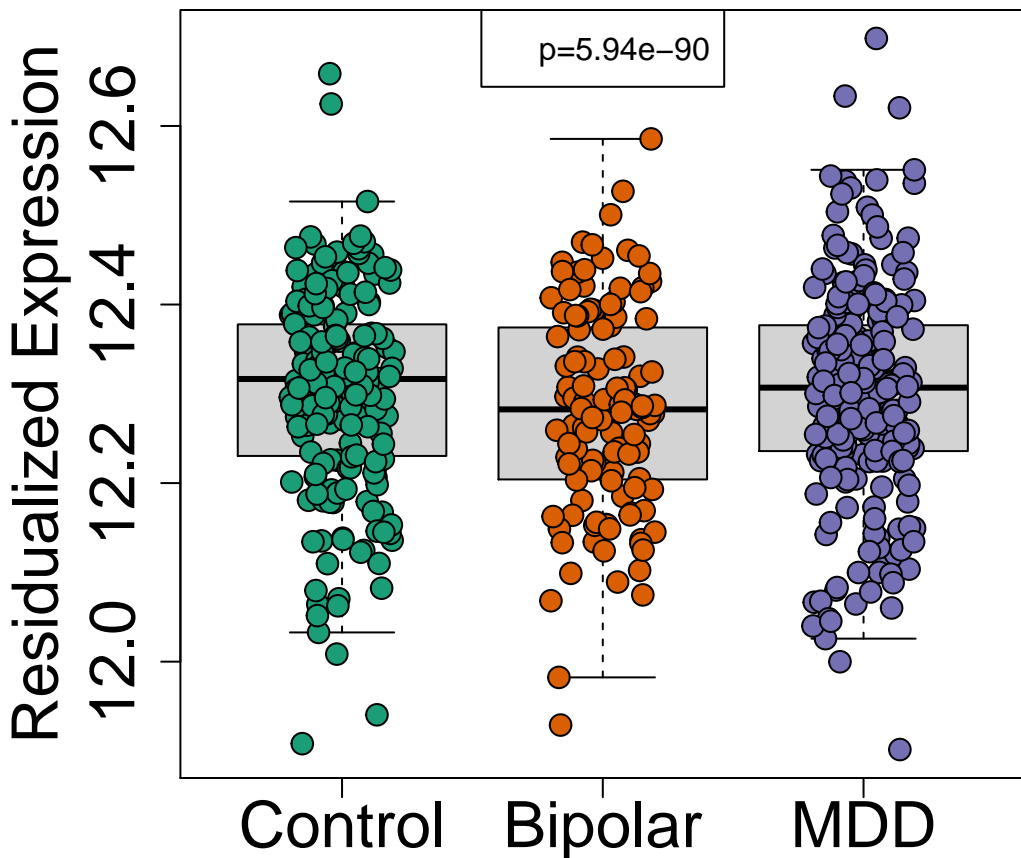


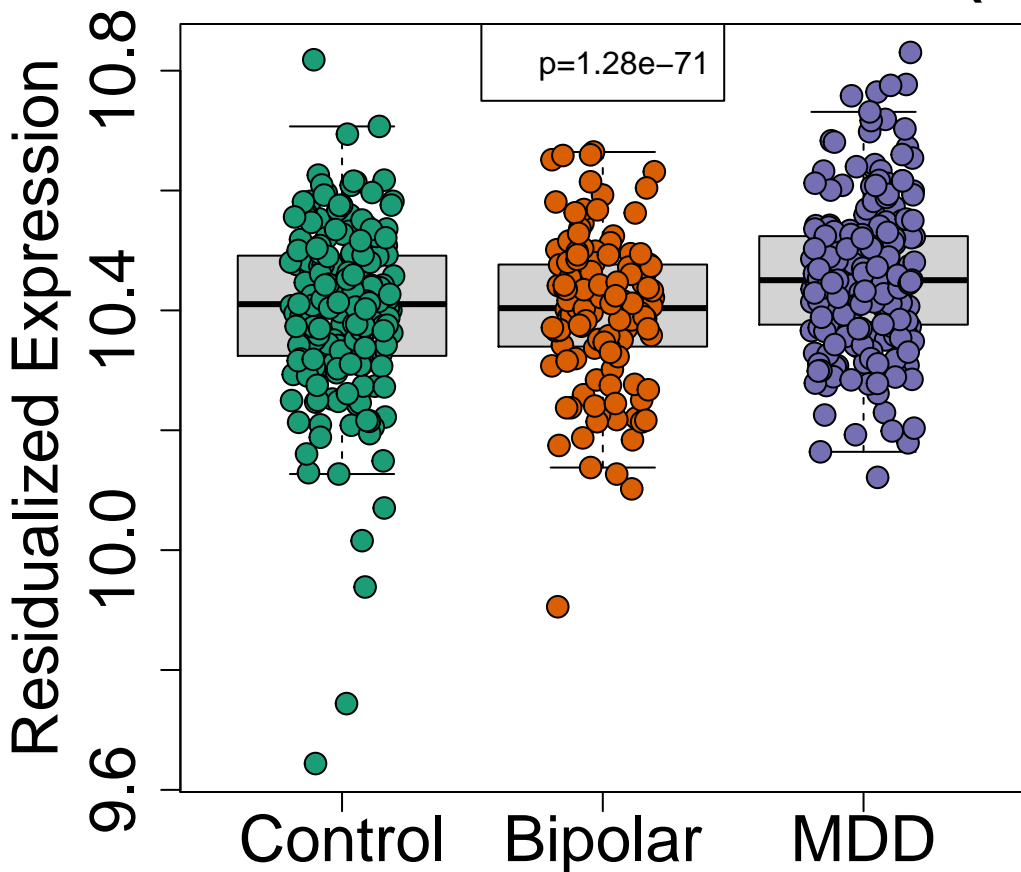
TMSB4X, InGen

chrX:12976362–12976776(+)



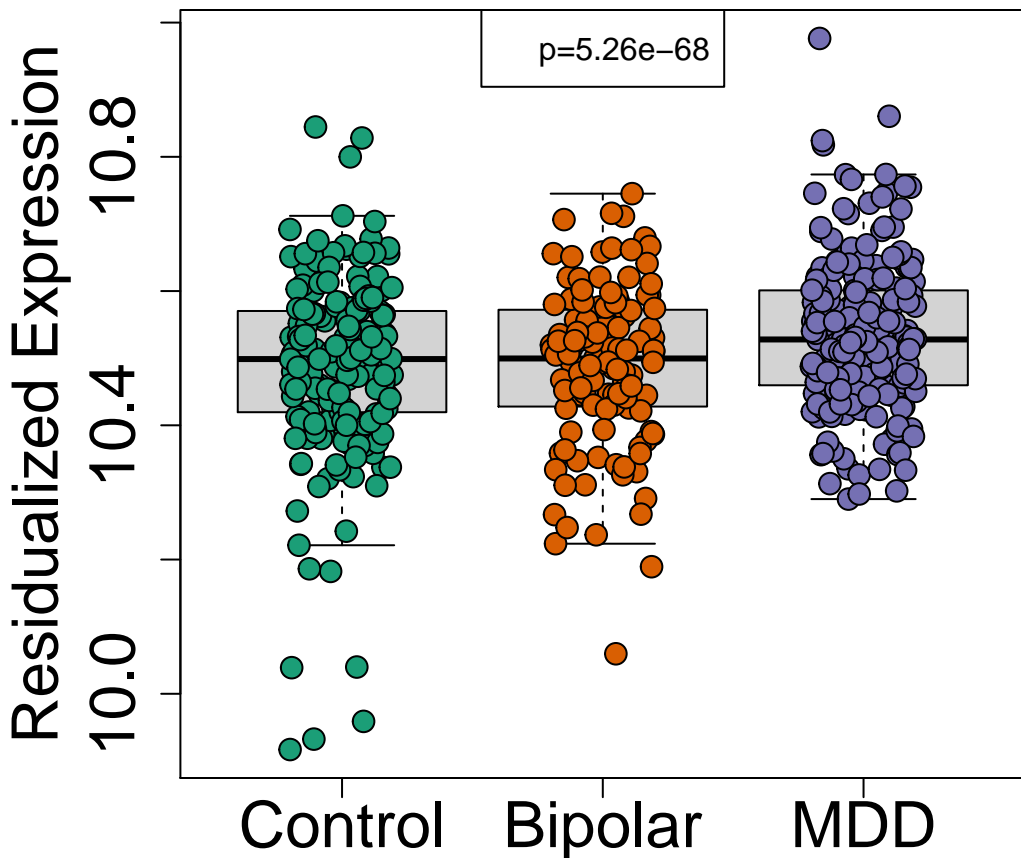
CALM3, InGen

chr19:46605858–46608196(+)



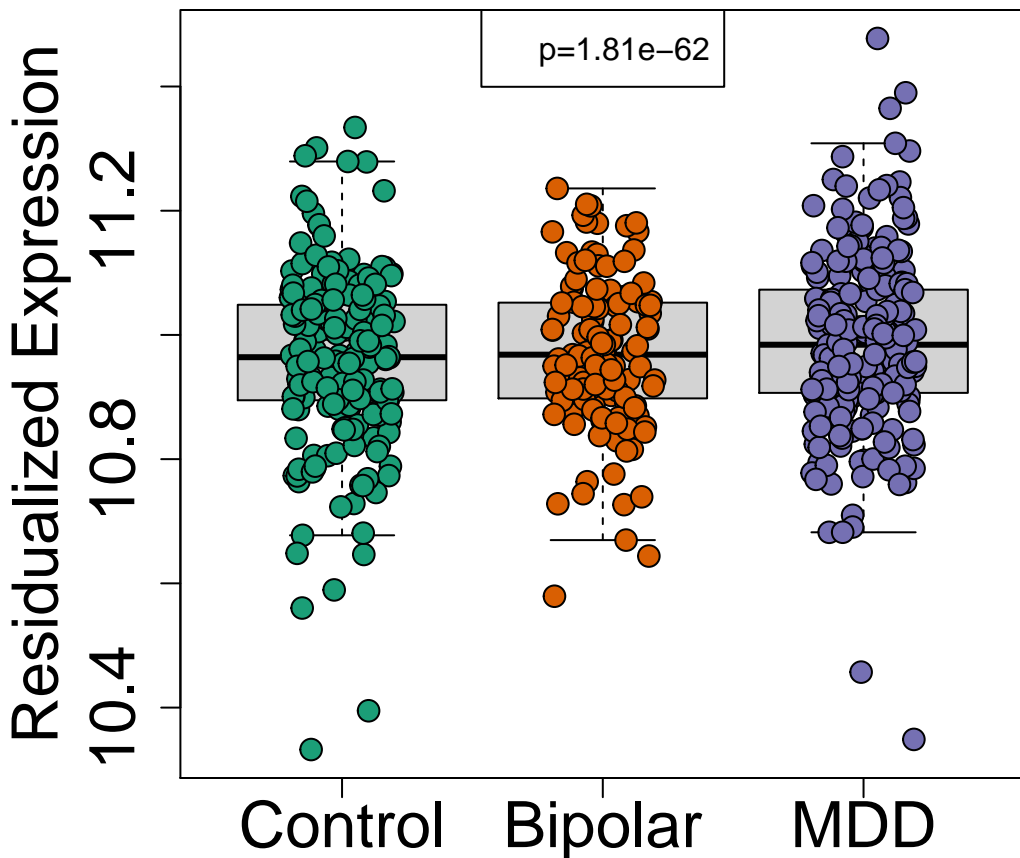
CALM3, InGen

chr19:46601438–46605826(+)



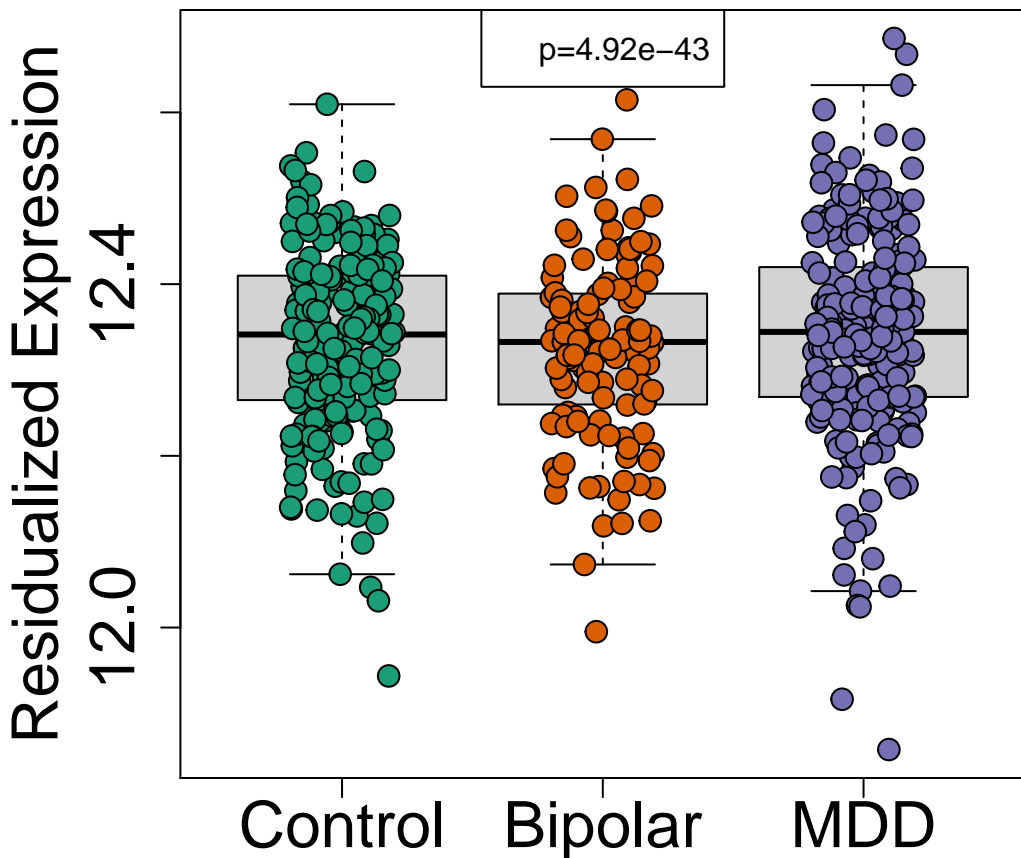
CALM3, InGen

chr19:46608982–46609124(+)



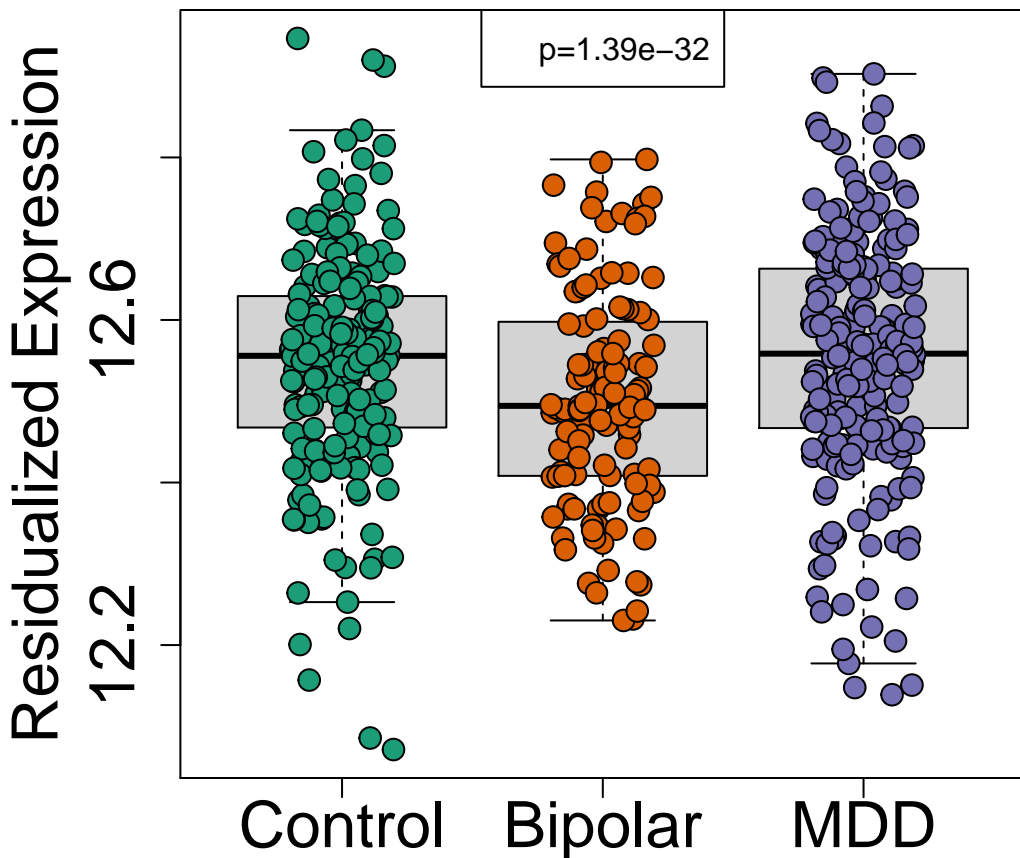
CALM2, InGen

chr2:47162663–47170733(–)



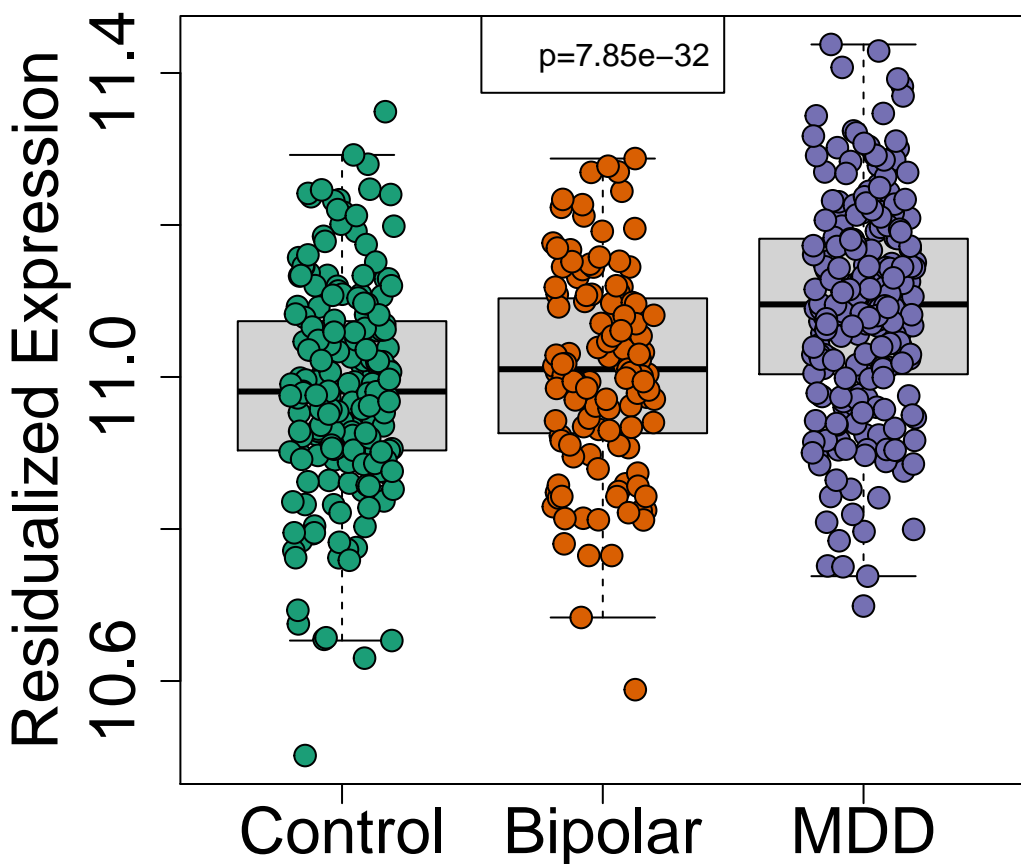
TMSB4X, InGen

chrX:12975169–12976245(+)



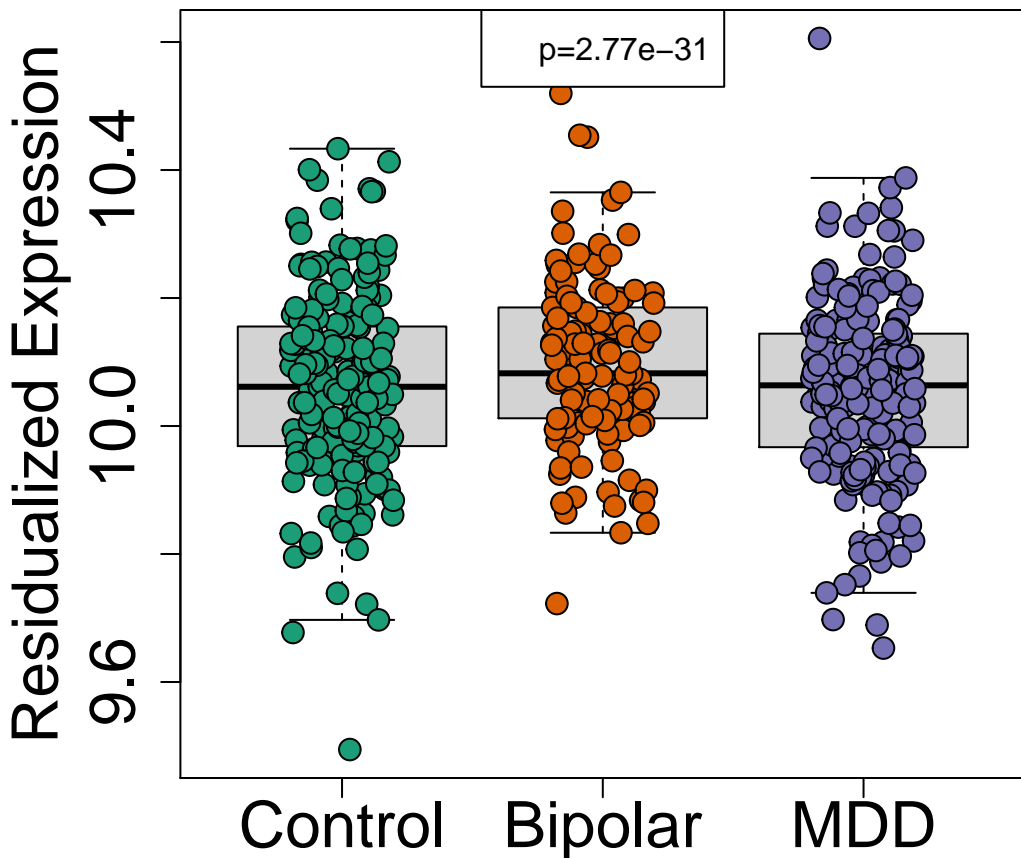
TMSB10, InGen

chr2:84906118–84906388(+)



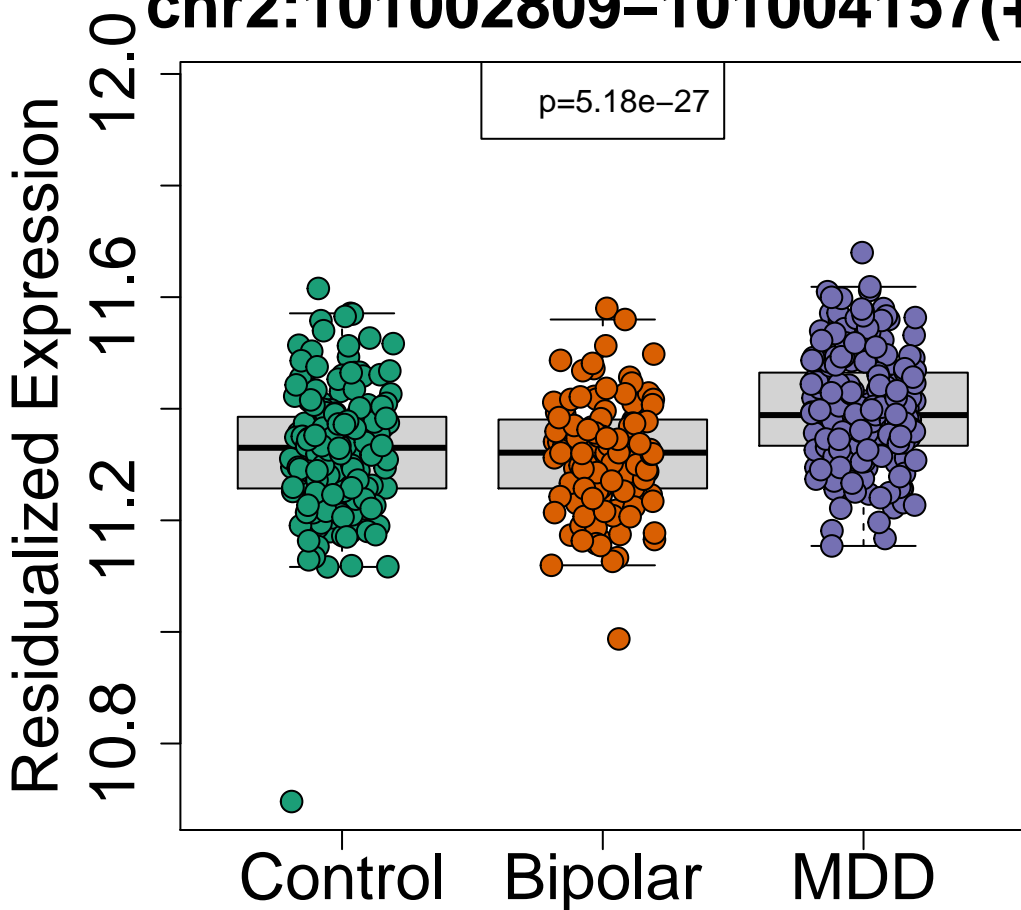
GNAS, InGen

chr20:58903792–58905382(+)



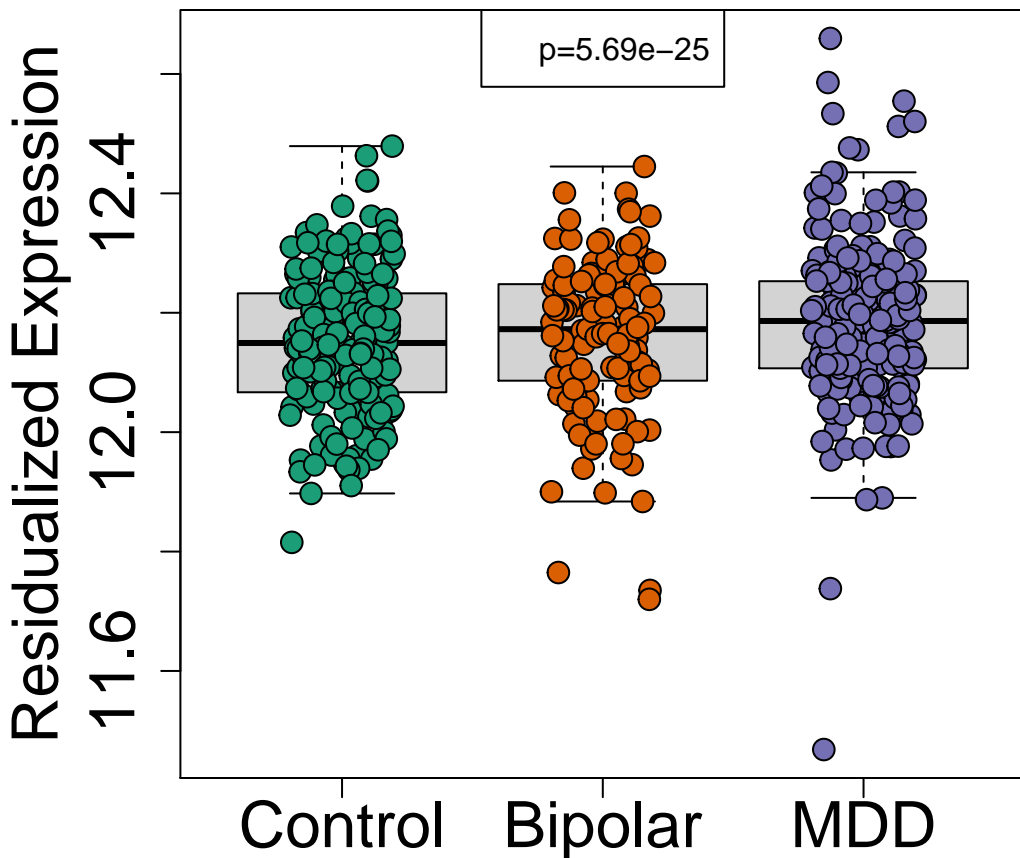
RPL31, InGen

chr2:101002809–101004157(+)



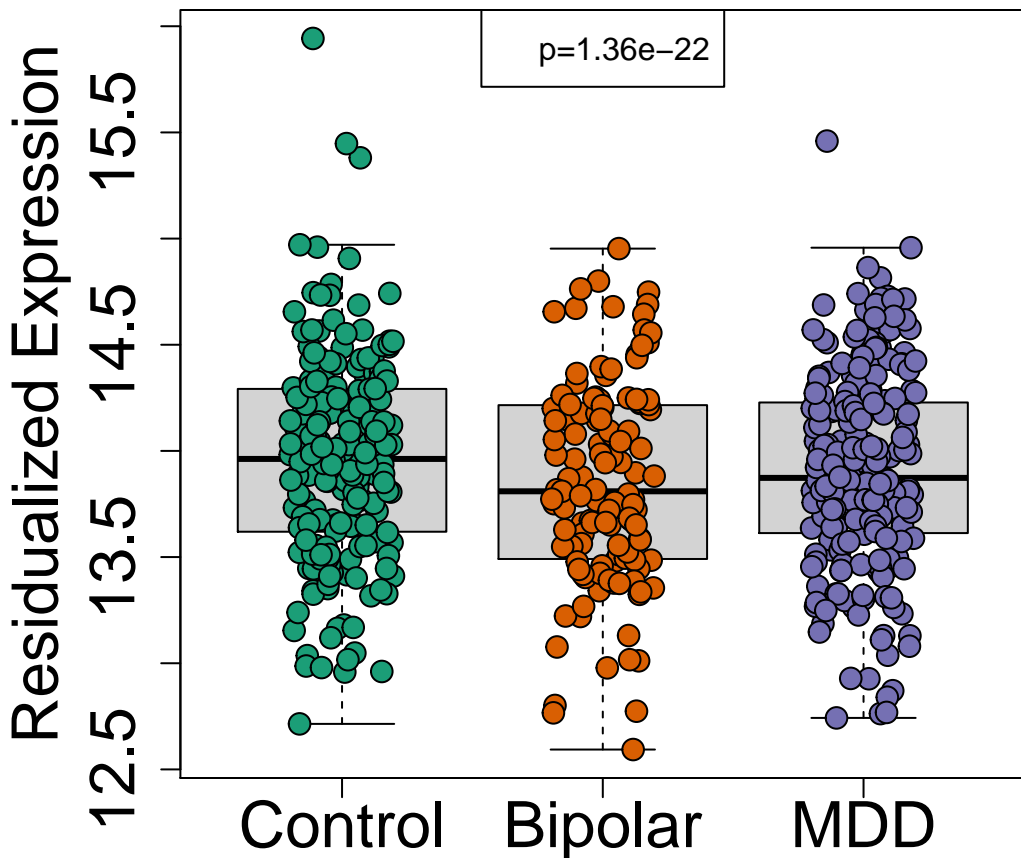
CALM2, InGen

chr2:47161859–47162285(-)



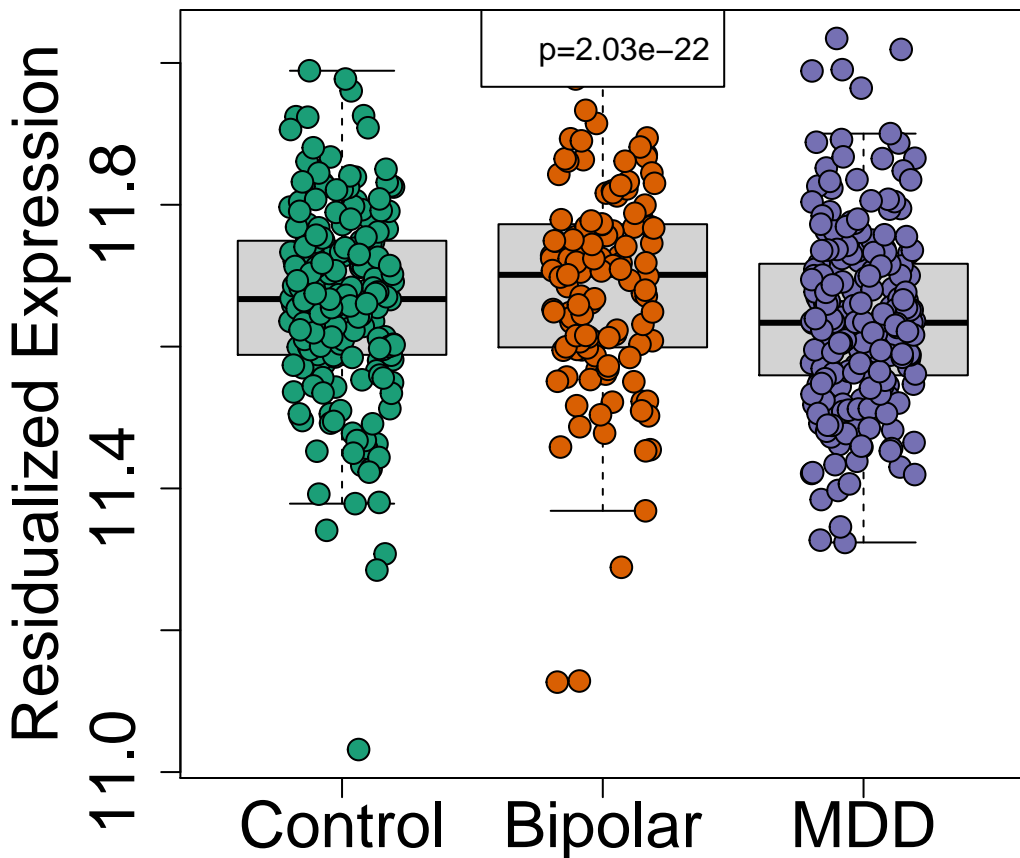
GFAP, InGen

chr17:44907389–44908063(-)



