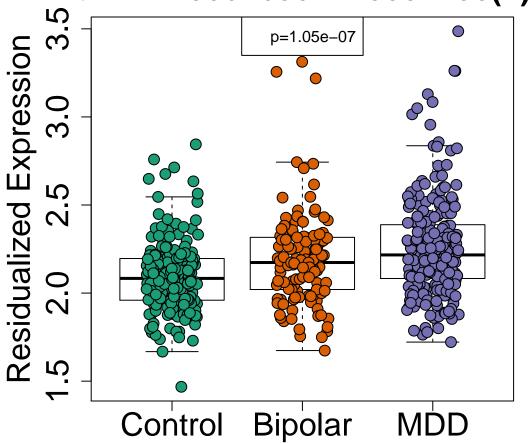
uc_338 chr12:53464720-53464901(+)



MSMO1 chr4:165341751-165343029(+)



HMGB2 chr4:173331695-173332238(-)



SFSWAP chr12:131711081–131714193(+)

