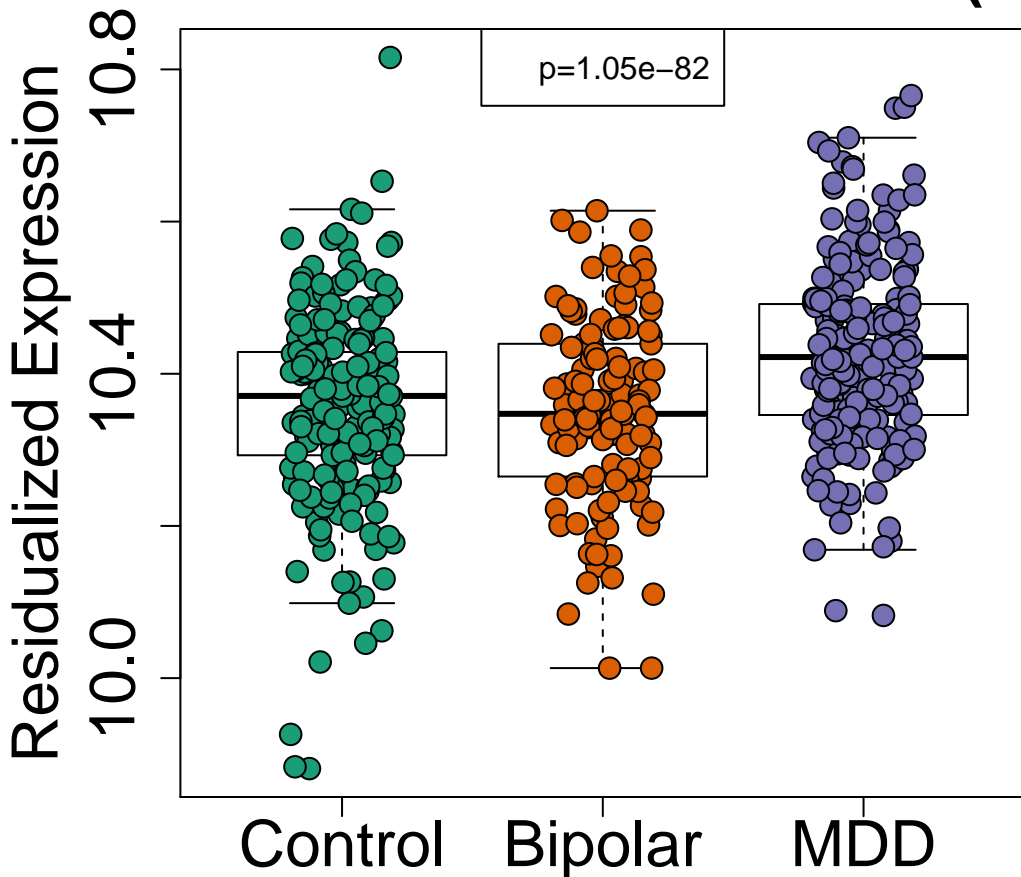


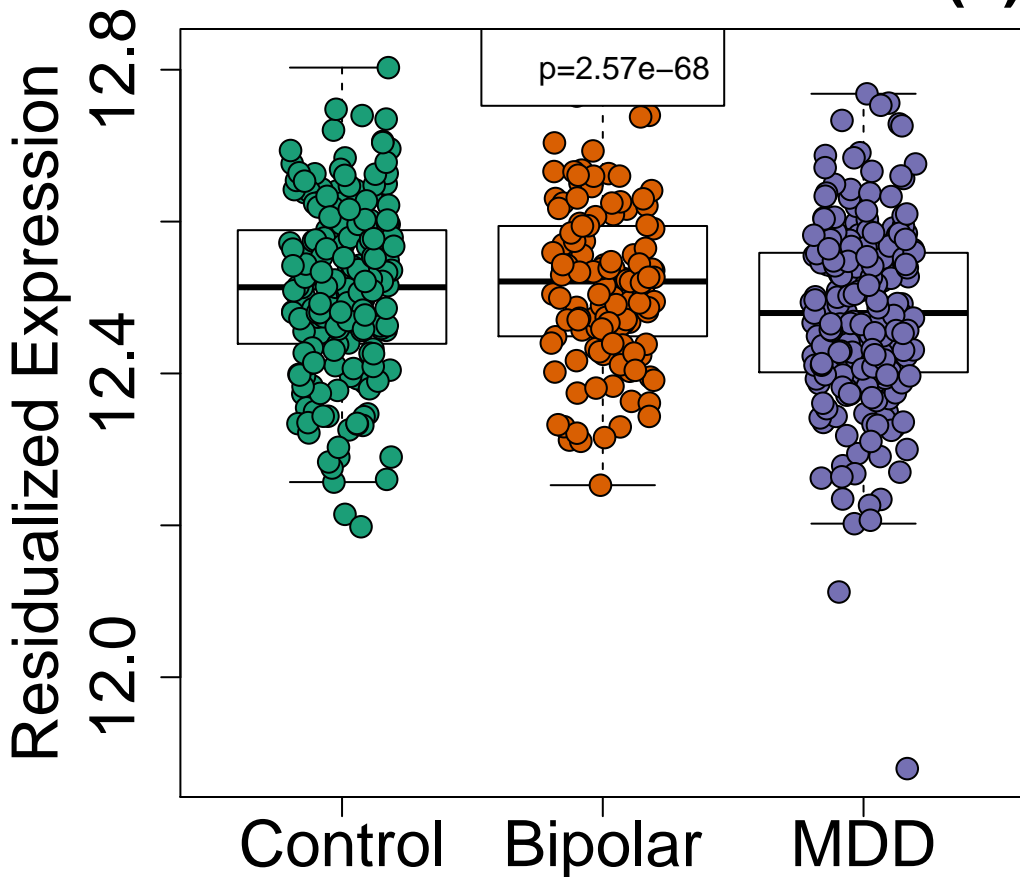
CALM3, InGen

chr19:46605858–46608196(+)



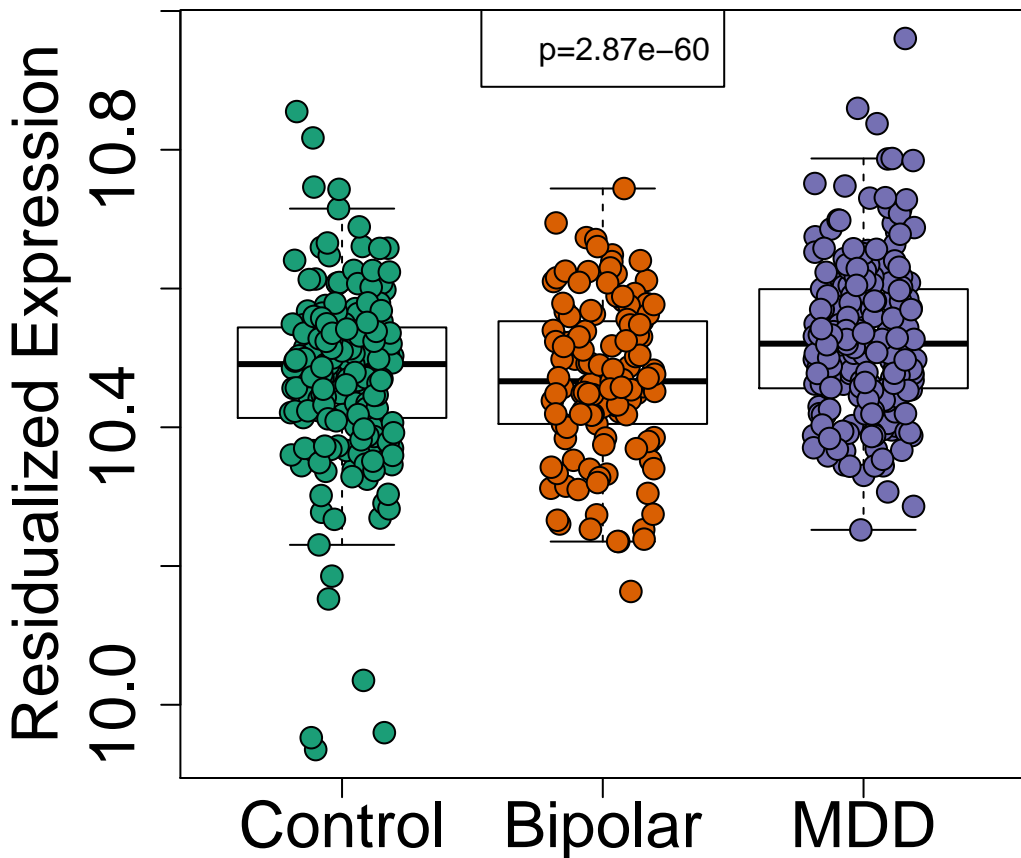
CALM2, InGen

chr2:47162393-47162518(-)



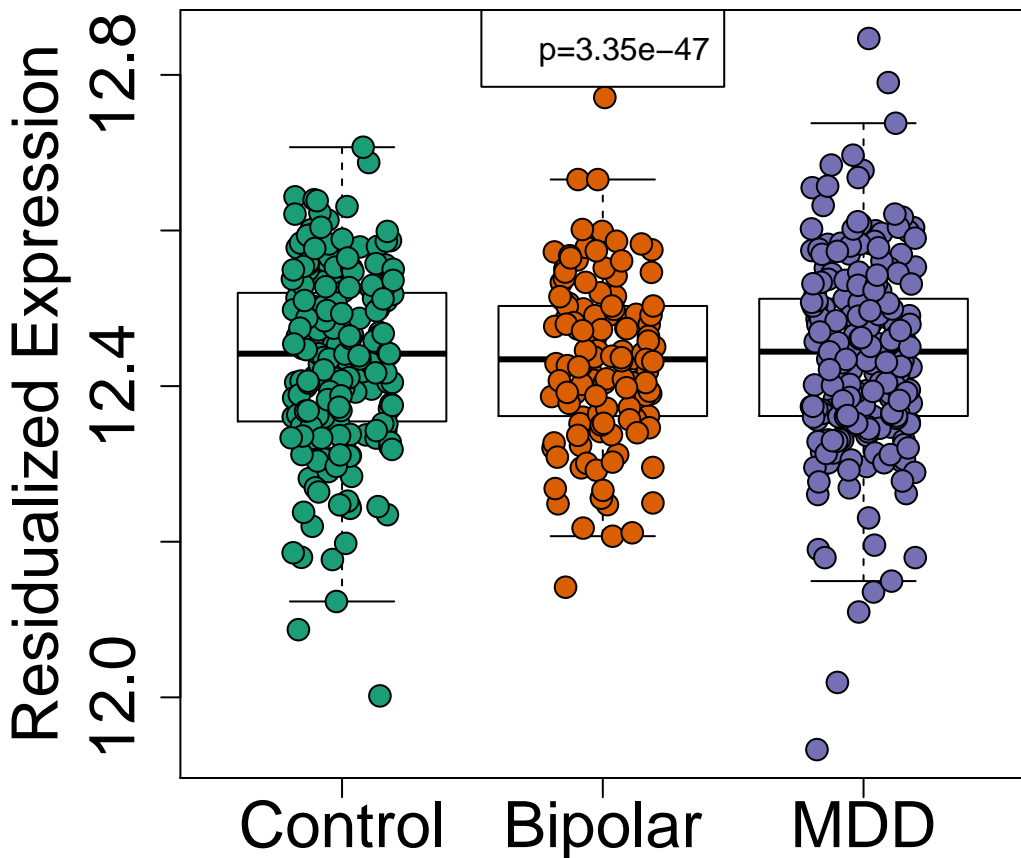
CALM3, InGen

chr19:46601438–46605826(+)



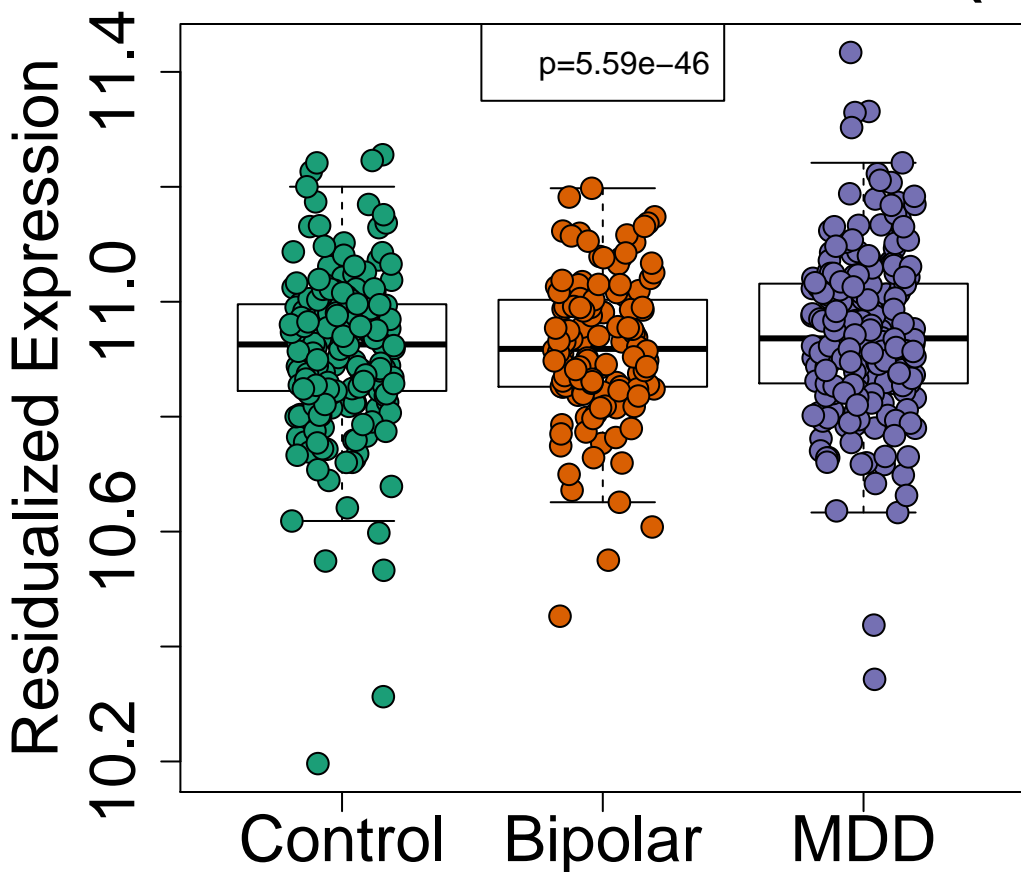
CALM2, InGen

chr2:47162663–47170733(-)

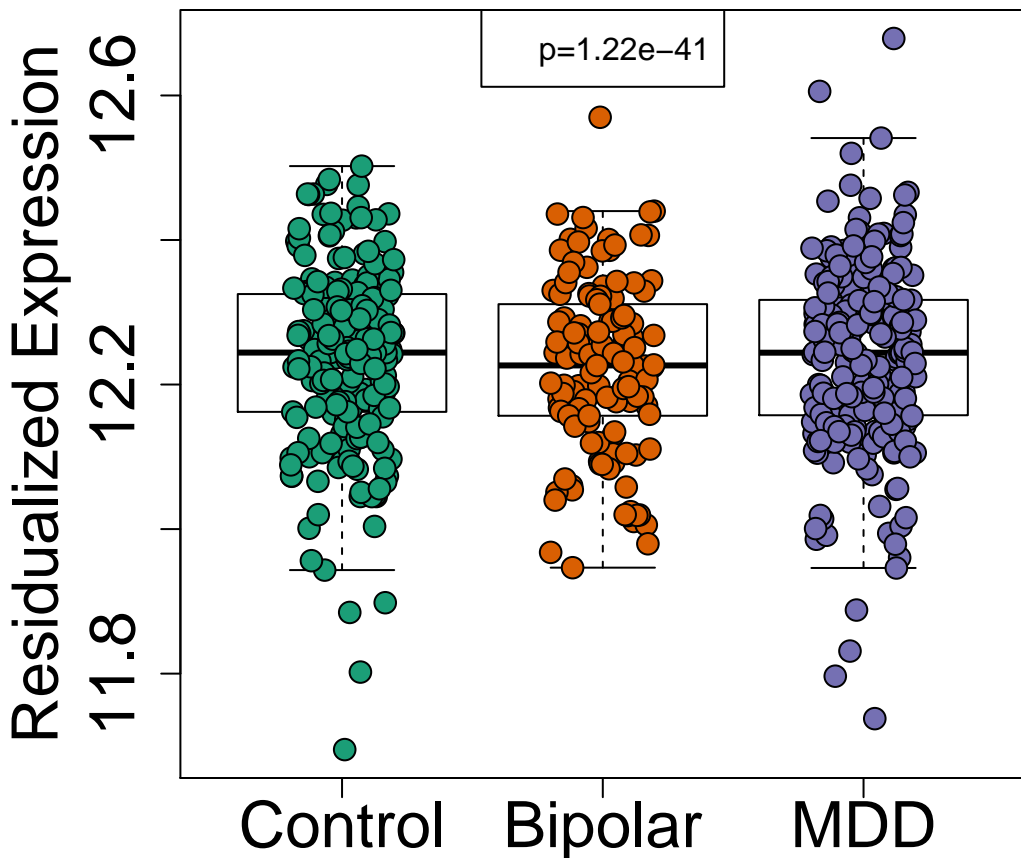


CALM3, InGen

chr19:46608982–46609124(+)