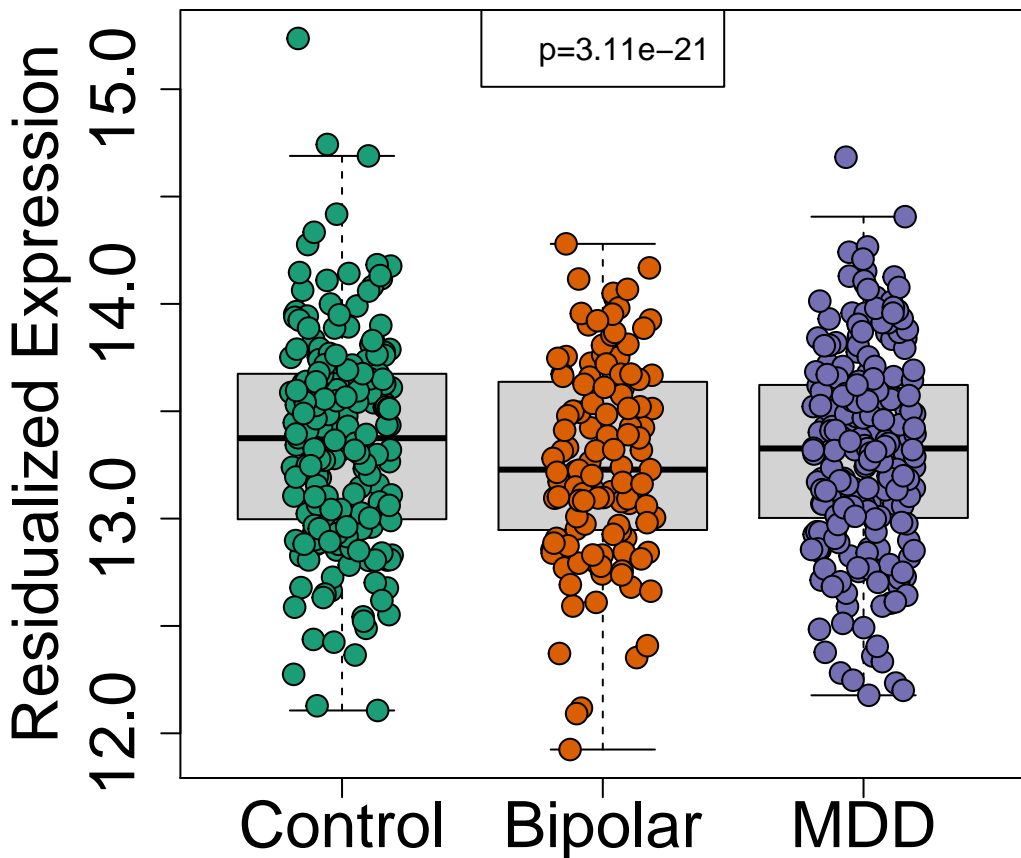
CALM1, InGen

chr14:90404515–90404688(+)



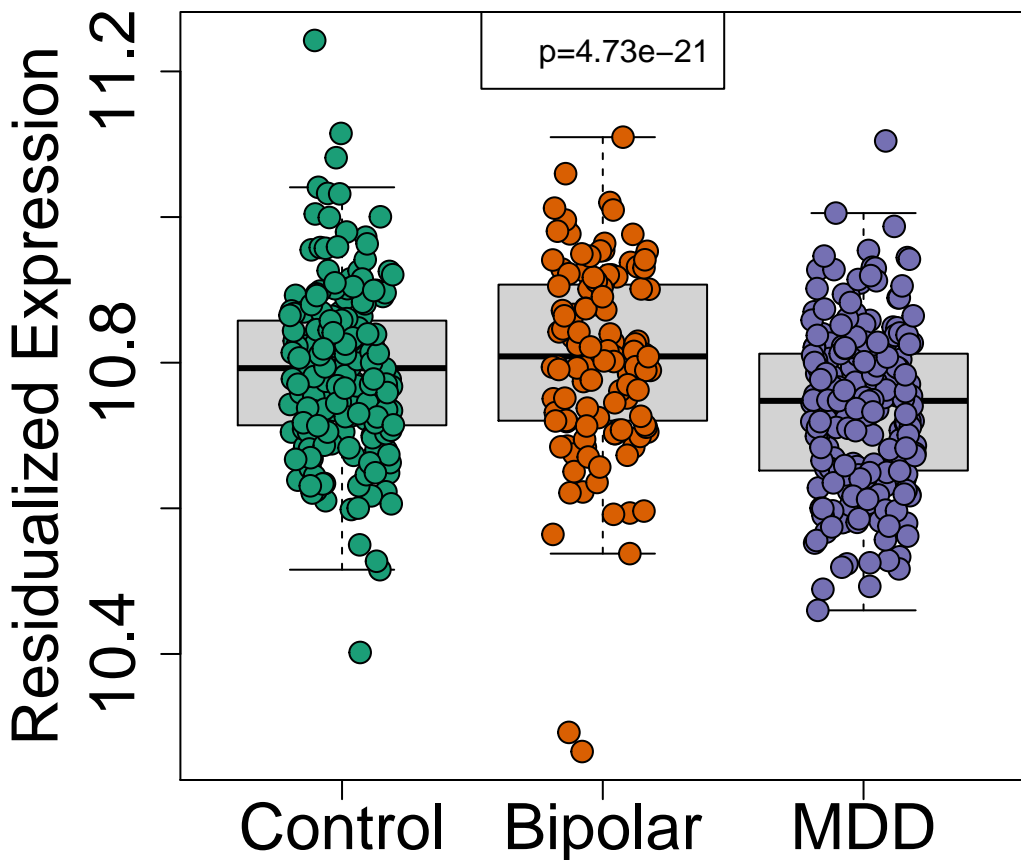
GFAP, InGen

chr17:44913431–44913727(-)



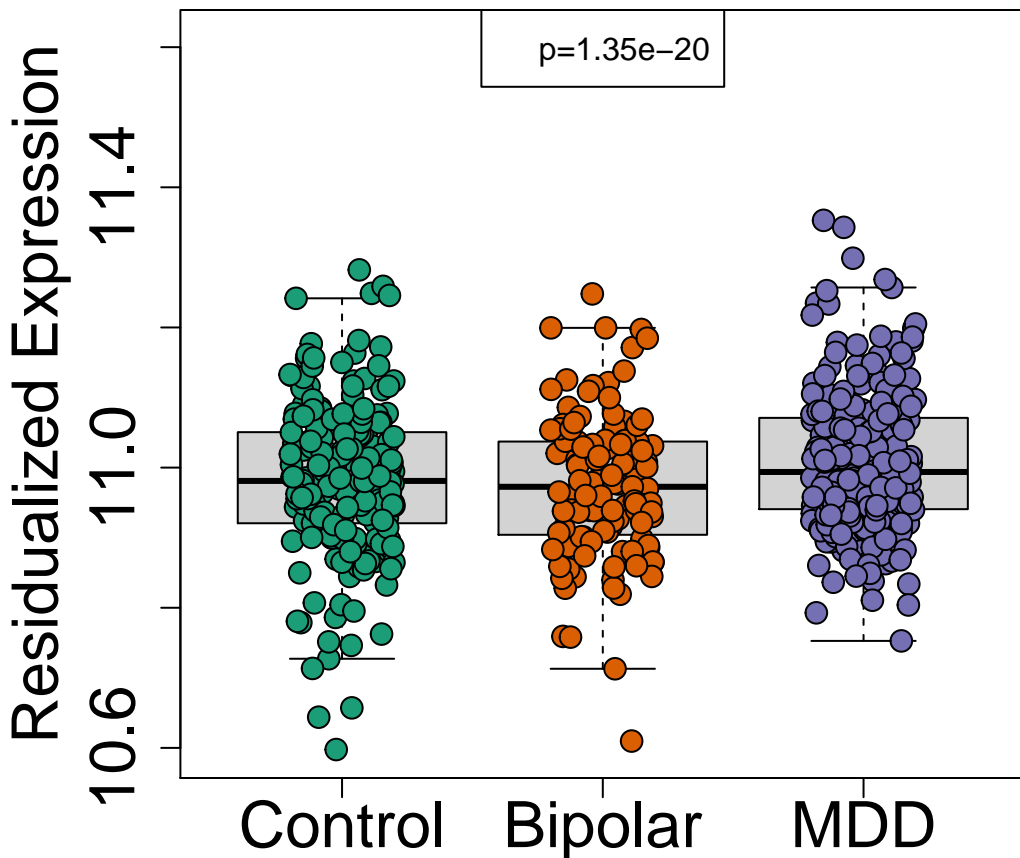
DPYSL2, InGen

chr8:26626679–26627214(+)



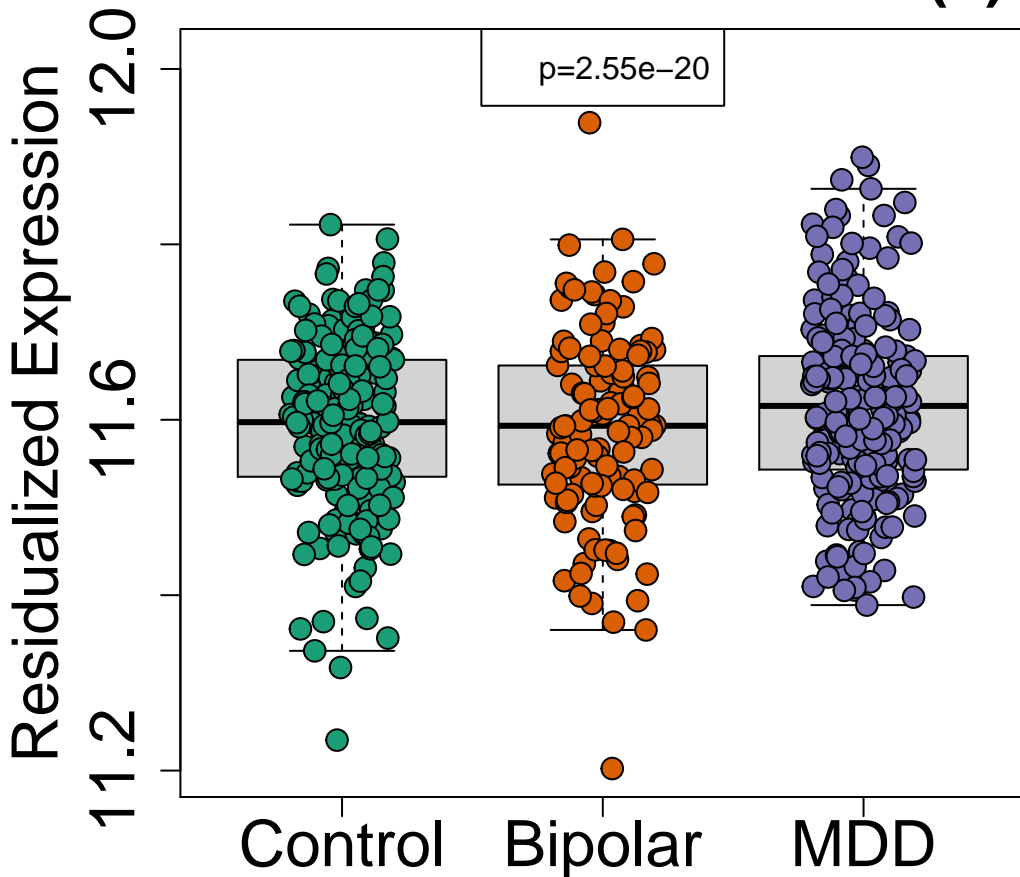
GAPDH, InGen

chr12:6537997–6538100(+)



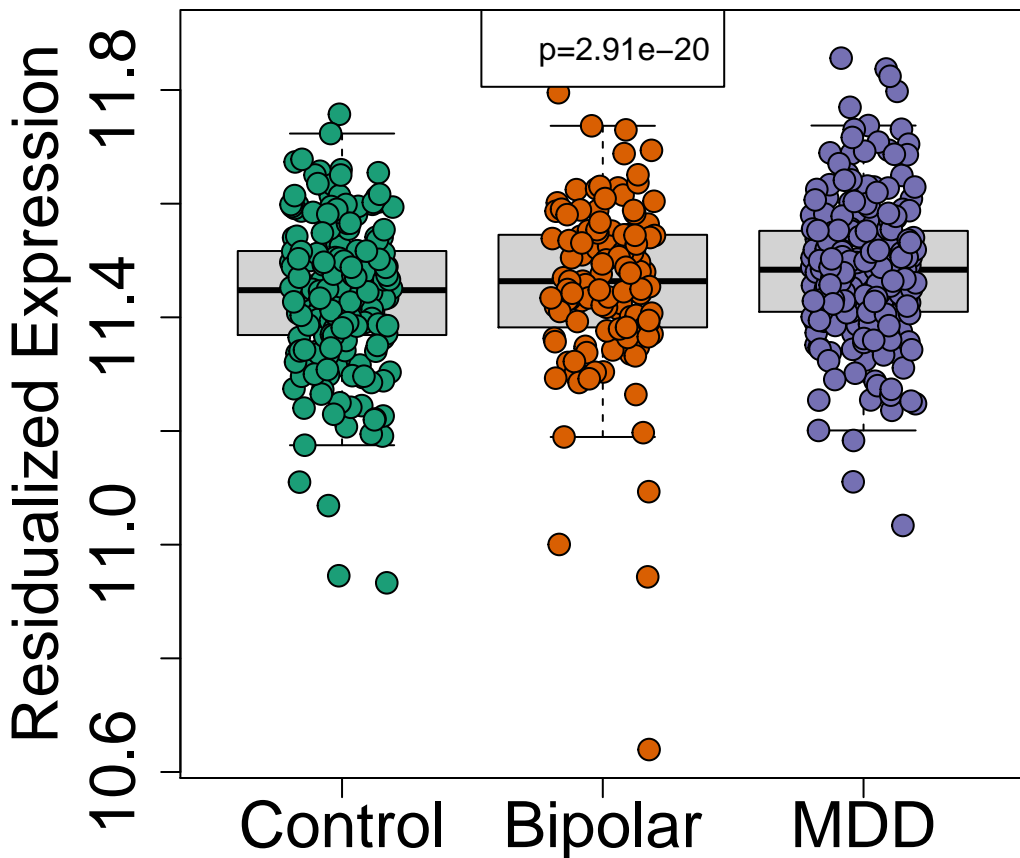
GAPDH, InGen

chr12:6537391–6537583(+)



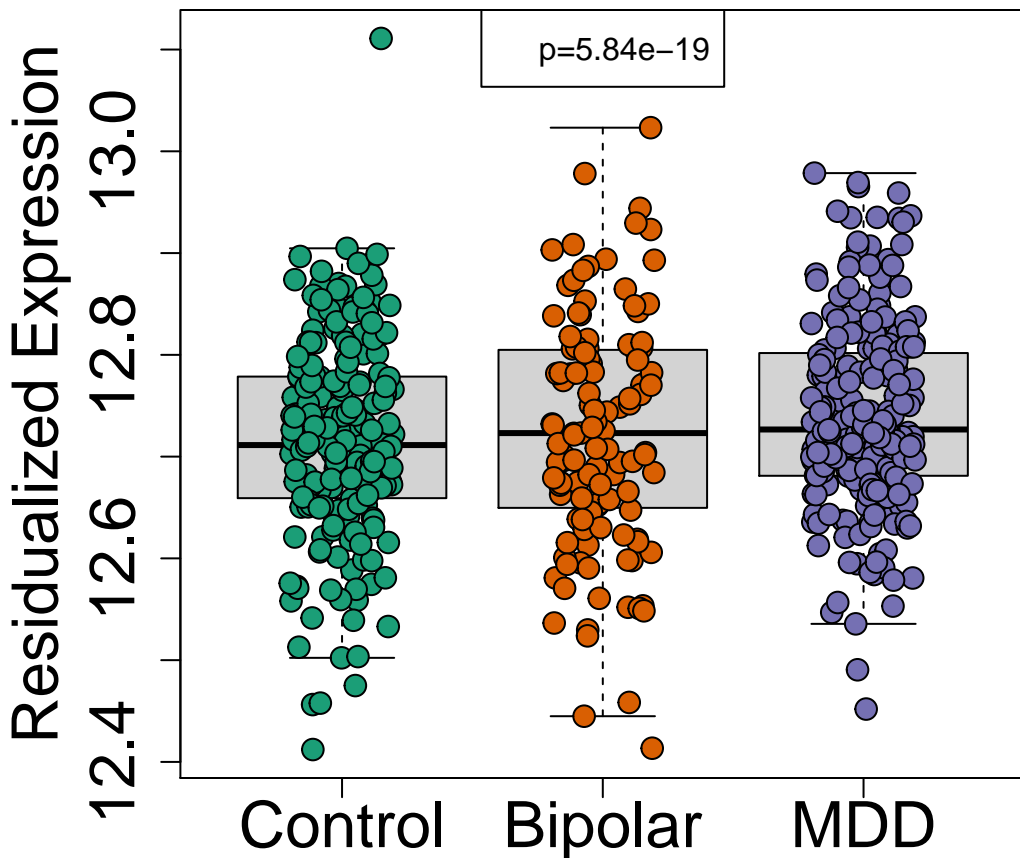
CALM1, InGen

chr14:90403969–90404378(+)



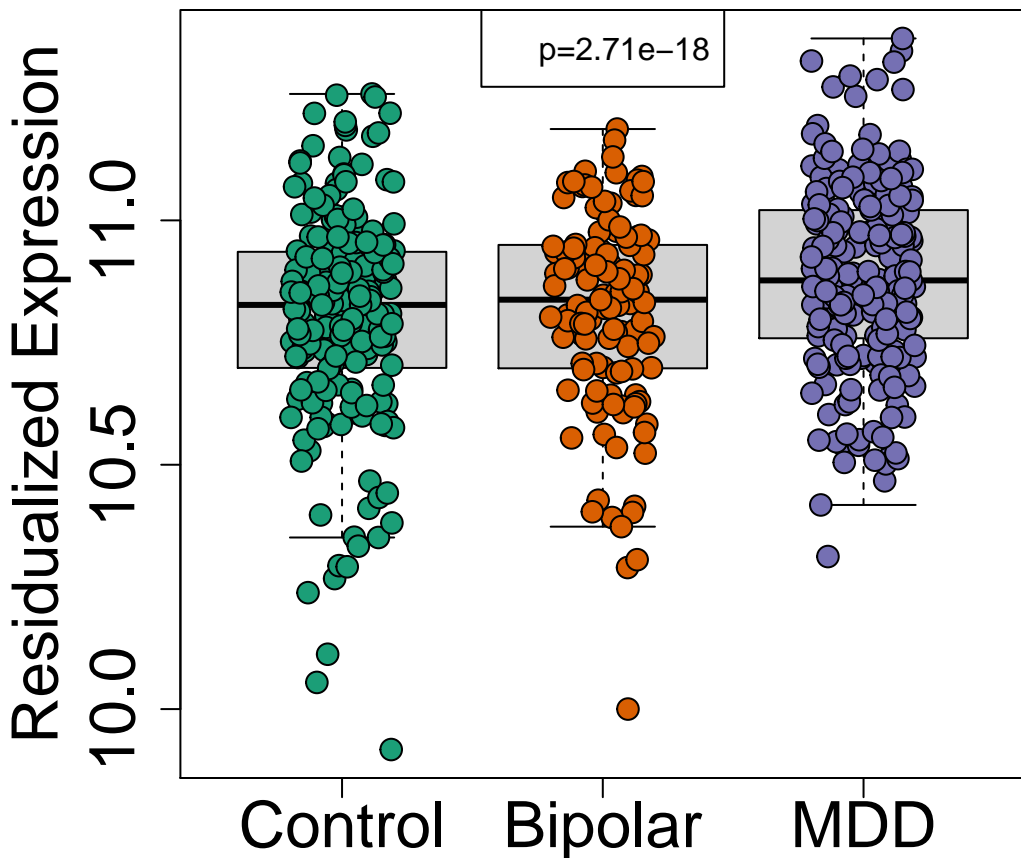
EEF1A1, InGen

chr6:73519229–73519336(–)



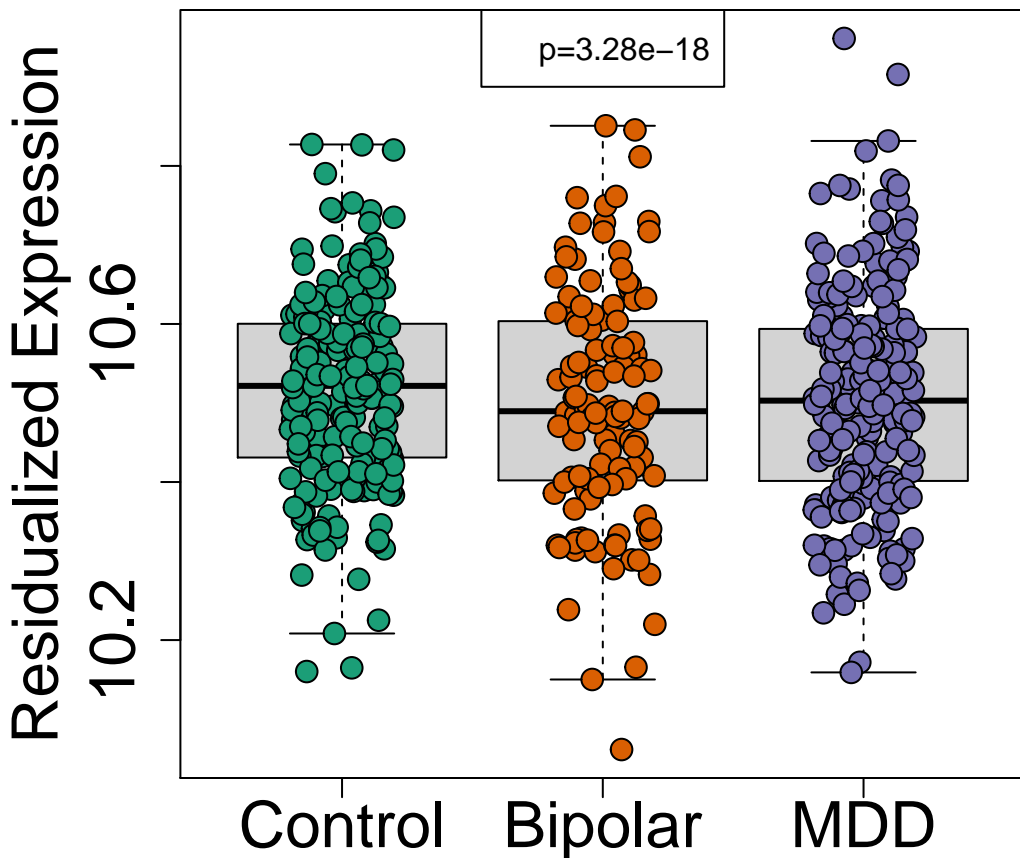
SPARCL1, InGen

chr4:87494599–87494980(–)



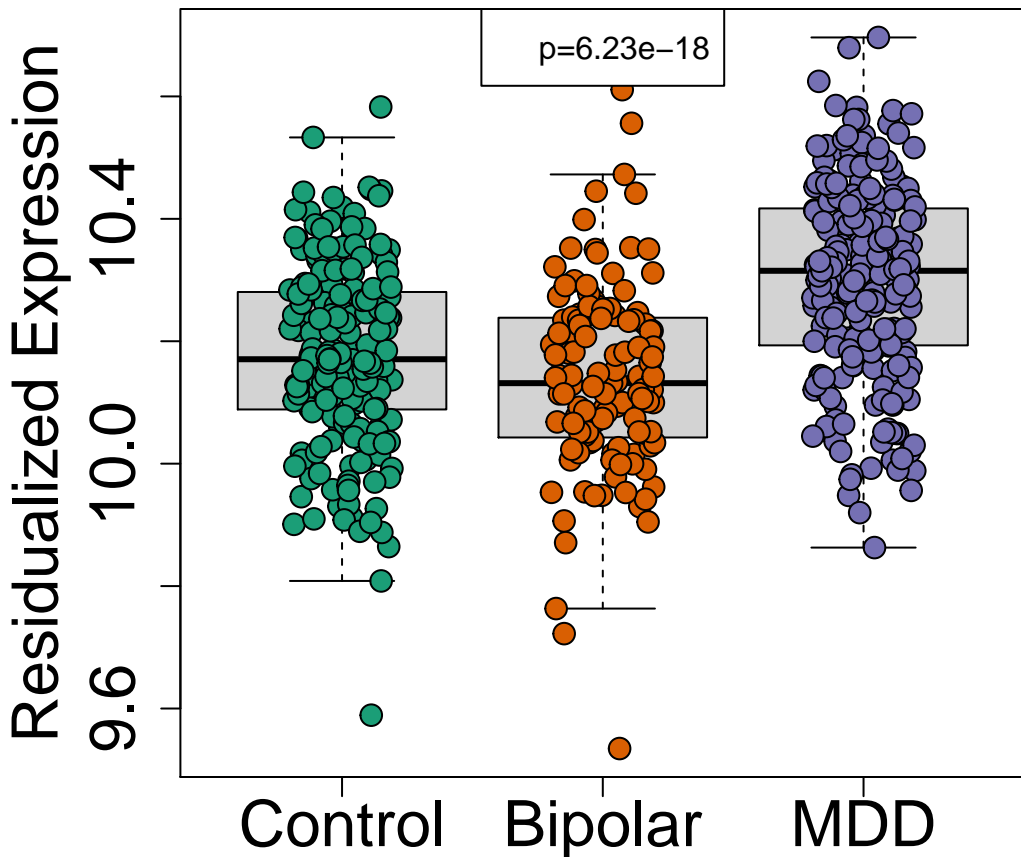
EIF4A2, InGen

chr3:186787265–186787494(+)



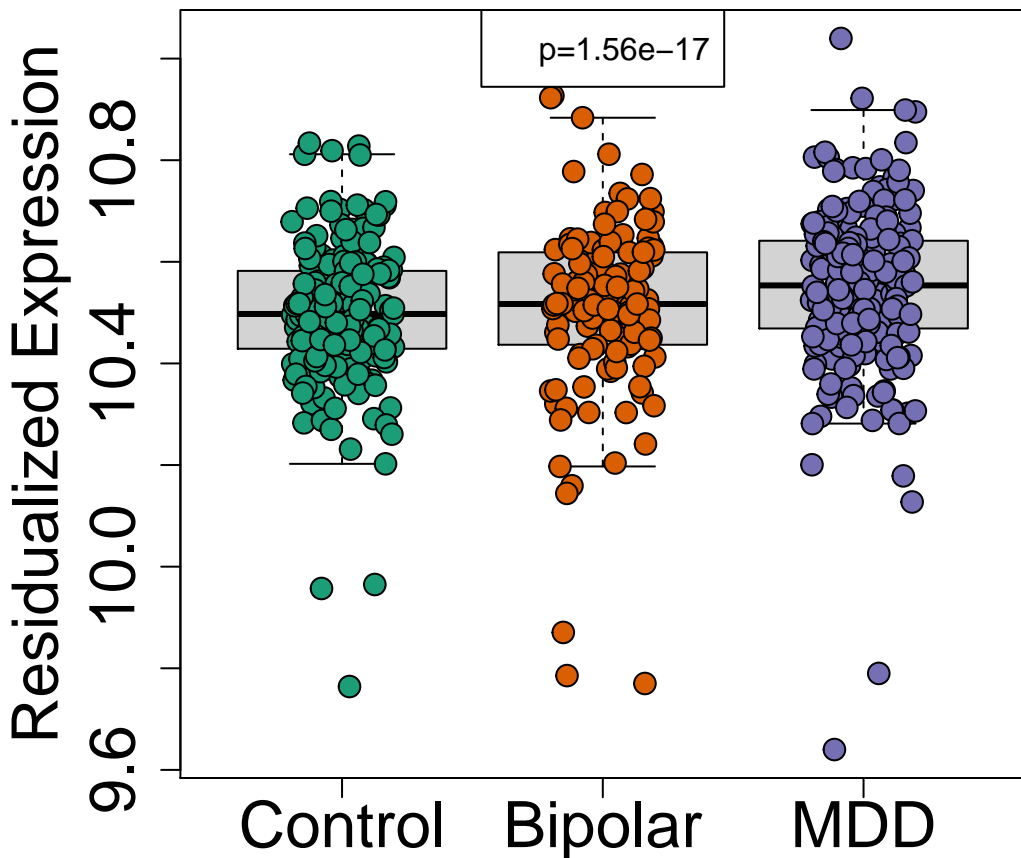
HNRNPA1, InGen

chr12:54280823–54281385(+)



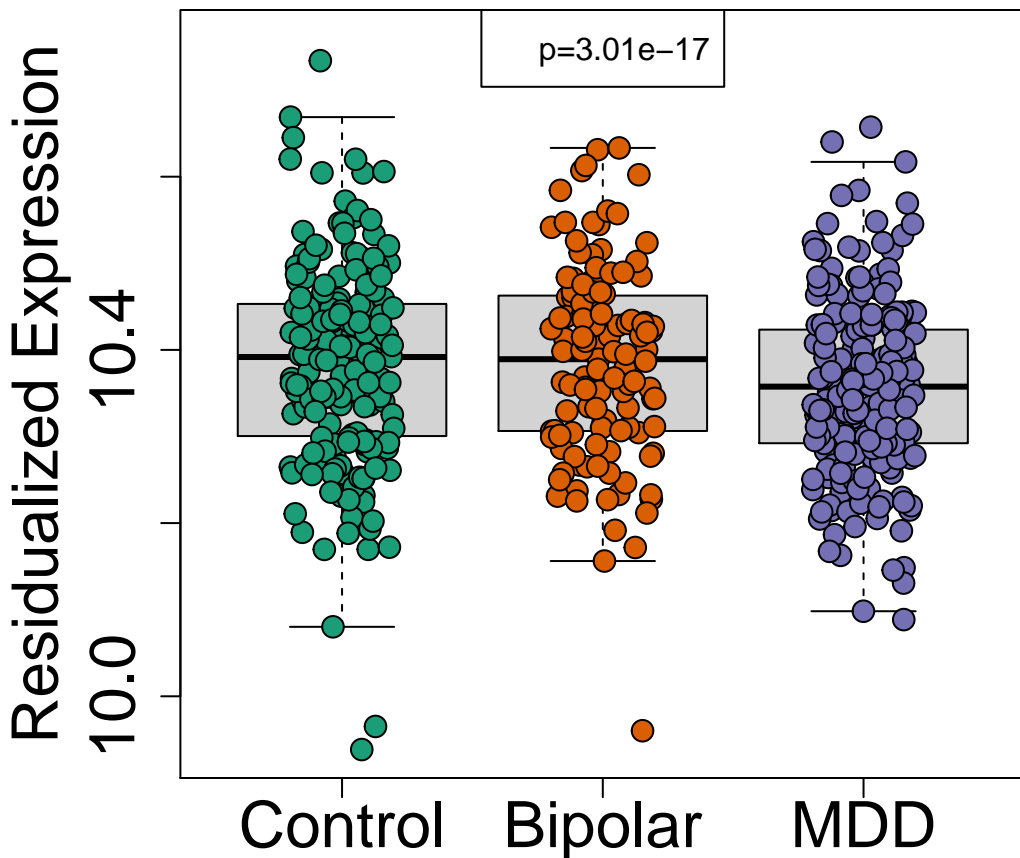
TUBA1B, InGen

chr12:49129391–49129499(-)



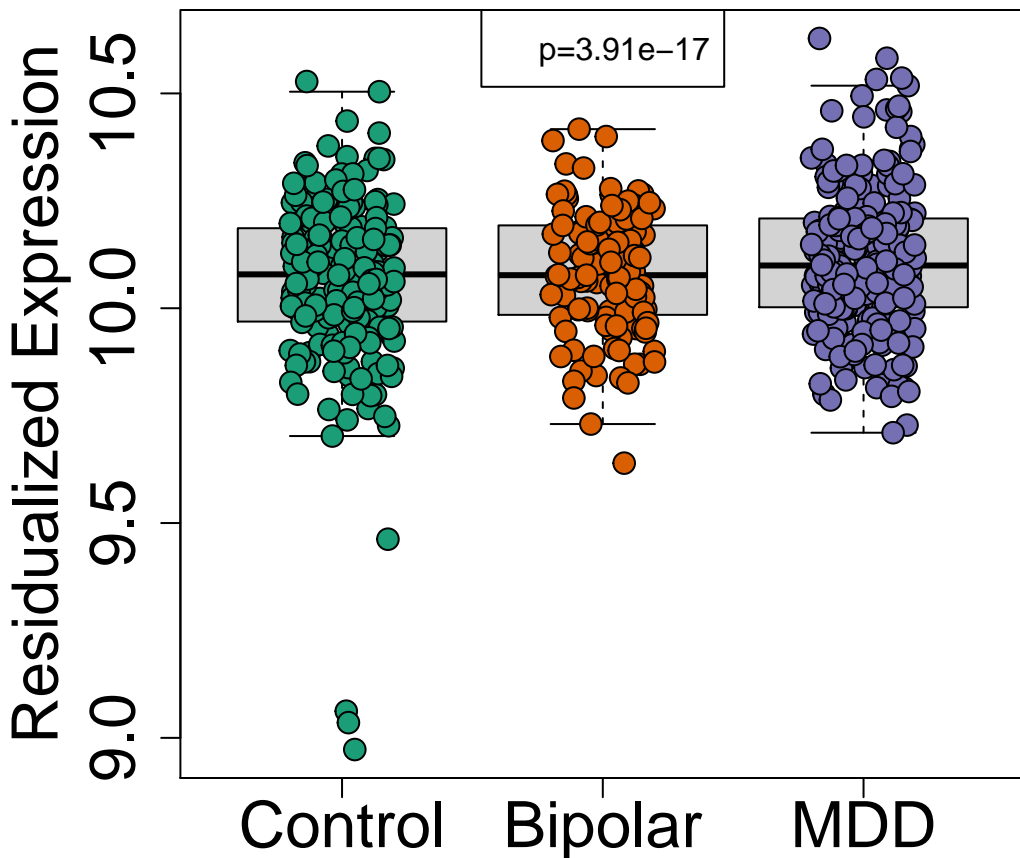
GNAS, InGen

chr20:58910082–58910333(+)



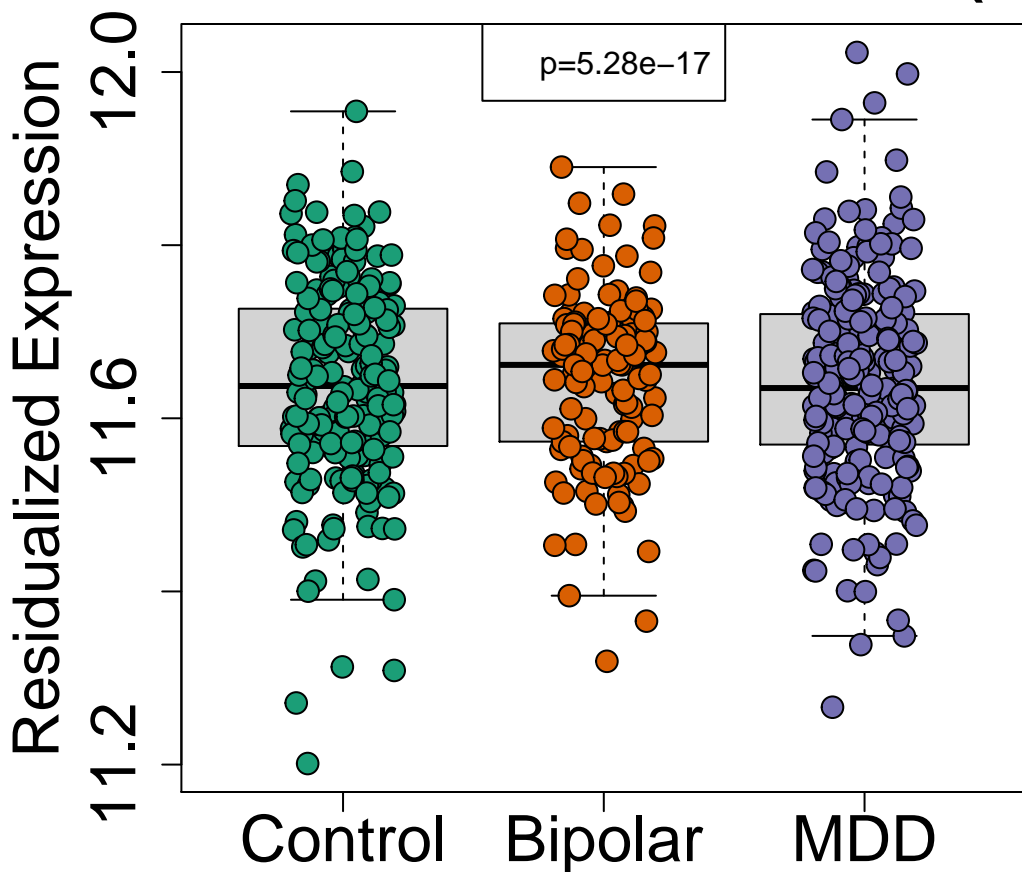
VSNL1, InGen

chr2:17540919–17592069(+)



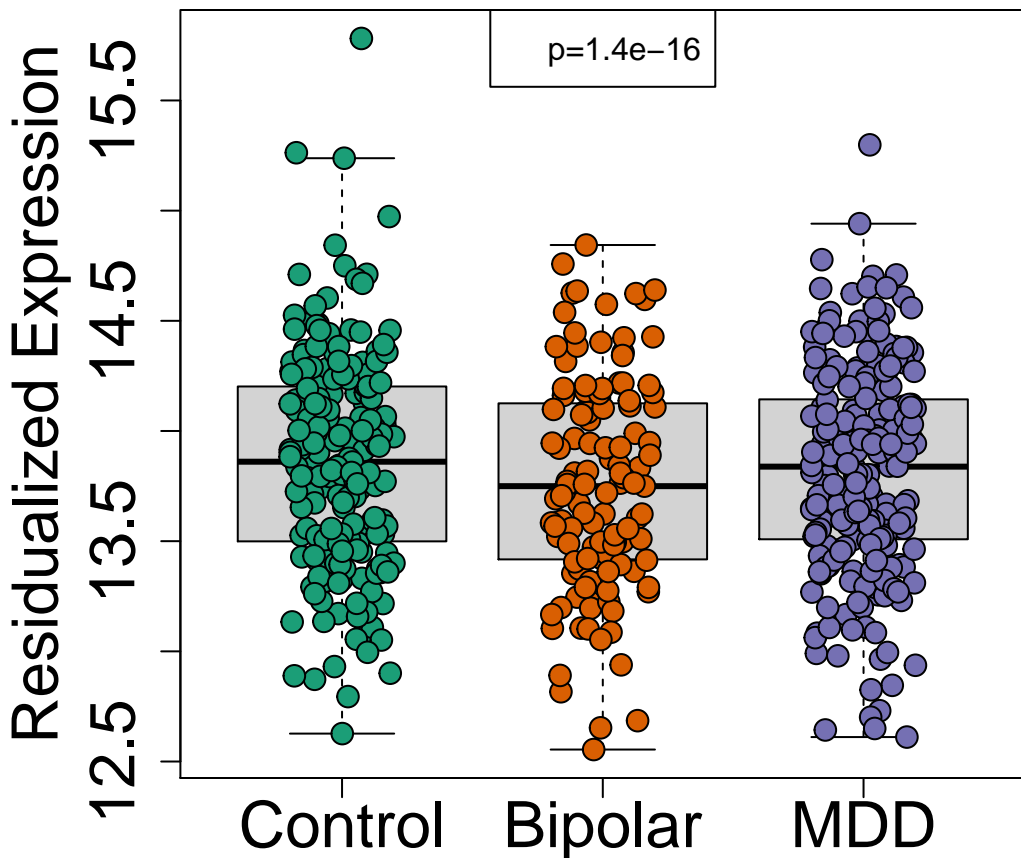
RPL38, InGen

chr17:74209310–74209803(+)



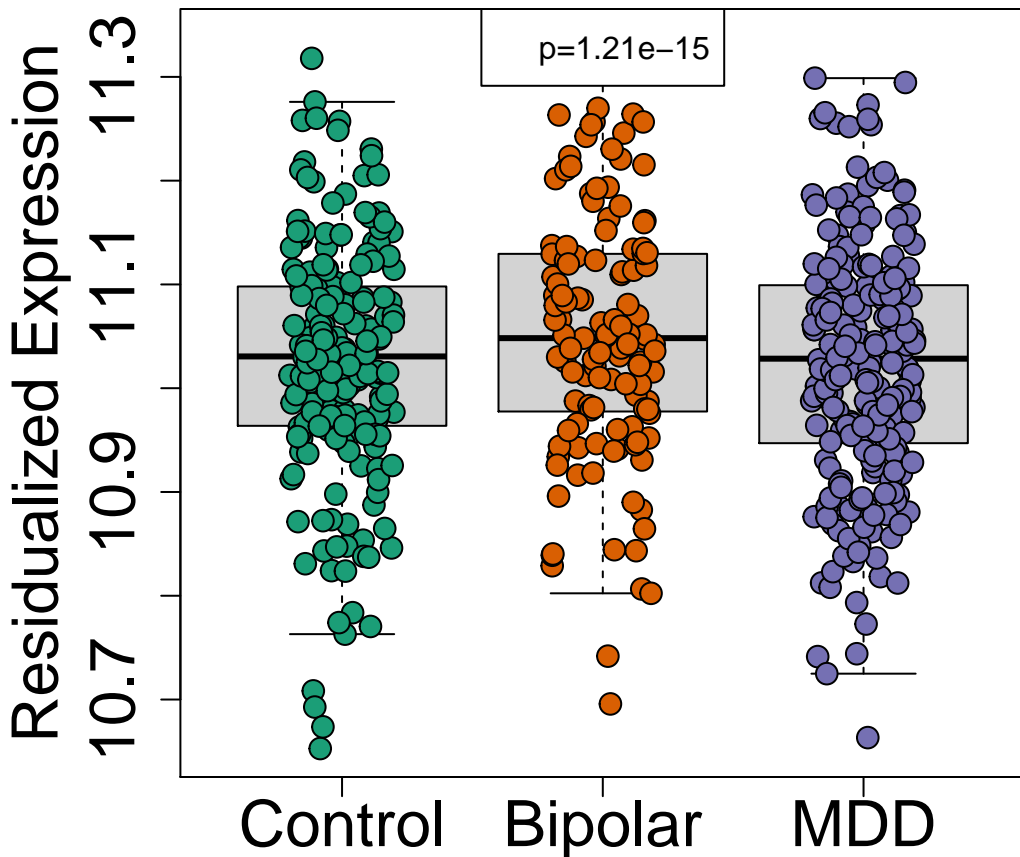
GFAP, InGen

chr17:44910659–44911235(-)



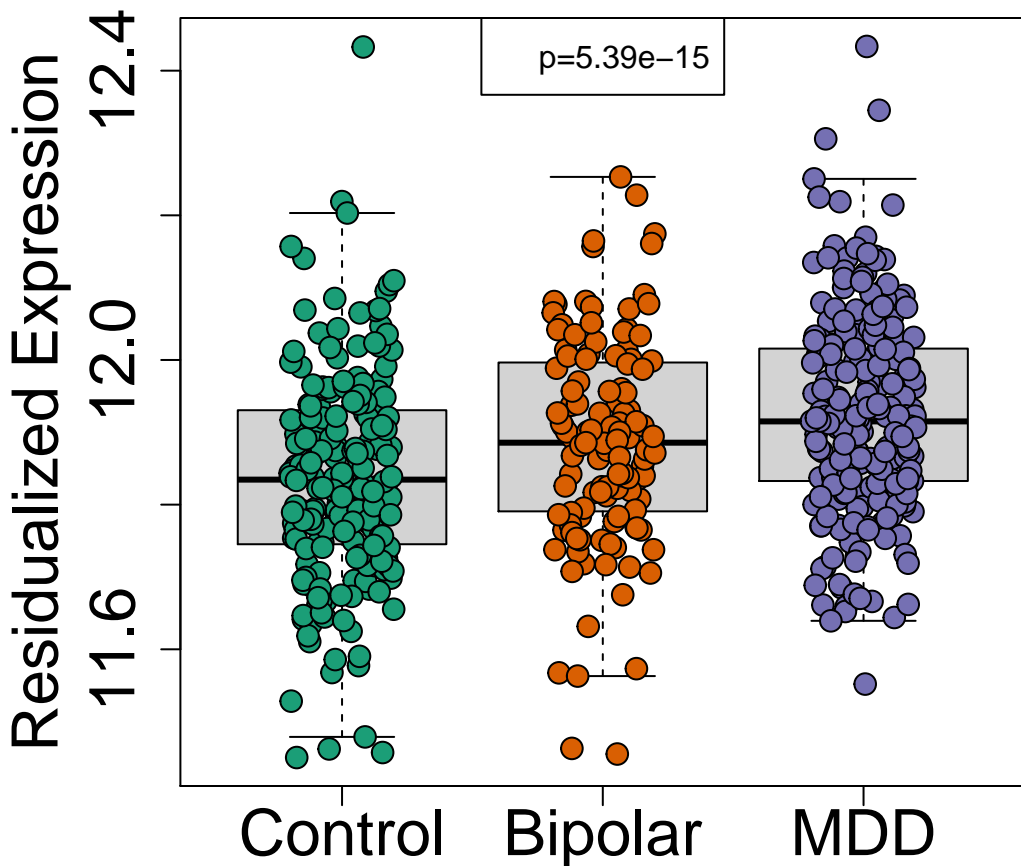
RPL37A, InGen

chr2:216499399–216499948(+)



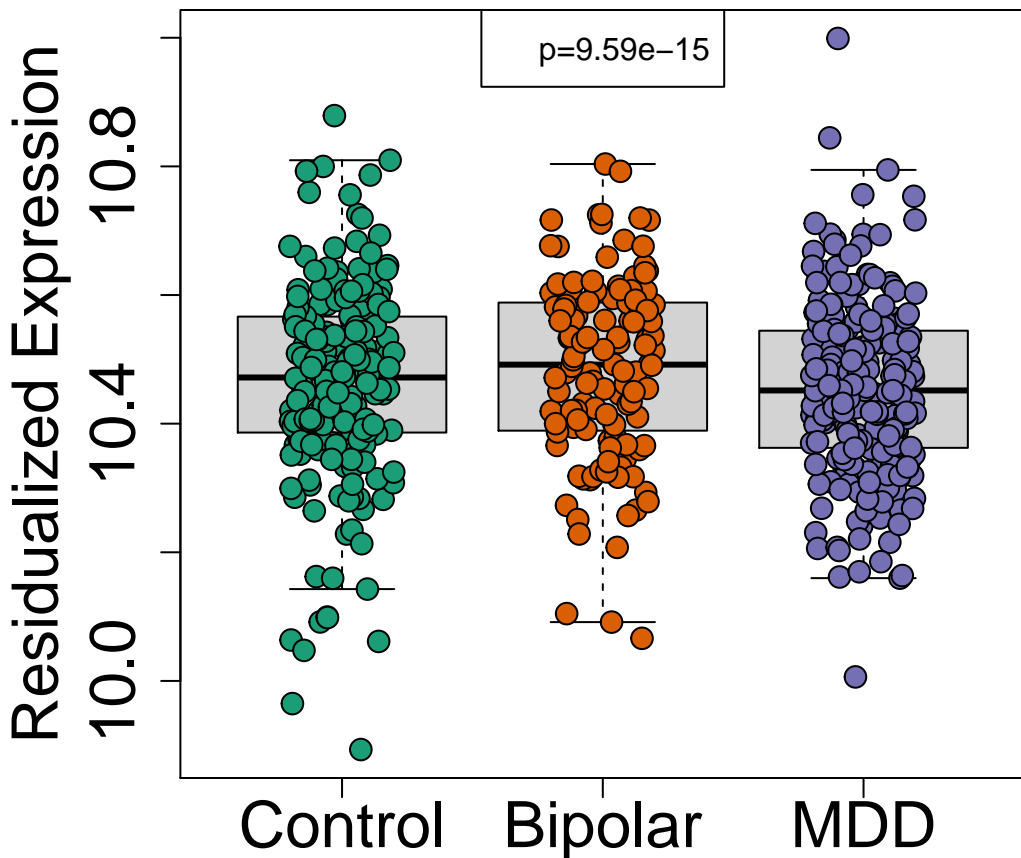
CLU, InGen

chr8:27606525–27608937(-)



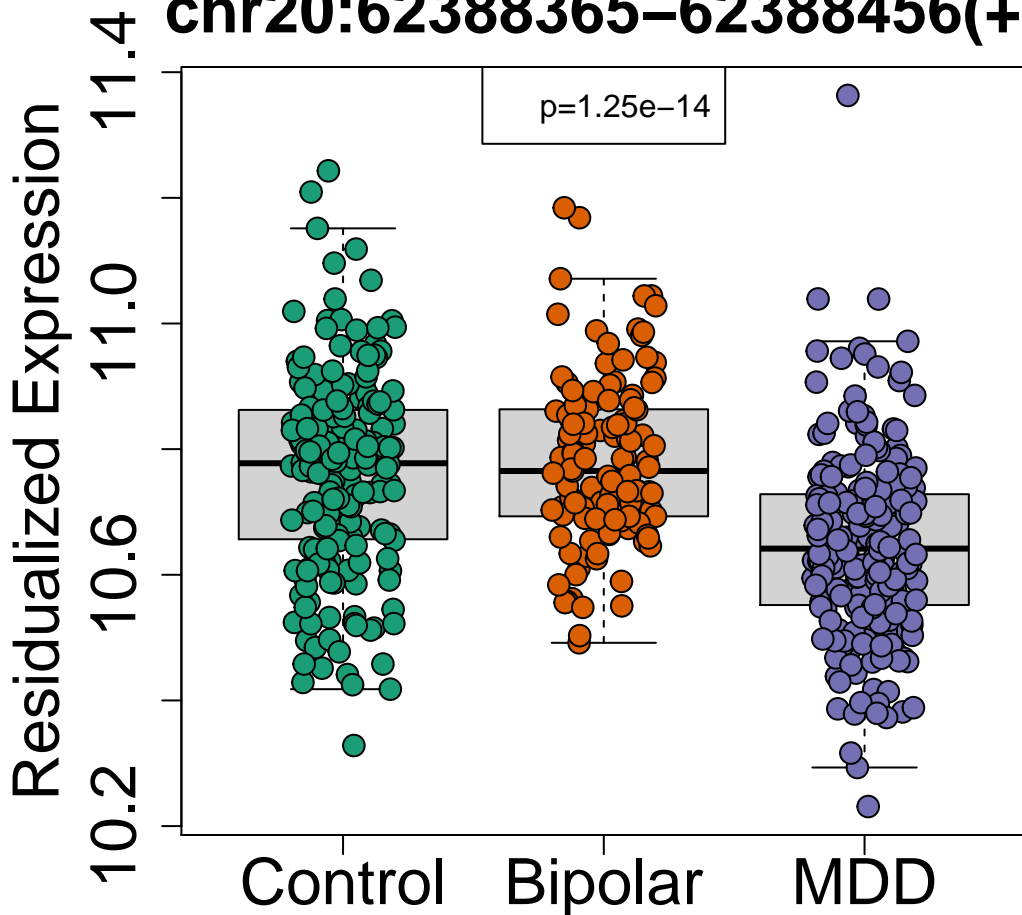
RTN3, InGen

chr11:63756171–63758155(+)



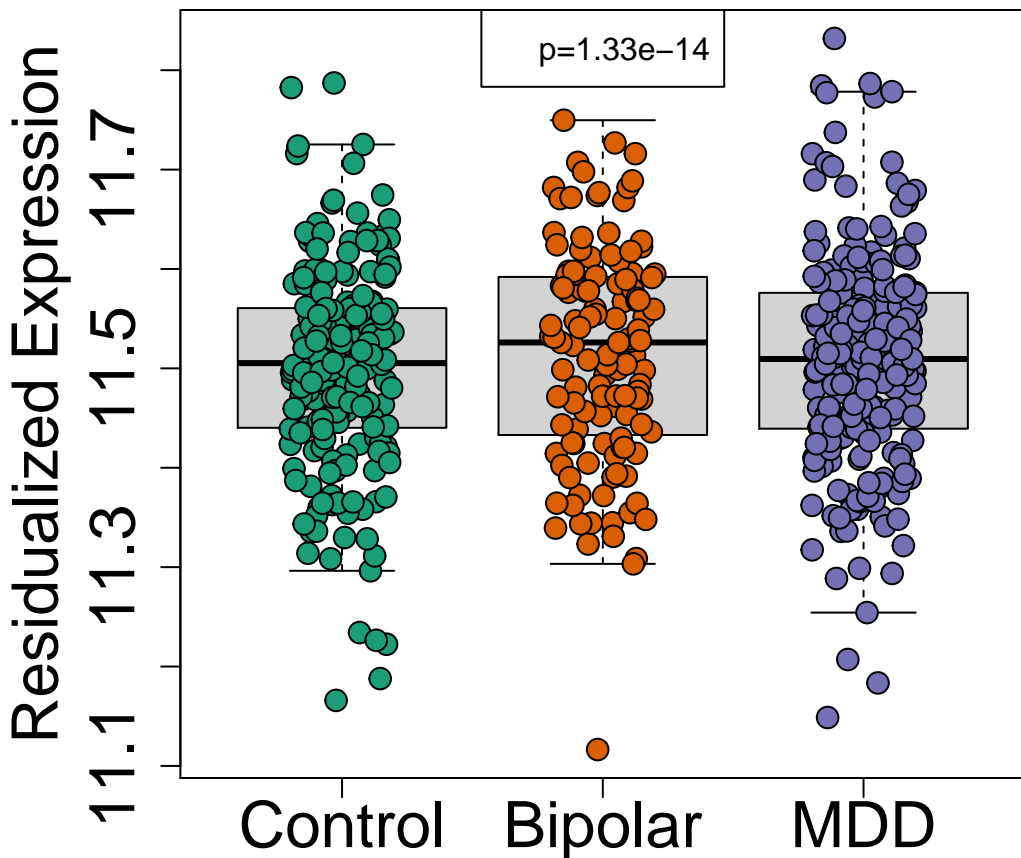
RPS21, InGen

chr20:62388365–62388456(+)



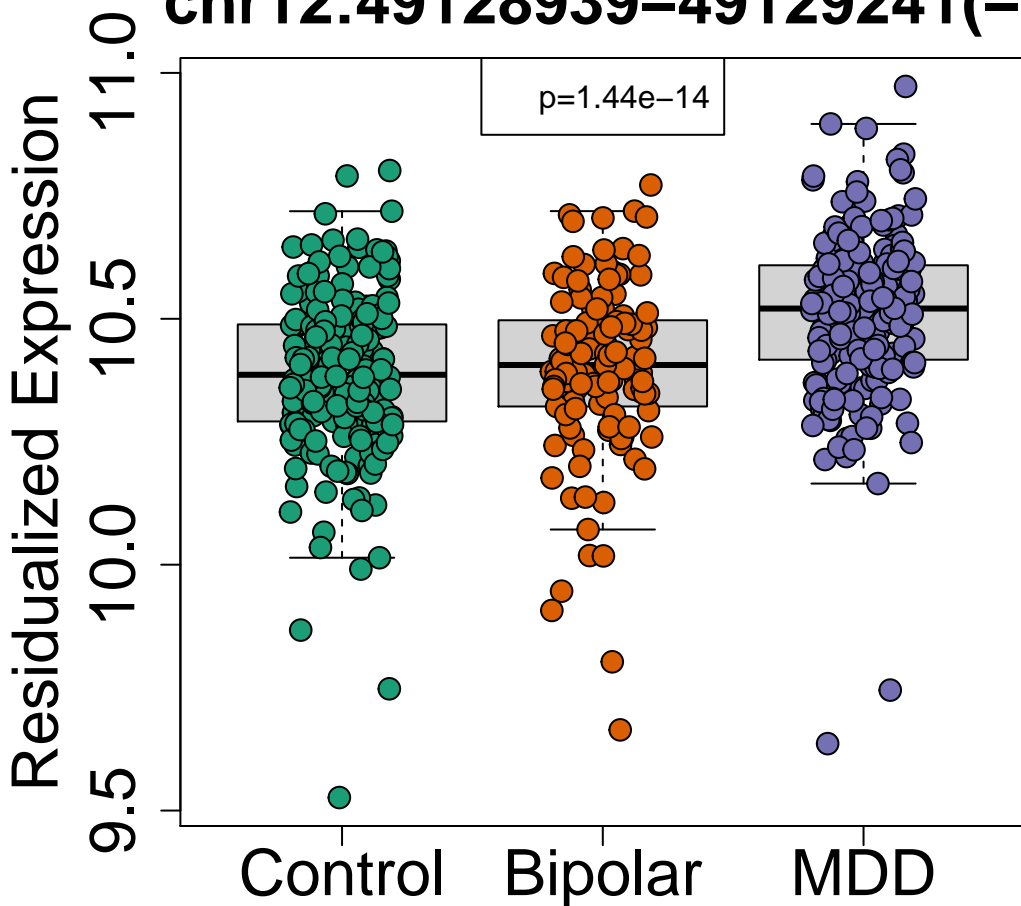
PPIA, InGen

chr7:44799875–44801286(+)



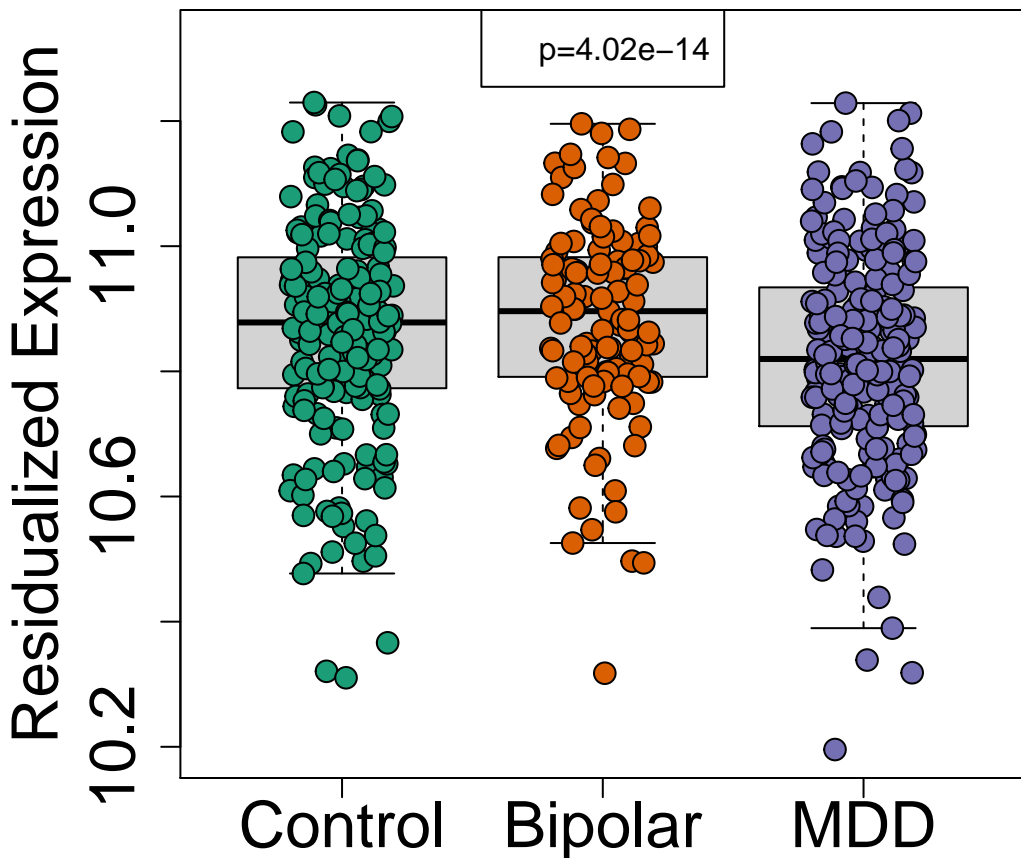
TUBA1B, InGen

chr12:49128939–49129241(-)



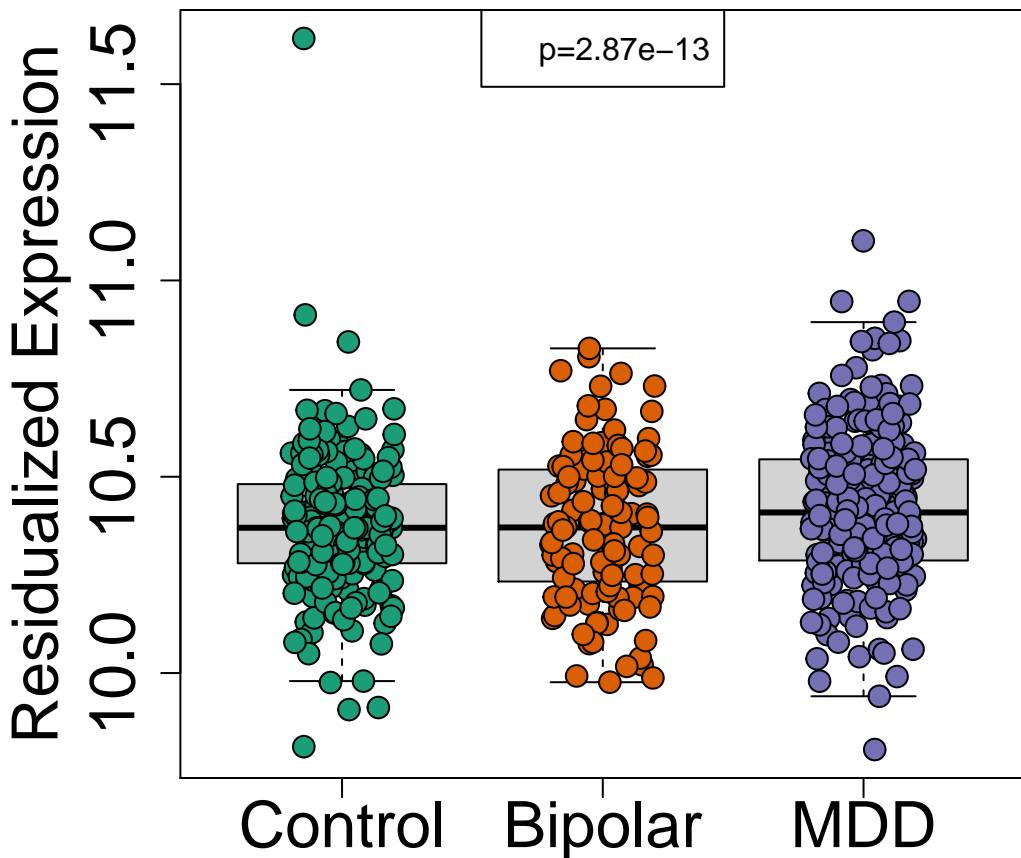
SNAP25, InGen

chr20:10297051–10299267(+)



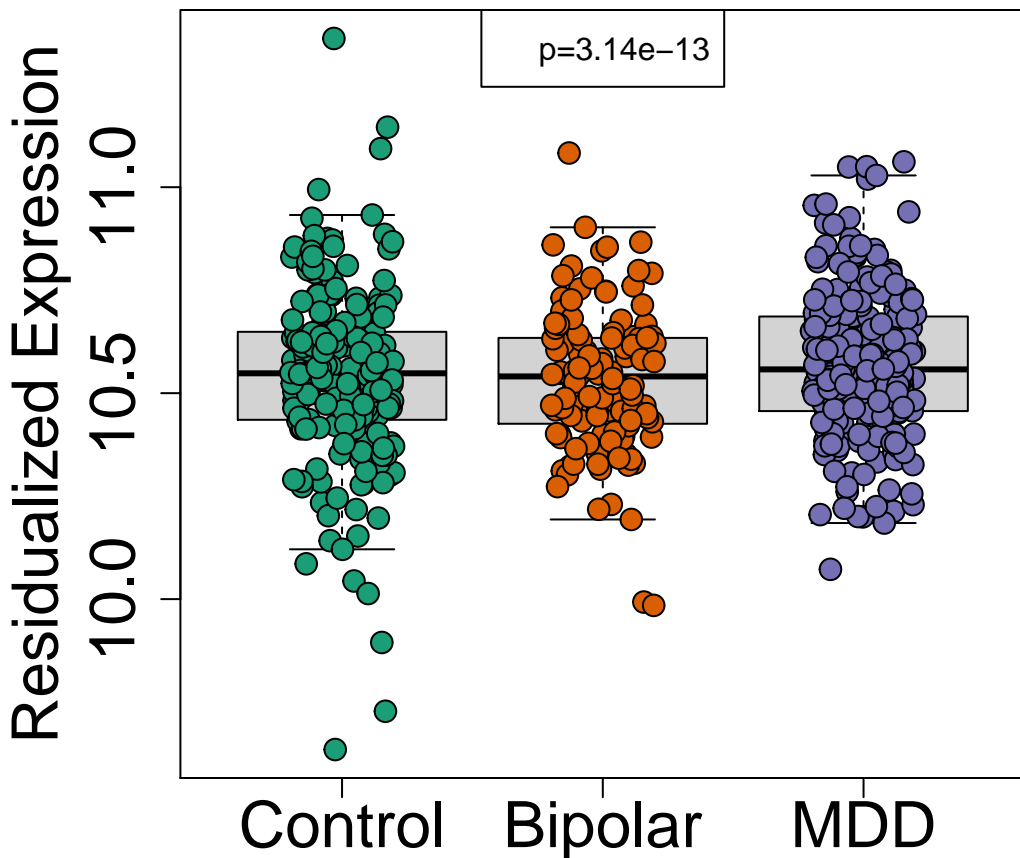
ARPP19, InGen

chr15:52552105–52557099(-)



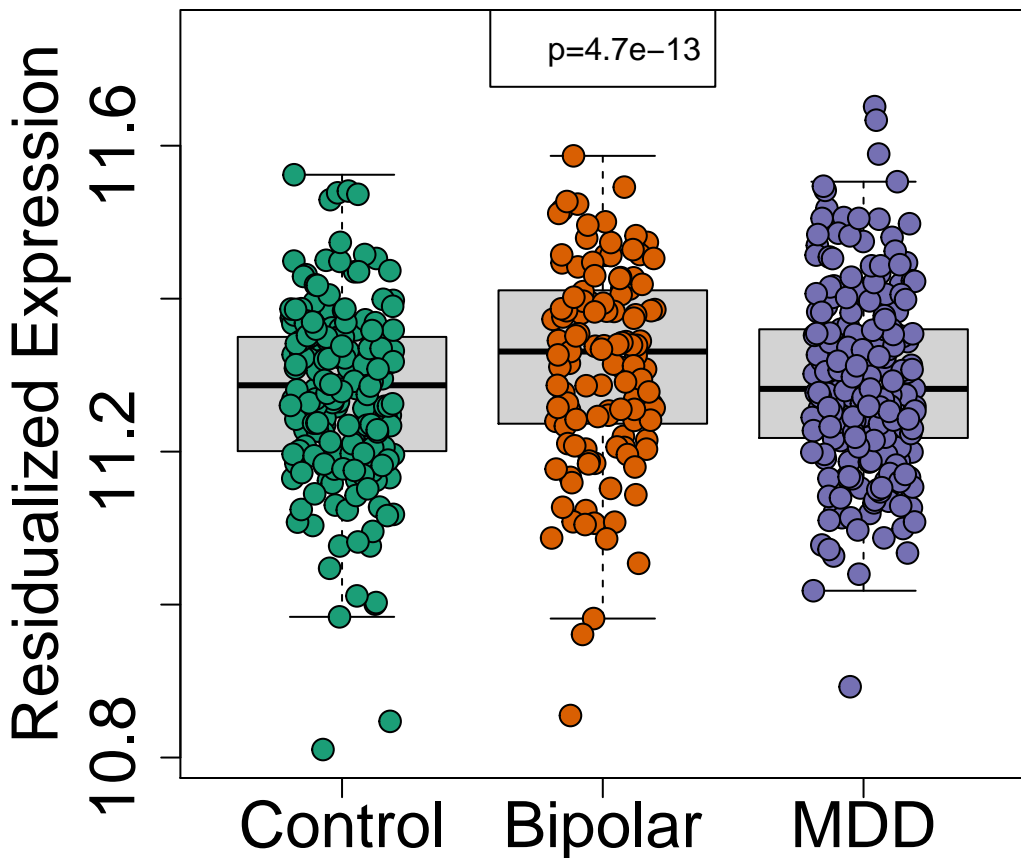
SNAP25, InGen

chr20:10293279–10296924(+)



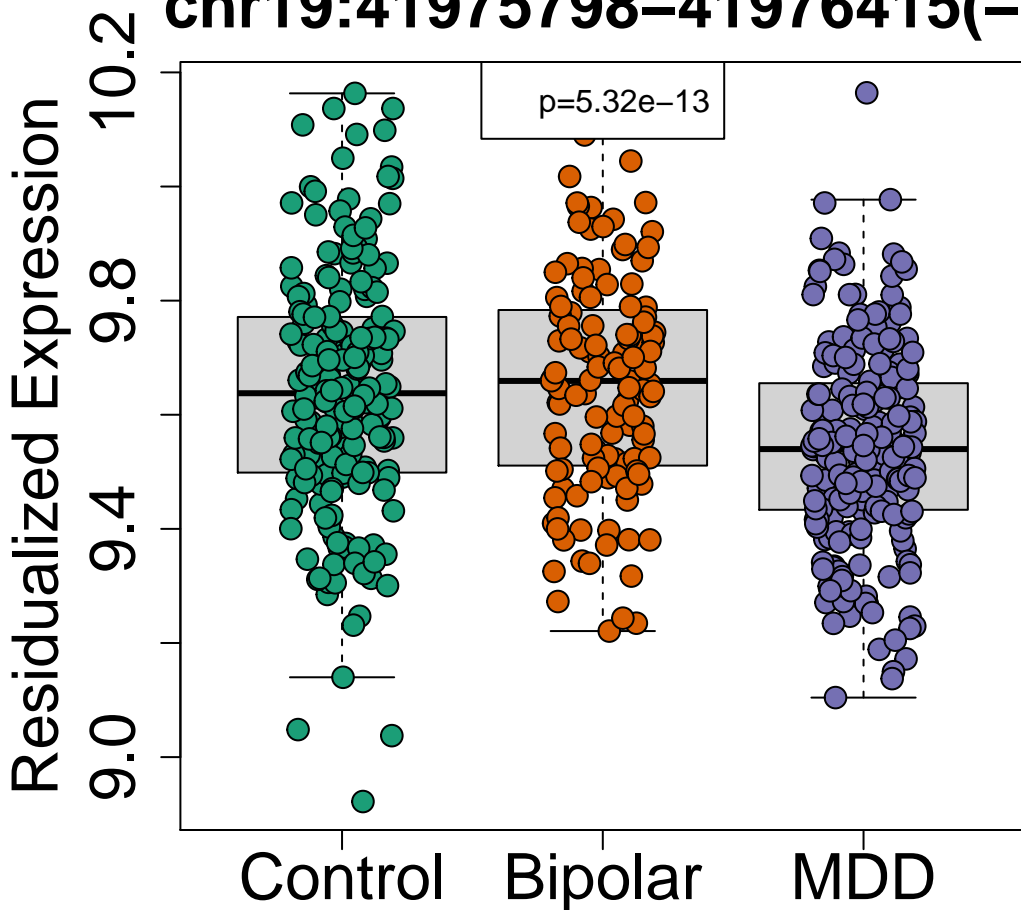
CPE, InGen

chr4:165484605–165487437(+)