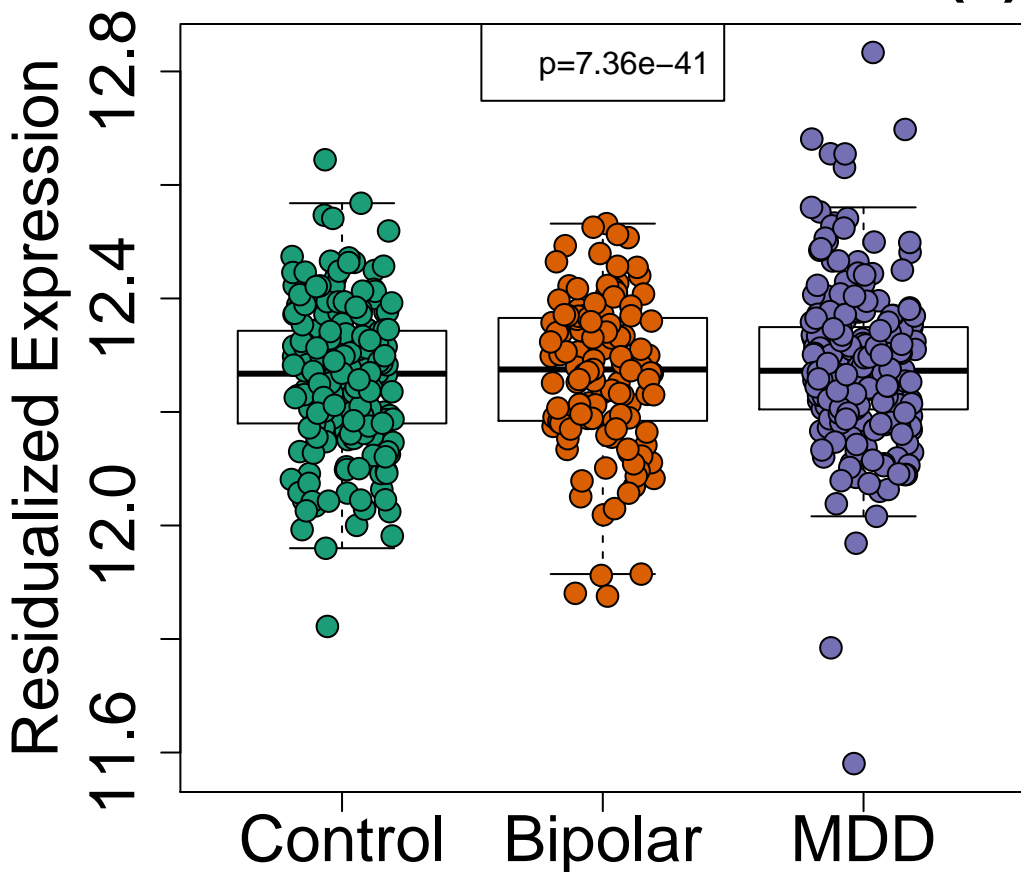


RP11-761B3.1, InGen
chr2:47170765-47176440(-)



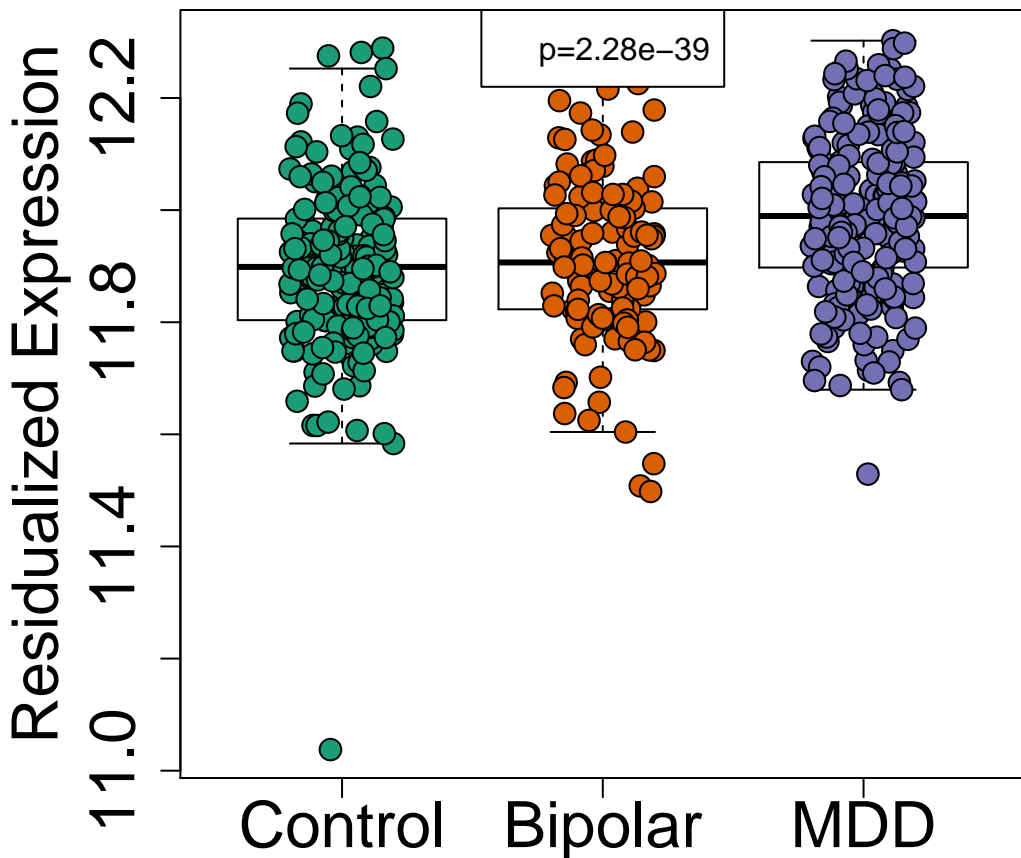
CALM2, InGen

chr2:47161859–47162285(-)



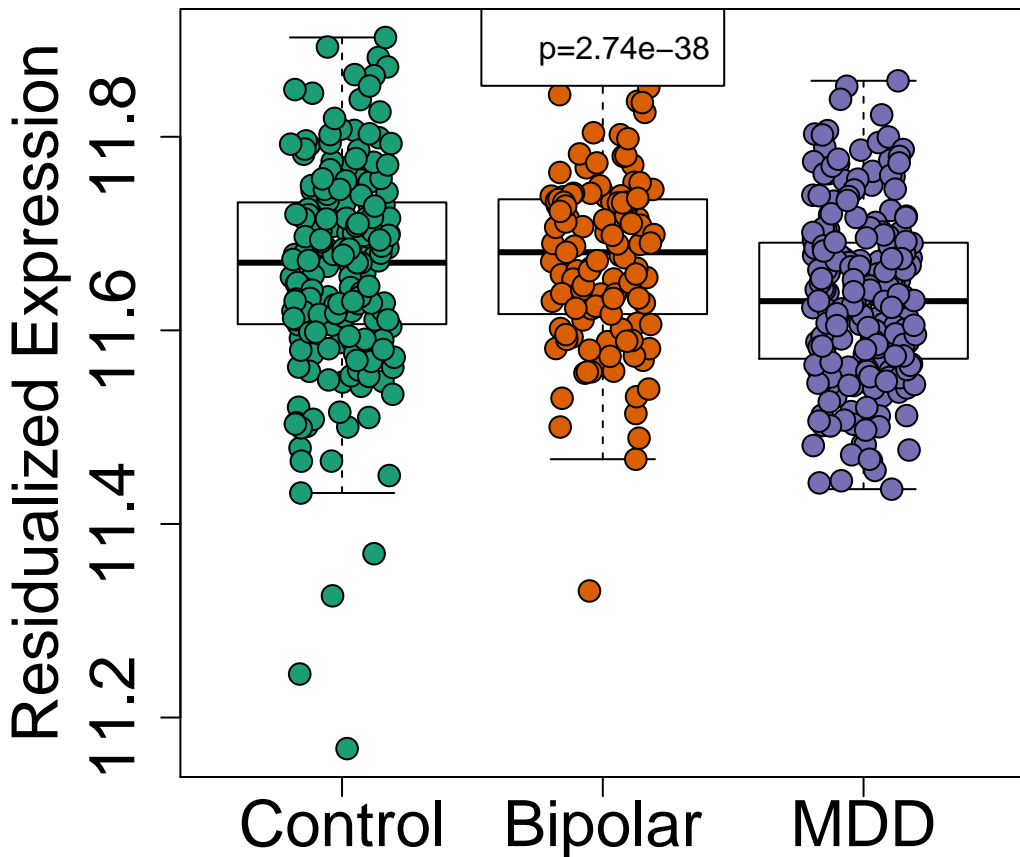
TMSB10, InGen

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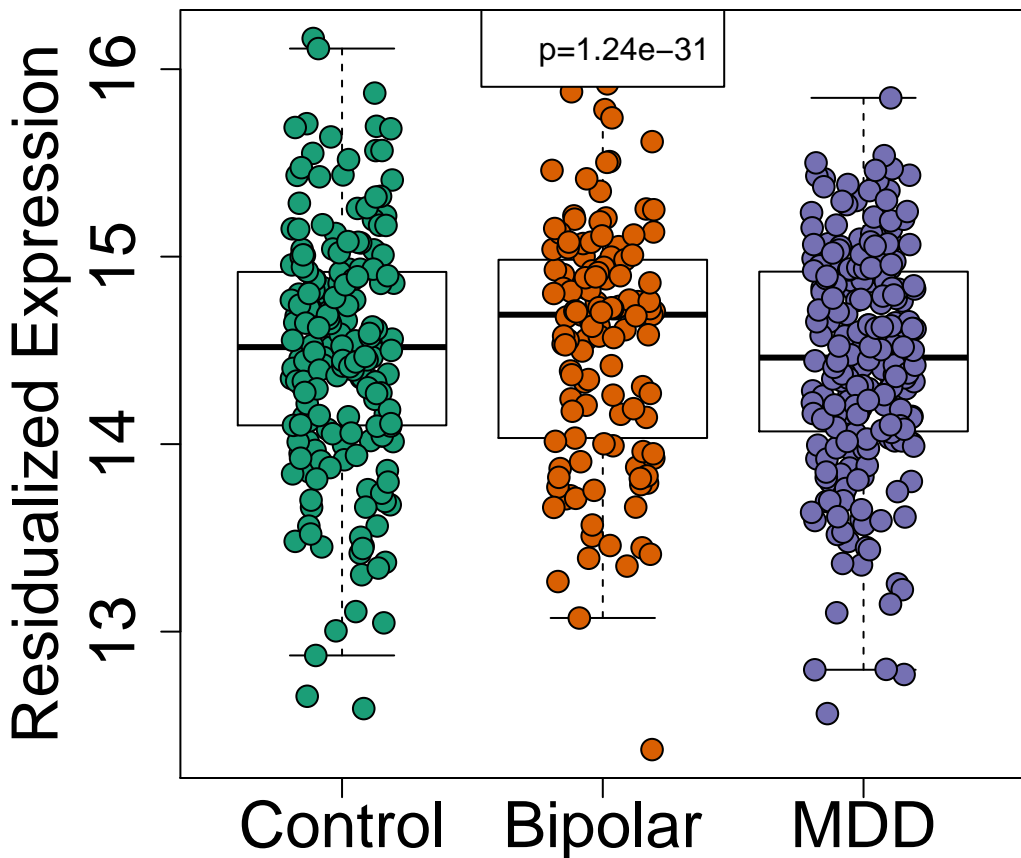
RPL37A, InGen

chr2:216500032–216501340(+)



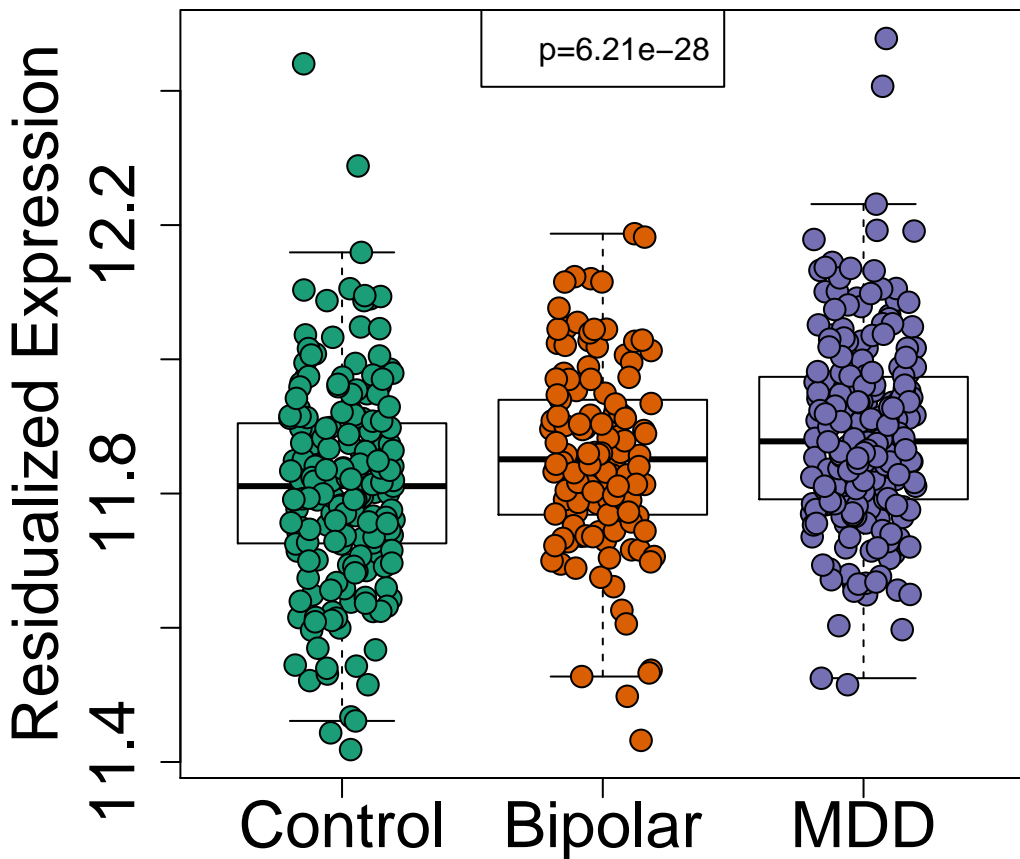
MBP, InGen

chr18:76988913–76989955(-)



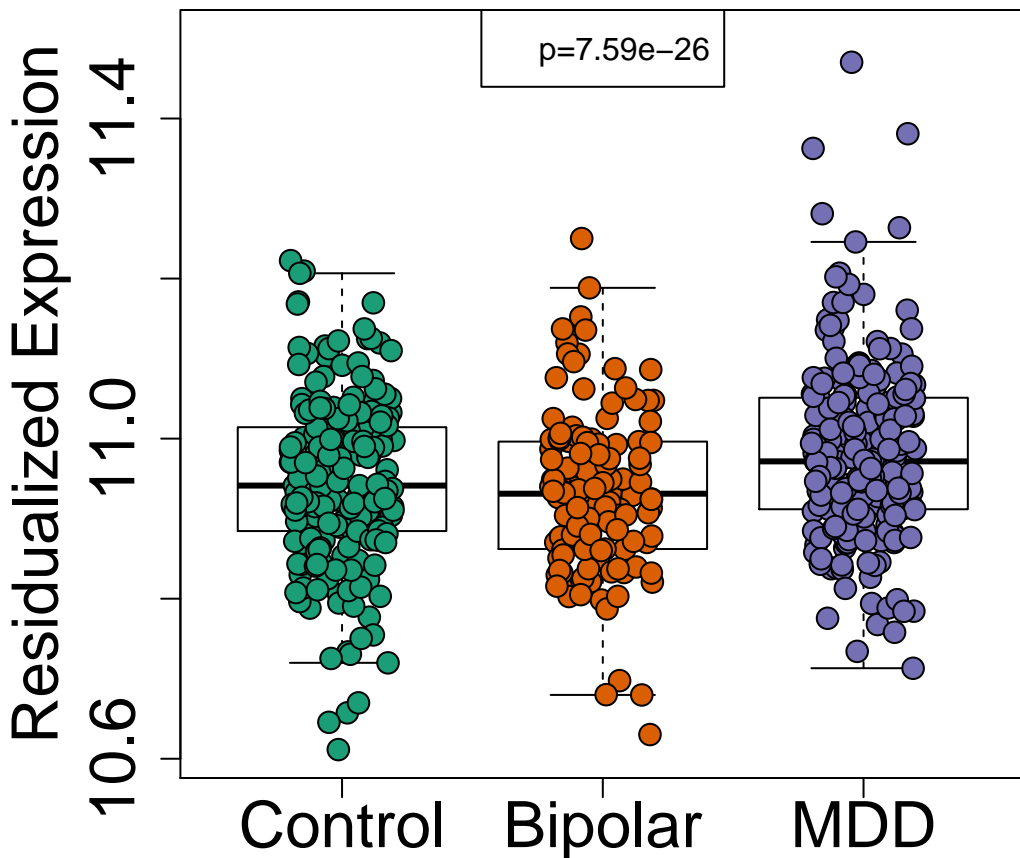
CLU, InGen

chr8:27606525–27608937(-)



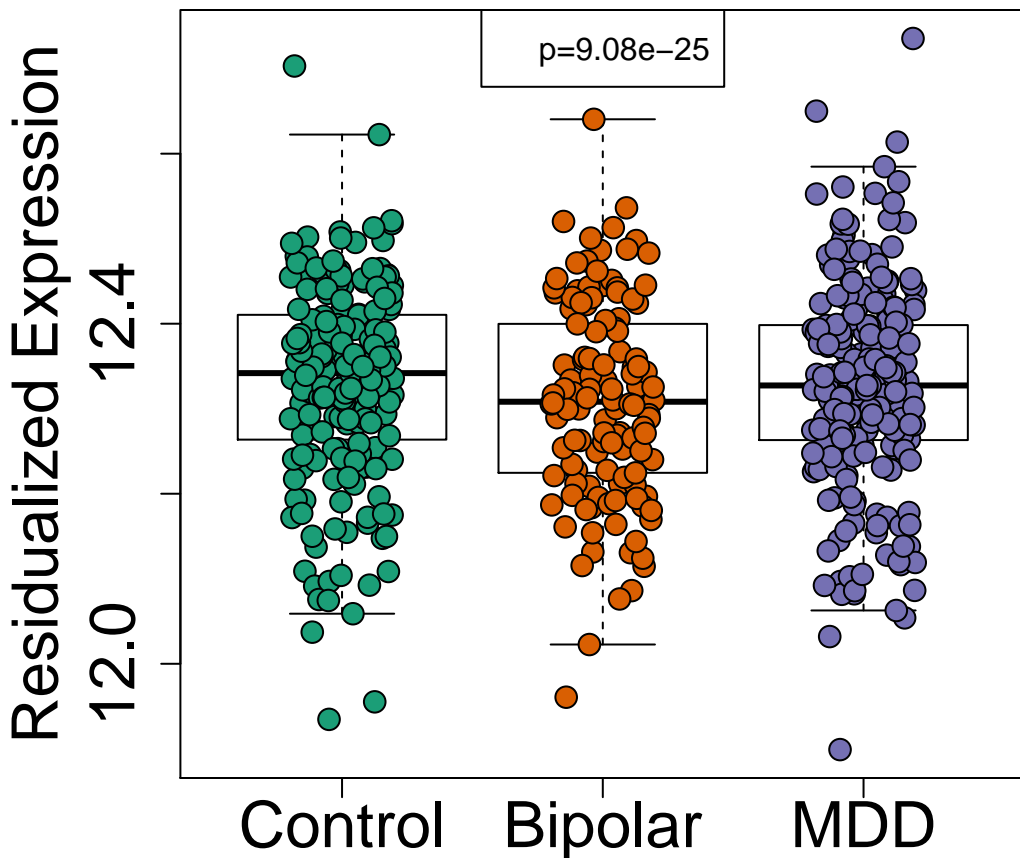
GAPDH, InGen

chr12:6537997–6538100(+)