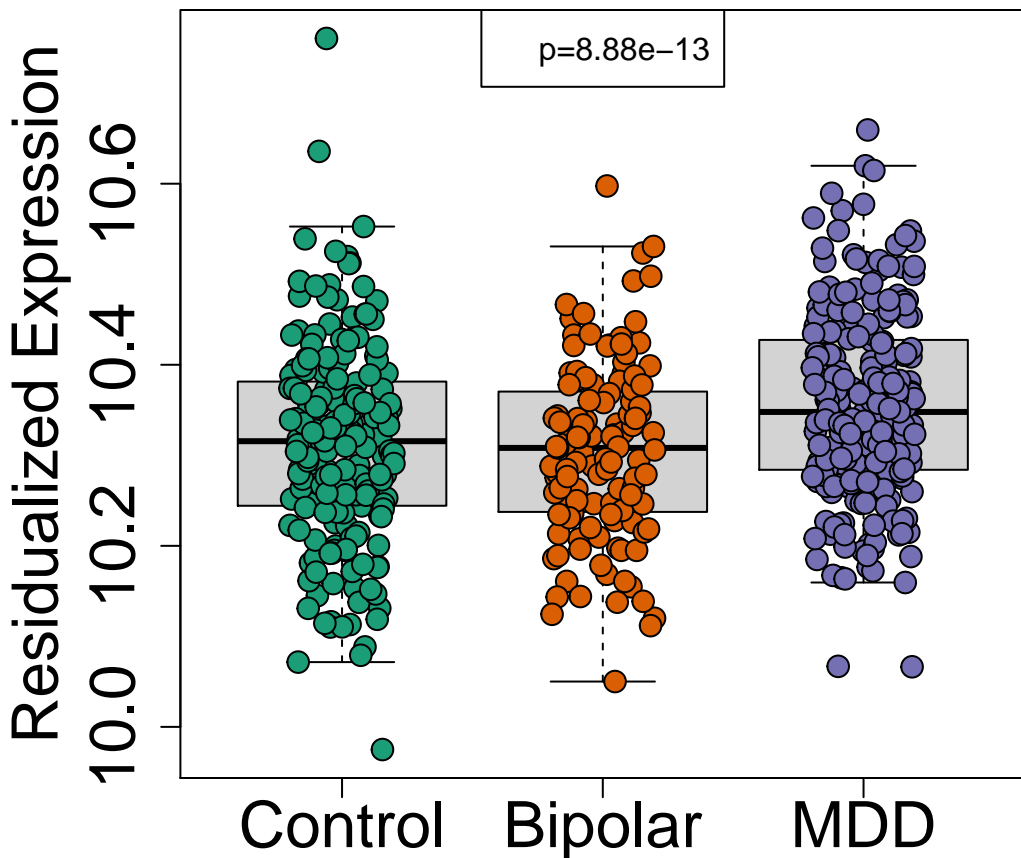
ATP1A3, InGen

chr19:41975798–41976415(-)



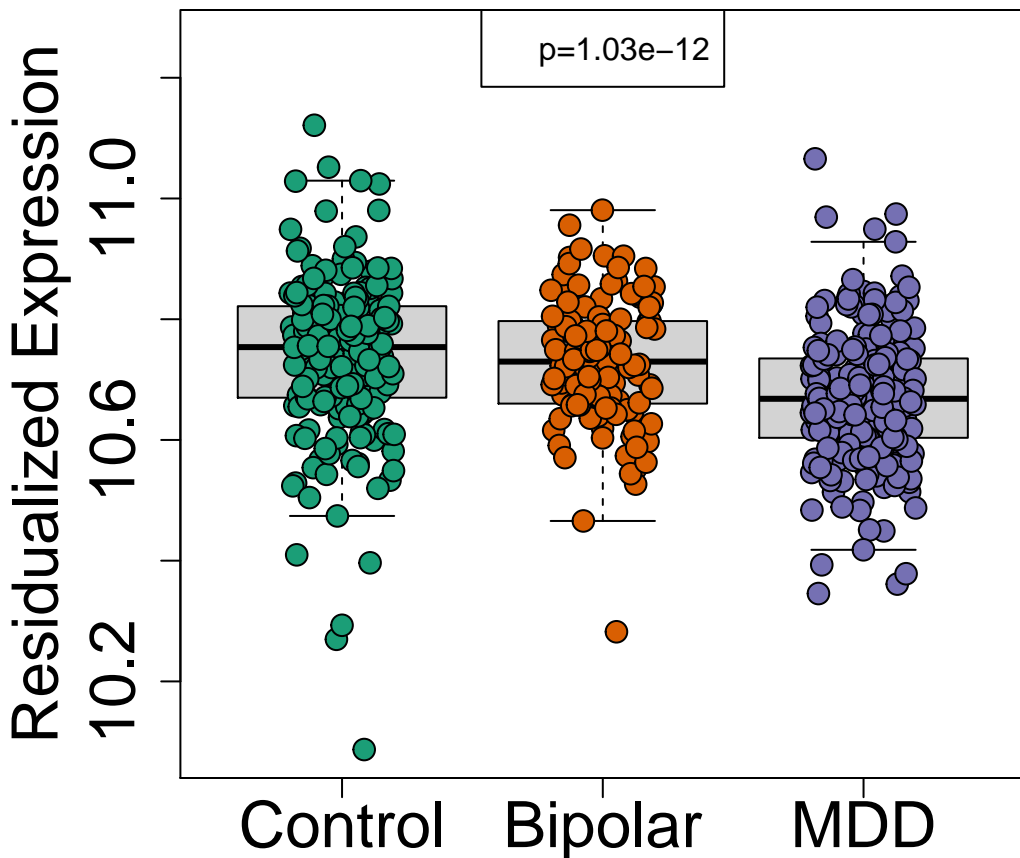
RPL11, InGen

chr1:23694792–23695797(+)



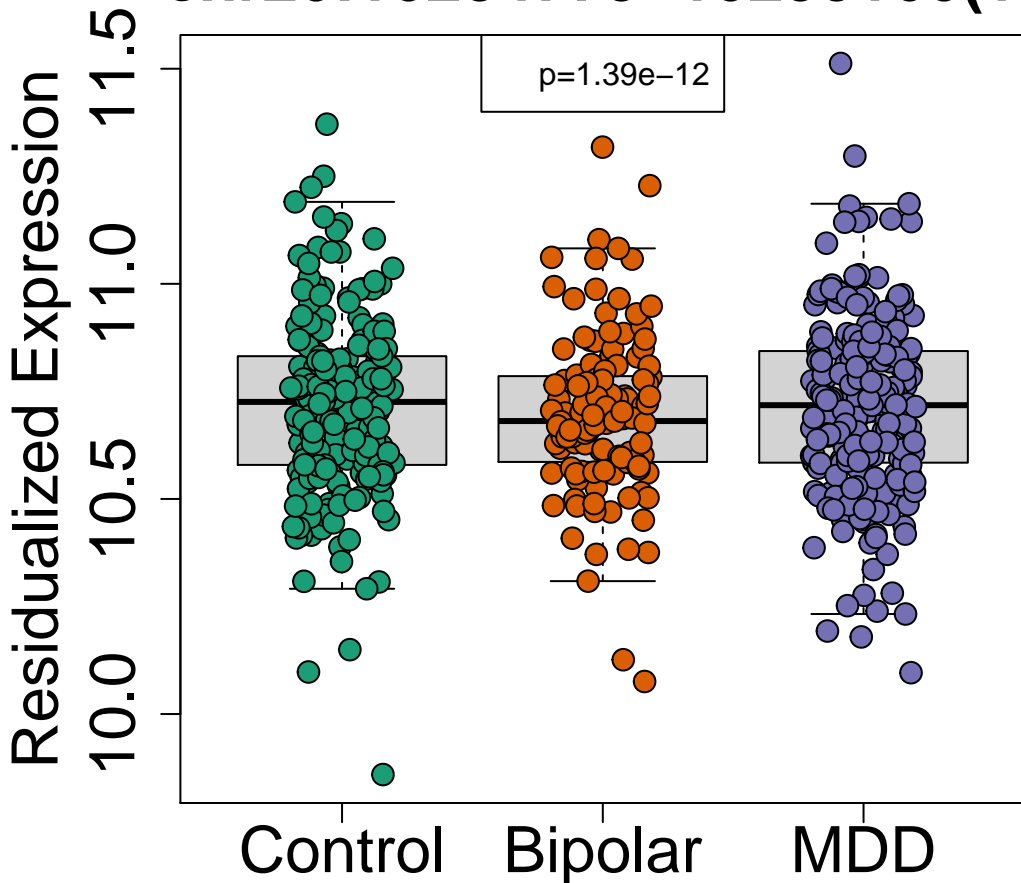
GNAS, InGen

chr20:58905481–58909161(+)



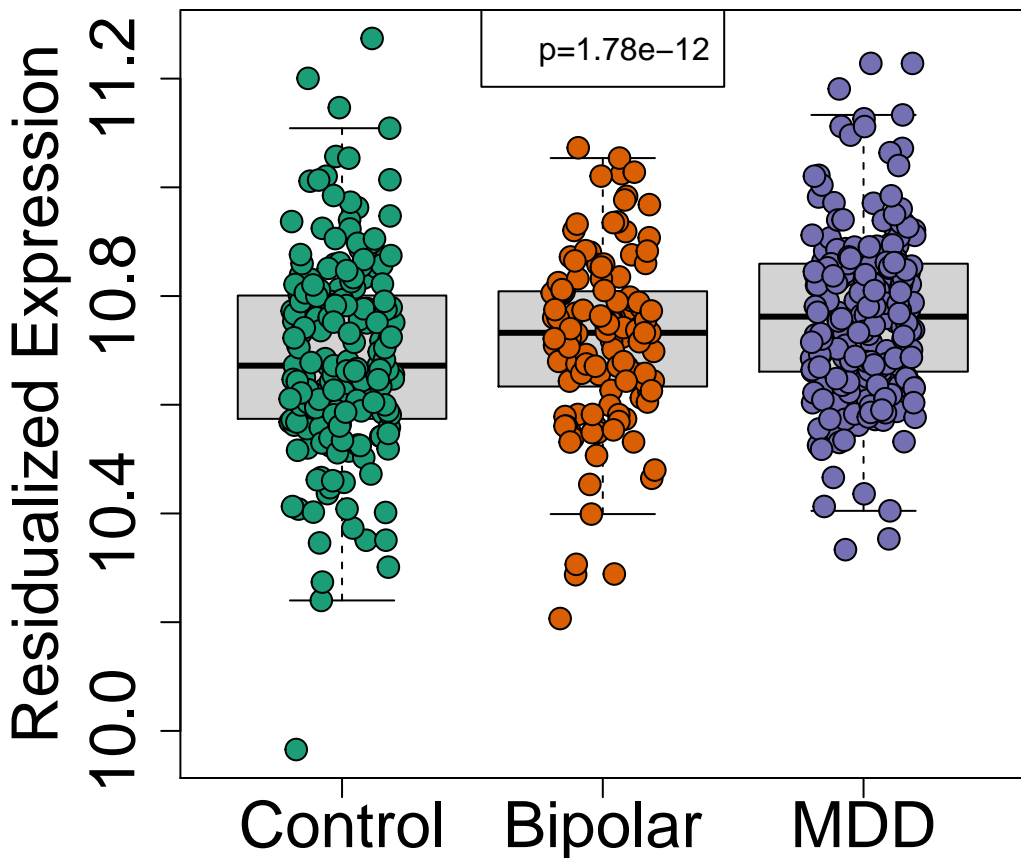
SNAP25, InGen

chr20:10284773–10293160(+)



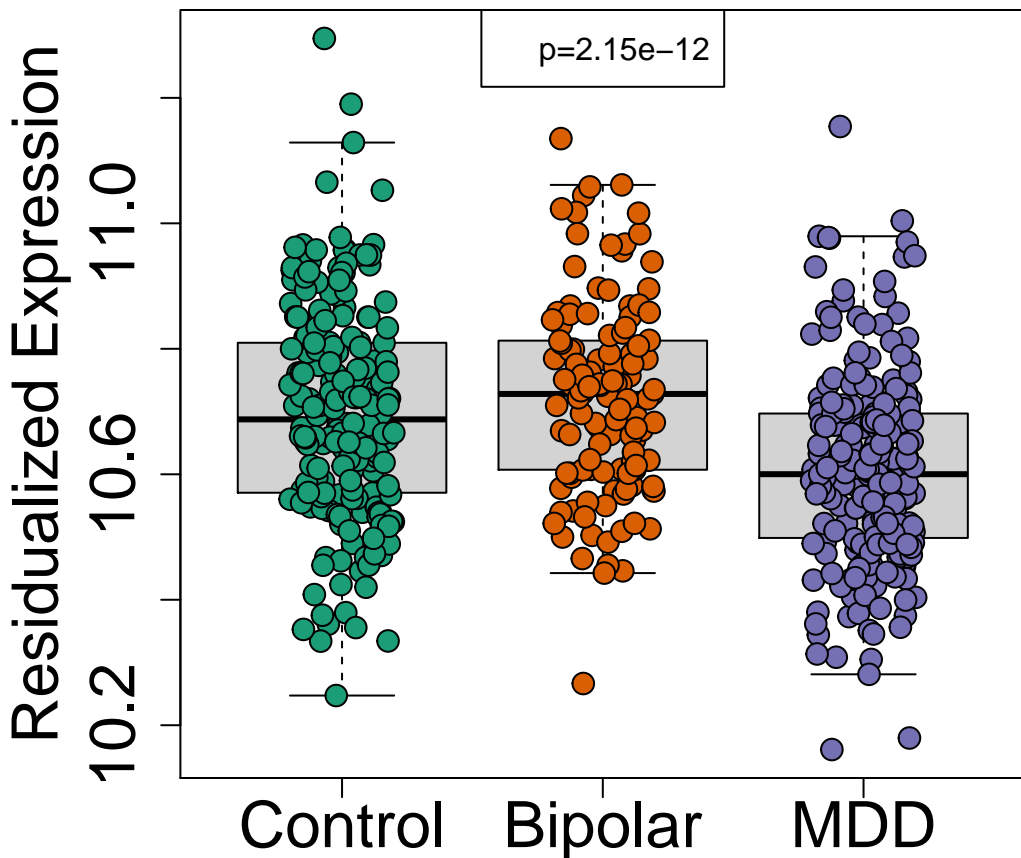
TSPAN7, InGen

chrX:38561628–38666120(+)



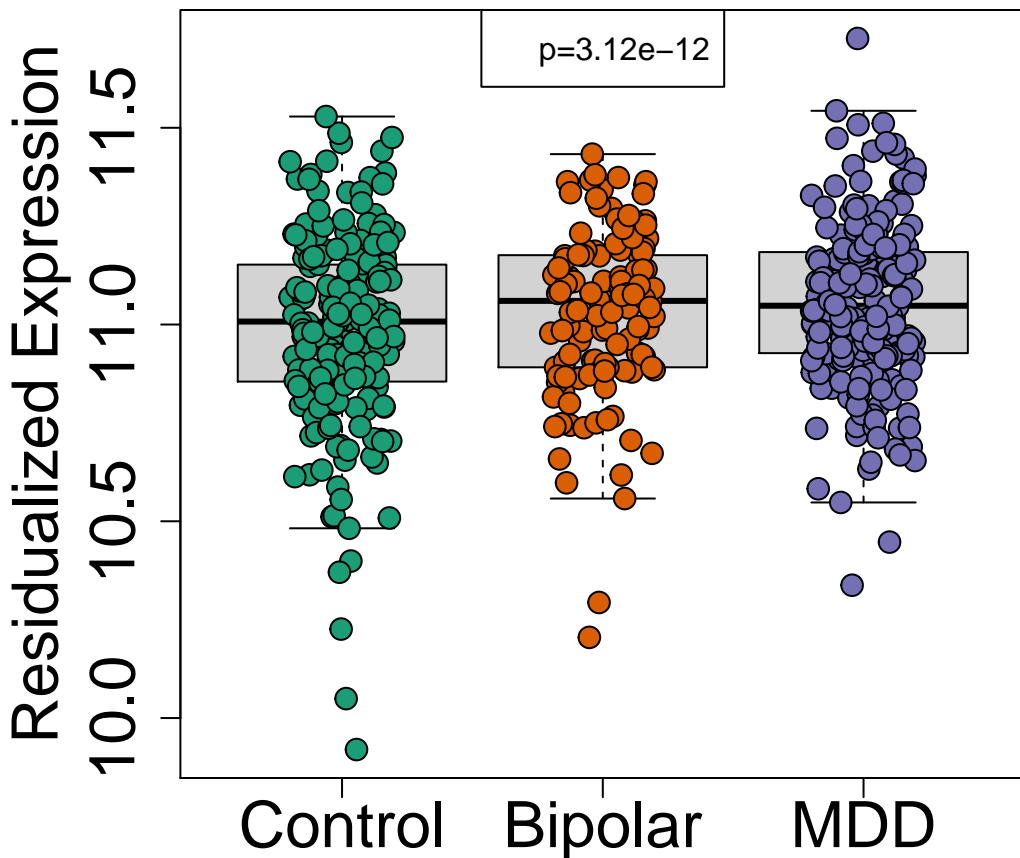
RPS21, InGen

chr20:62387915–62388308(+)



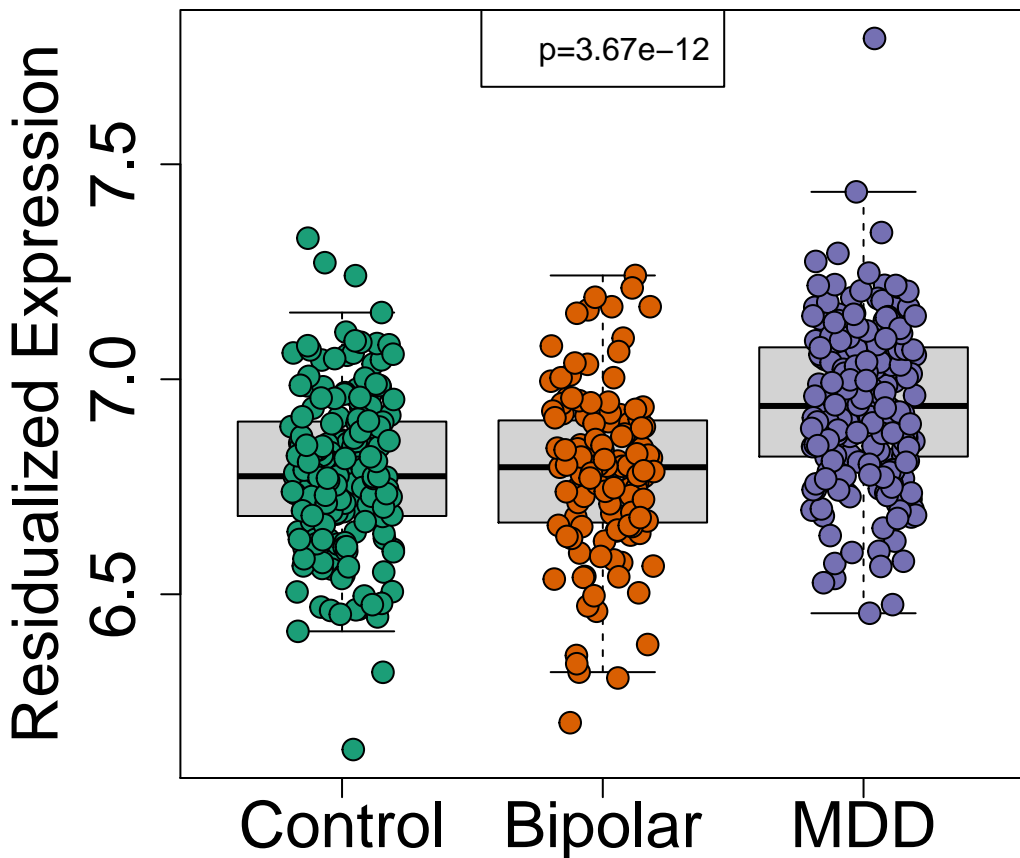
SPARCL1, InGen

chr4:87491691–87493581(-)



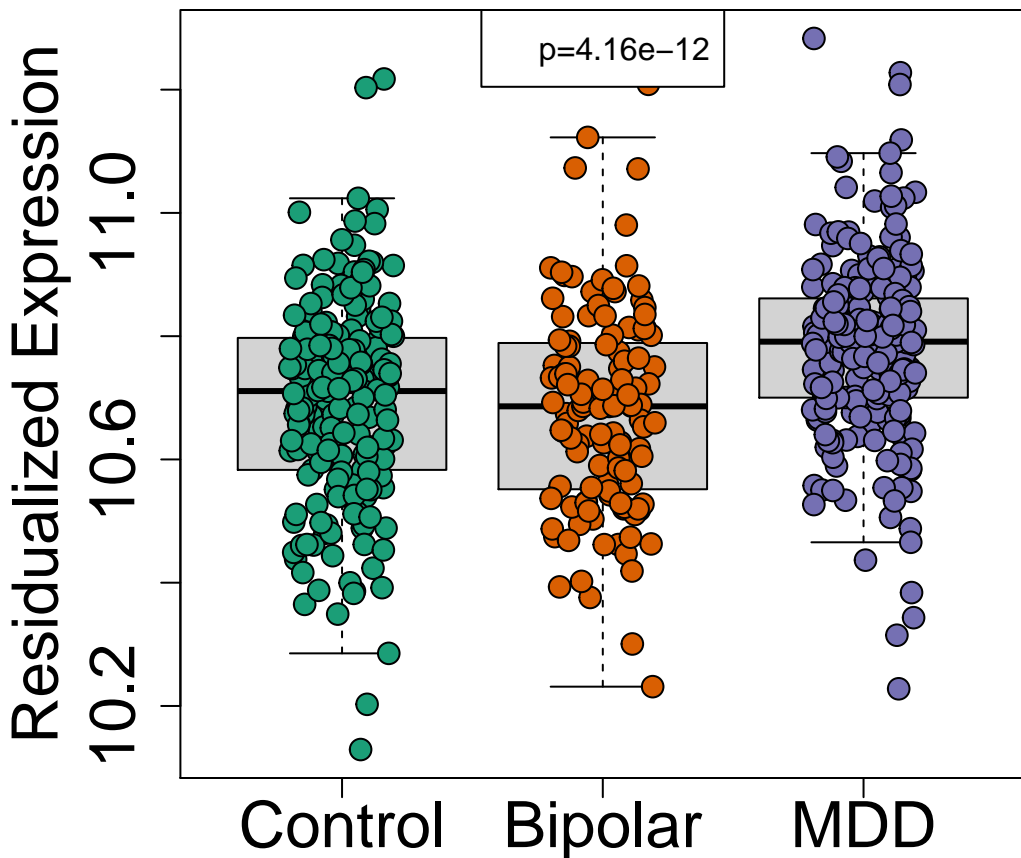
MAPT, InGen

chr17:45894687–45962320(+)



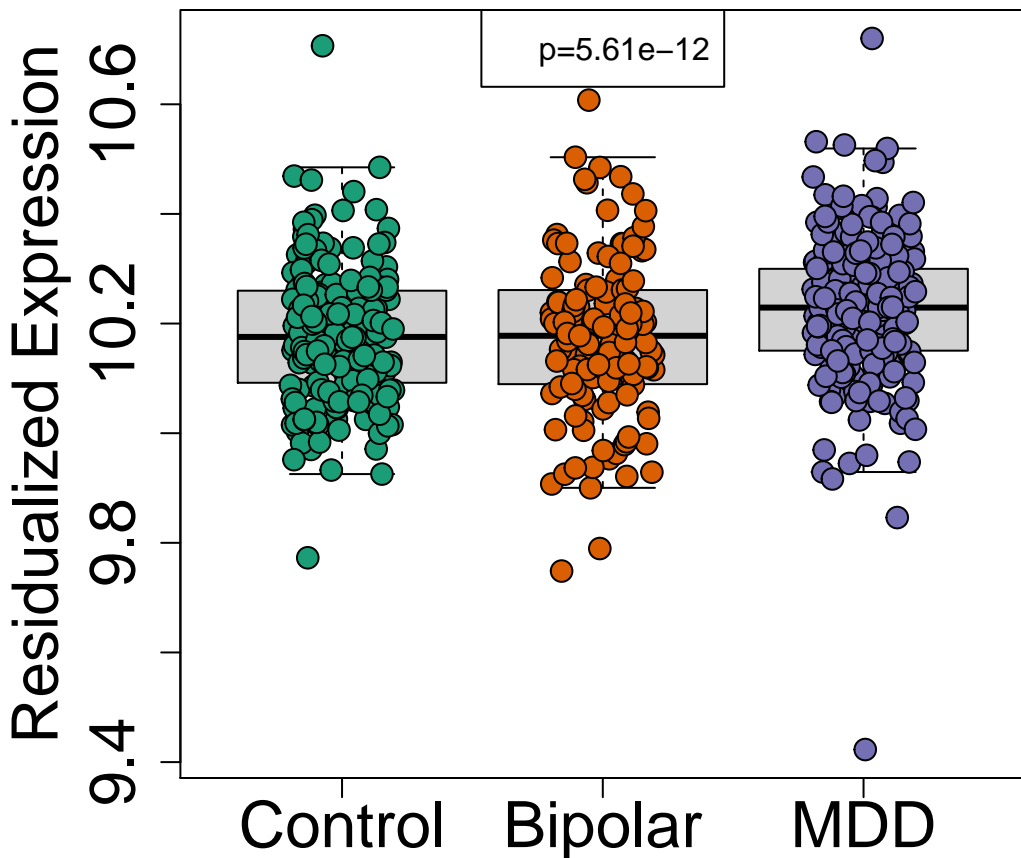
TUBA1A, InGen

chr12:49185991–49186309(-)



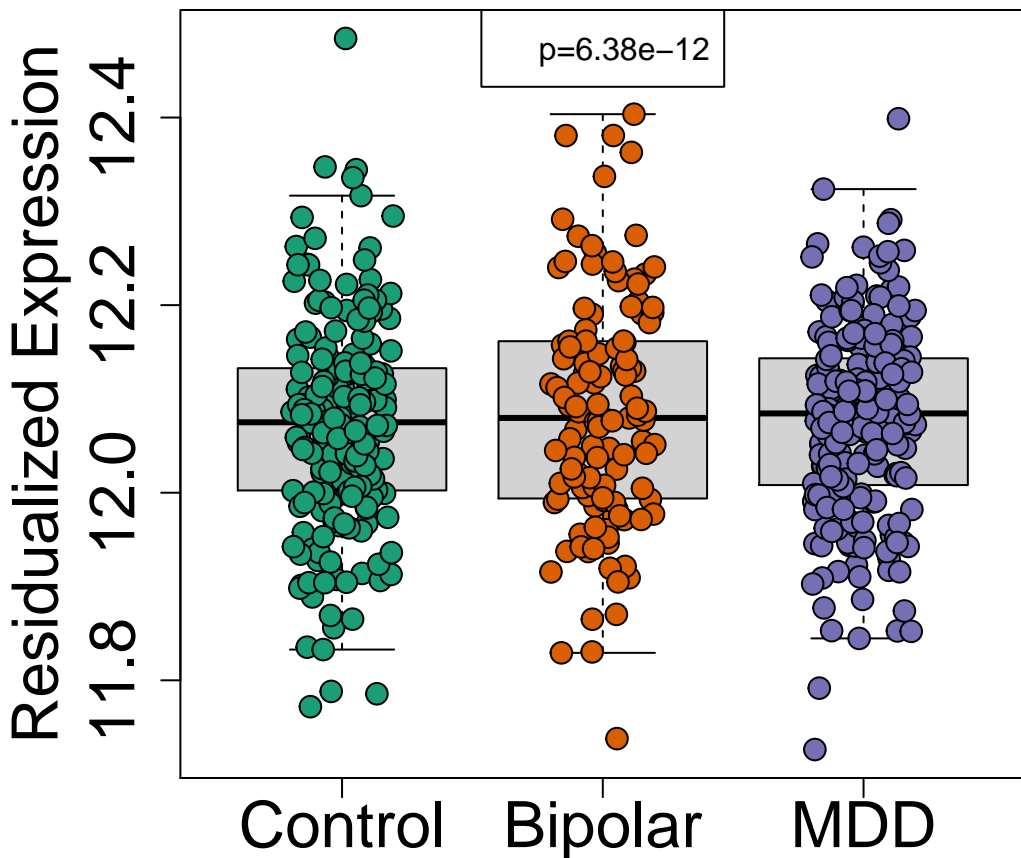
HSPA8, InGen

chr11:123059262–123059472(-)



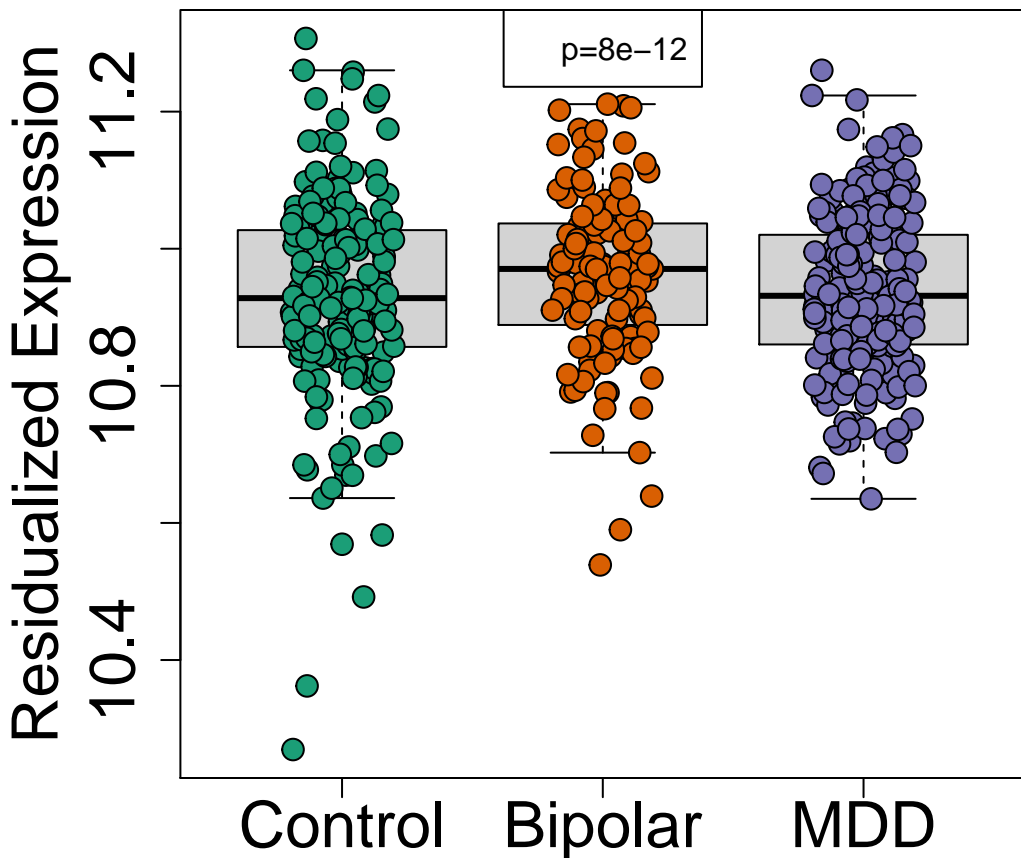
EEF1A1, InGen

chr6:73518849–73518931(–)



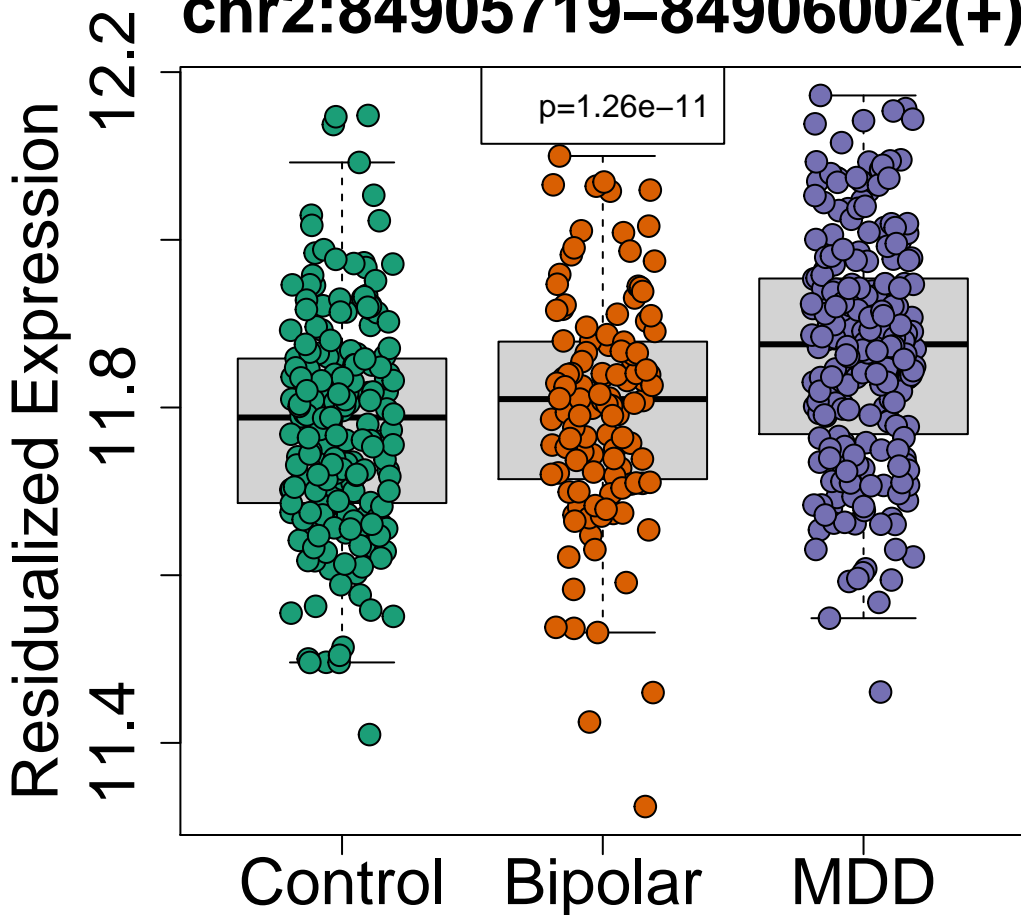
CALM3, InGen

chr19:46608341–46608481(+)