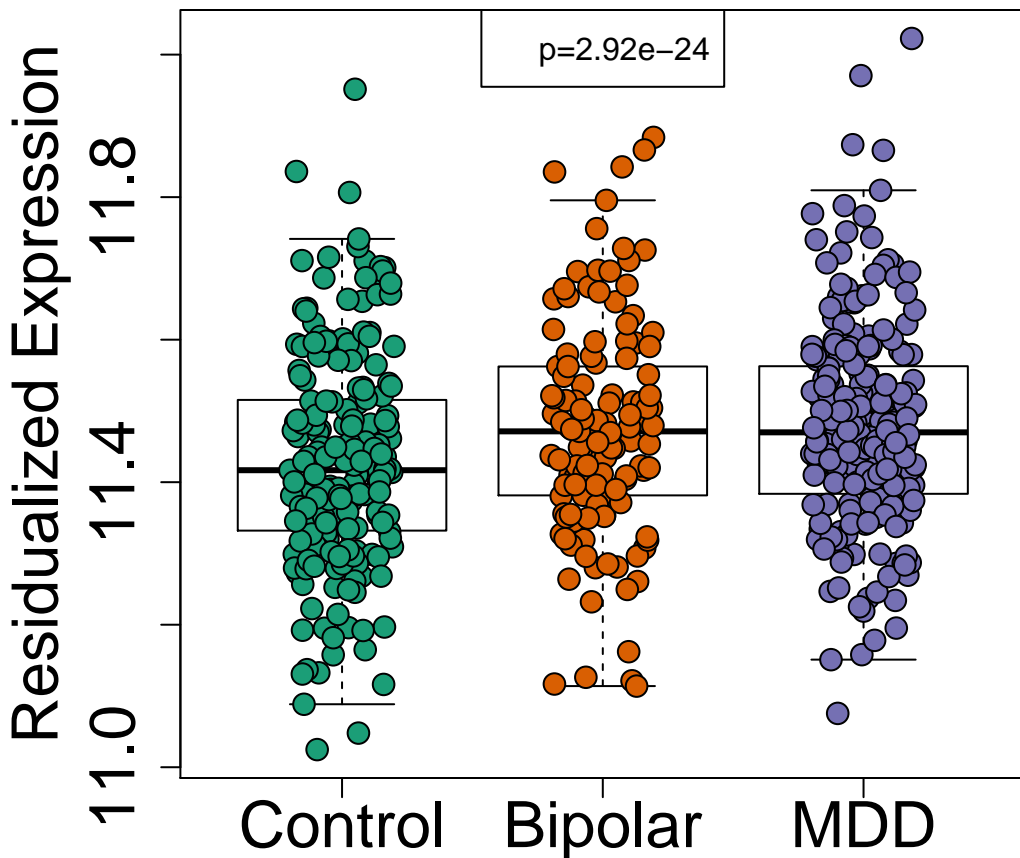
TMSB4X, InGen

chrX:12976362–12976776(+)



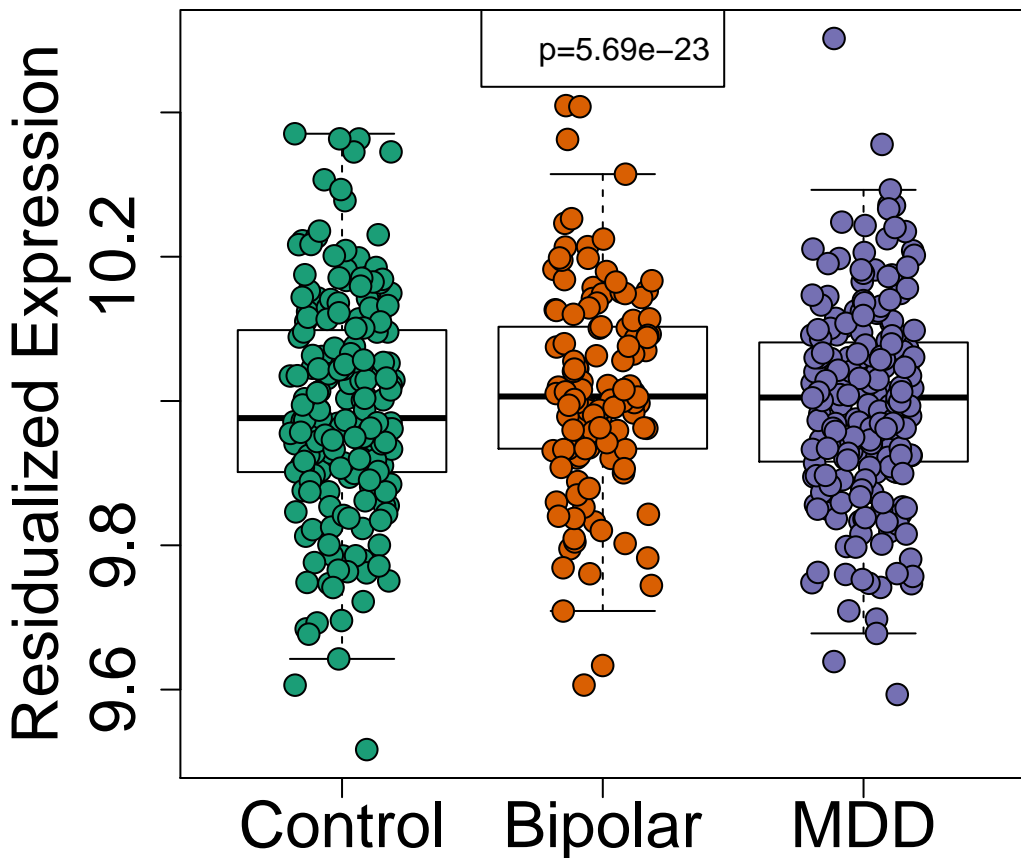
CLU, InGen

chr8:27609087-27610474(-)



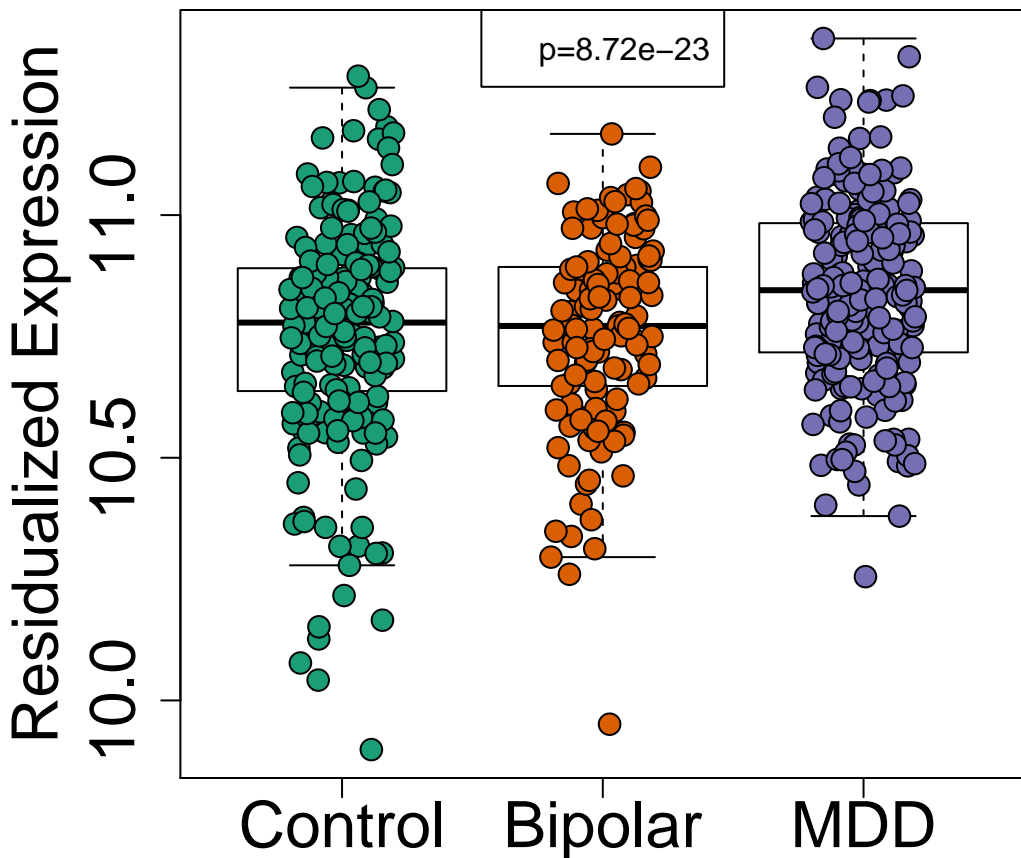
GNAS, InGen

chr20:58903792–58905382(+)



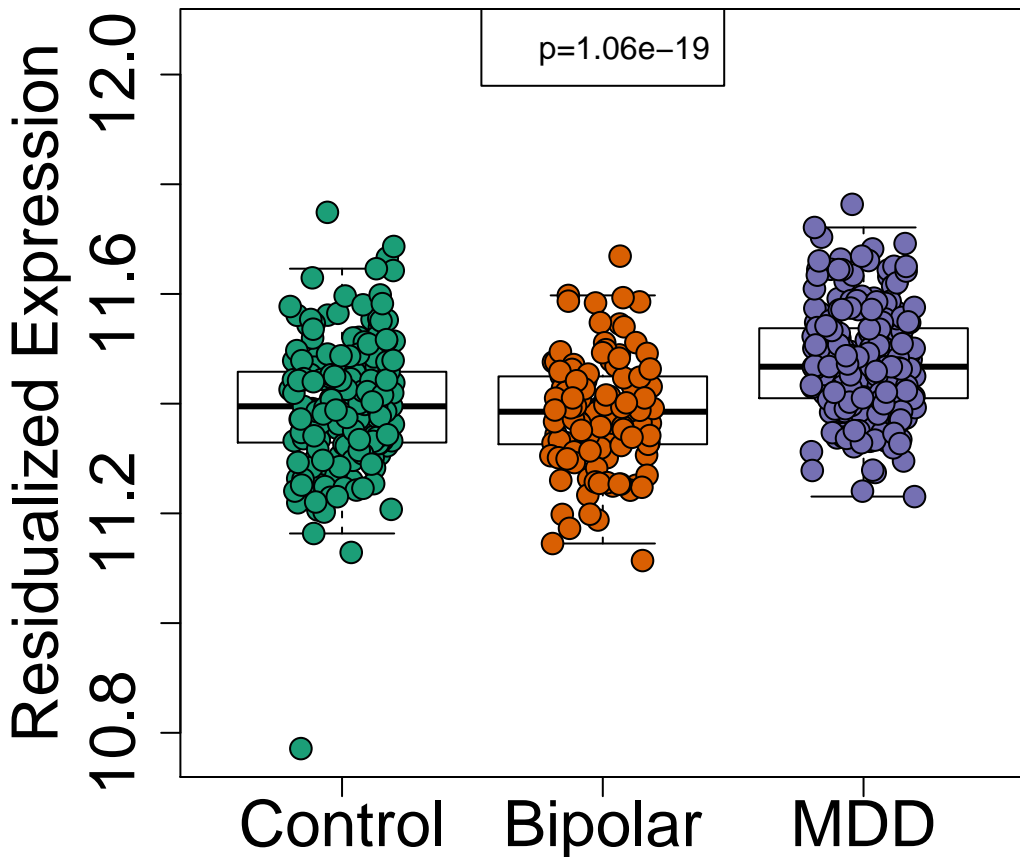
SPARCL1, InGen

chr4:87494599–87494980(–)



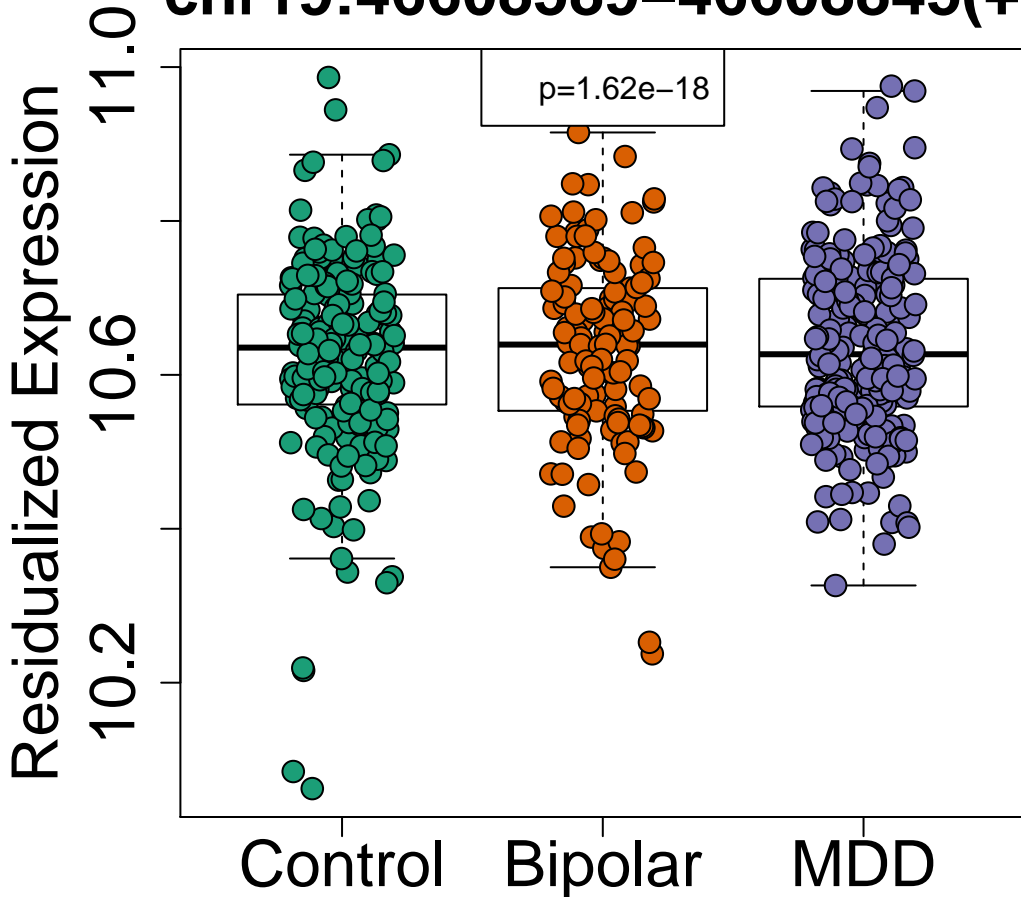
RPL31, InGen

chr2:101002809–101004157(+)



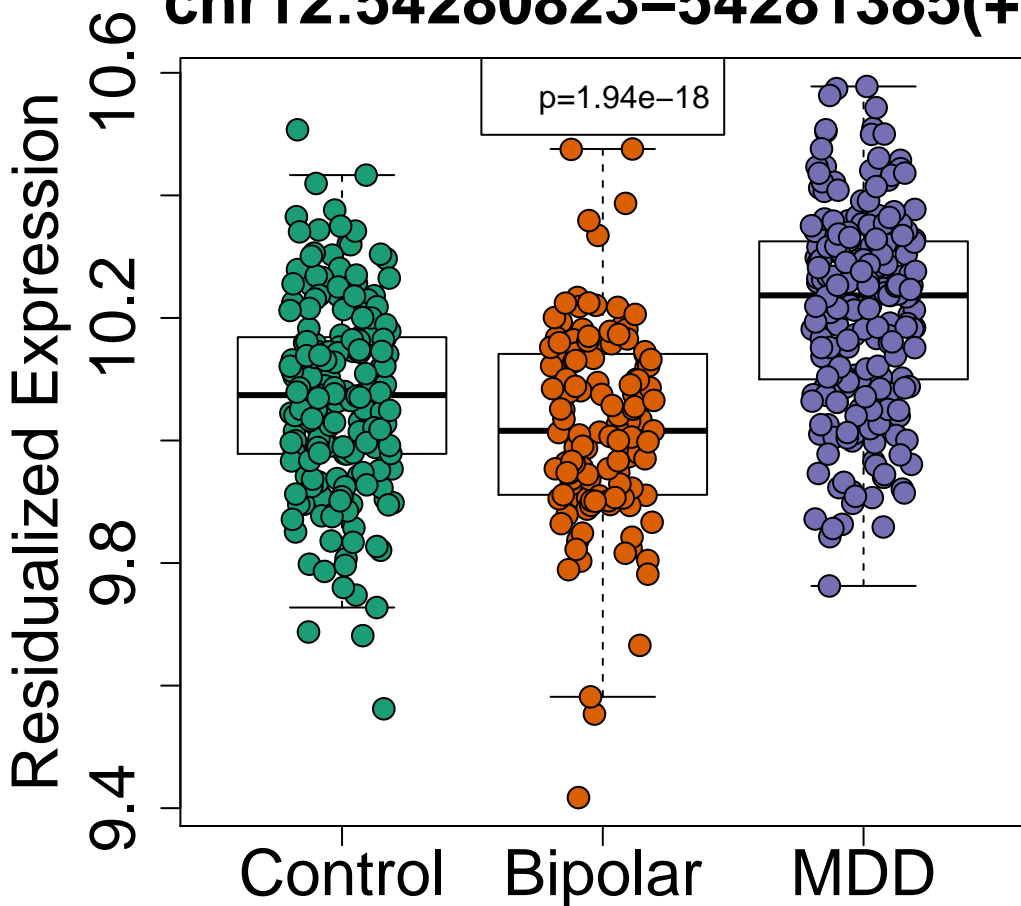
CALM3, InGen

chr19:46608589–46608845(+)