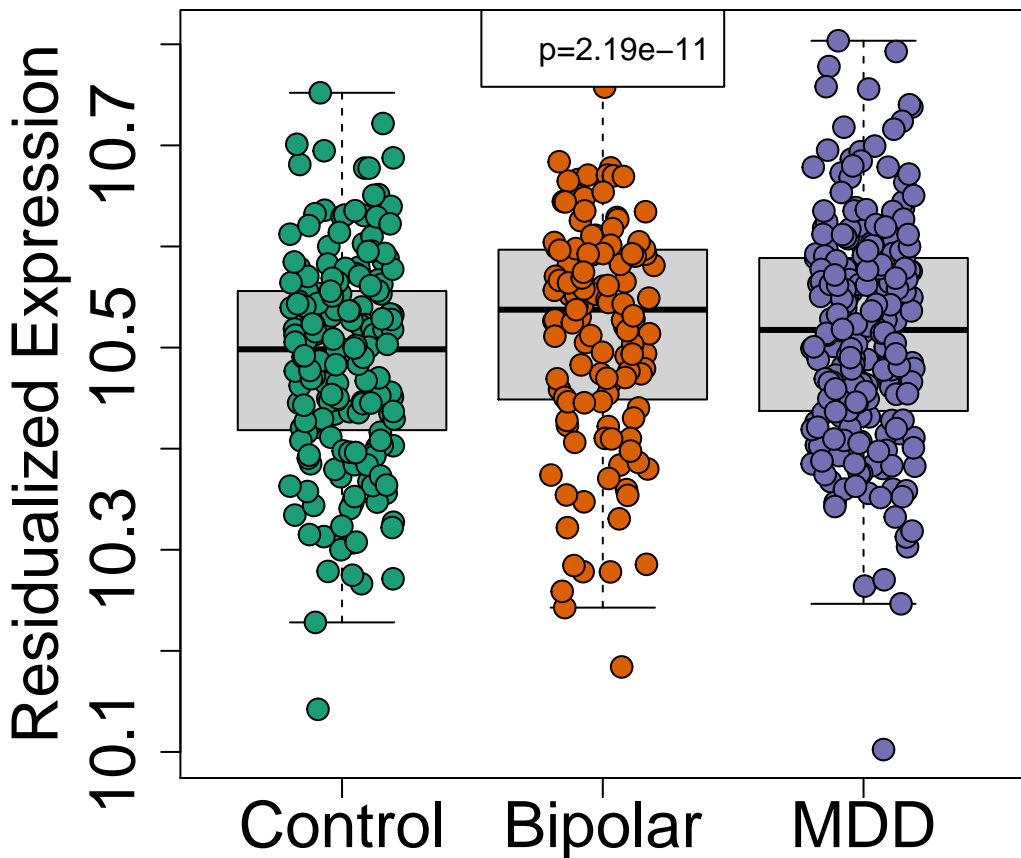
TMSB10, InGen

chr2:84905719–84906002(+)



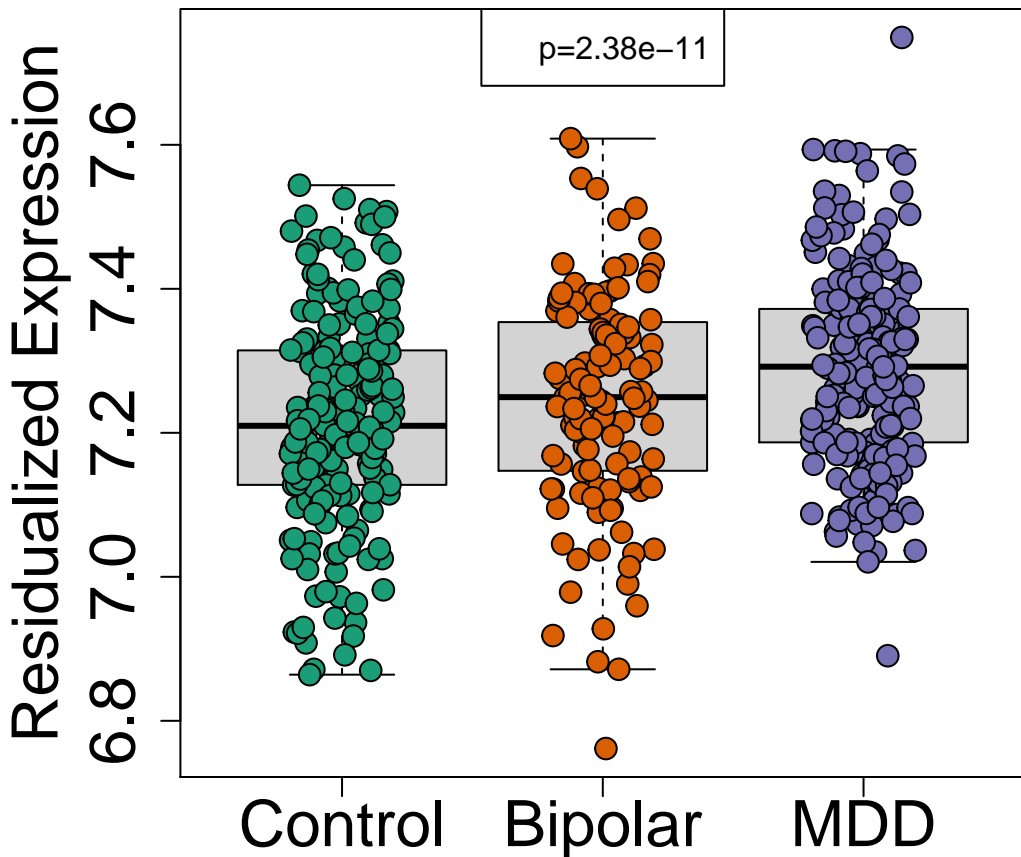
SKP1, InGen

chr5:134158596–134160986(–)



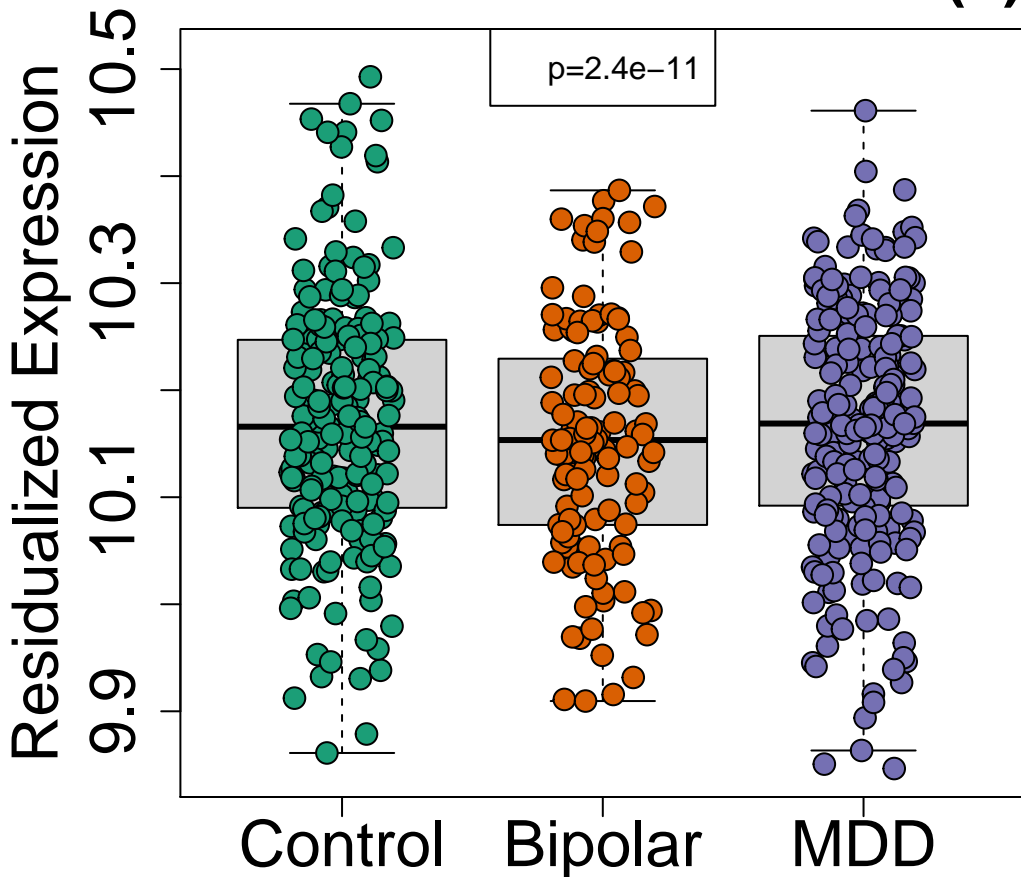
FKBP8, InGen

chr19:18532256–18532663(-)



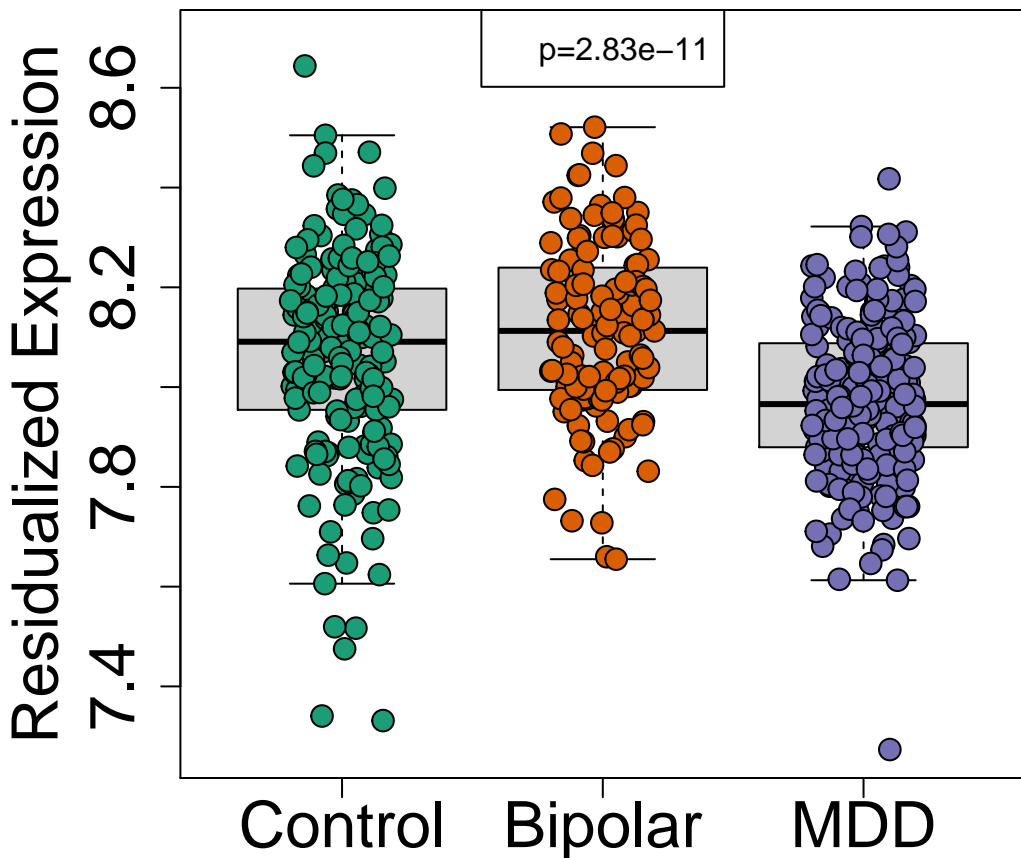
RPS6, InGen

chr9:19378919–19379486(-)



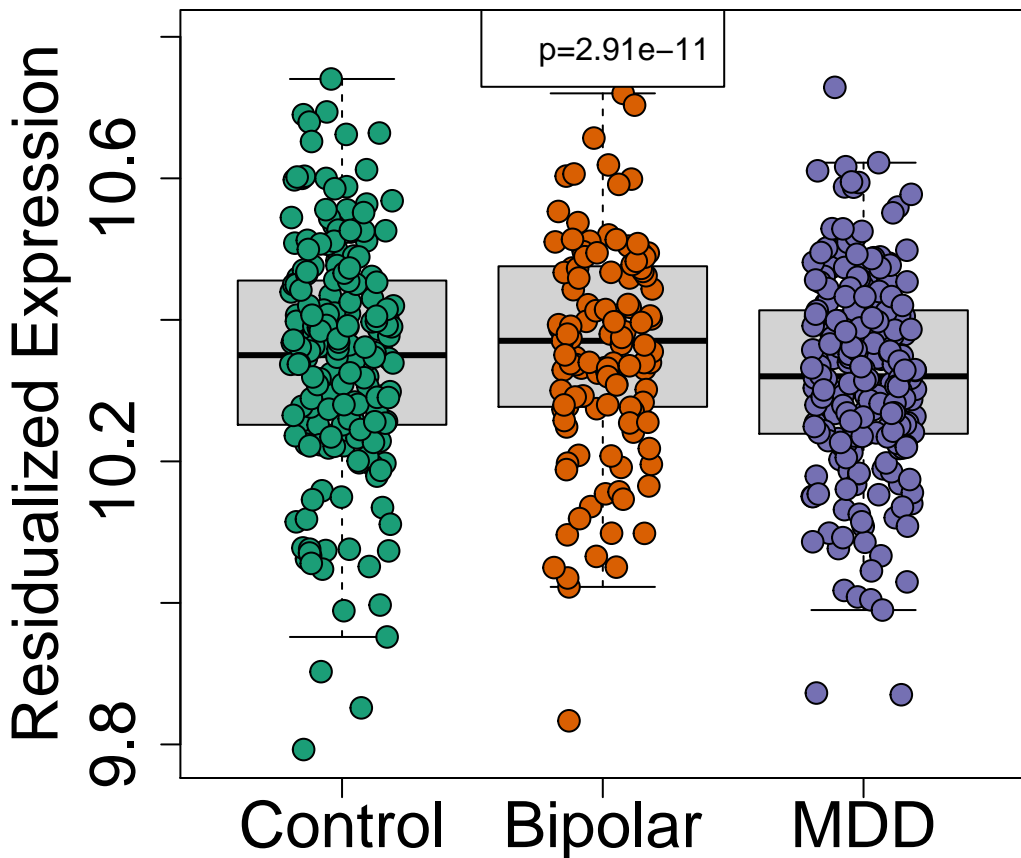
ATP1A3, InGen

chr19:41985424–41985863(-)



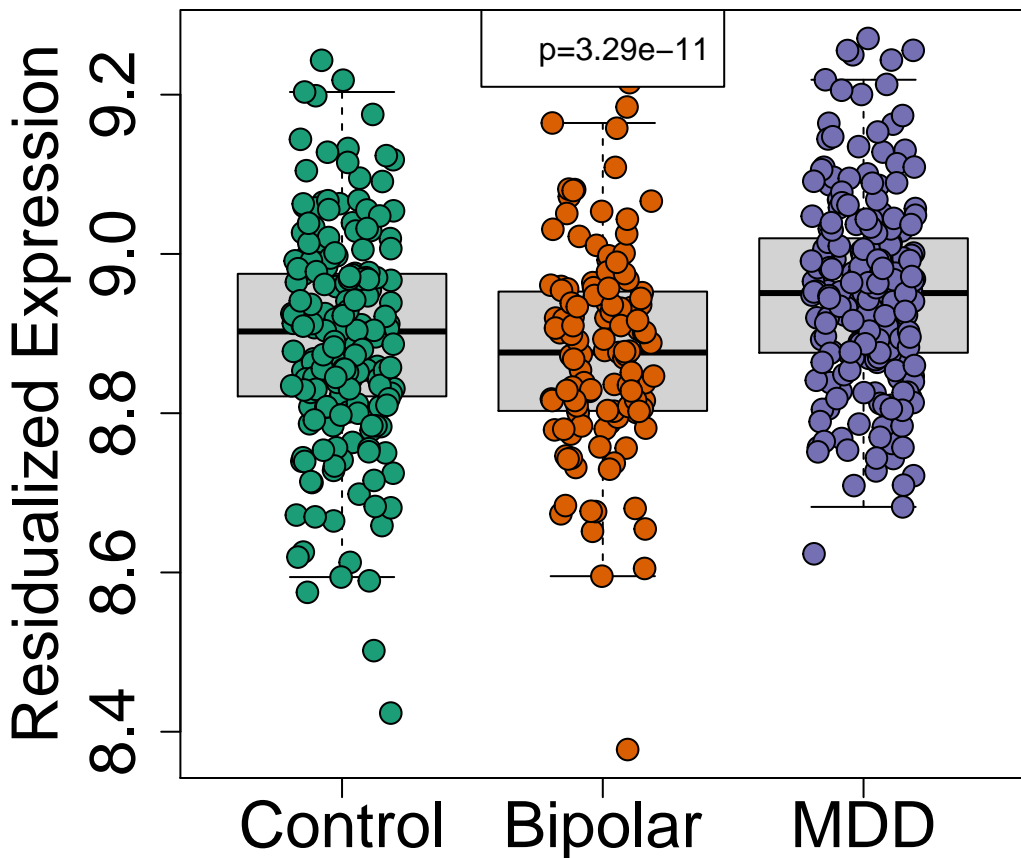
APP, InGen

chr21:26022043–26050999(-)



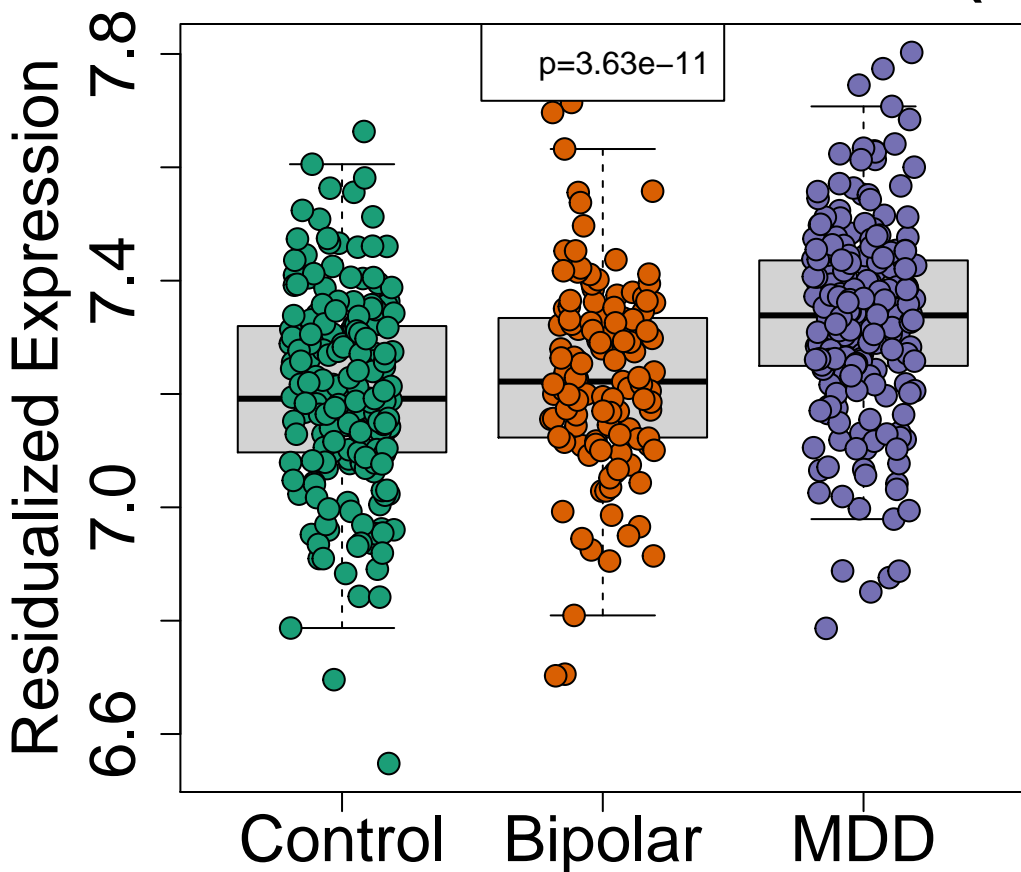
RPL18, InGen

chr19:48616203–48616725(-)



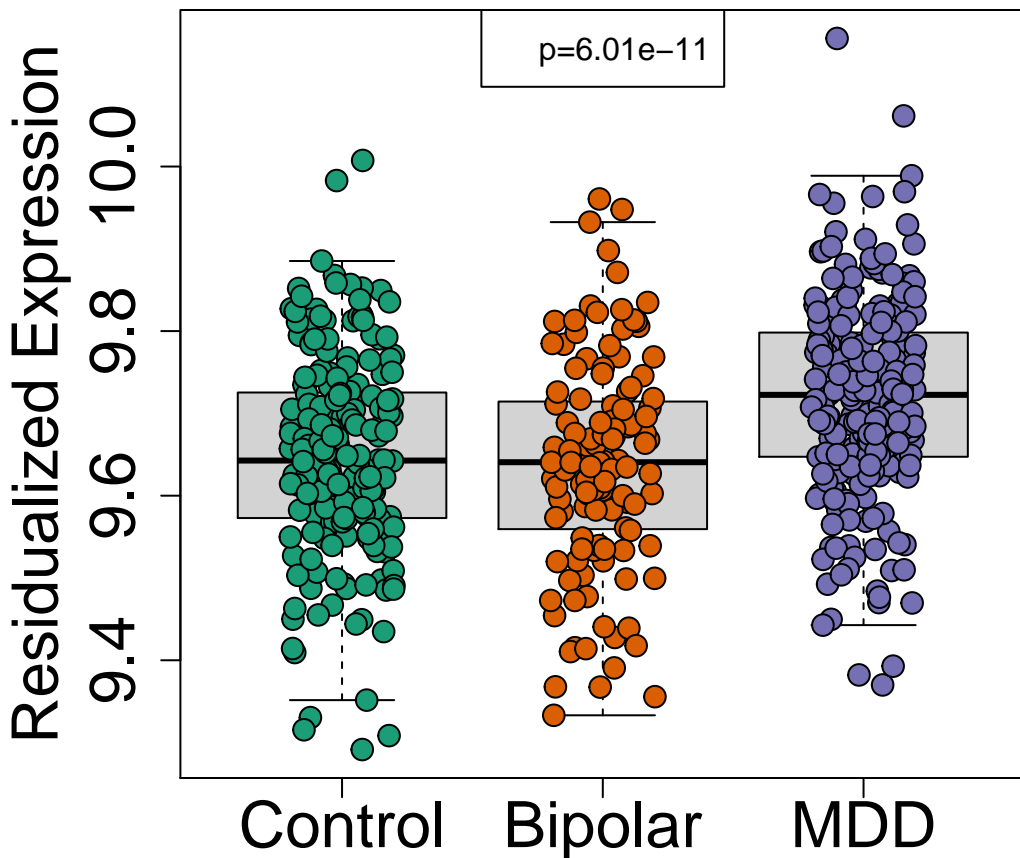
EIF3L, InGen

chr22:37870348–37874369(+)



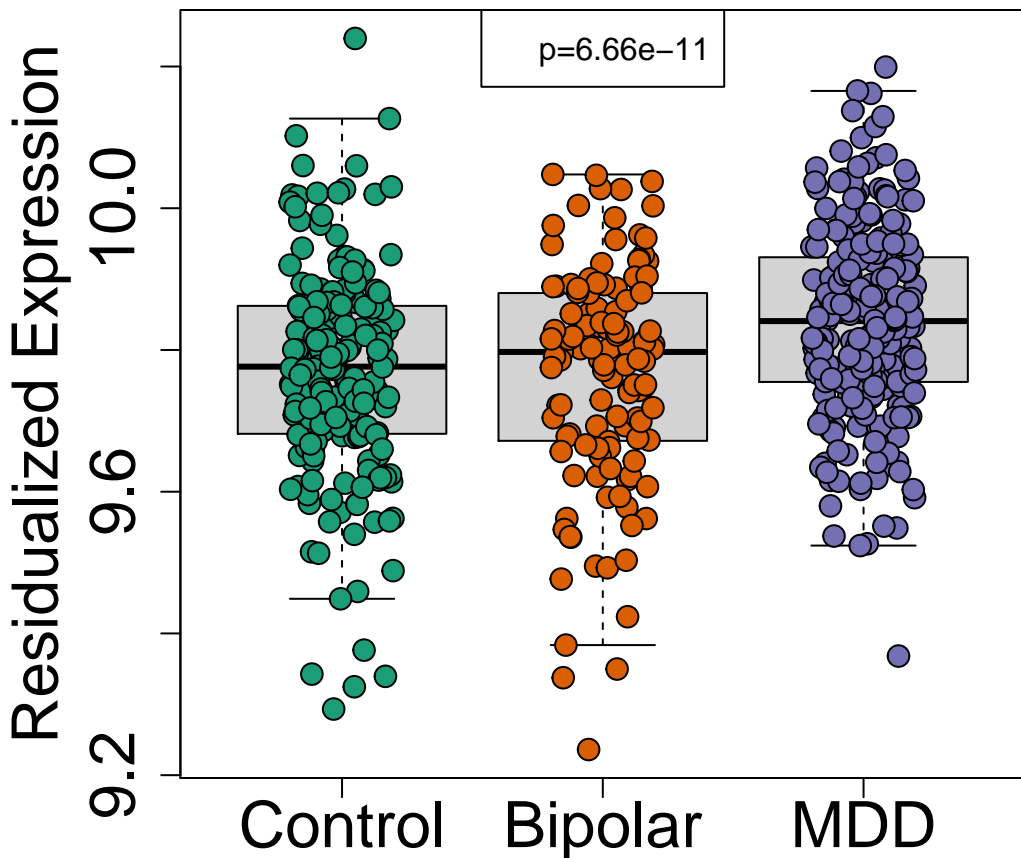
ENO1, InGen

chr1:8866502–8867116(–)



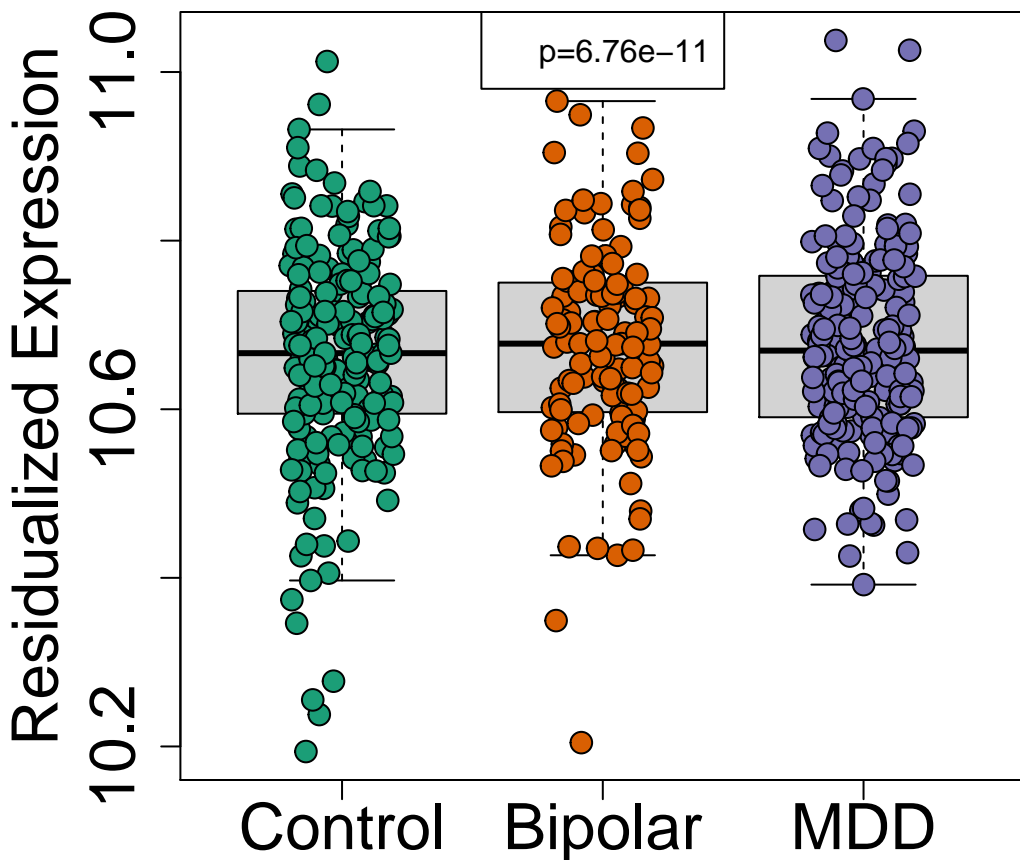
GABARAPL1, InGen

chr12:10220559–10221786(+)



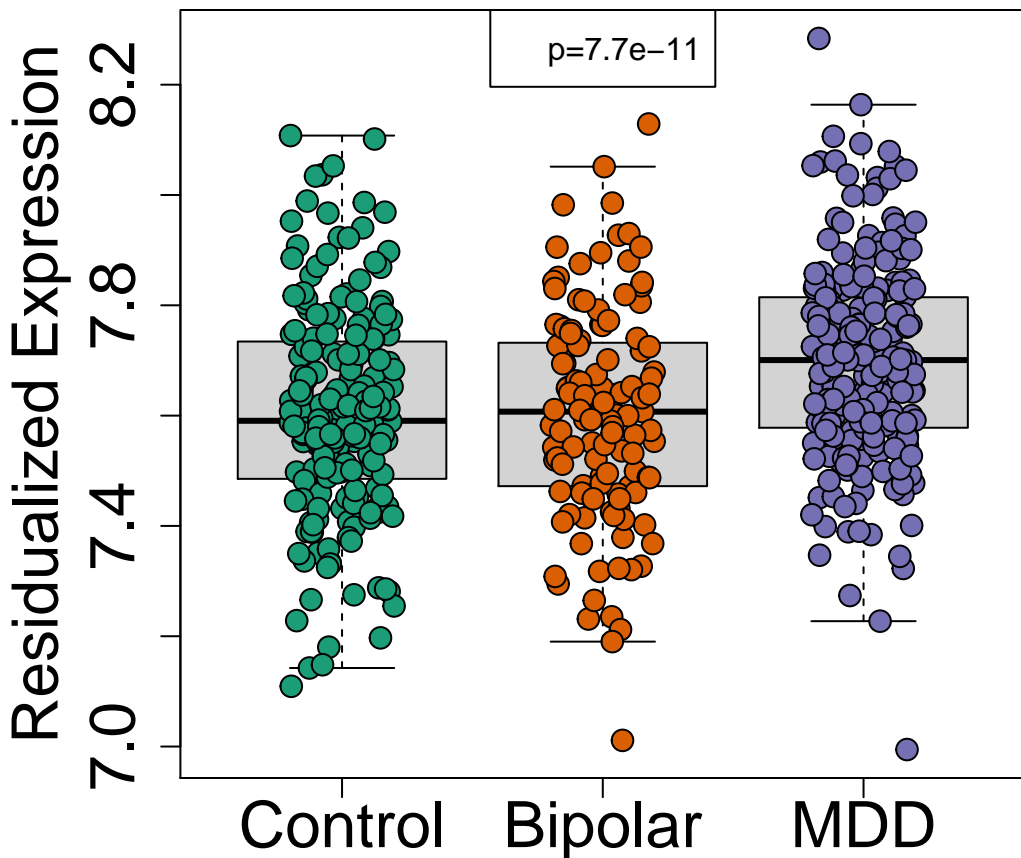
CALM3, InGen

chr19:46608589–46608845(+)



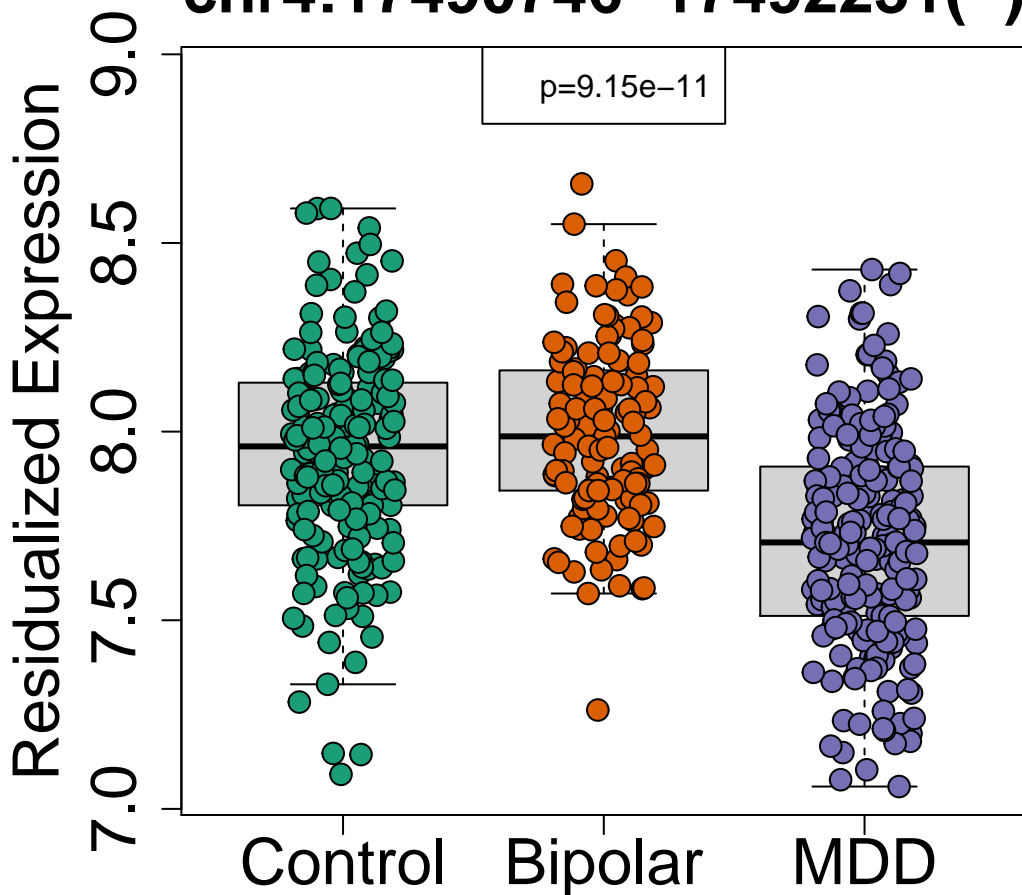
GPM6B, InGen

chrX:13785809–13816843(-)



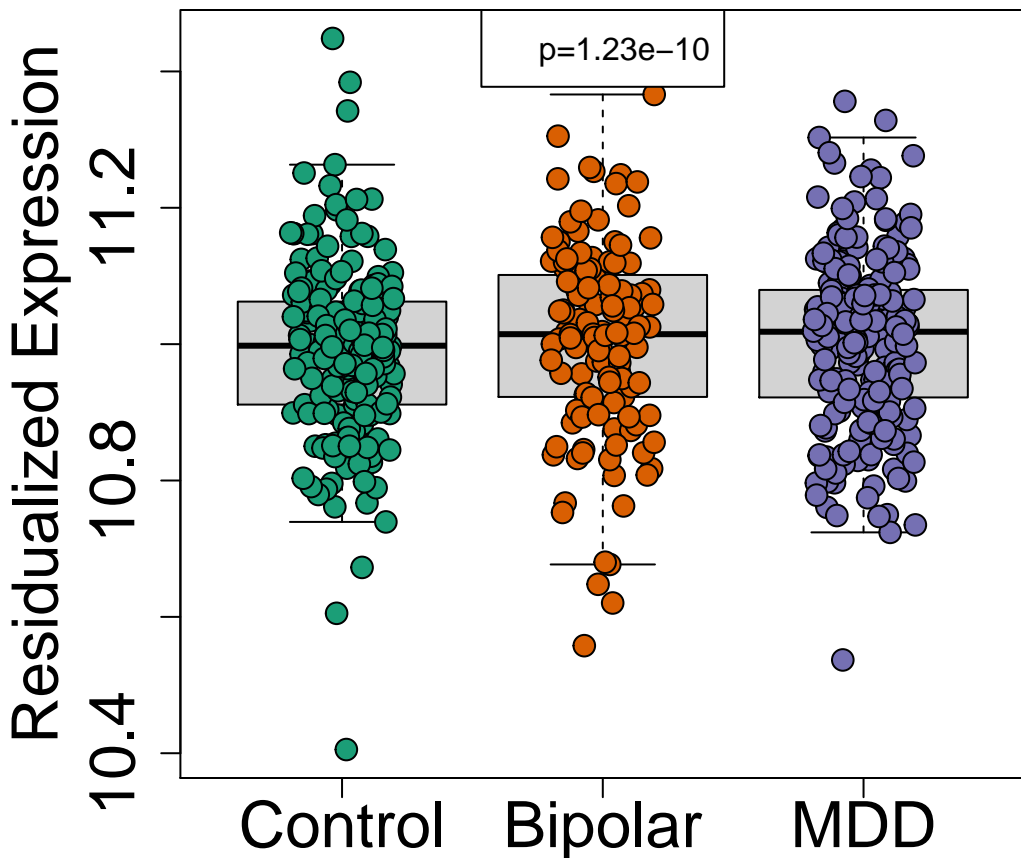
QDPR, InGen

chr4:17490746–17492231(–)