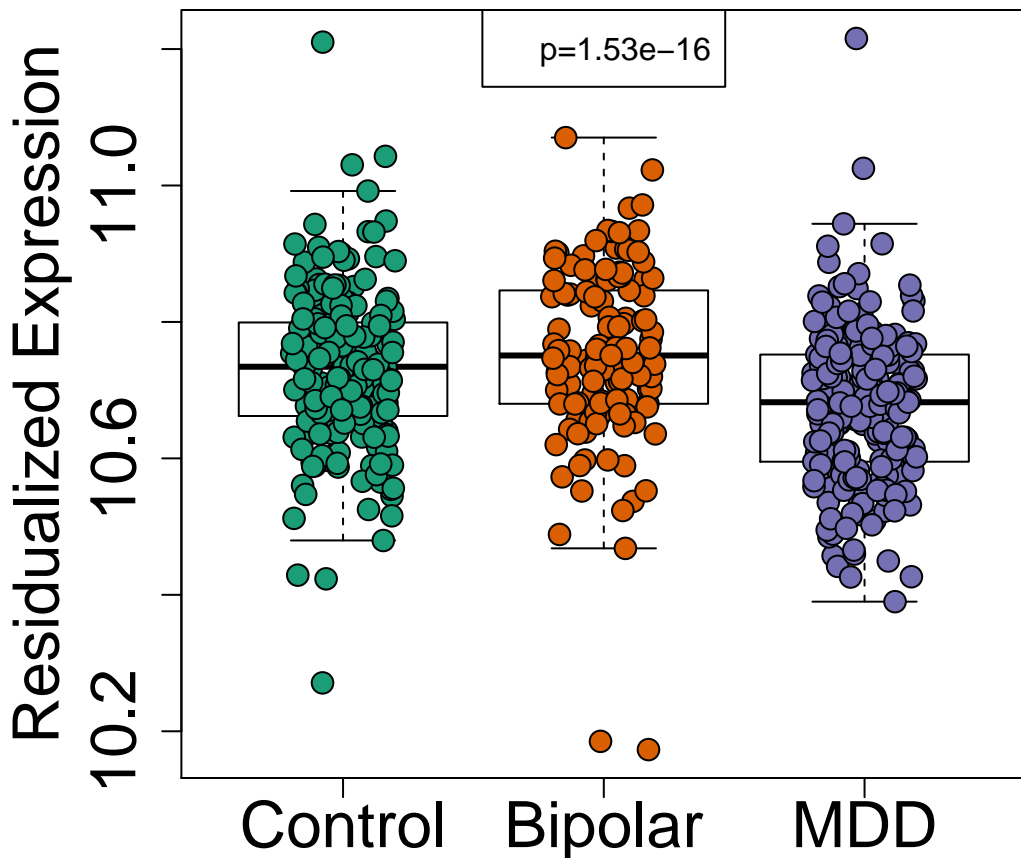
HNRNPA1, InGen

chr12:54280823–54281385(+)



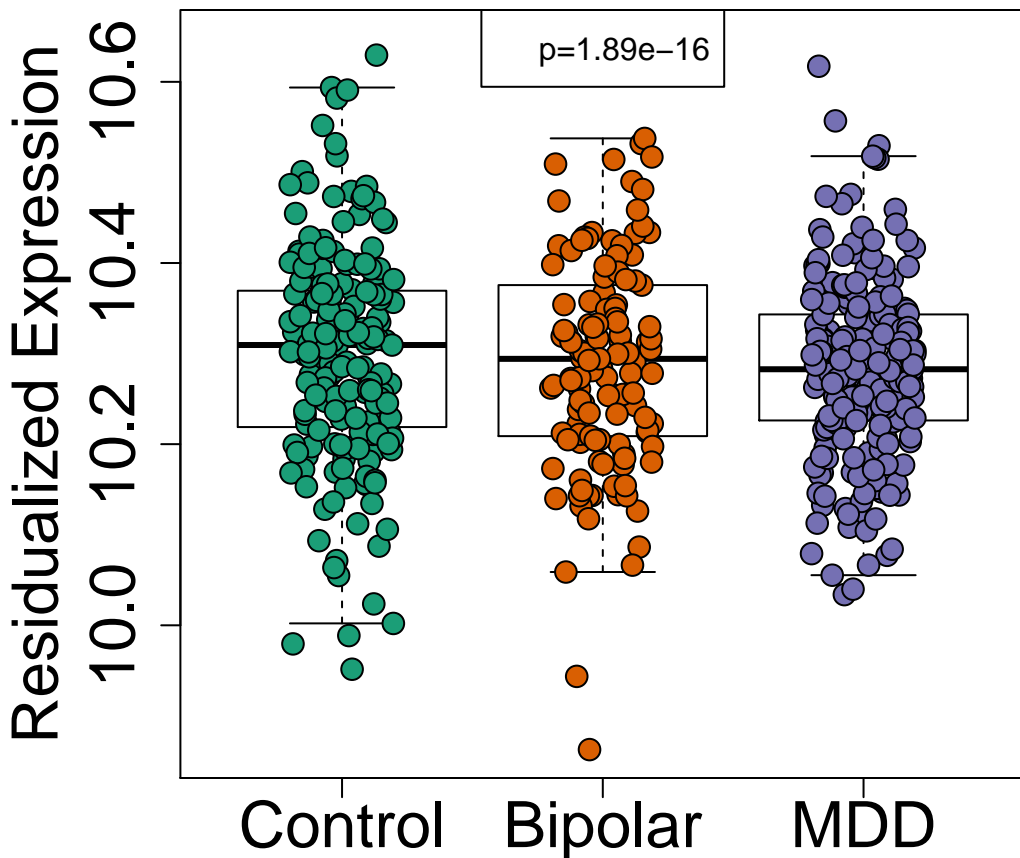
DPYSL2, InGen

chr8:26626679–26627214(+)



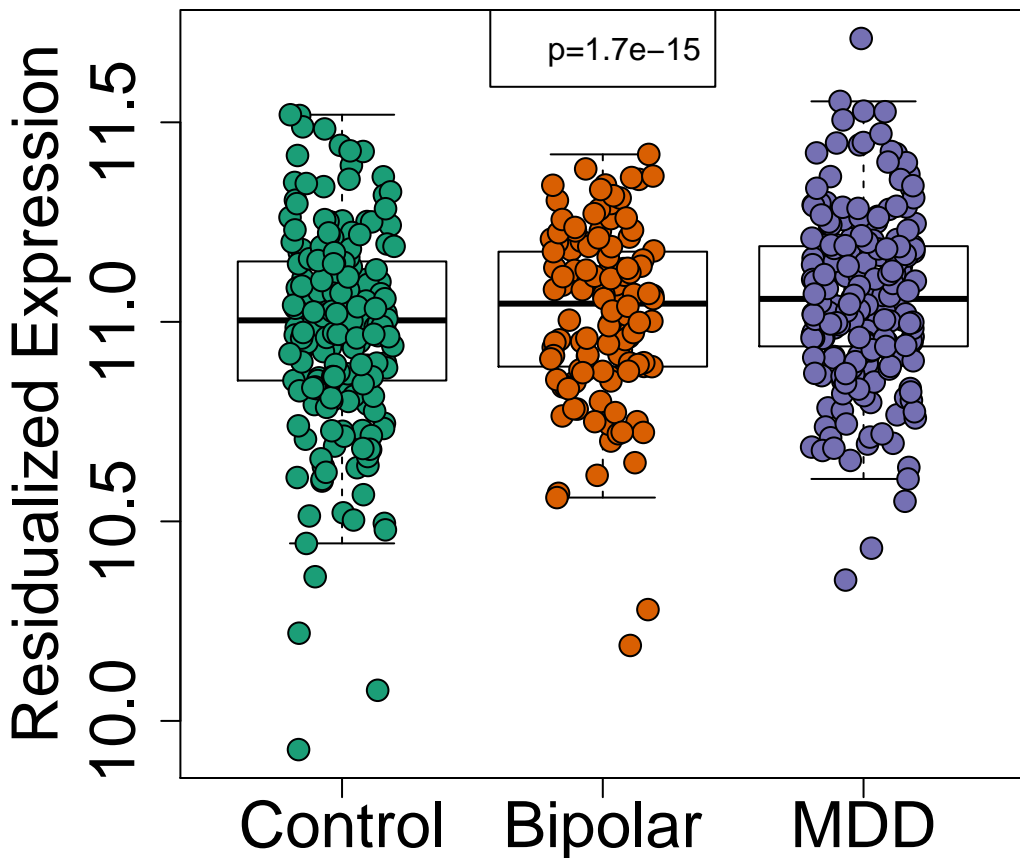
GNAS, InGen

chr20:58910082–58910333(+)



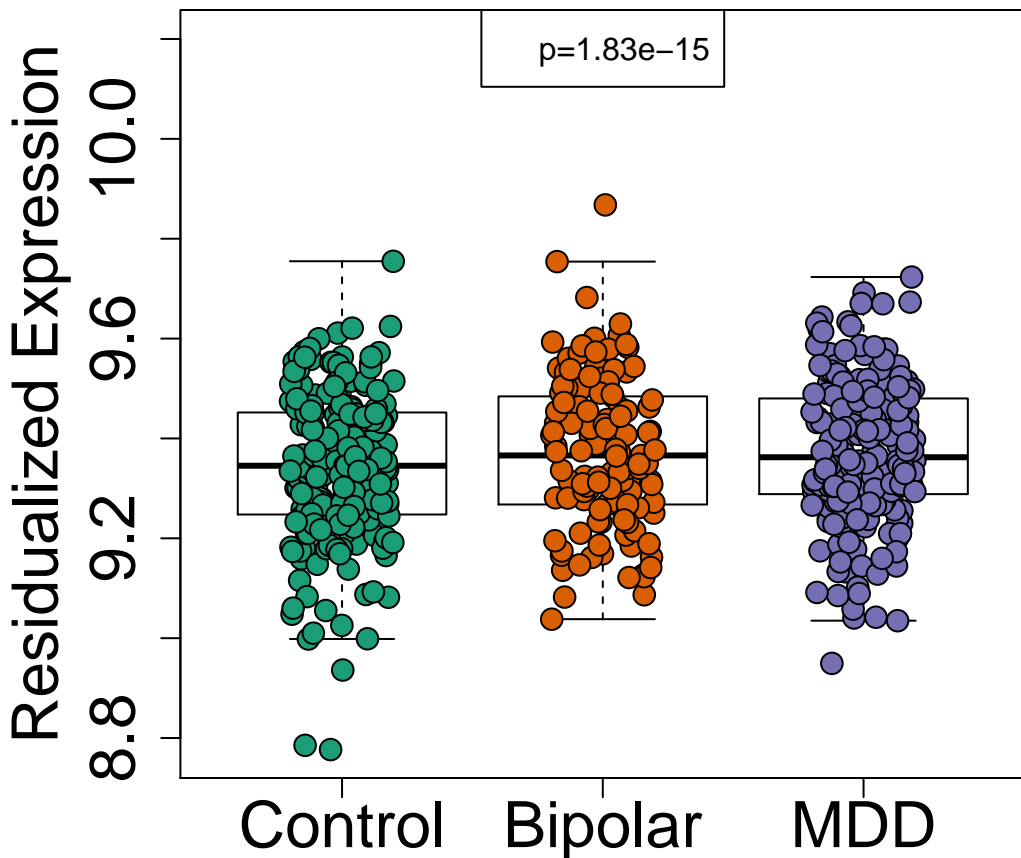
SPARCL1, InGen

chr4:87491691–87493581(-)



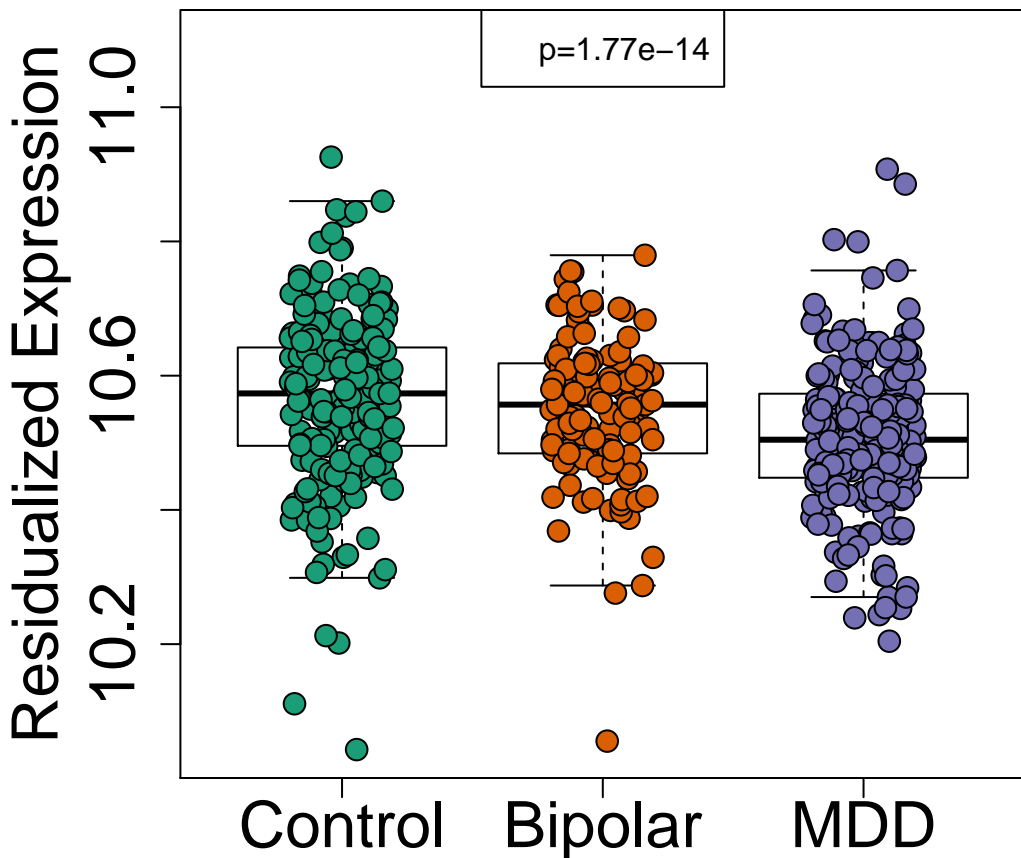
HBB, InGen

chr11:5226800–5226929(-)



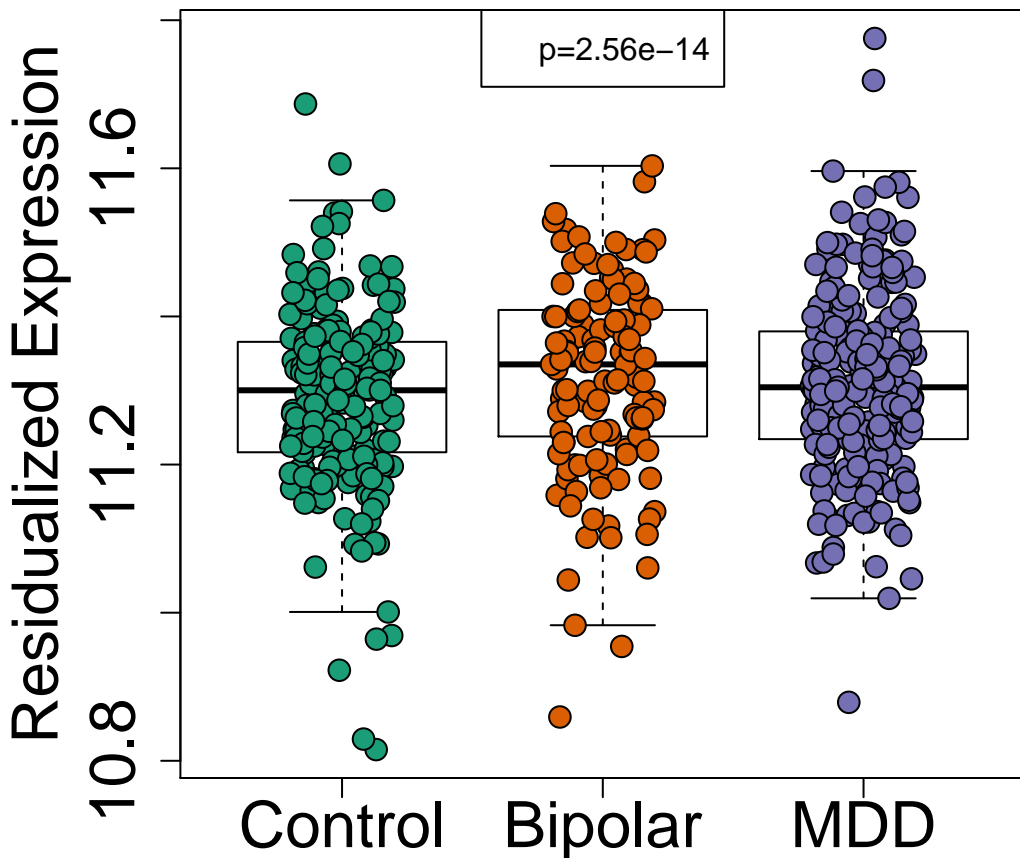
GNAS, InGen

chr20:58905481–58909161(+)