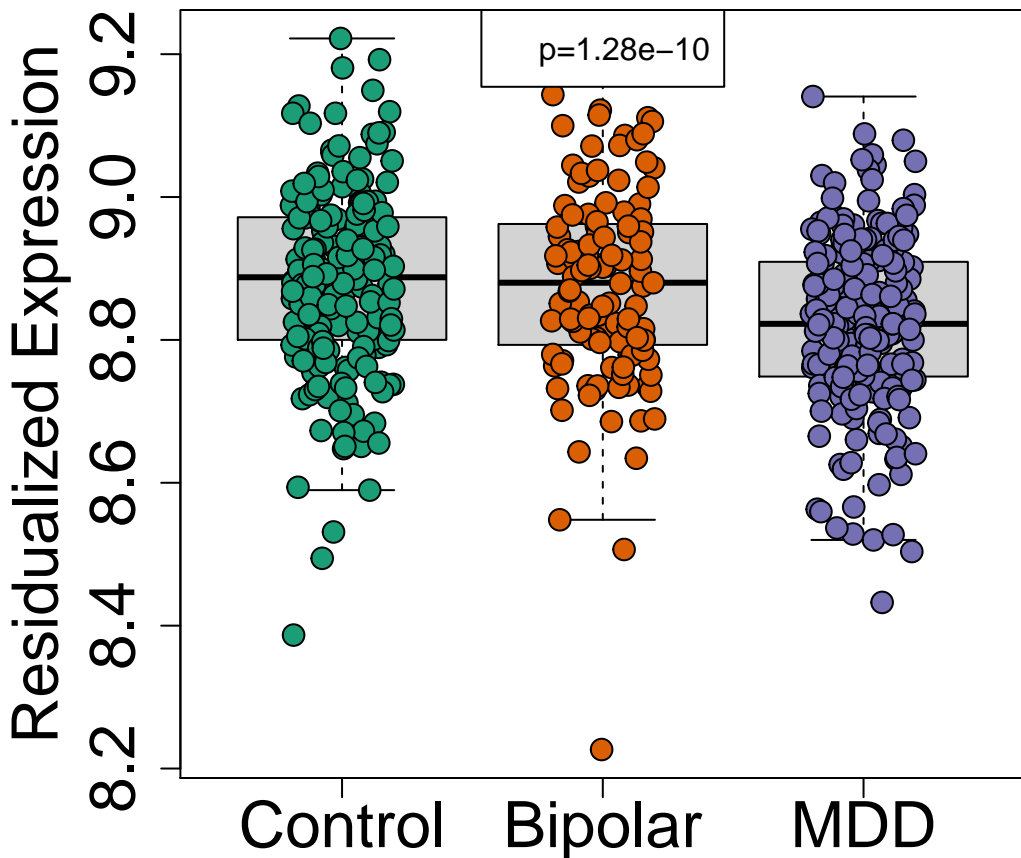
CPE, InGen

chr4:165487578–165493170(+)



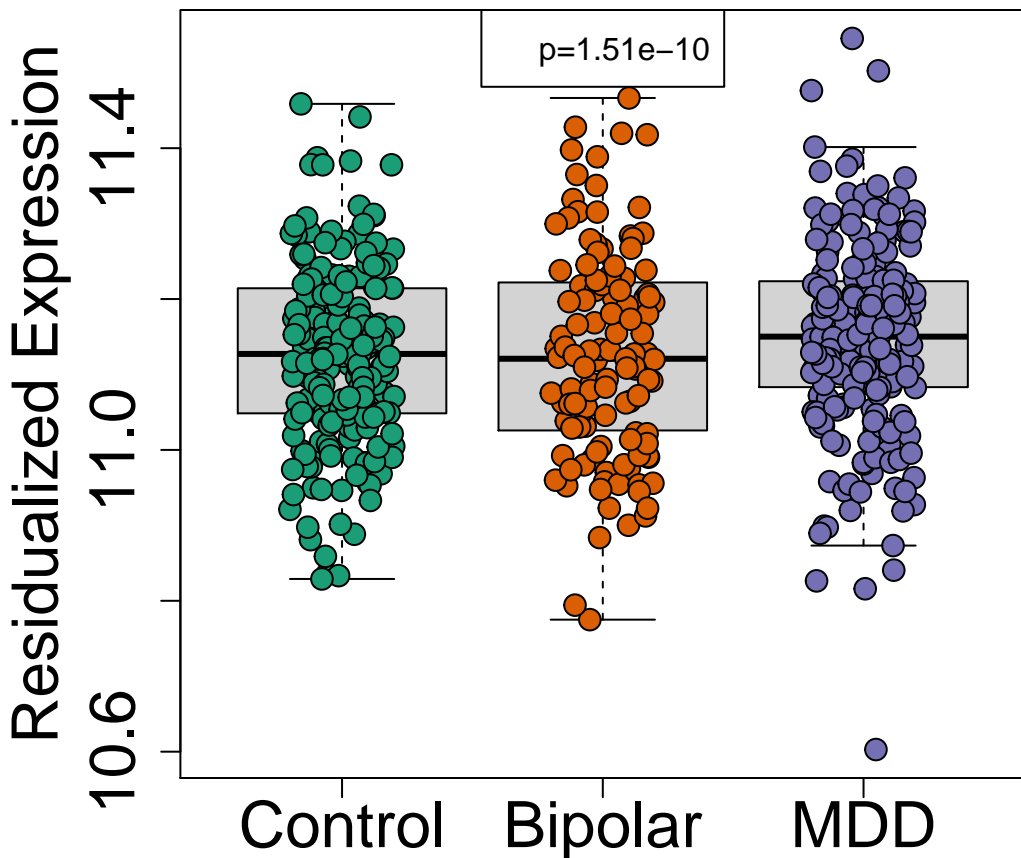
SPTBN1, InGen

chr2:54647262–54648985(+)



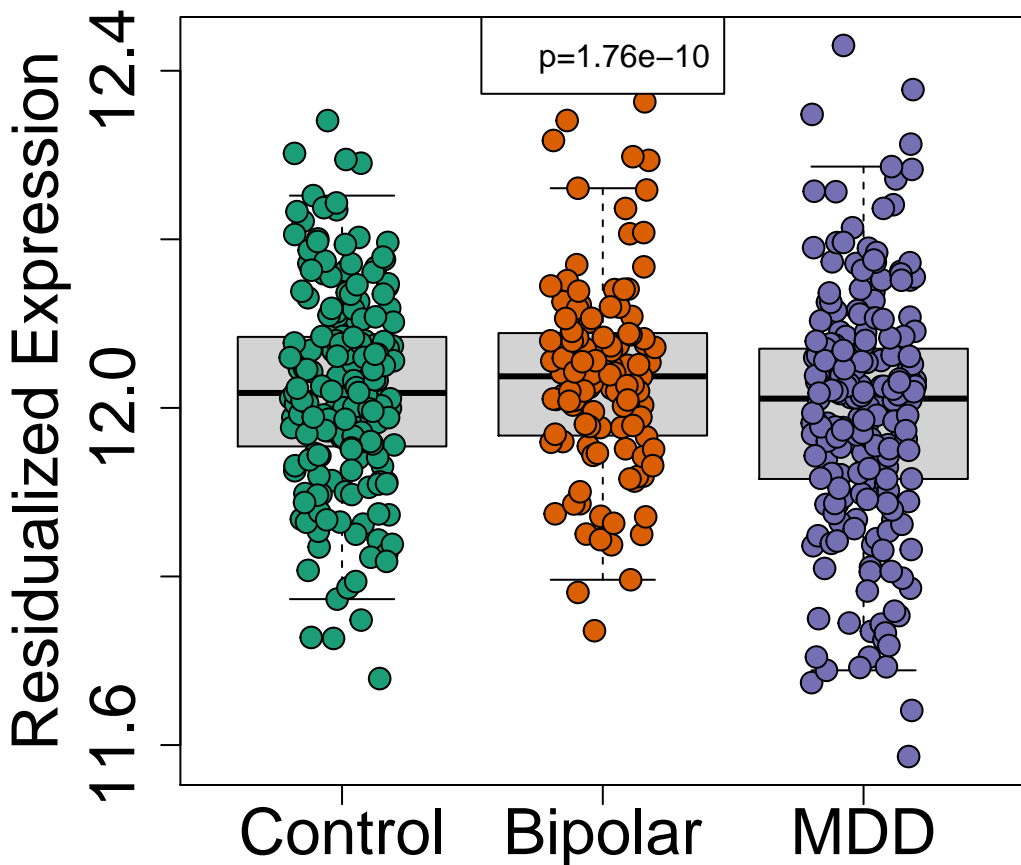
RPL13A, InGen

chr19:49489923–49490231(+)



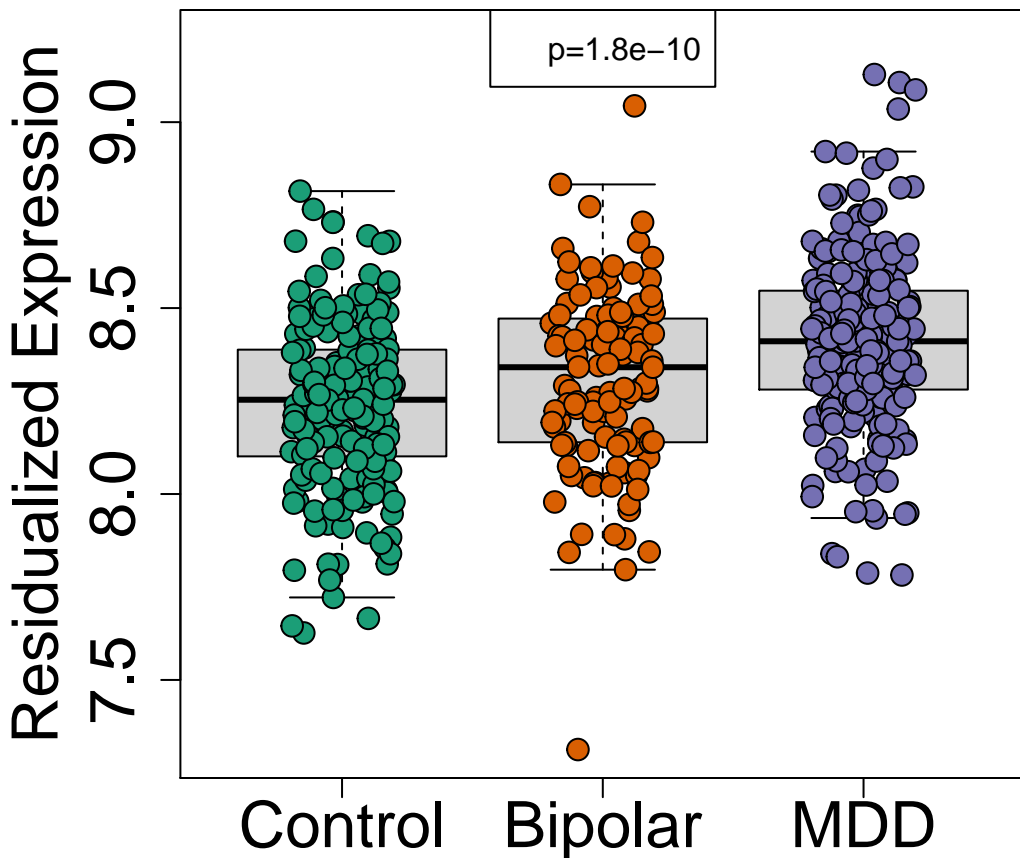
UBB, InGen

chr17:16381185–16381901(+)



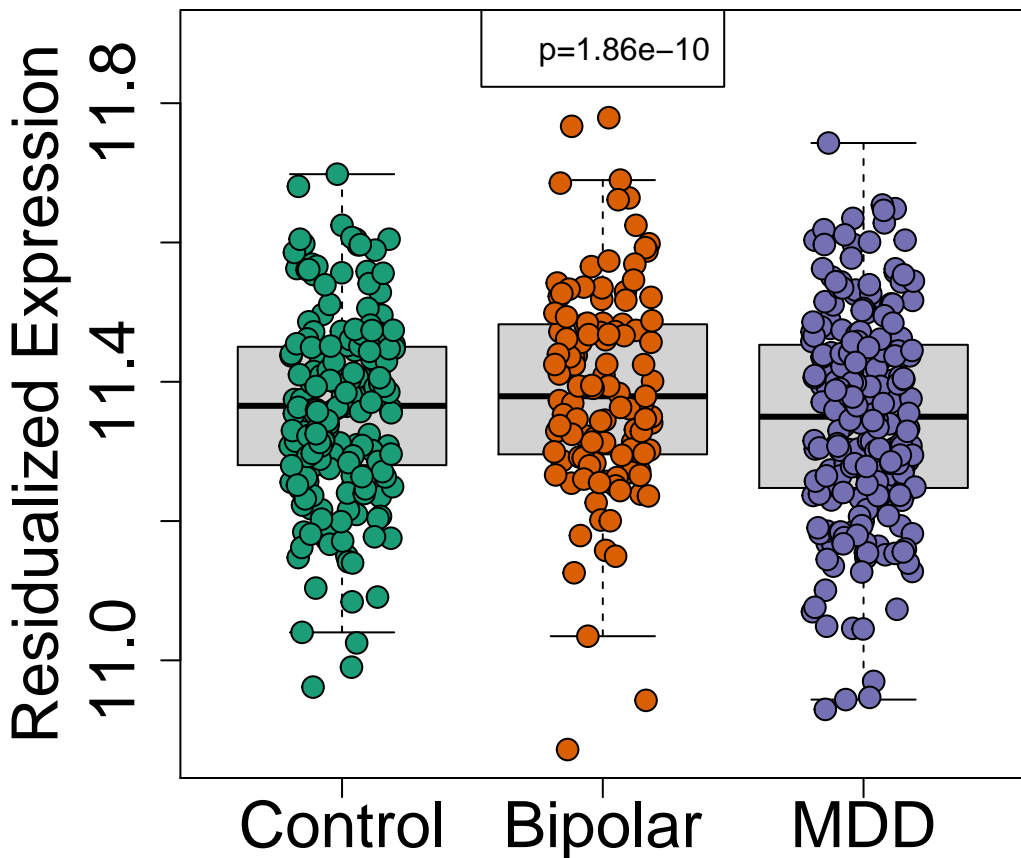
HMGN2, InGen

chr1:26474668–26475112(+)



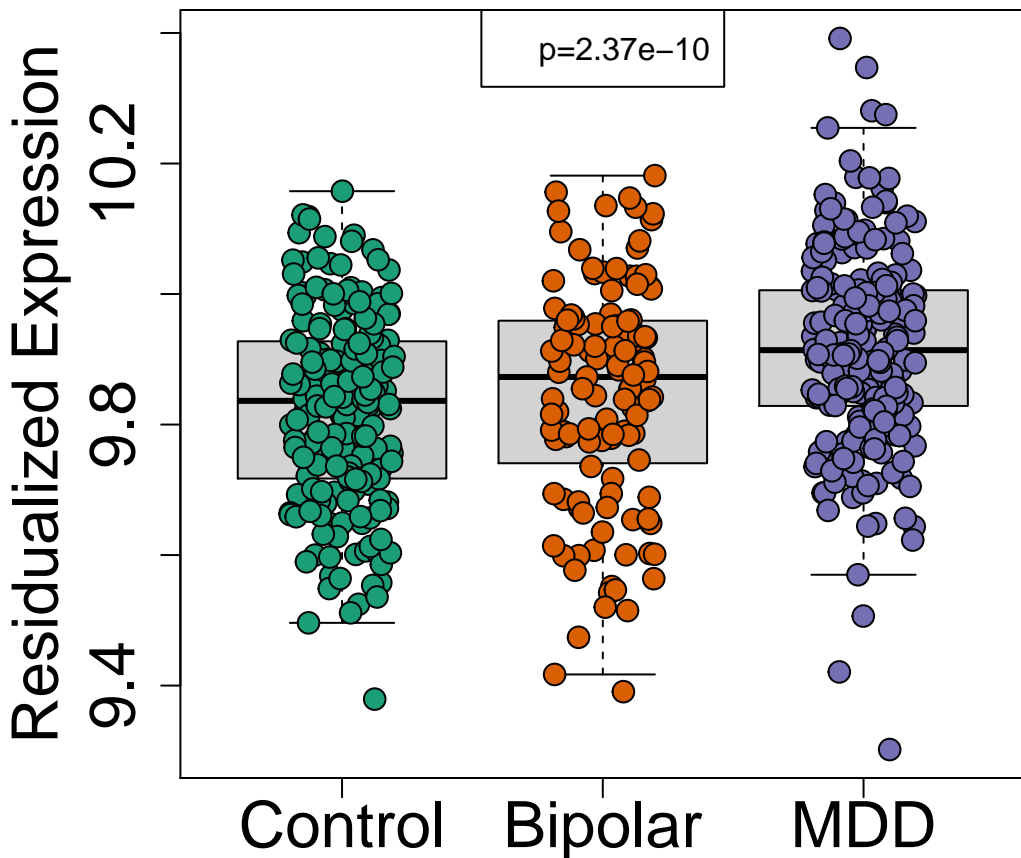
CPE, InGen

chr4:165495678–165497511(+)



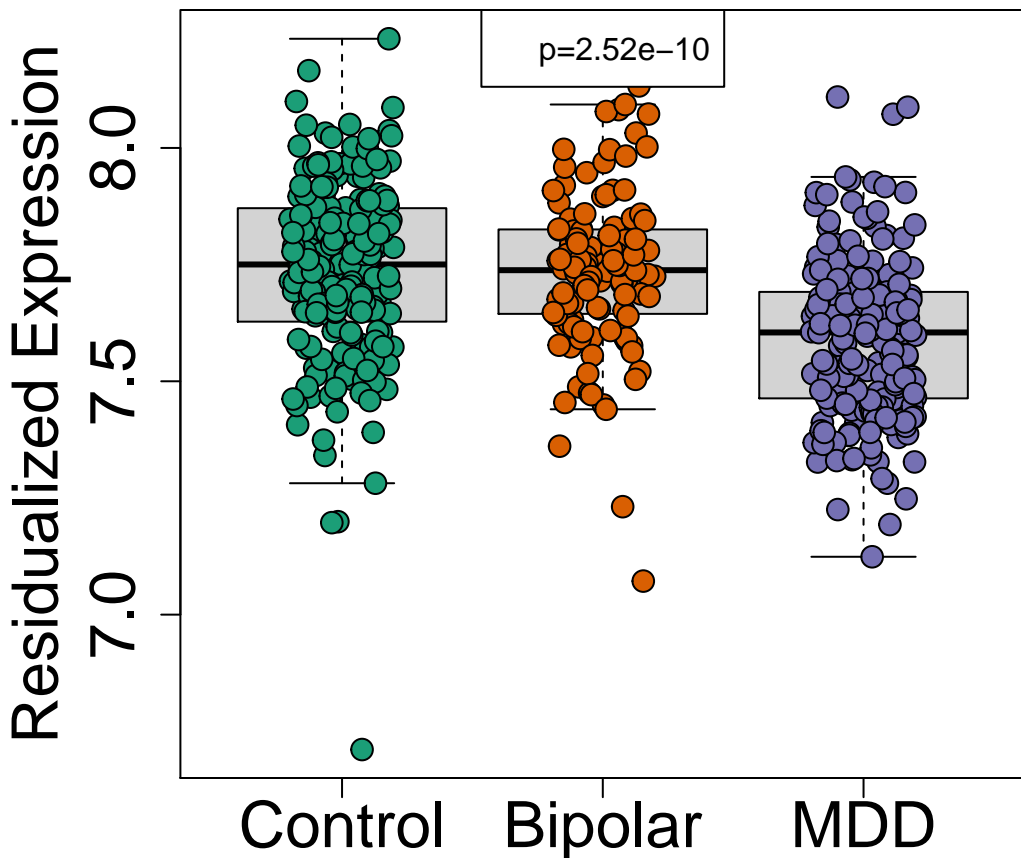
GLUD1, InGen

chr10:87060826-87060914(-)



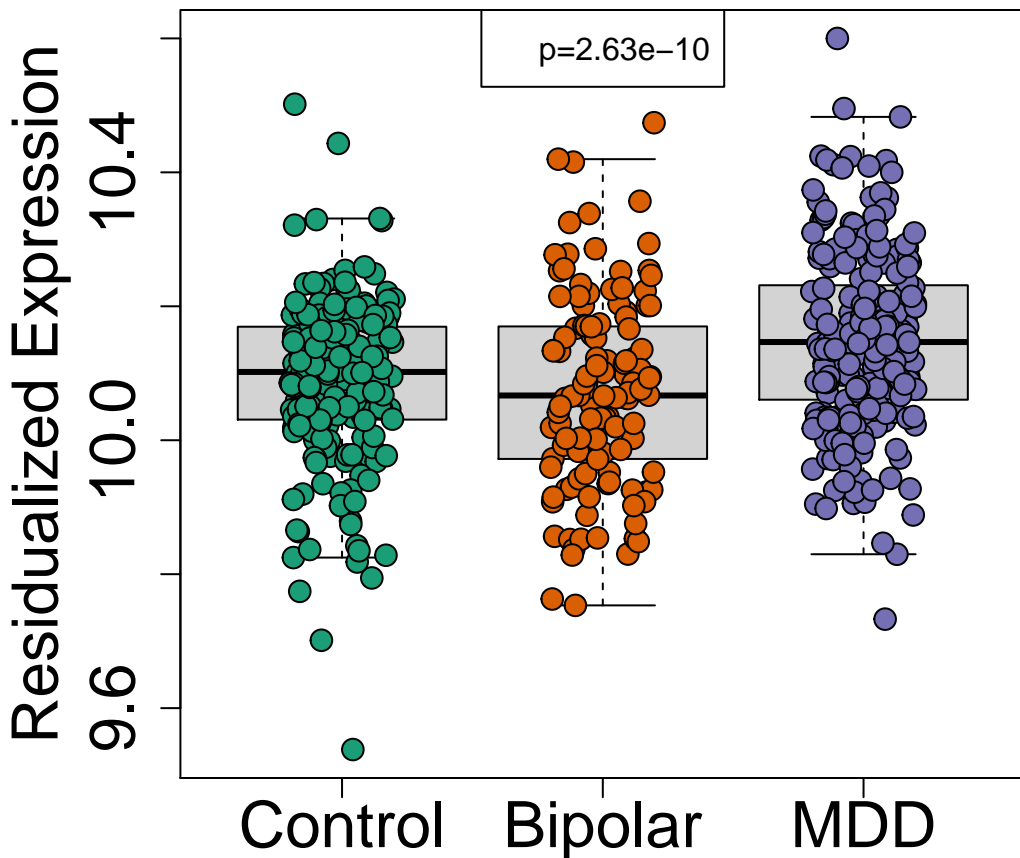
RP11-793H13.8, InGen

chr12:53464853-53465931(+)



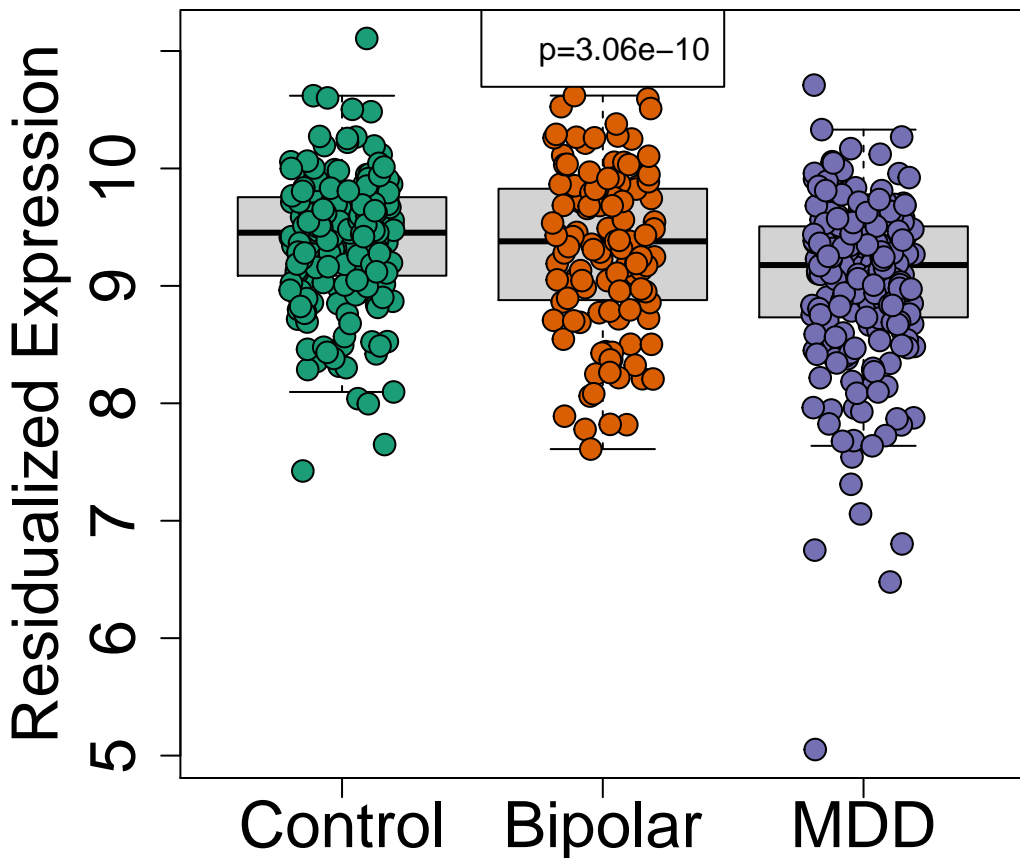
ENO1, InGen

chr1:8874918–8878579(-)



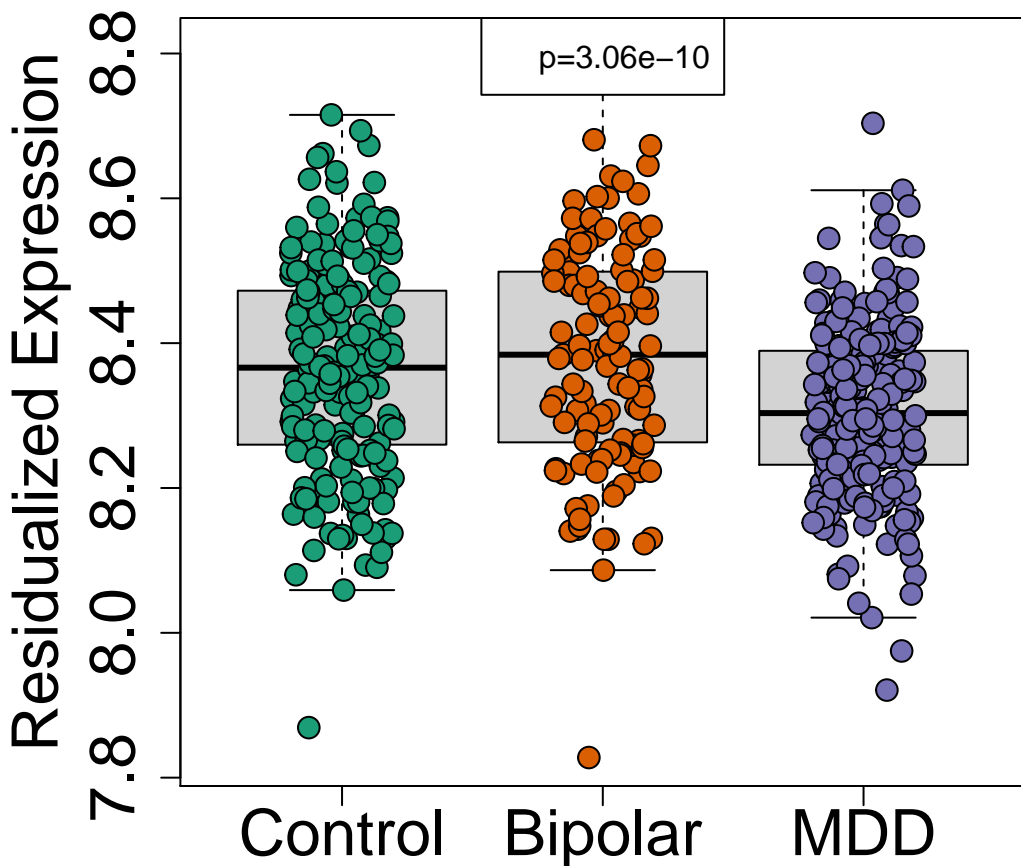
ANLN, InGen

chr7:36415885–36417079(+)



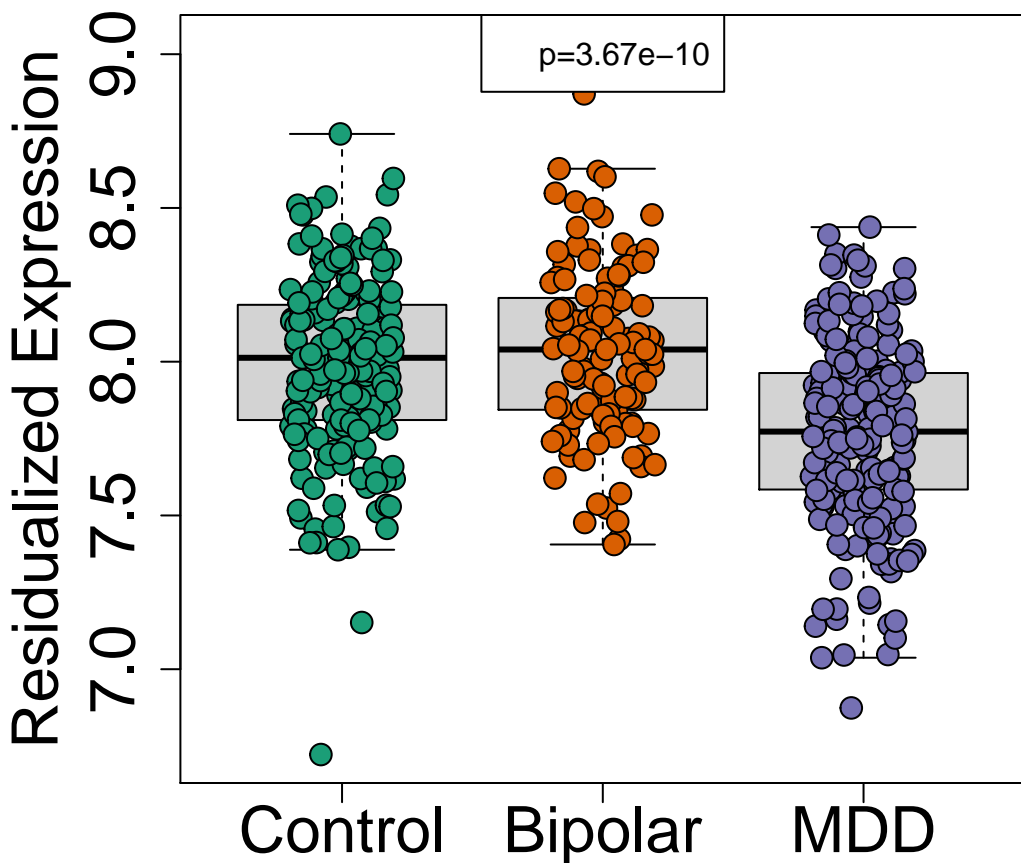
ATP9A, InGen

chr20:51629073–51639342(-)



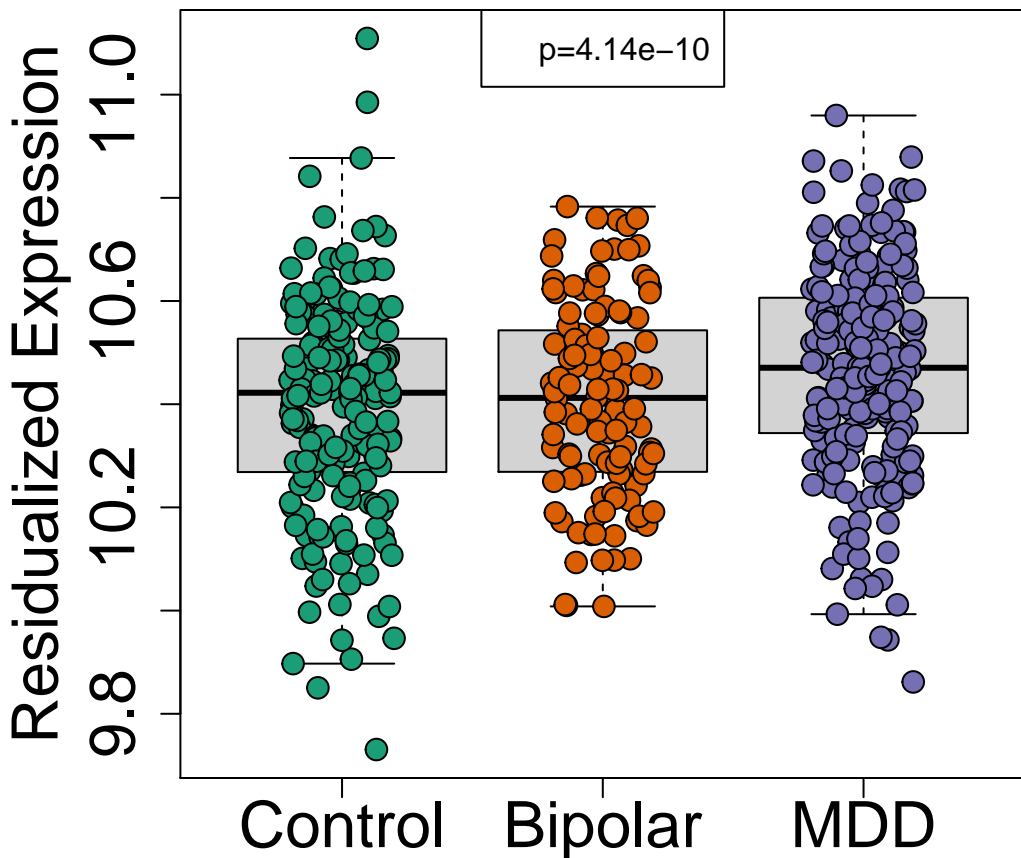
QDPR, InGen

chr4:17501860–17504378(-)