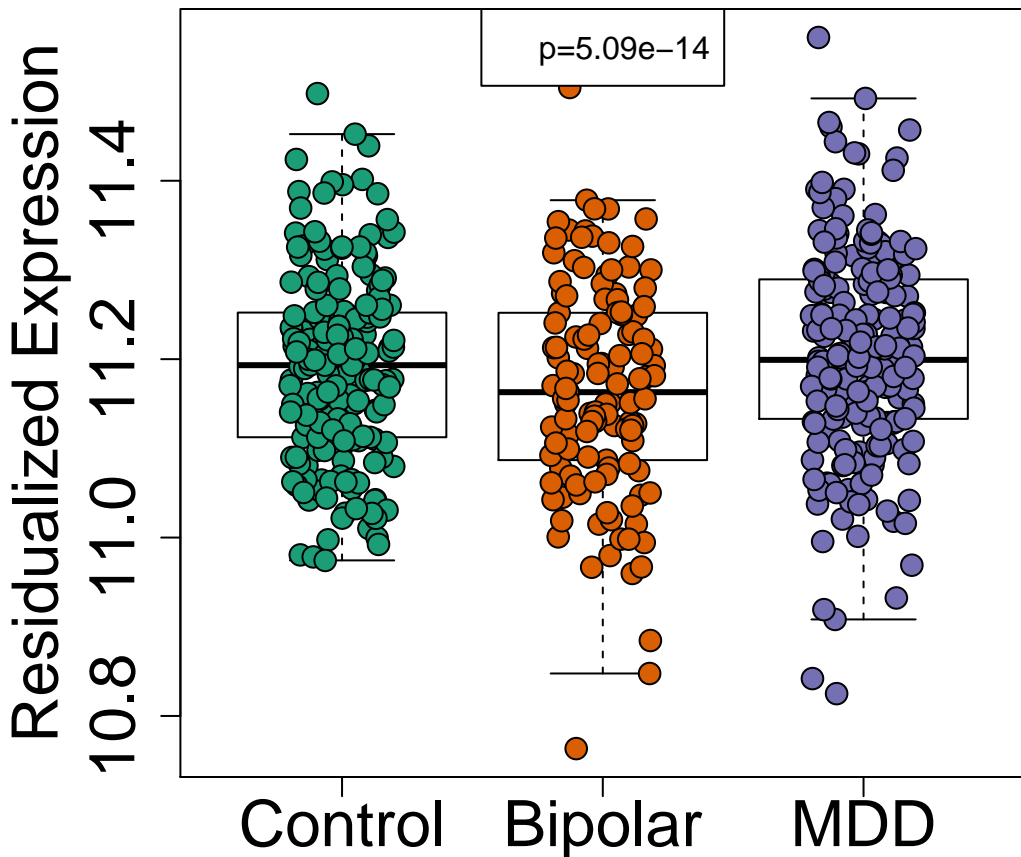
CPE, InGen

chr4:165484605–165487437(+)



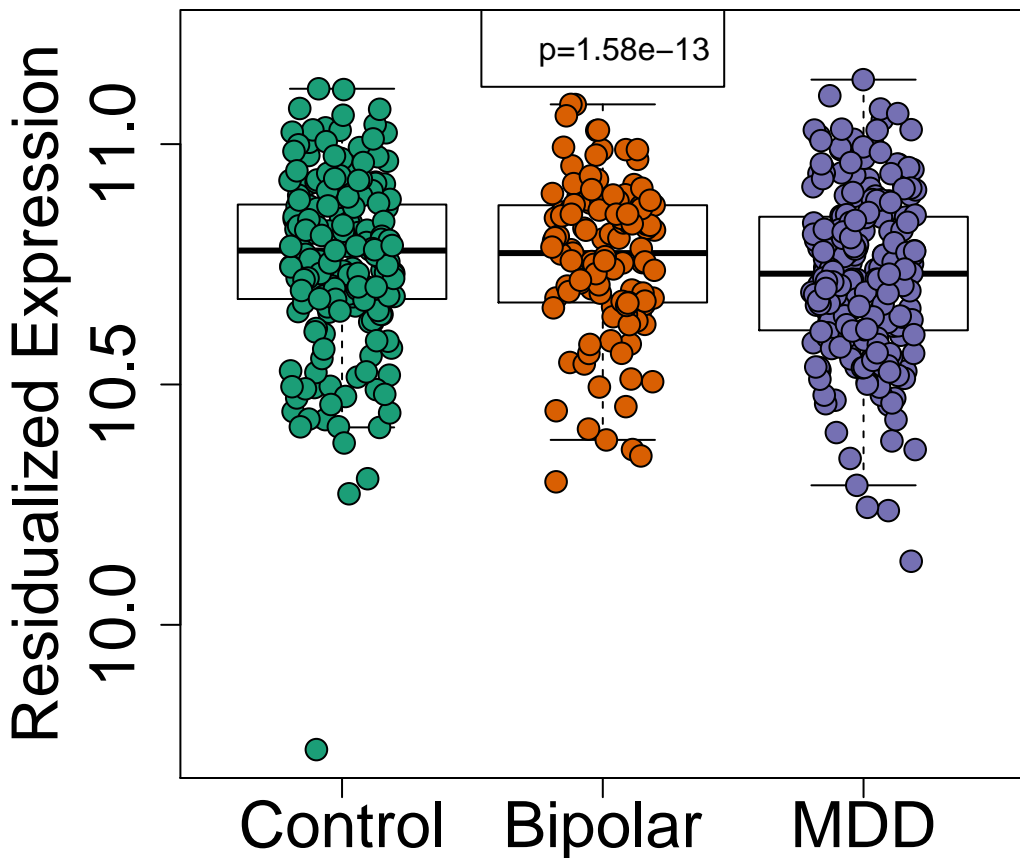
TPT1, InGen

chr13:45338777-45339496(-)



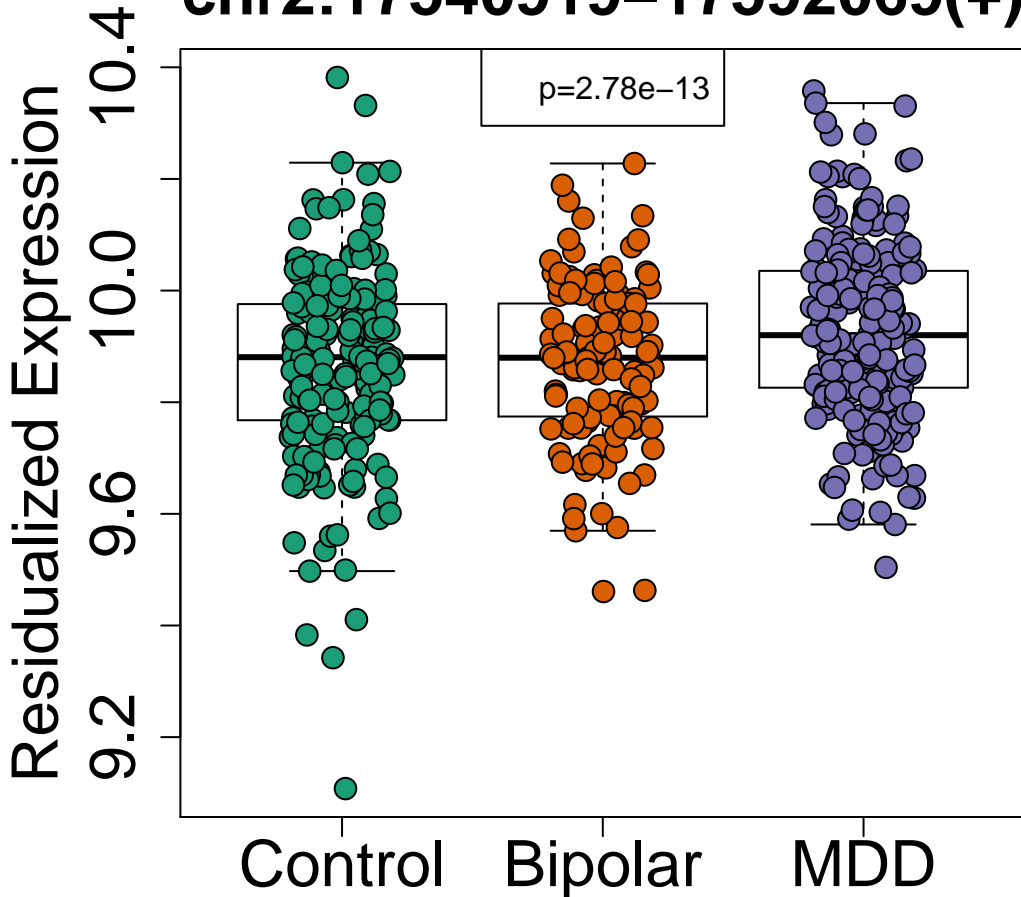
SNAP25, InGen

chr20:10297051–10299267(+)



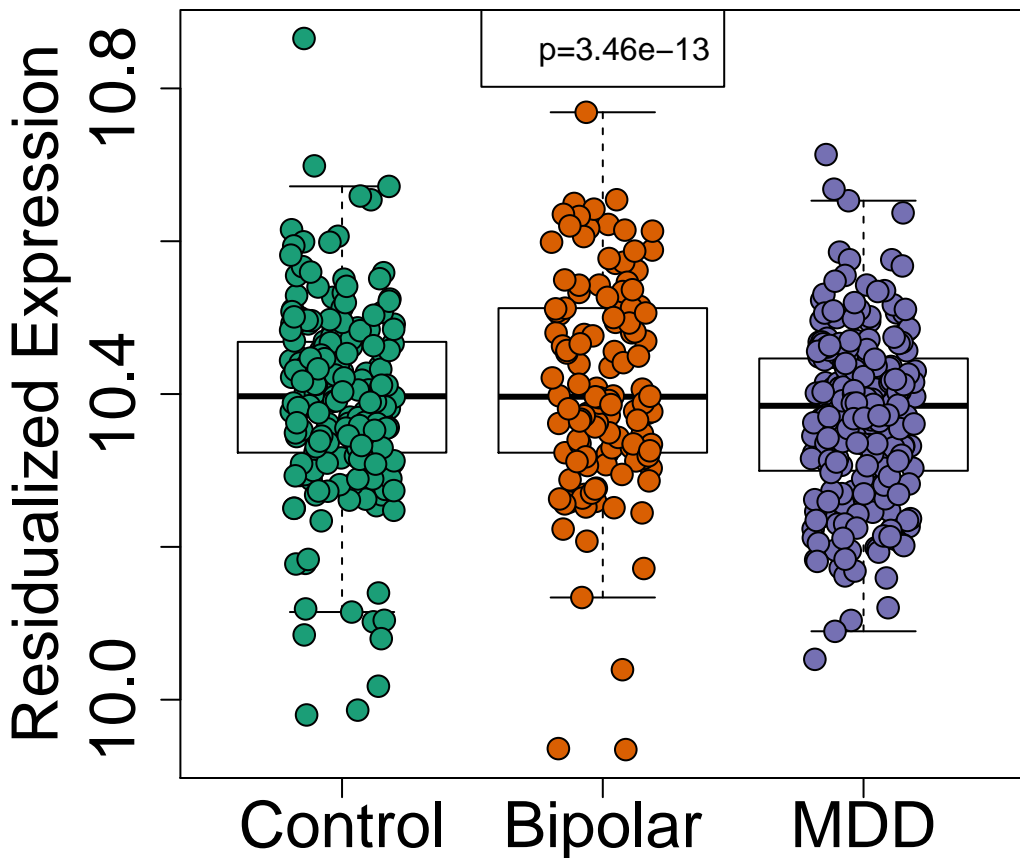
VSNL1, InGen

chr2:17540919–17592069(+)



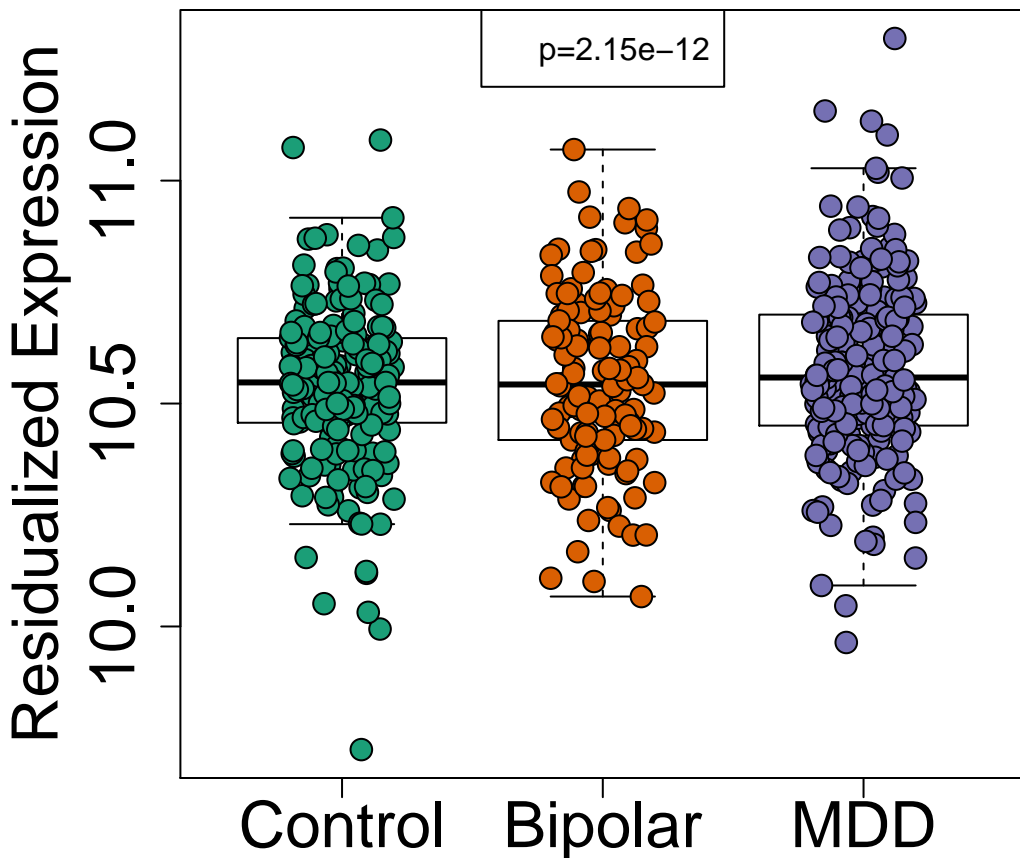
ACTB, InGen

chr7:5528720–5529160(-)



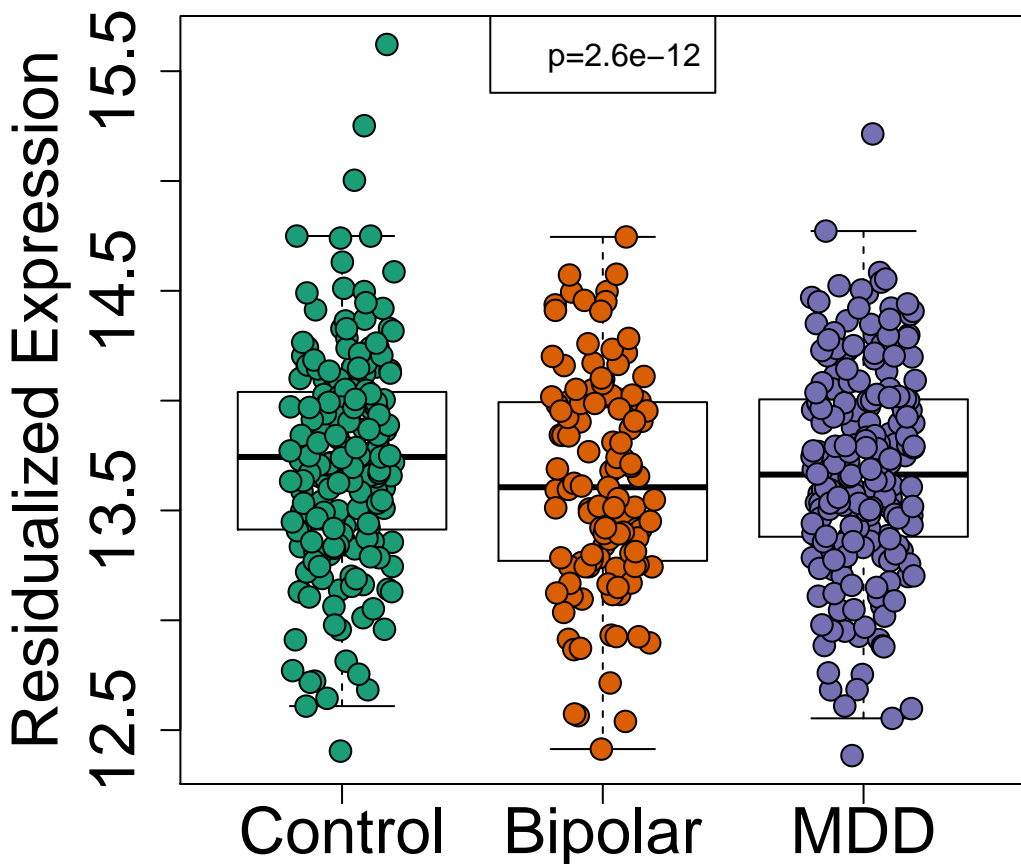
ARPP19, InGen

chr15:52552105–52557099(-)