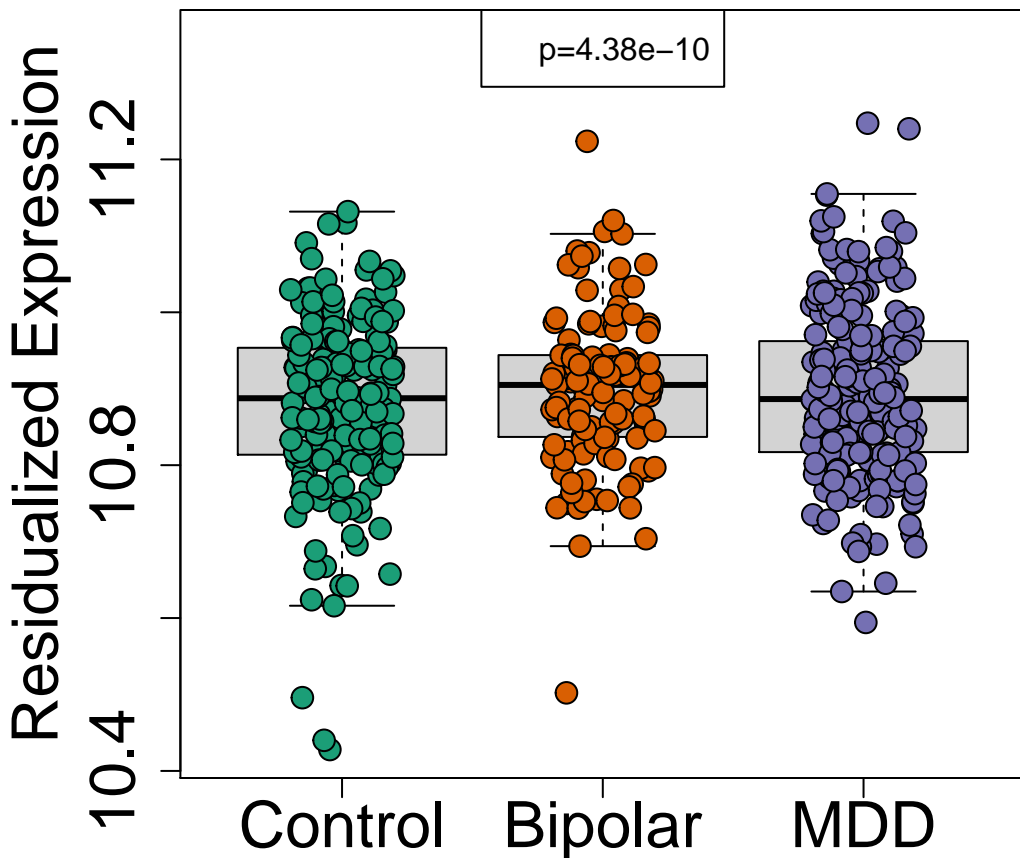
UCHL1, InGen

chr4:41257127–41257608(+)



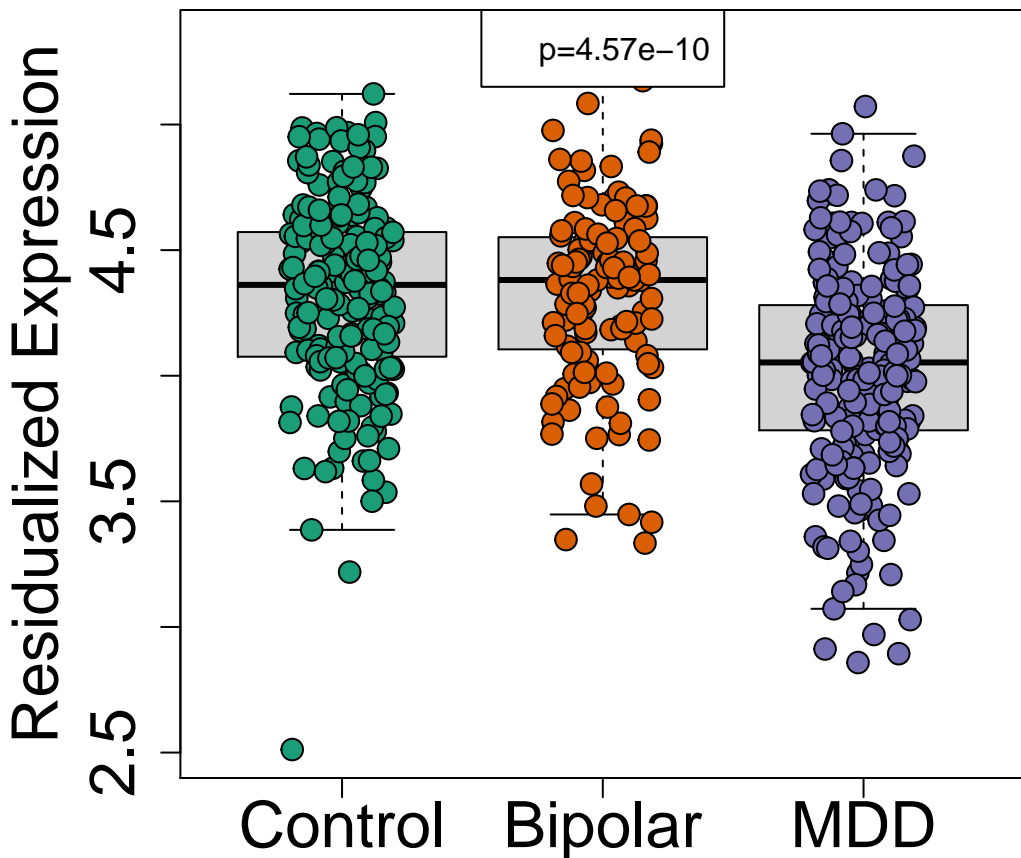
HSP90AB1, InGen

chr6:44249584–44249674(+)



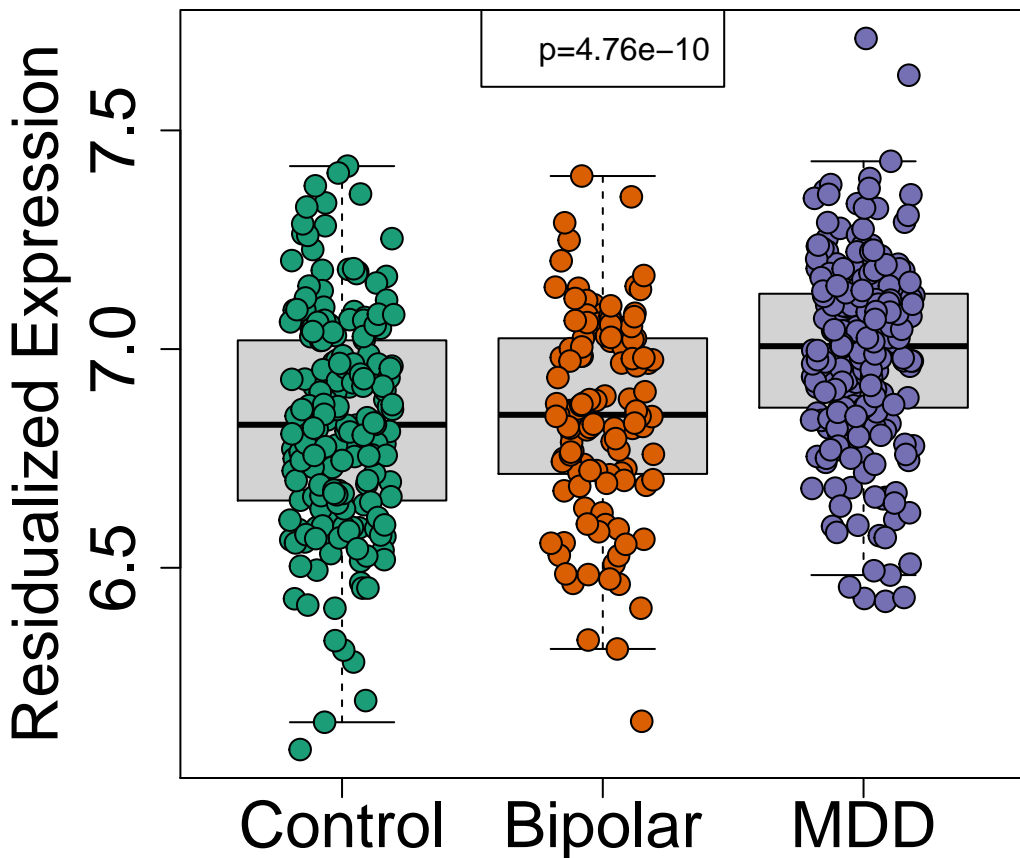
PPP4C, InGen

chr16:30076476–30081258(+)



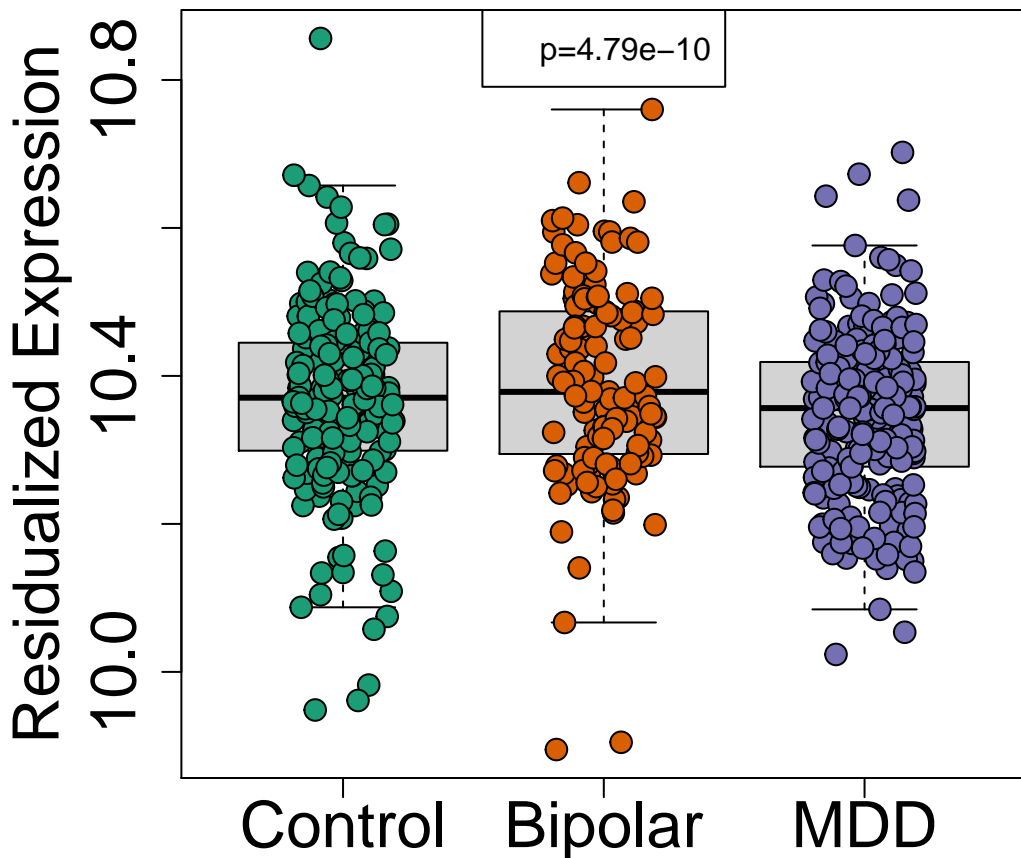
ATP1A1, InGen

chr1:116401007–116401129(+)



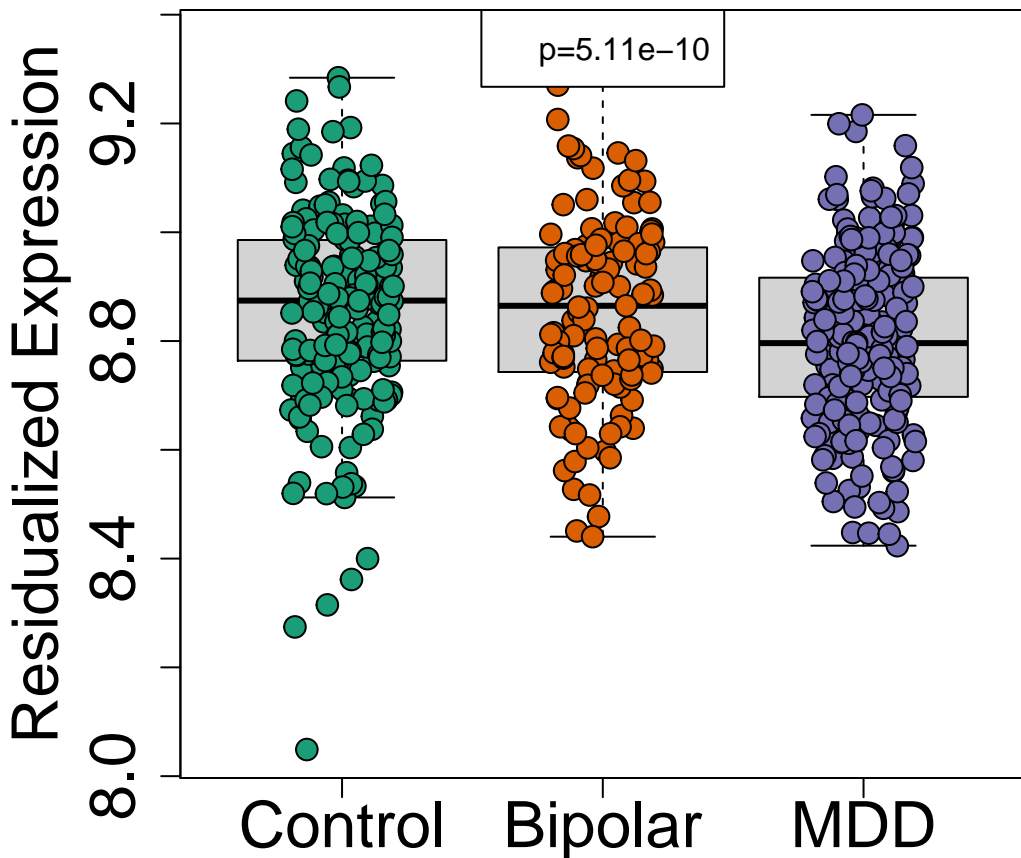
ACTB, InGen

chr7:5528720–5529160(-)



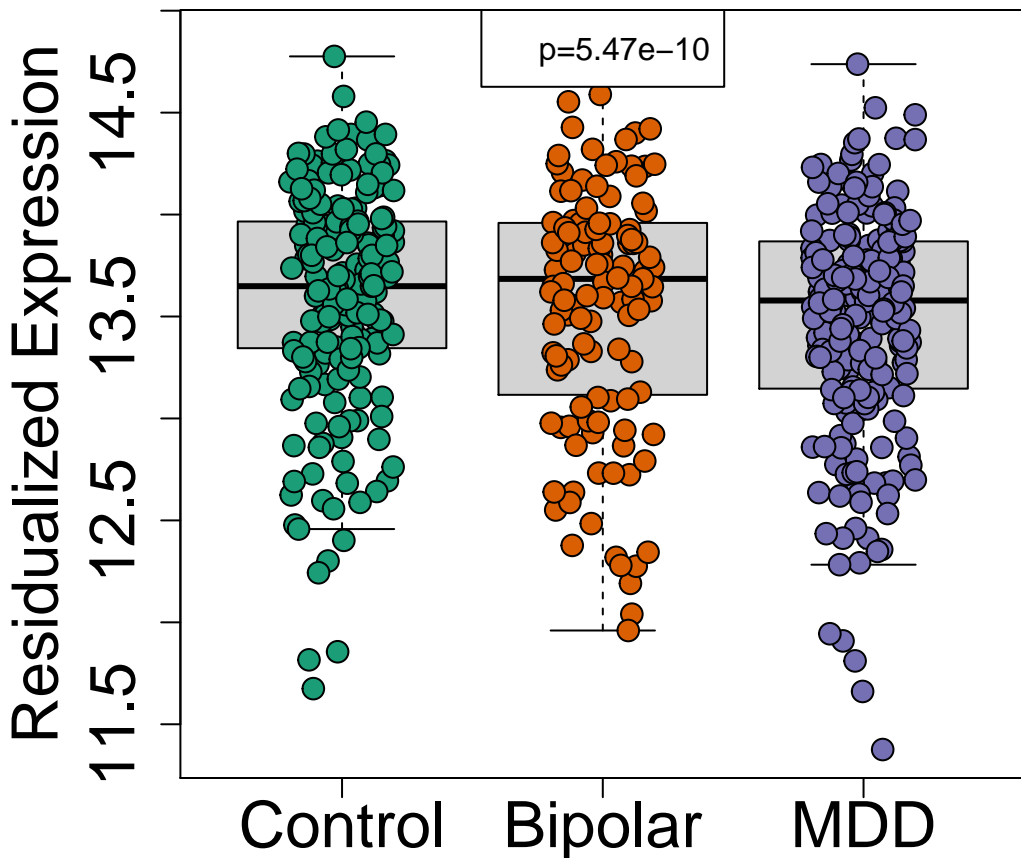
ATP1A3, InGen

chr19:41978799–41981501(-)



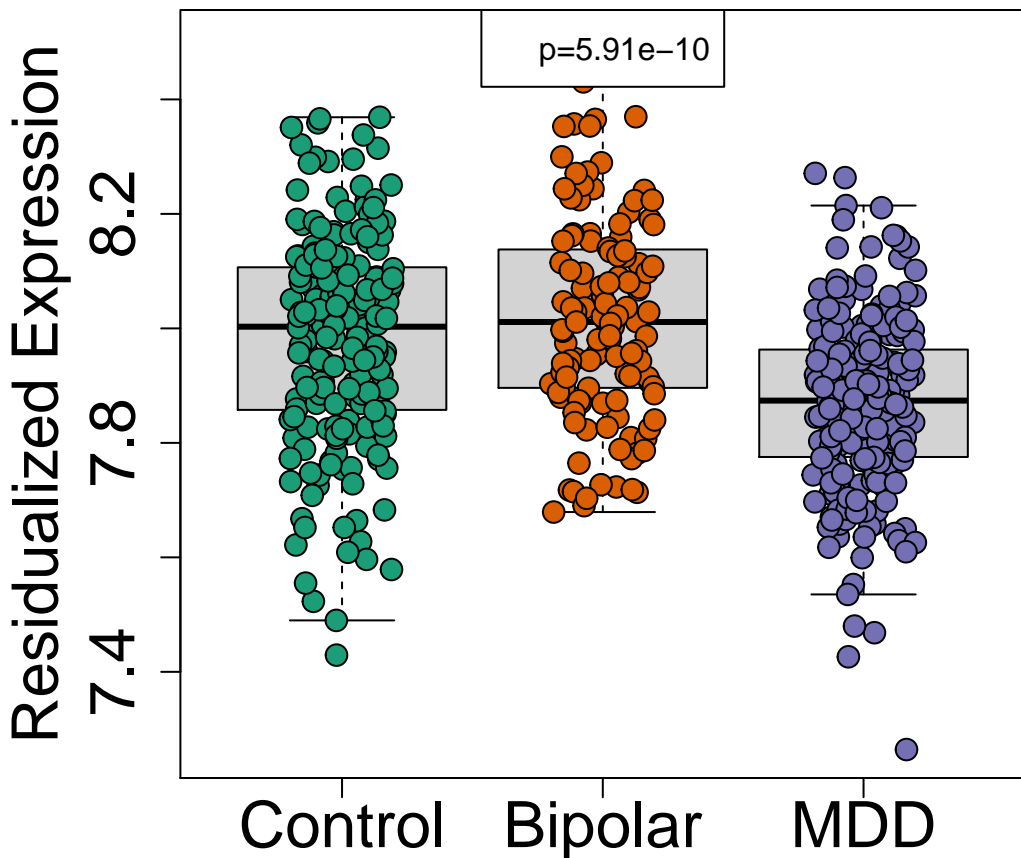
PLP1, InGen

chrX:103787967-103788436(+)



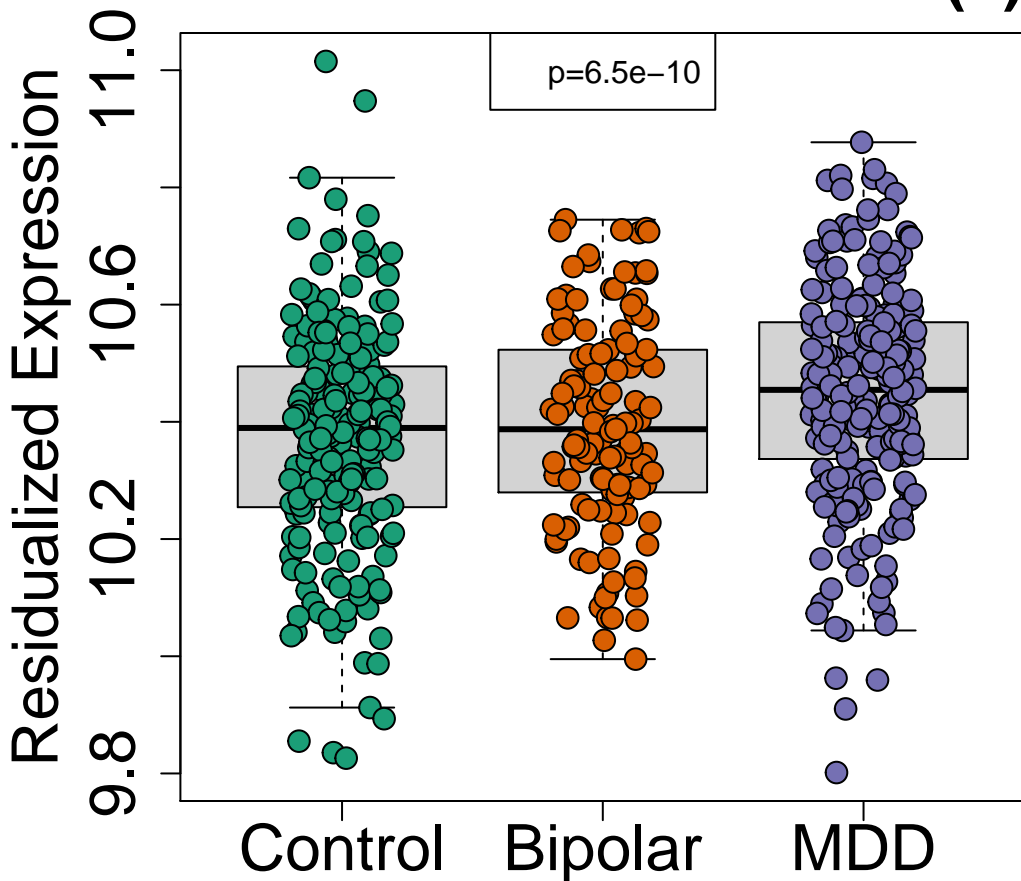
ATP2A2, InGen

chr12:110339722–110340658(+)



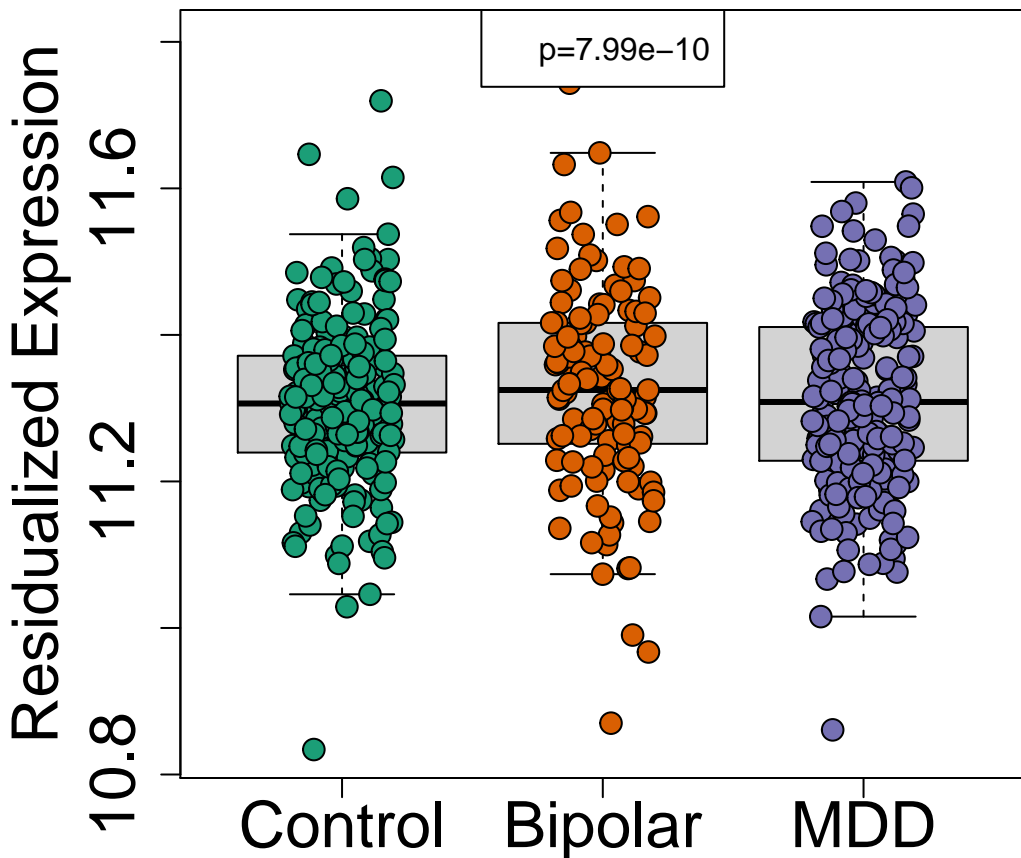
UCHL1, InGen

chr4:41257010–41257114(+)



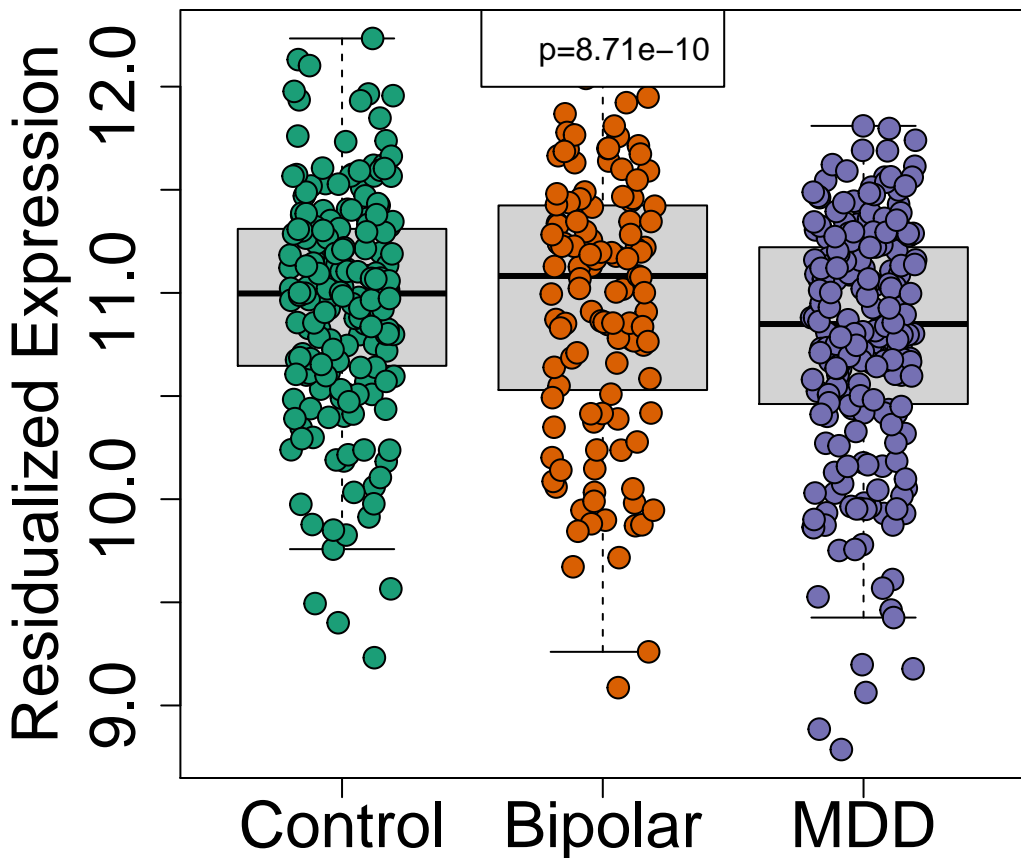
CPE, InGen

chr4:165493271–165495558(+)



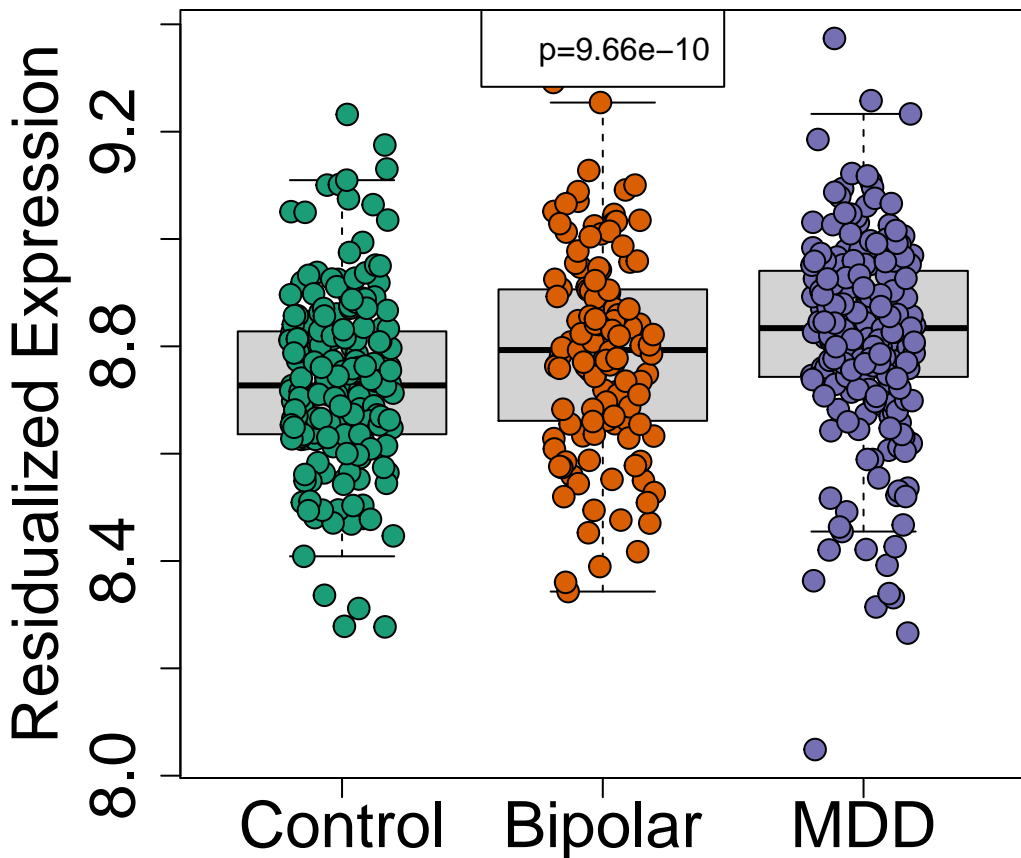
TF, InGen

chr3:133759330–133764181(+)



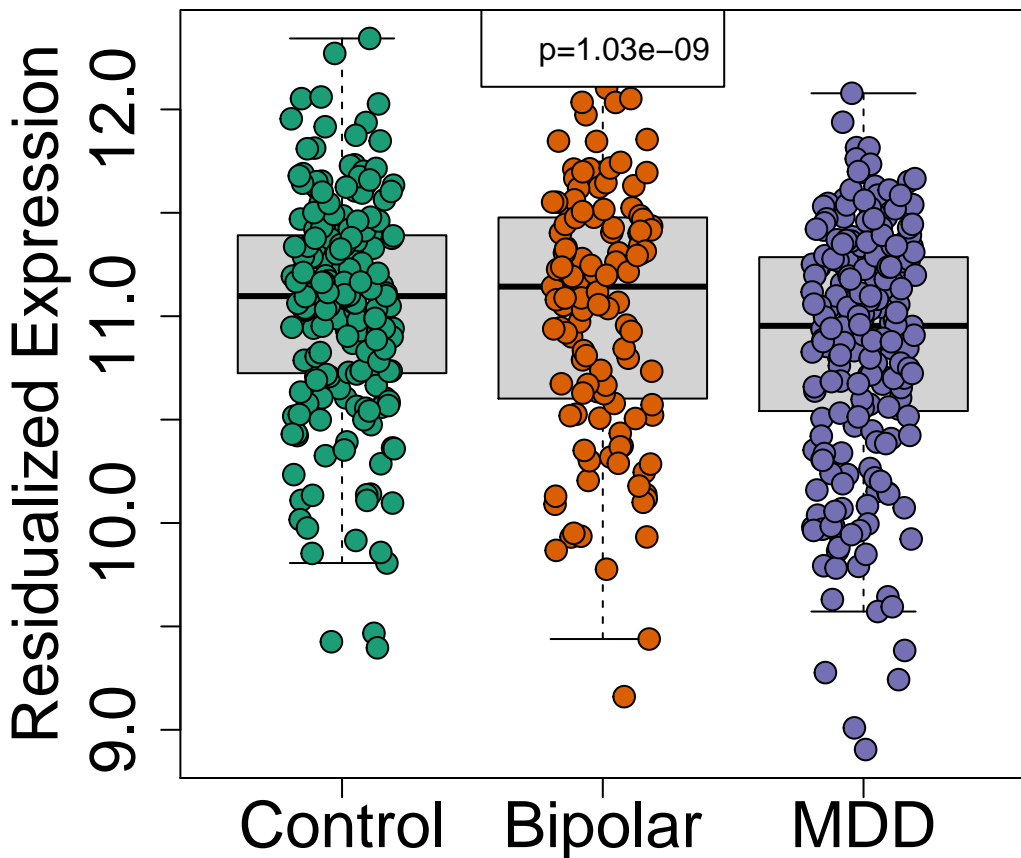
PKIA, InGen

chr8:78598536–78601741(+)



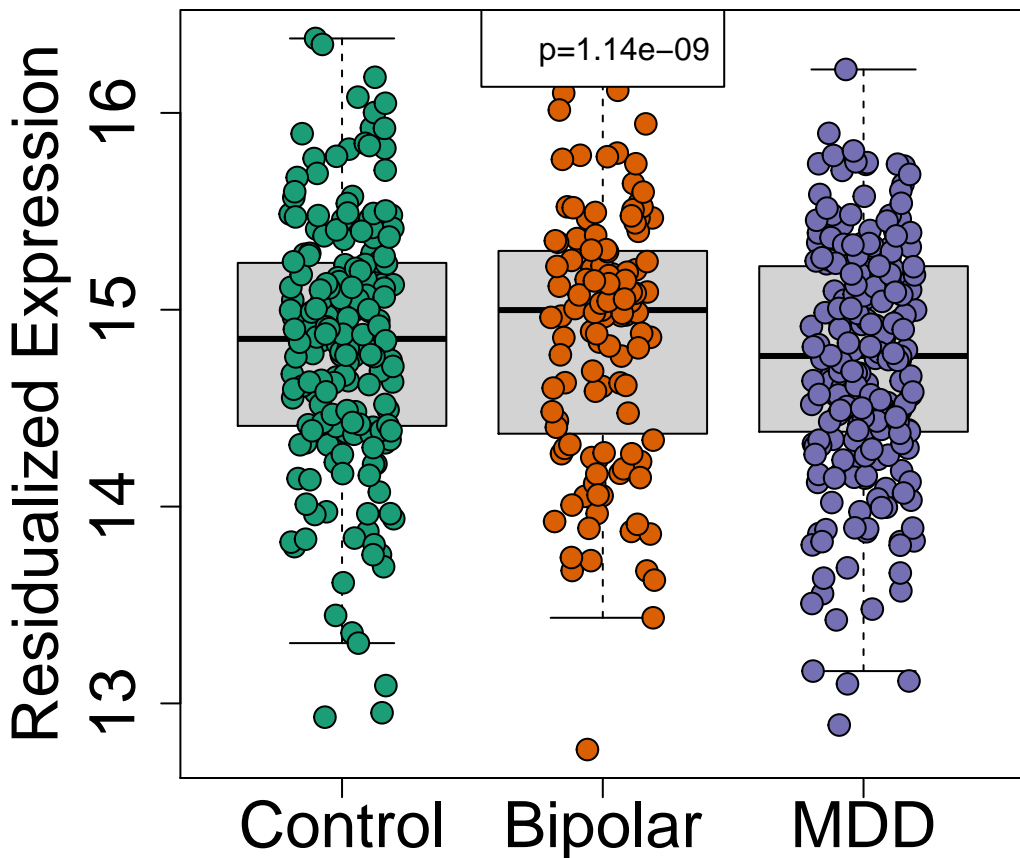
TF, InGen

chr3:133764276–133764874(+)



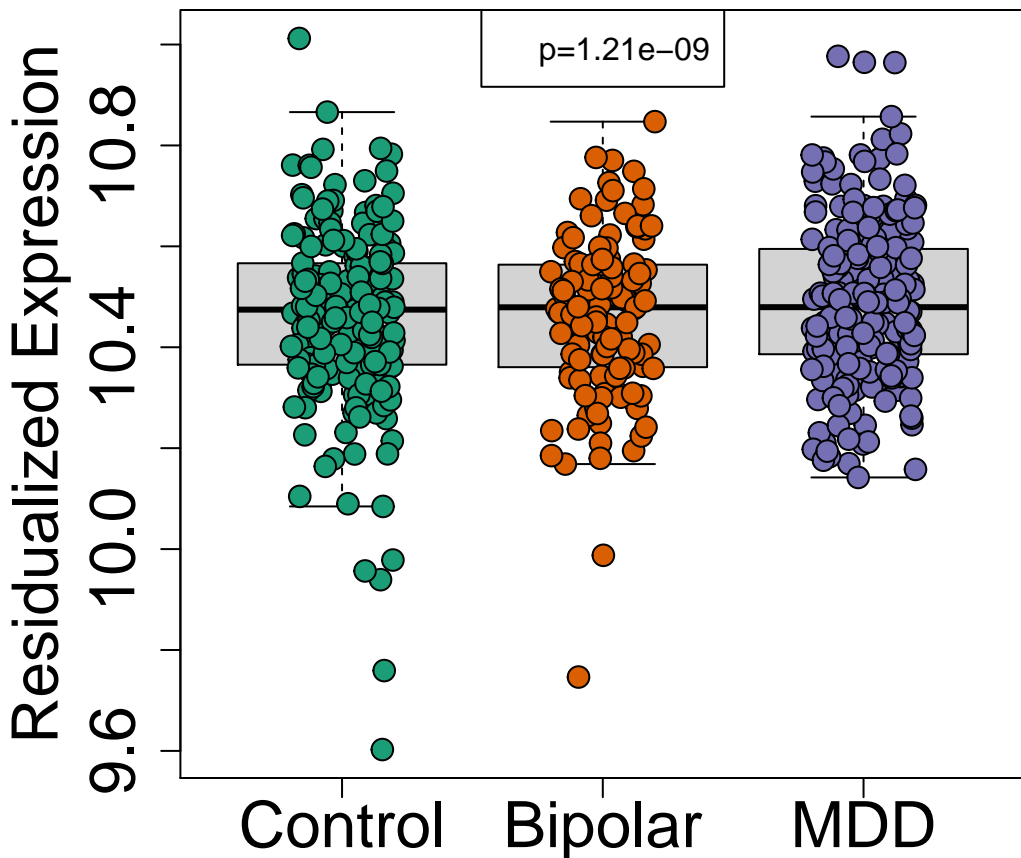
MBP, InGen

chr18:76988913–76989955(-)



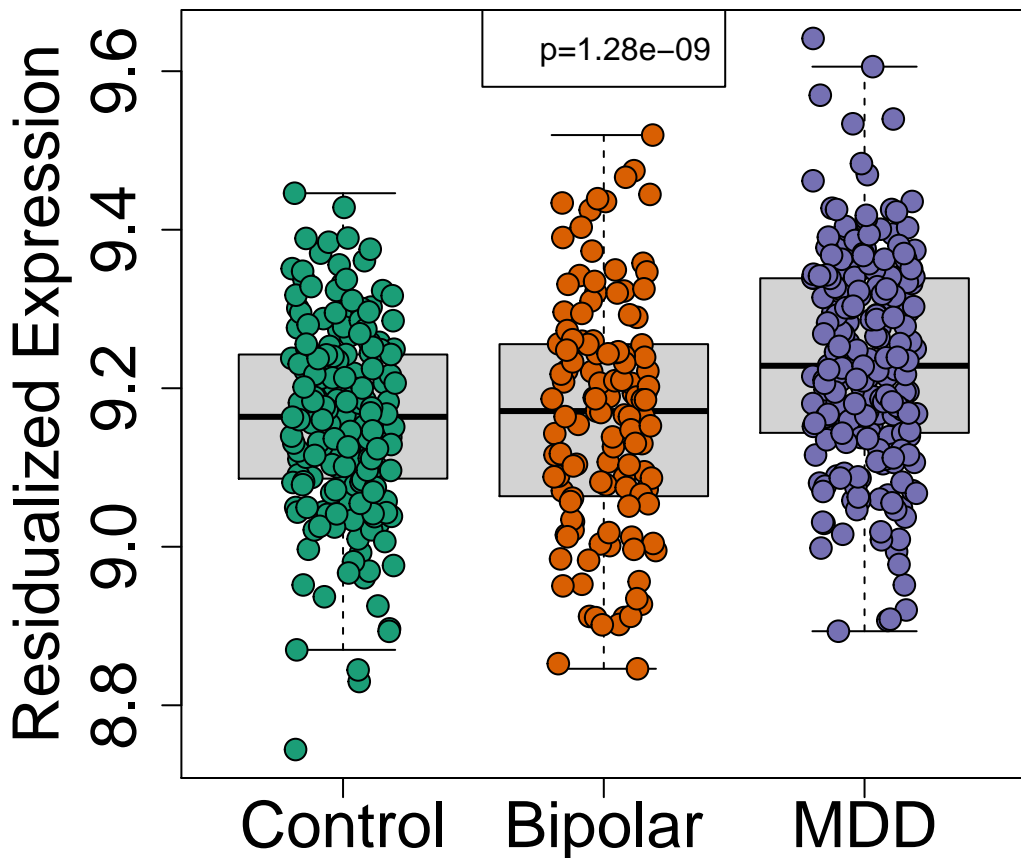
SNAP25, InGen

chr20:10218978–10275428(+)



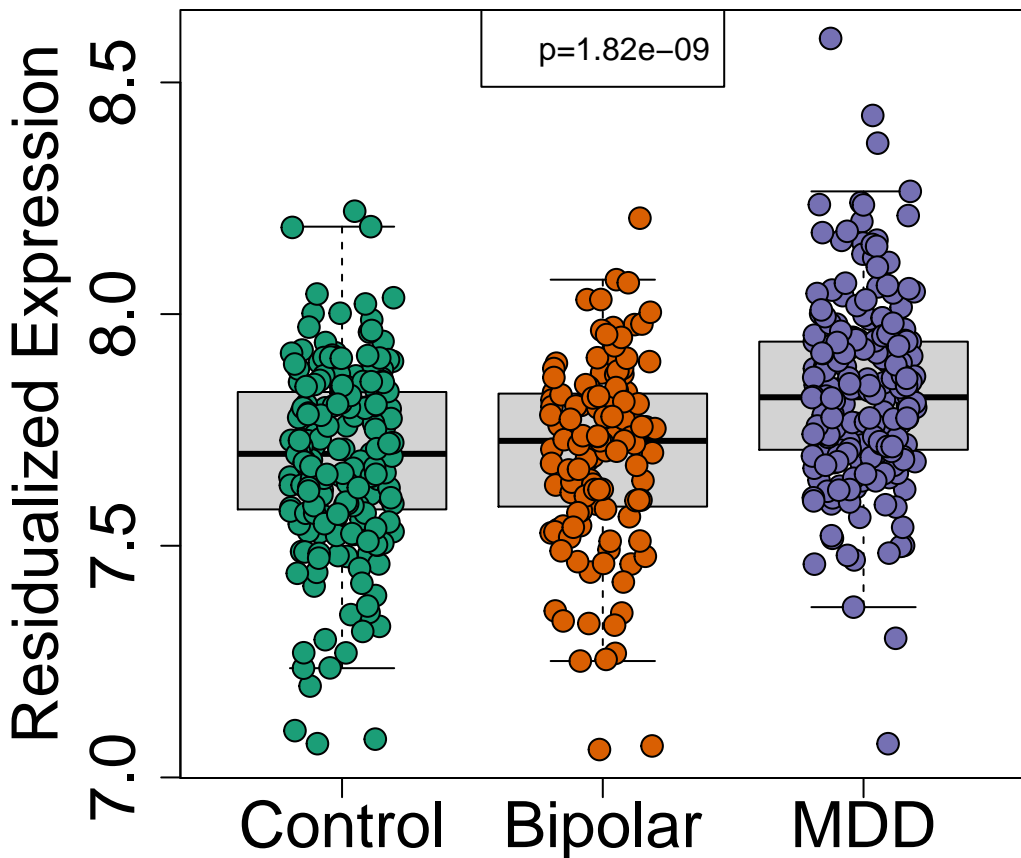
ENO1, InGen

chr1:8862946–8863234(-)



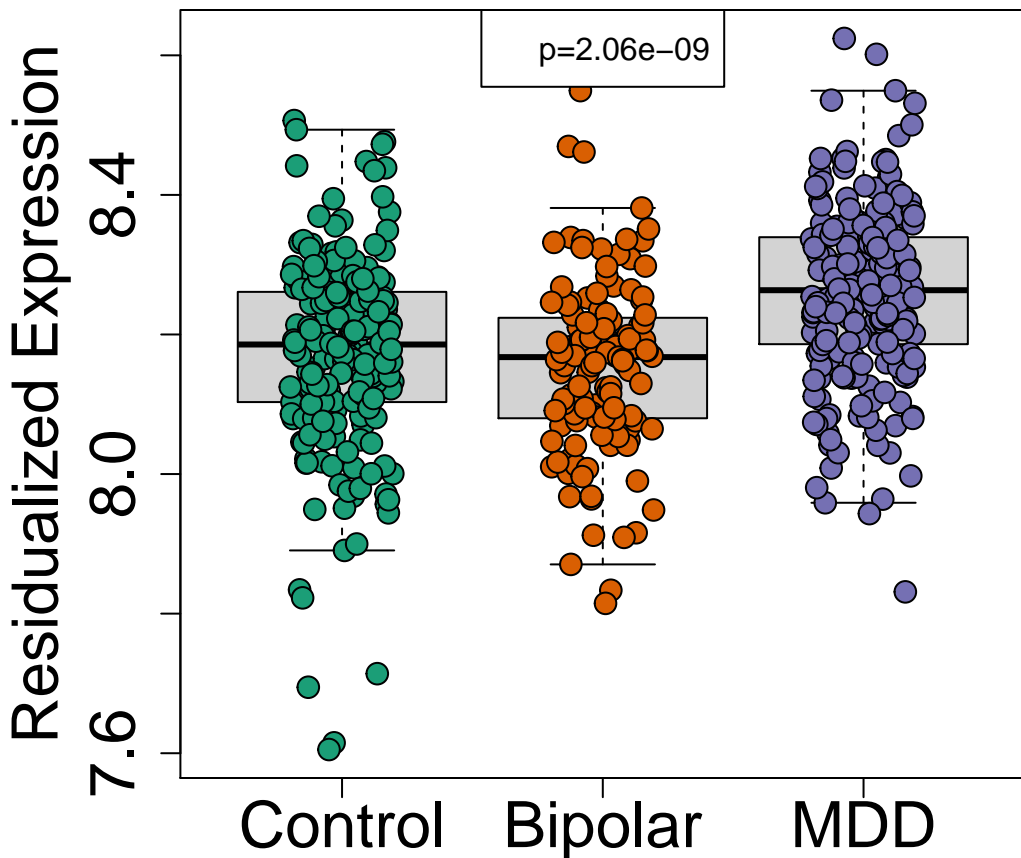
GNG3, InGen

chr11:62708395–62708677(+)



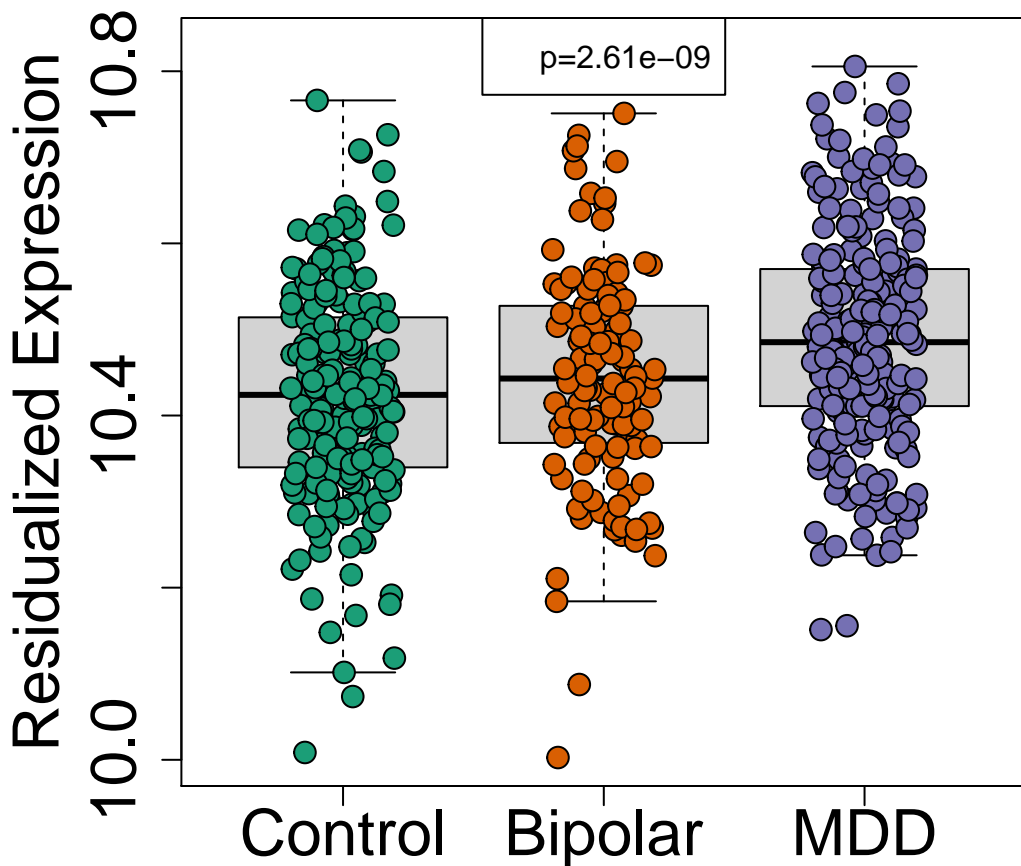
RAN, InGen

chr12:130875783–130875881(+)



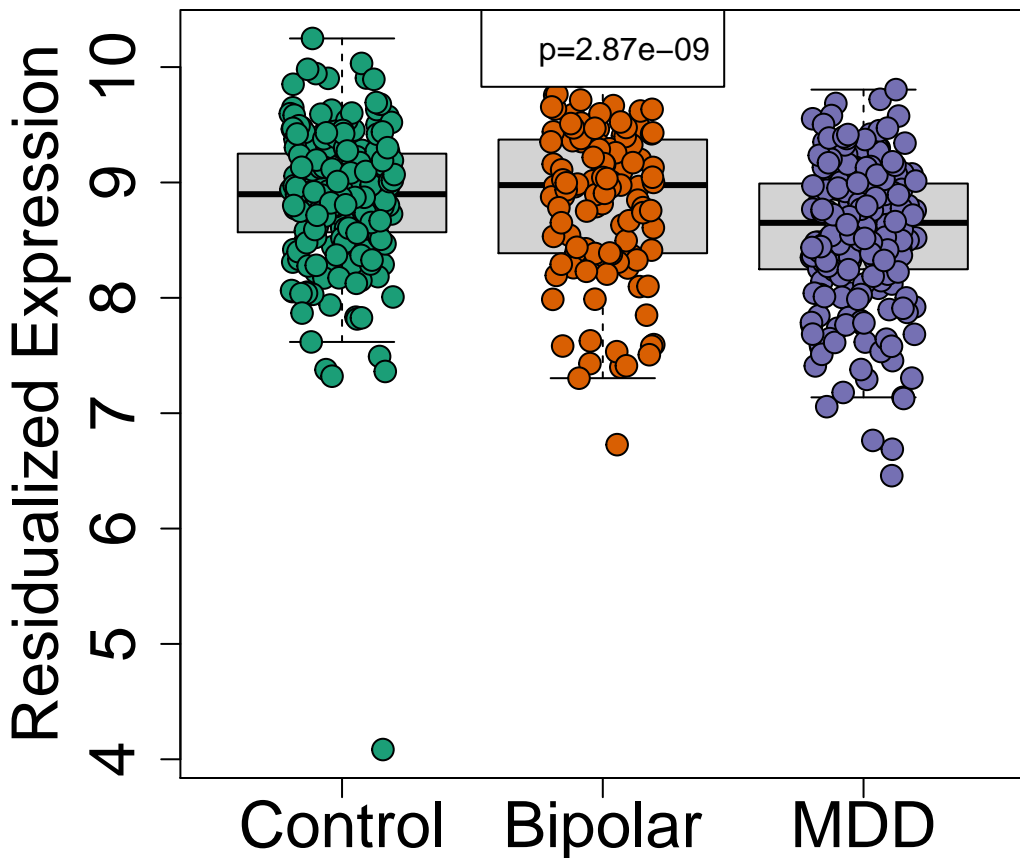
RPL22, InGen

chr1:6193055–6197651(-)



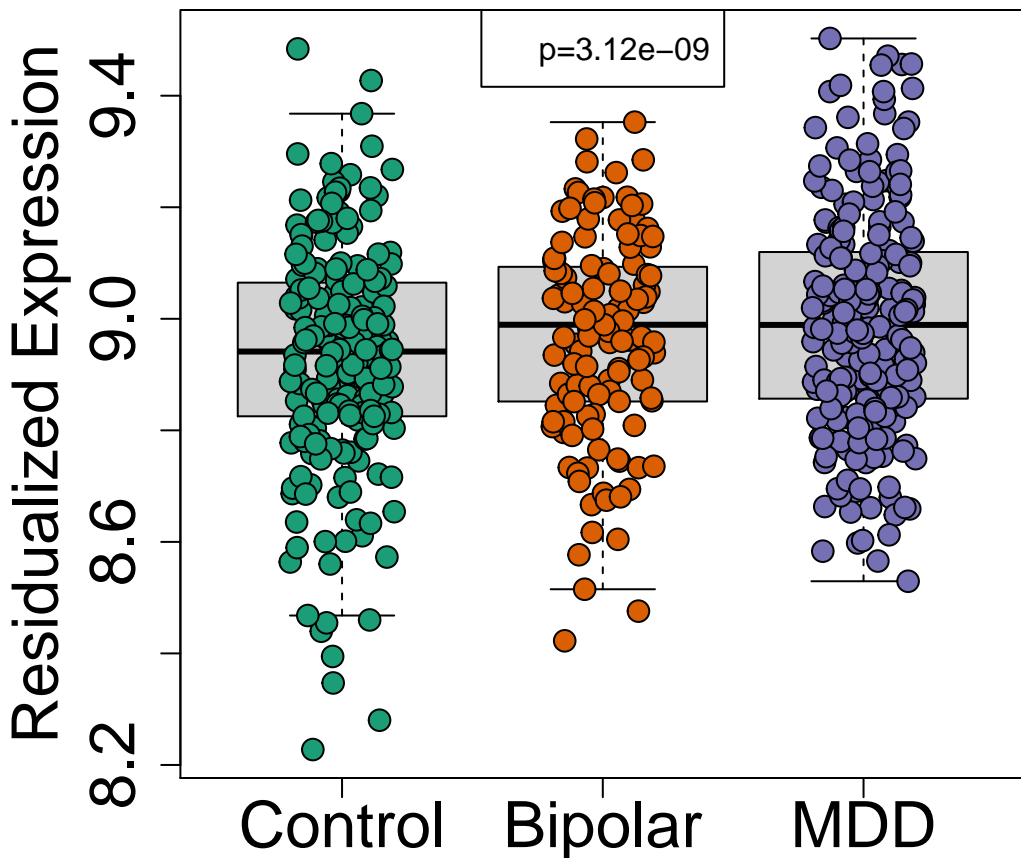
ANLN, InGen

chr7:36420315–36420596(+)



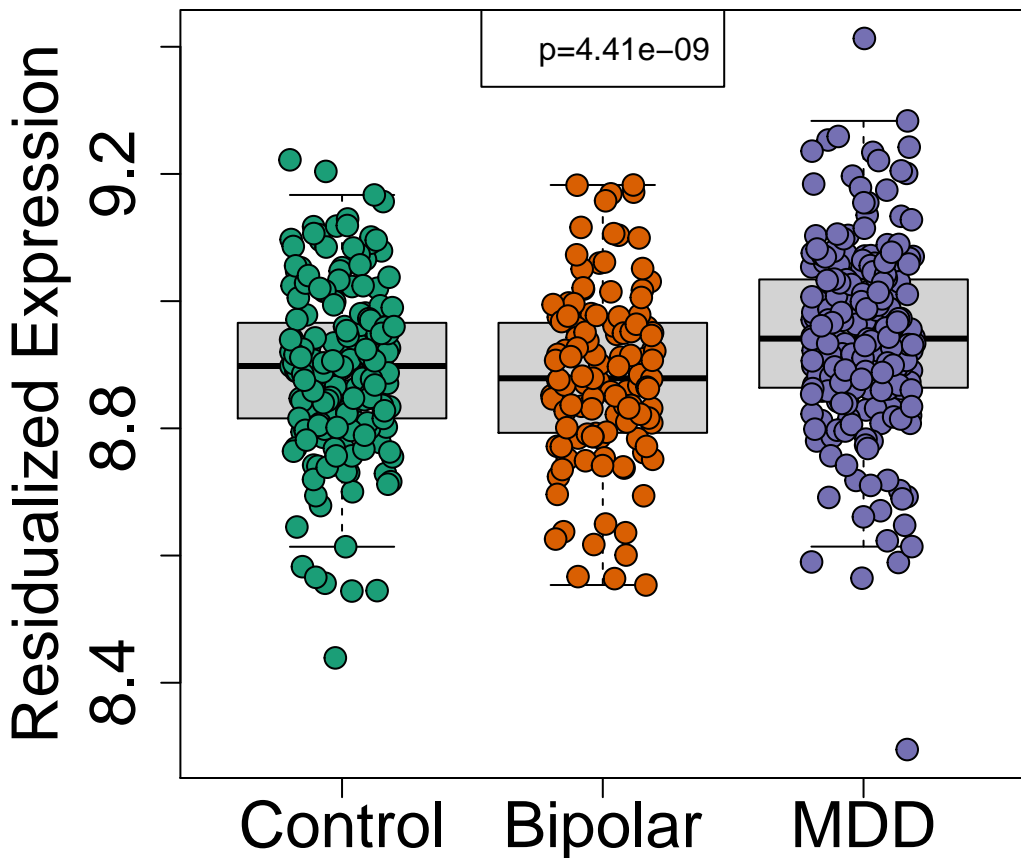
SNCA, InGen

chr4:89822389–89828142(-)



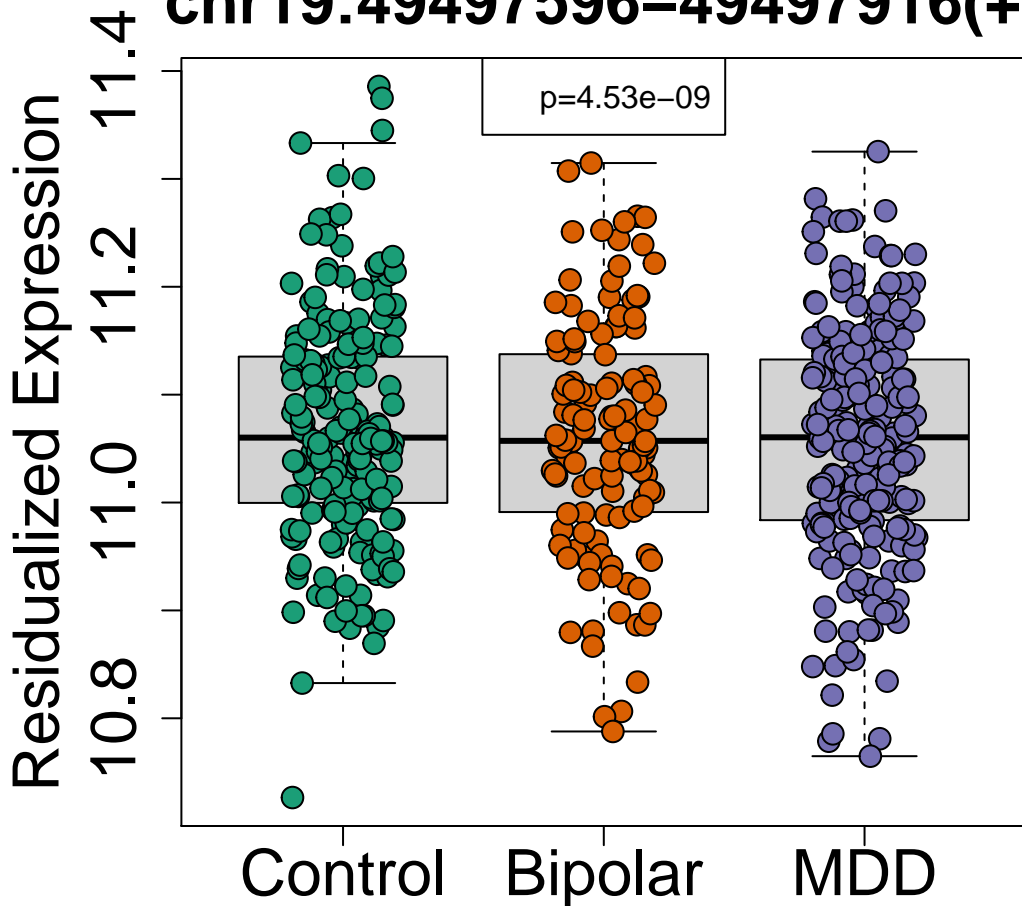
PGK1, InGen

chrX:78109918–78113743(+)



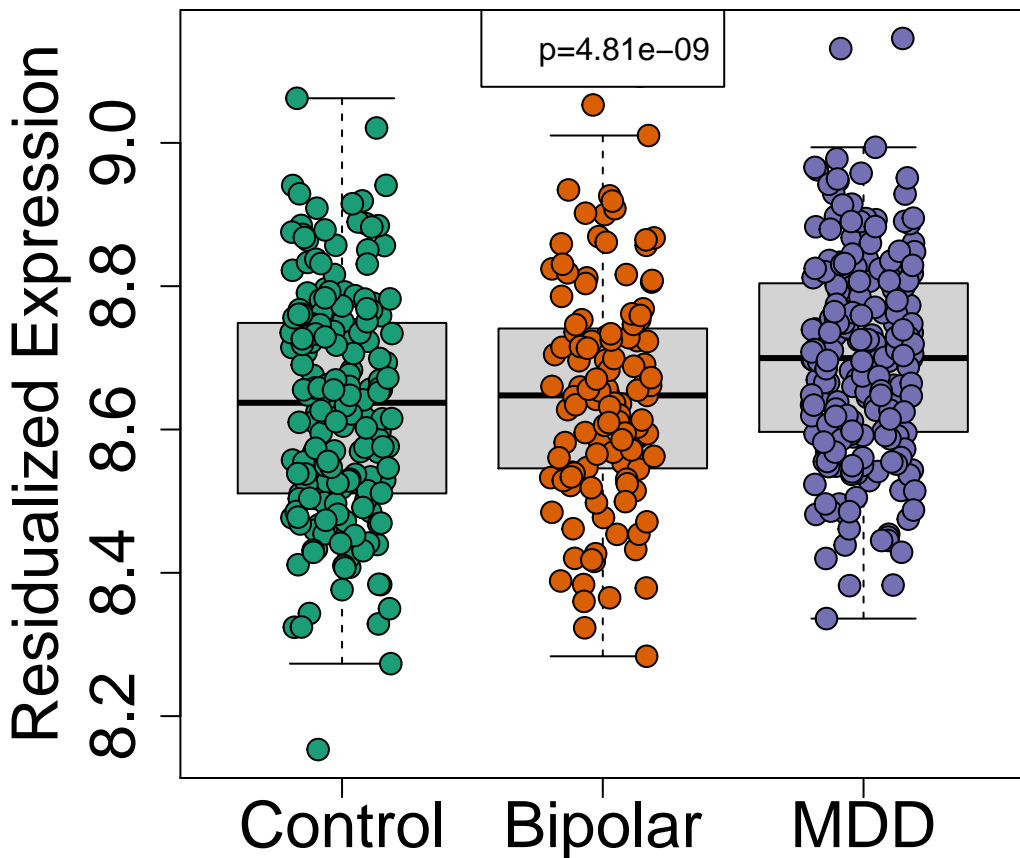
RPS11, InGen

chr19:49497596–49497916(+)



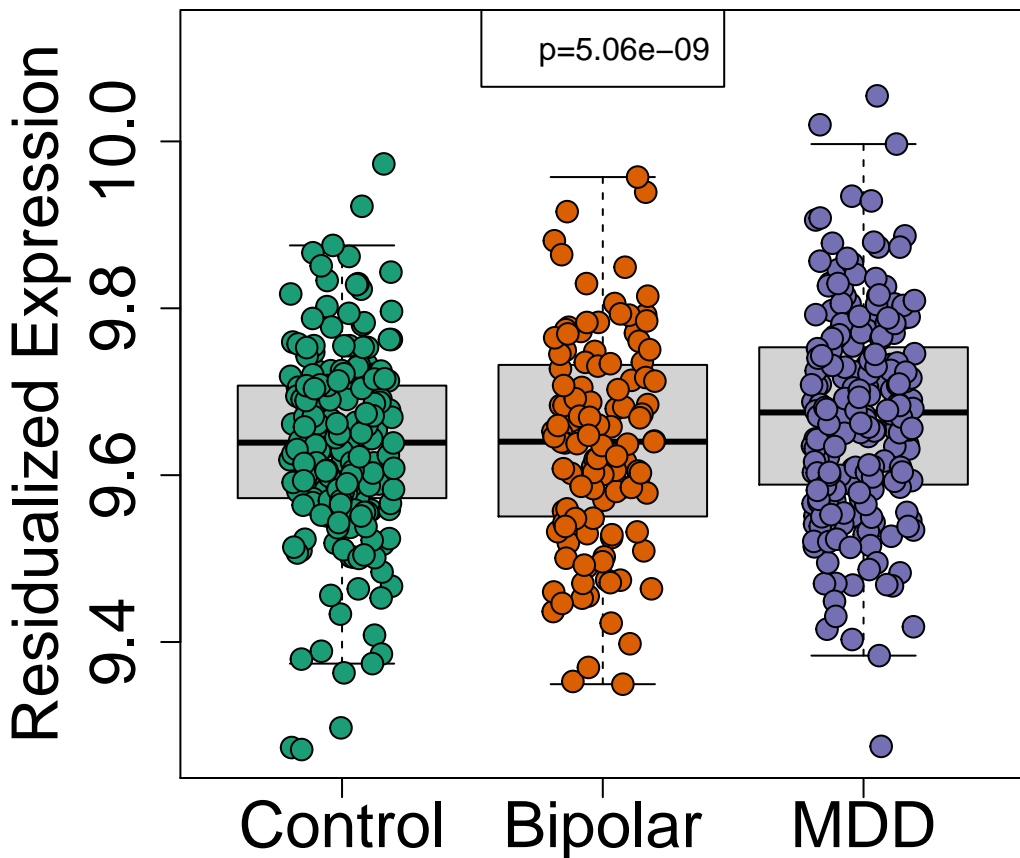
AZIN1, InGen

chr8:102828679–102829271(-)



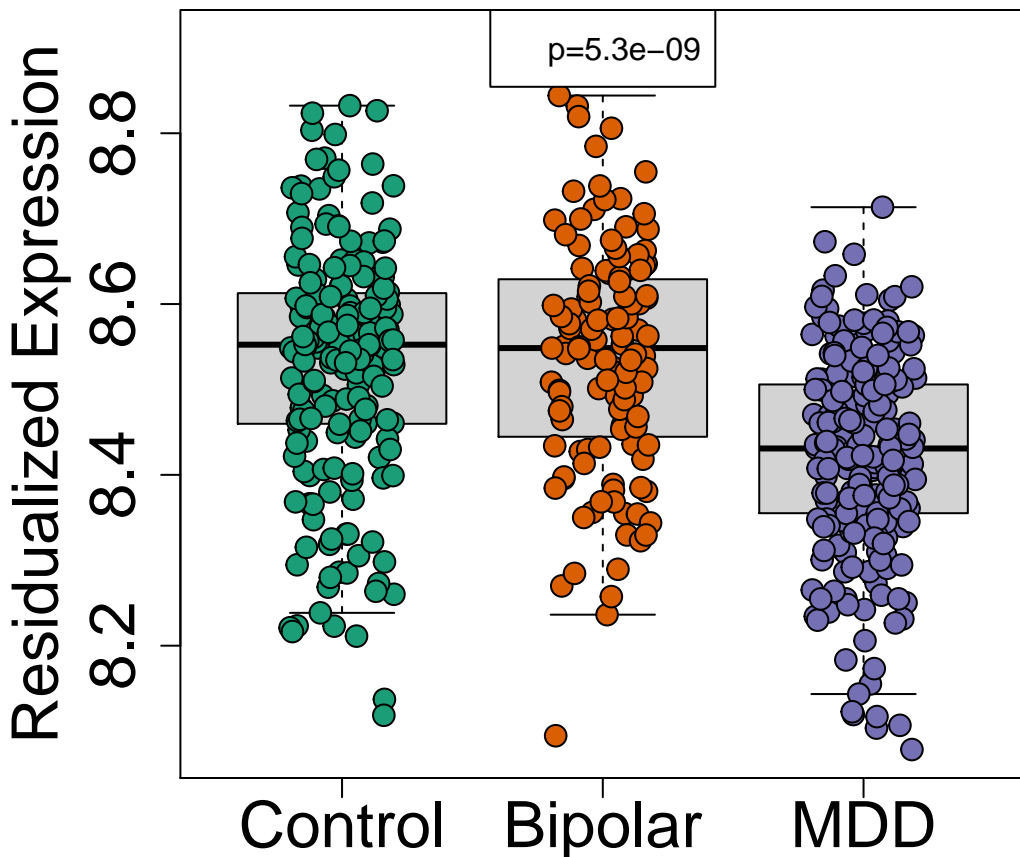
ENO1, InGen

chr1:8871987–8874823(-)



DYNC1H1, InGen

chr14:102005237-102005887(+)



ATP1B1, InGen

chr1:169125040–169127223(+)