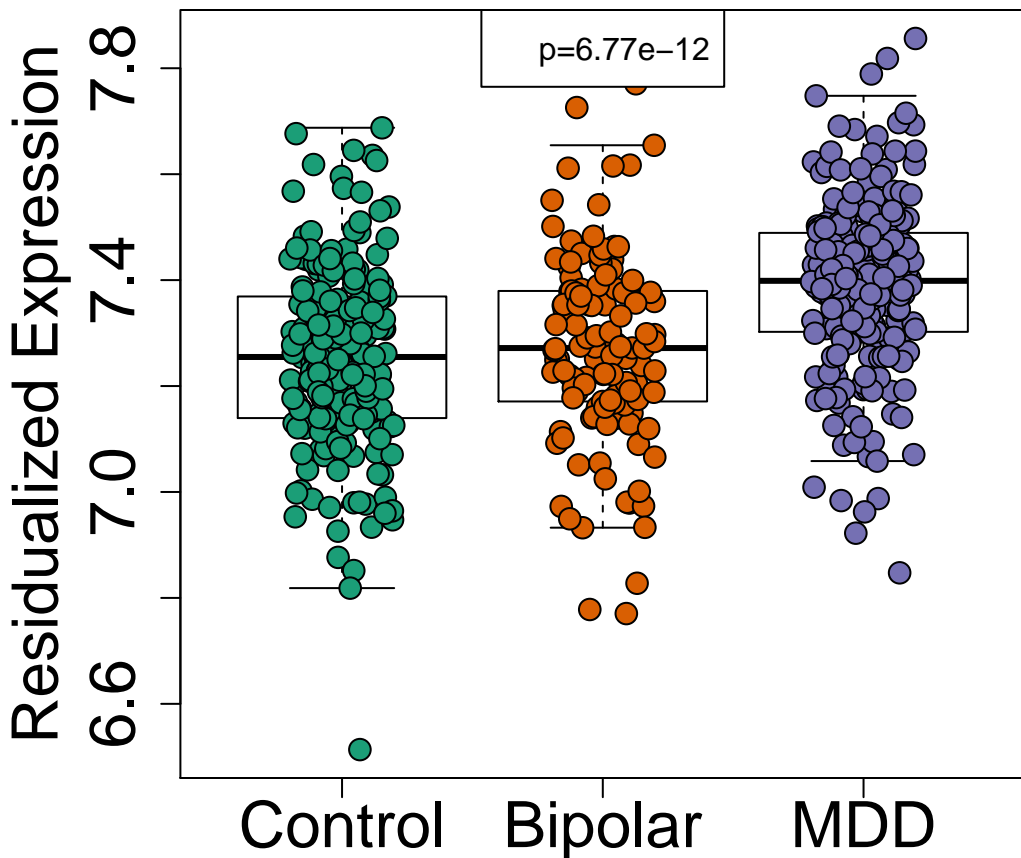
GFAP, InGen

chr17:44907389–44908063(-)



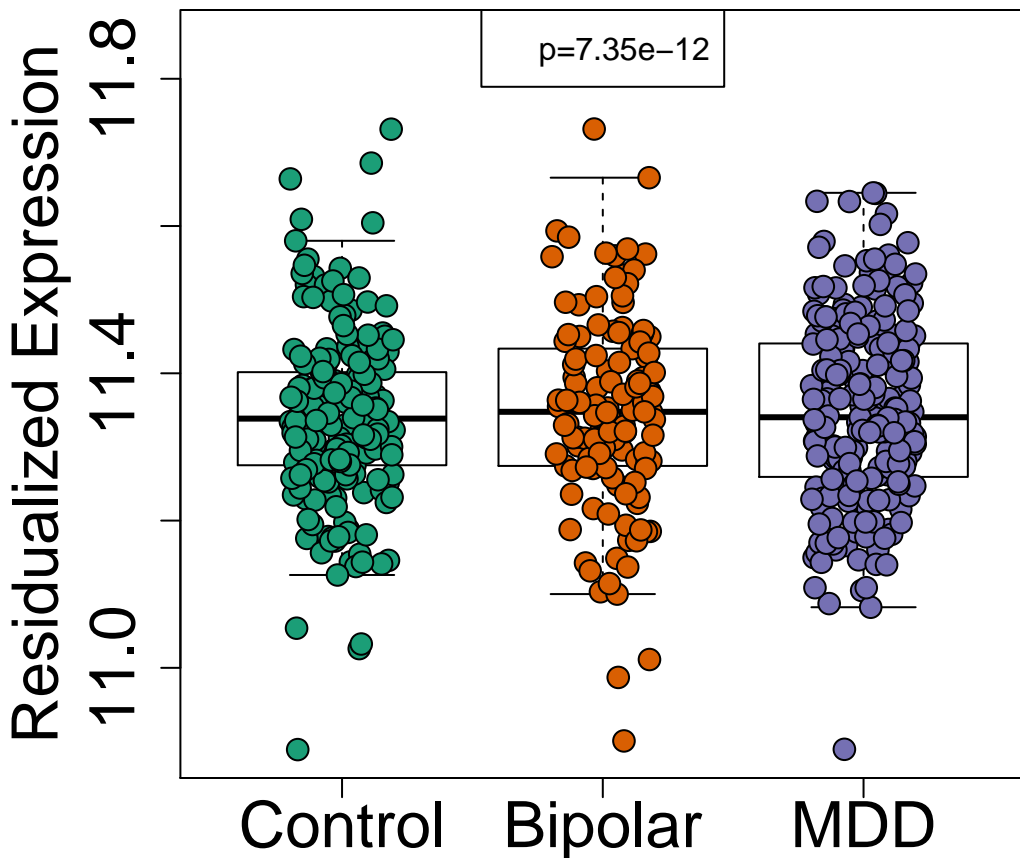
EIF3L, InGen

chr22:37870348–37874369(+)



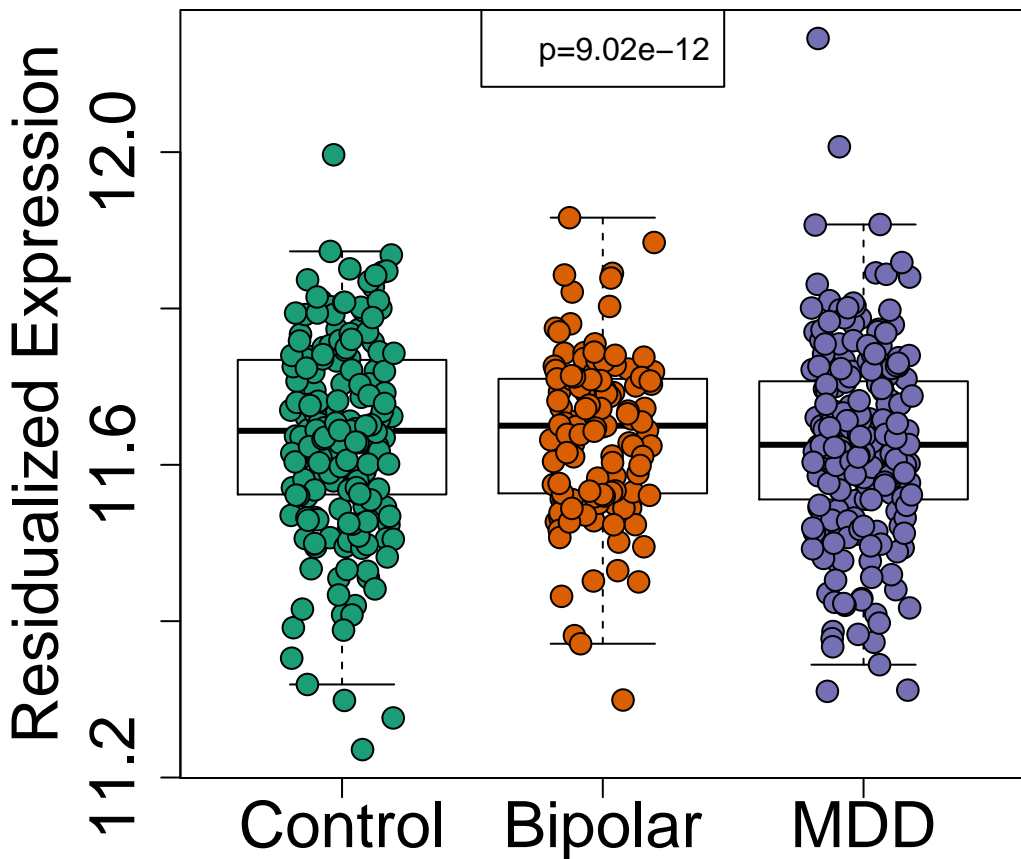
CPE, InGen

chr4:165493271–165495558(+)



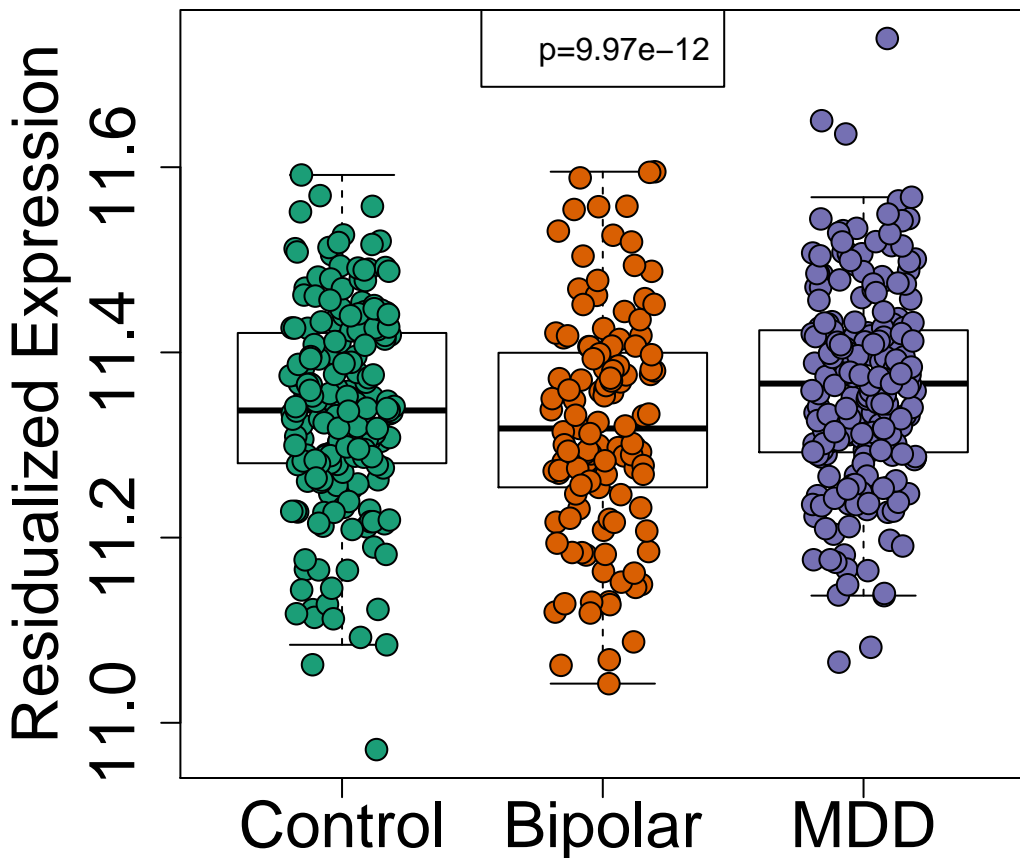
RPL38, InGen

chr17:74209310–74209803(+)



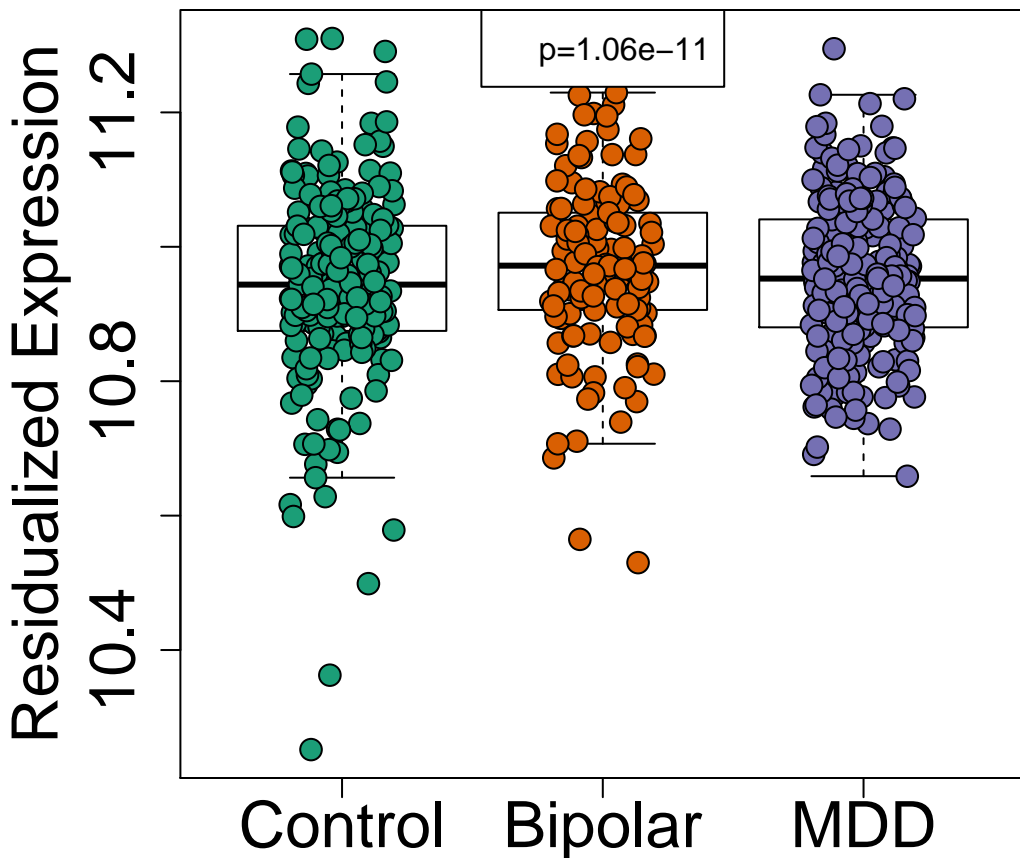
UQCRB, InGen

chr8:96231941–96233155(–)



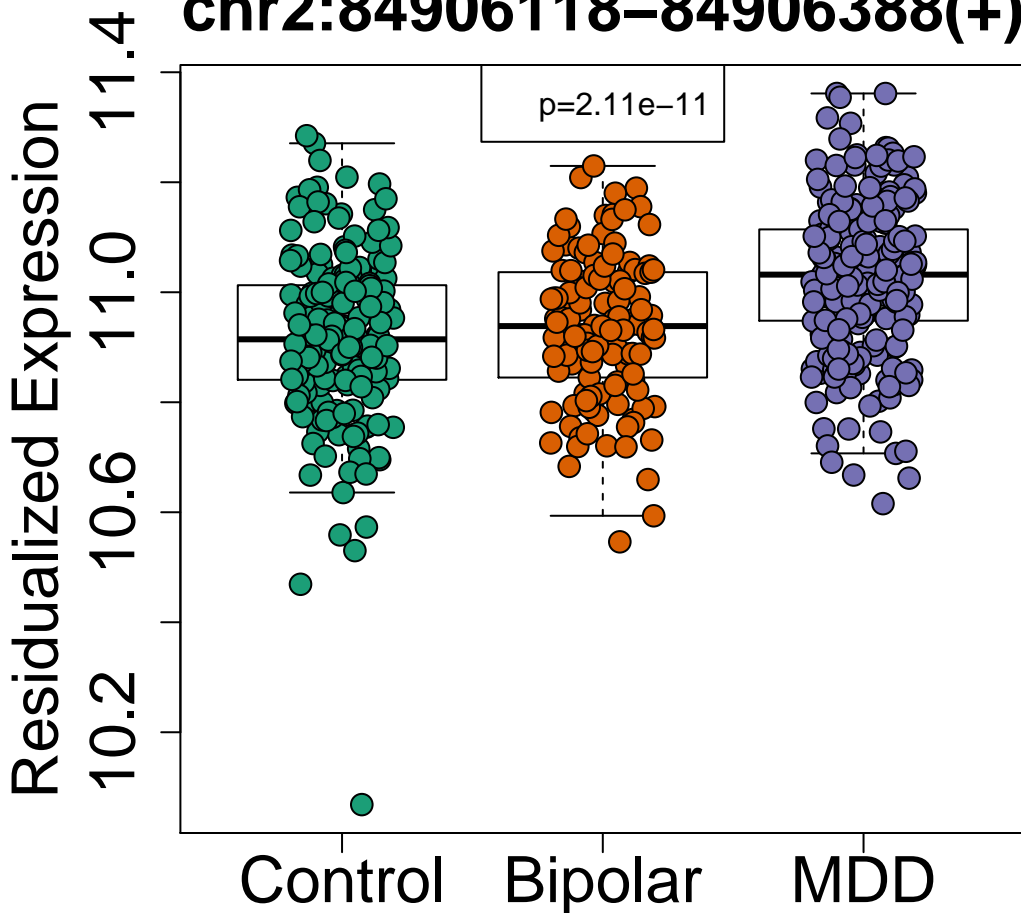
CALM3, InGen

chr19:46608341–46608481(+)



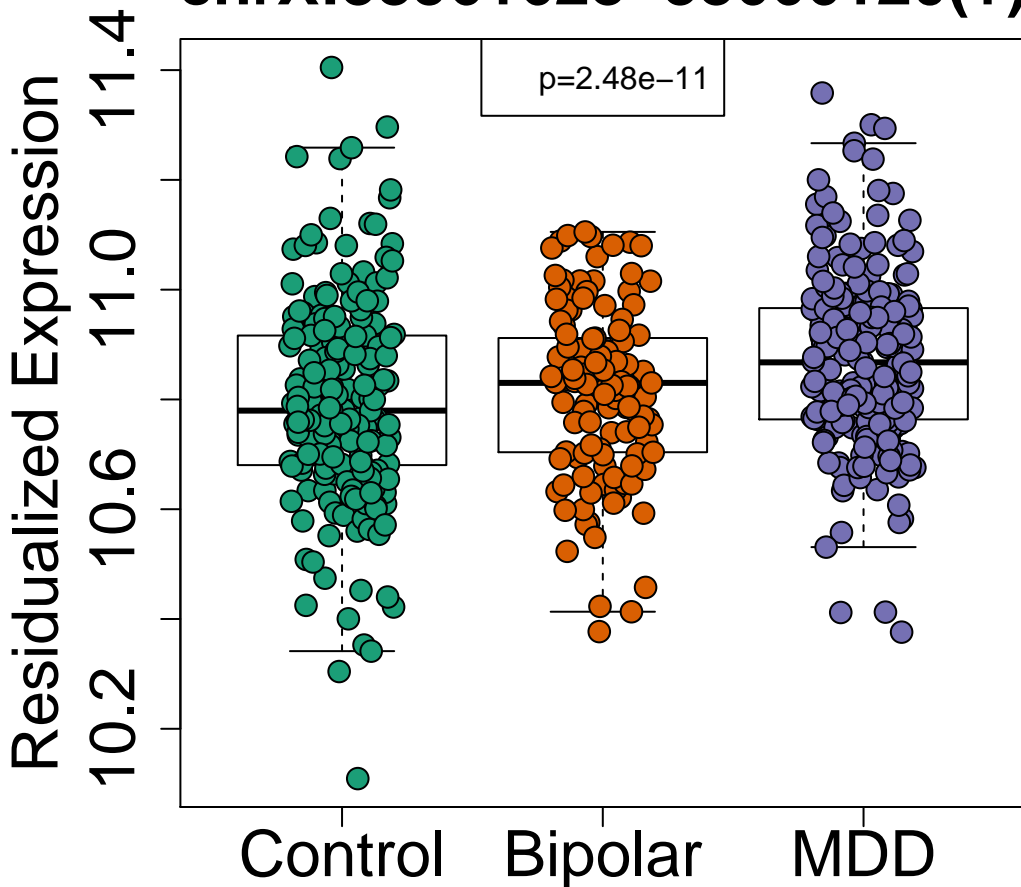
TMSB10, InGen

chr2:84906118–84906388(+)



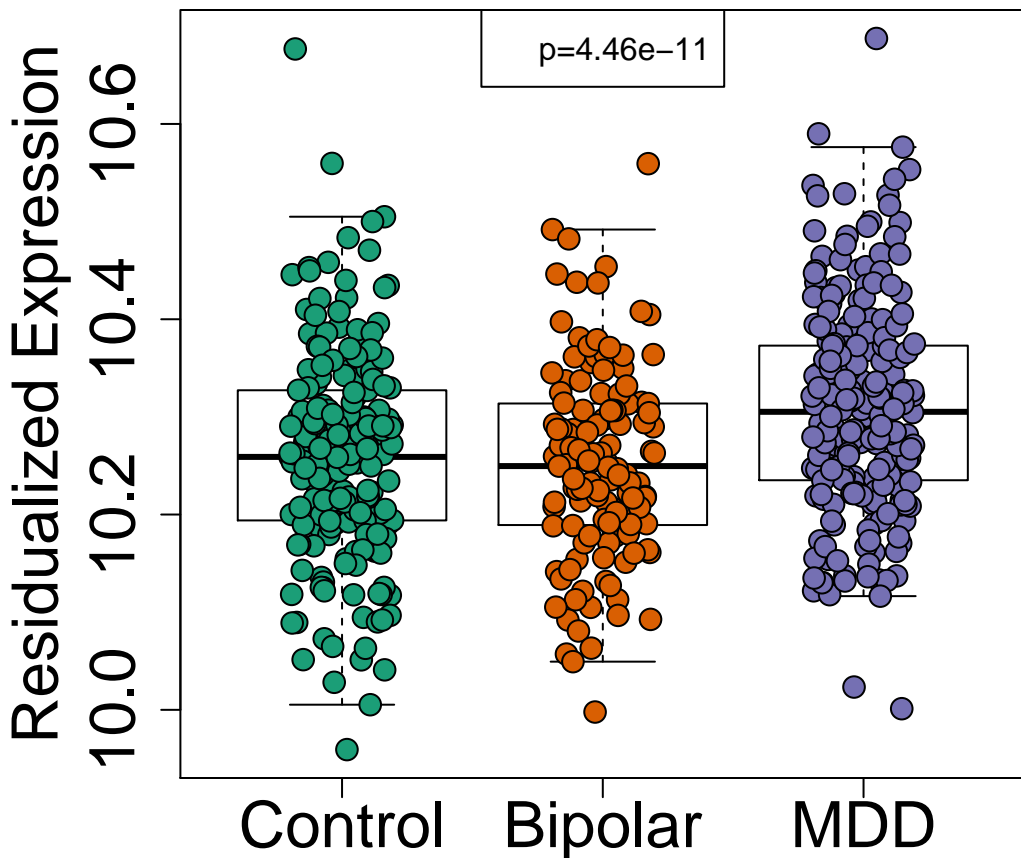
TSPAN7, InGen

chrX:38561628–38666120(+)



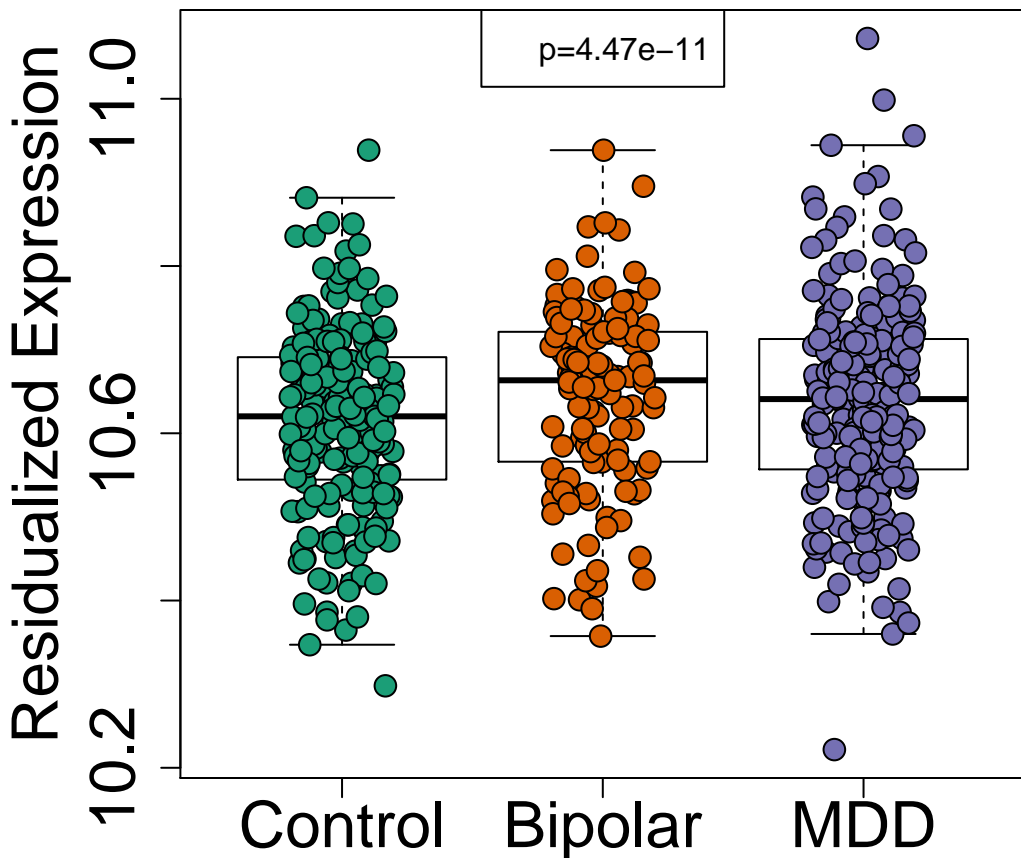
RPL11, InGen

chr1:23694792–23695797(+)



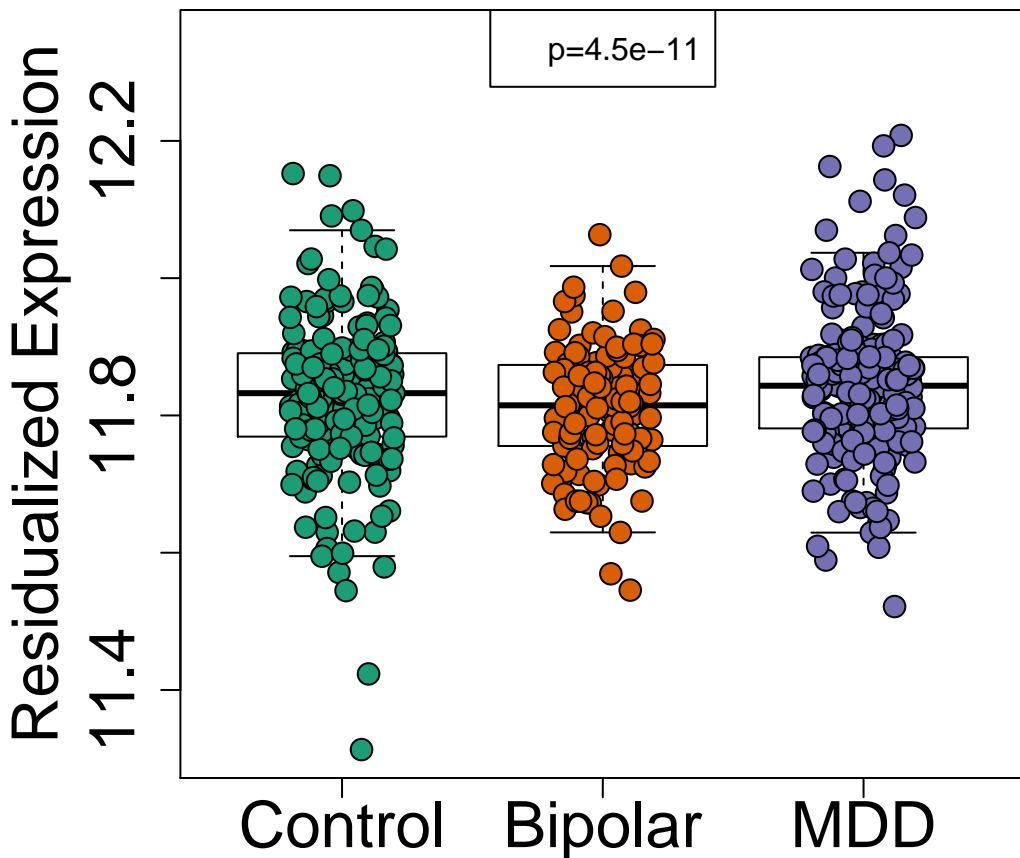
SKP1, InGen

chr5:134158596–134160986(-)



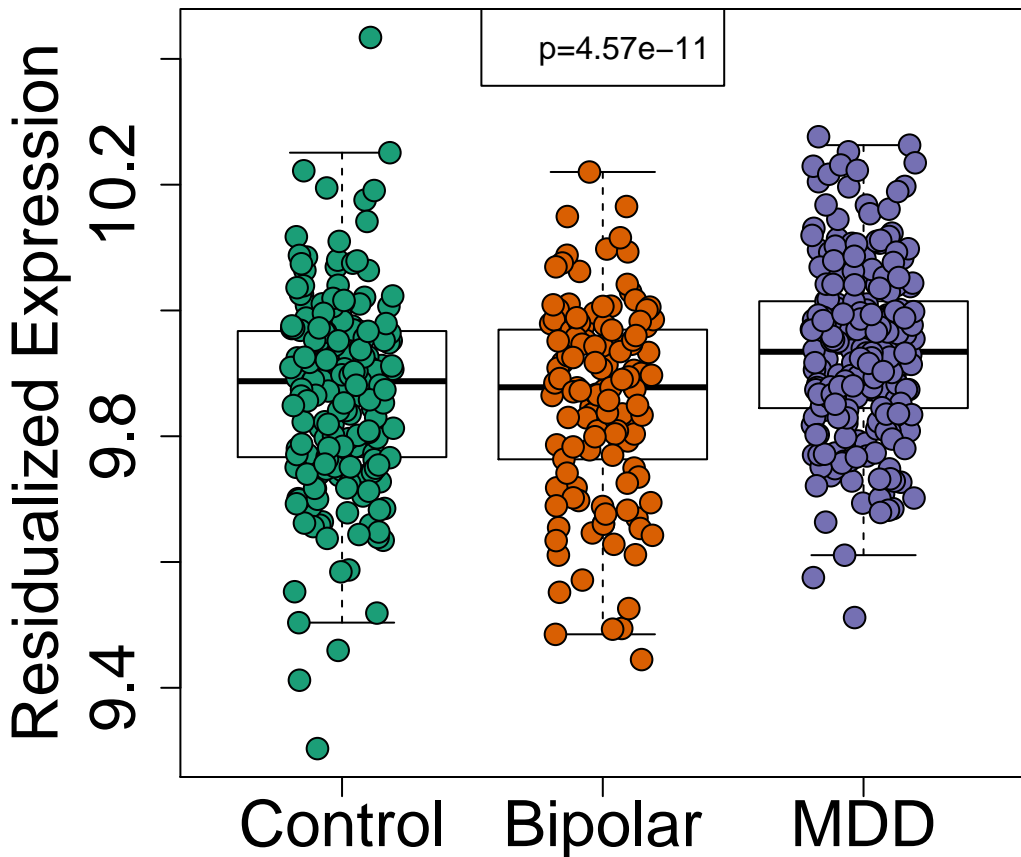
GAPDH, InGen

chr12:6534862–6536493(+)



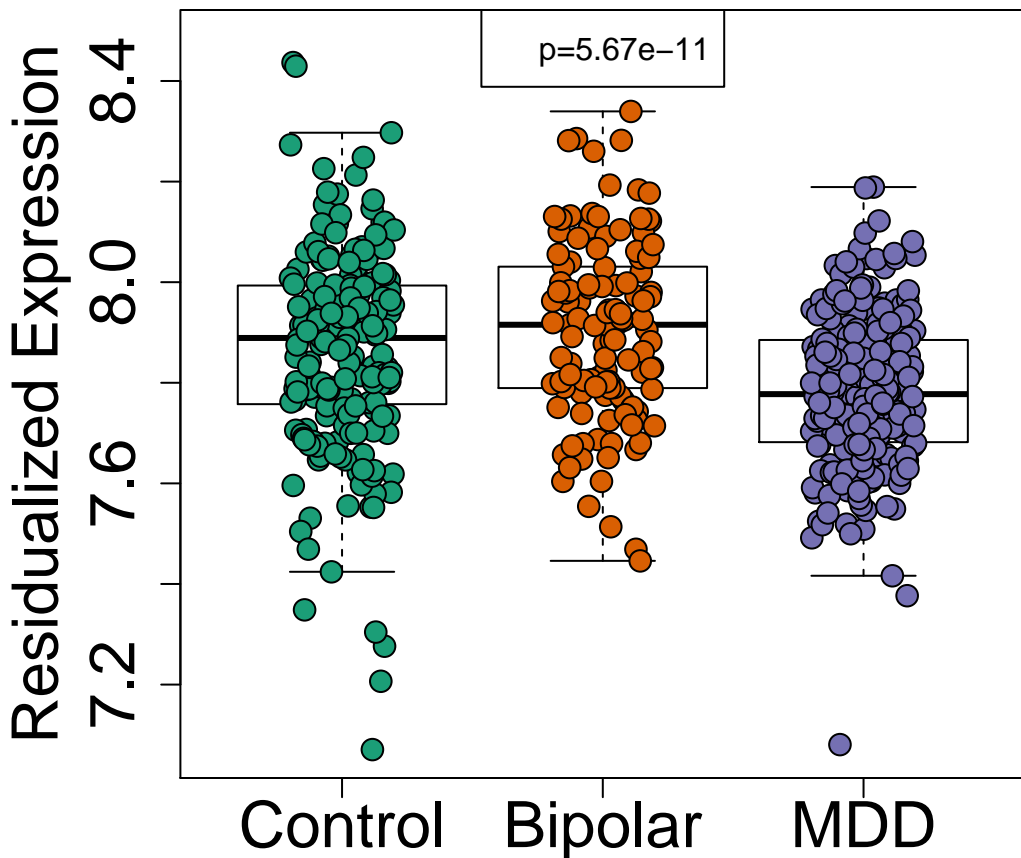
GABARAPL1, InGen

chr12:10220559–10221786(+)



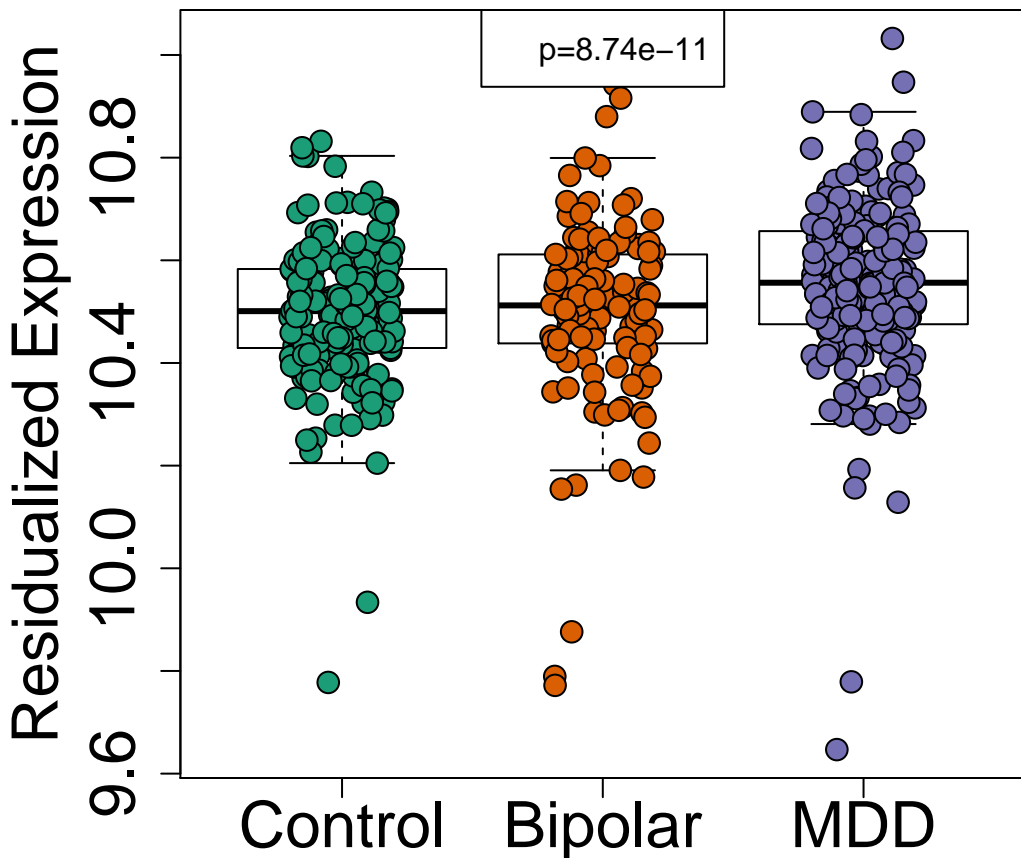
ATP1A3, InGen

chr19:41985424–41985863(-)



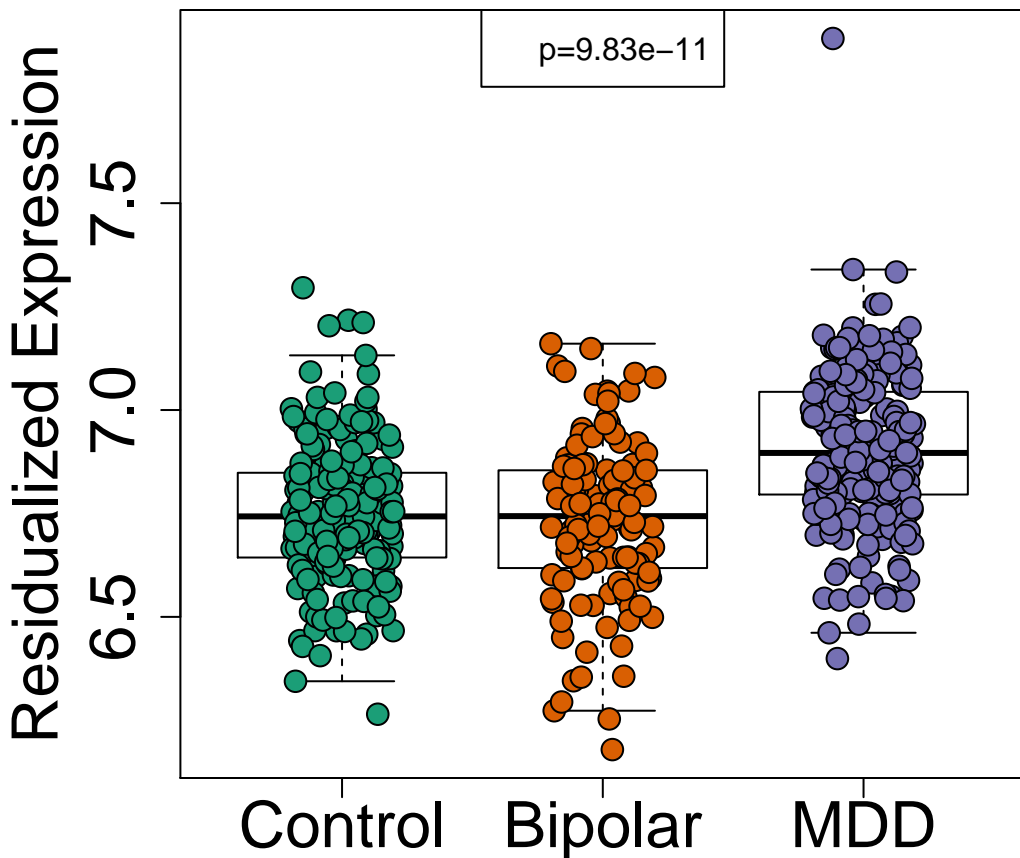
TUBA1B, InGen

chr12:49129391–49129499(–)



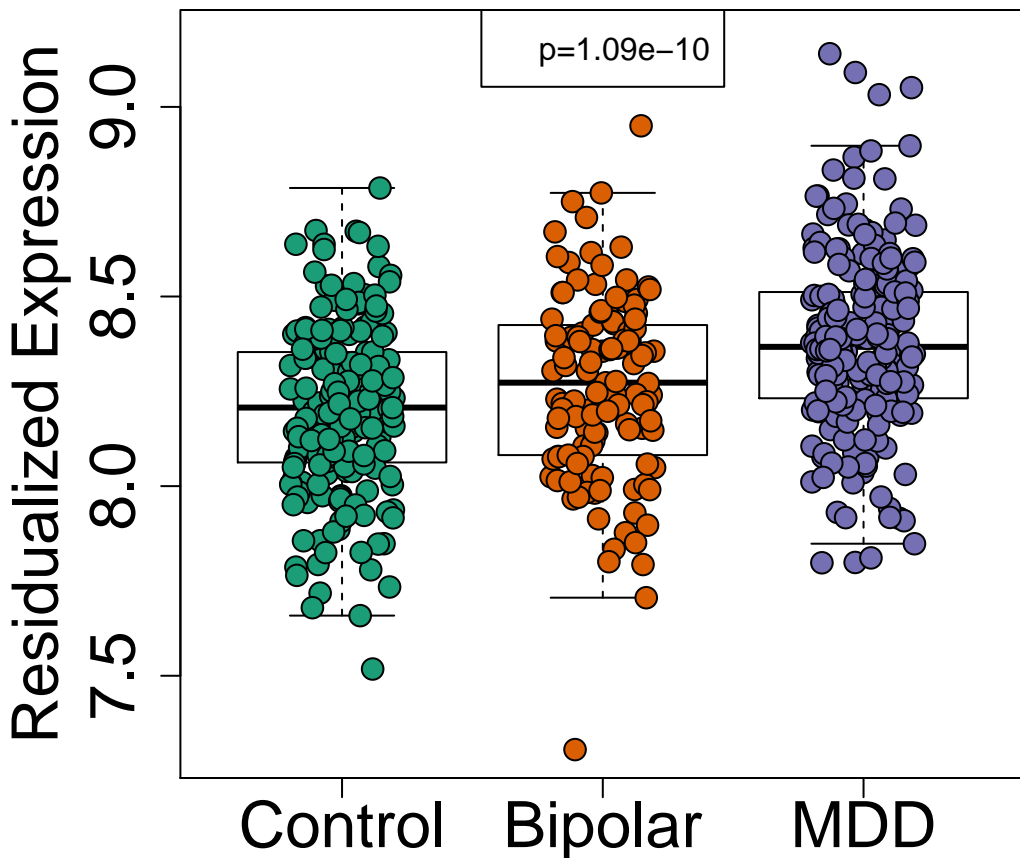
MAPT, InGen

chr17:45894687–45962320(+)



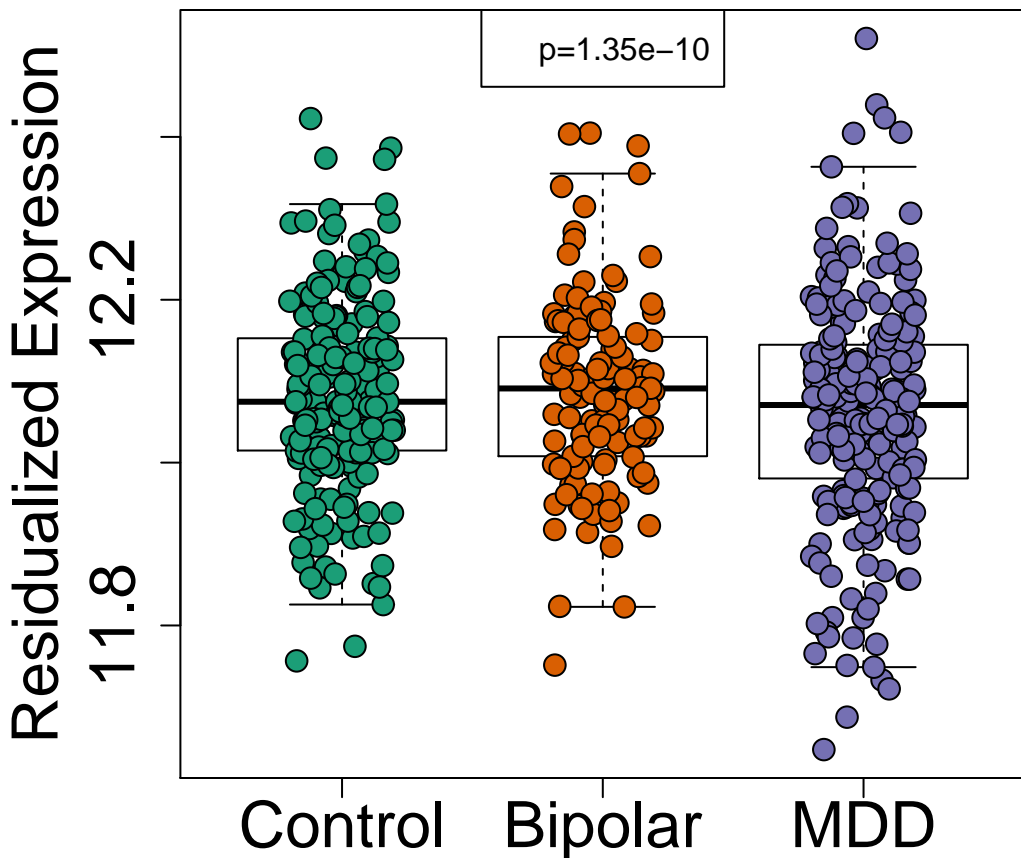
HMGN2, InGen

chr1:26474668–26475112(+)



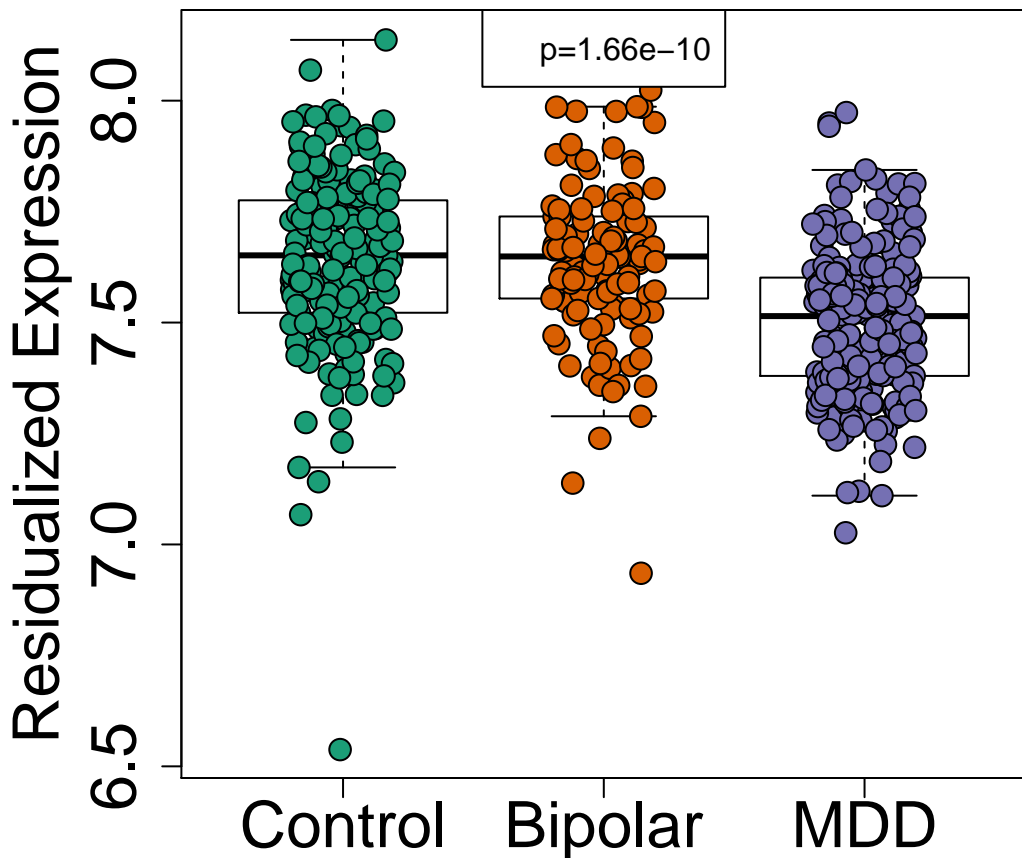
UBB, InGen

chr17:16381185–16381901(+)



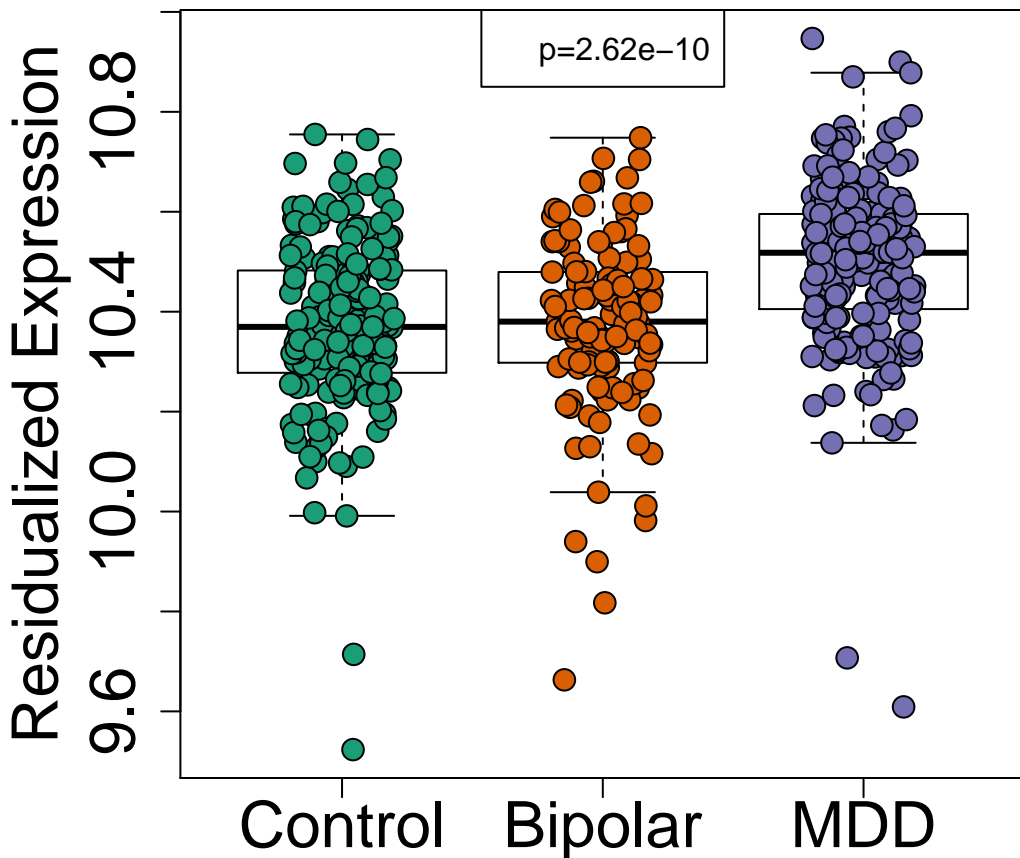
RP11-793H13.8, InGen

chr12:53464853-53465931(+)



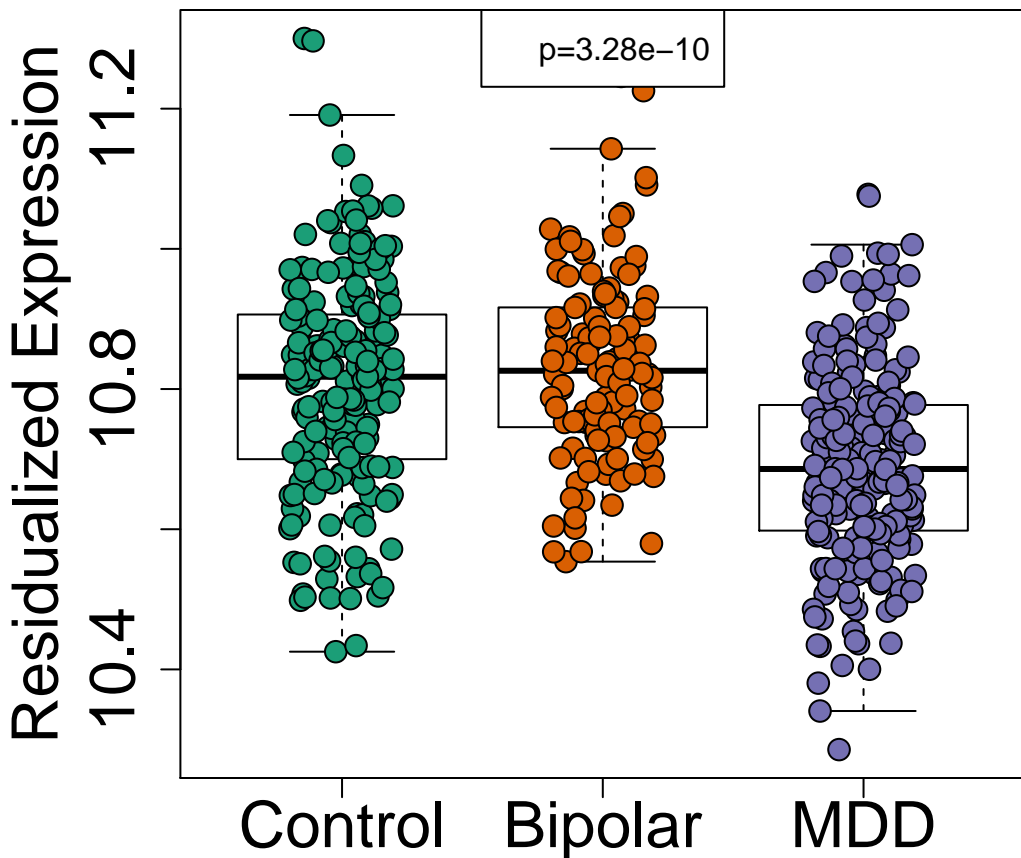
TUBA1B, InGen

chr12:49128939–49129241(-)



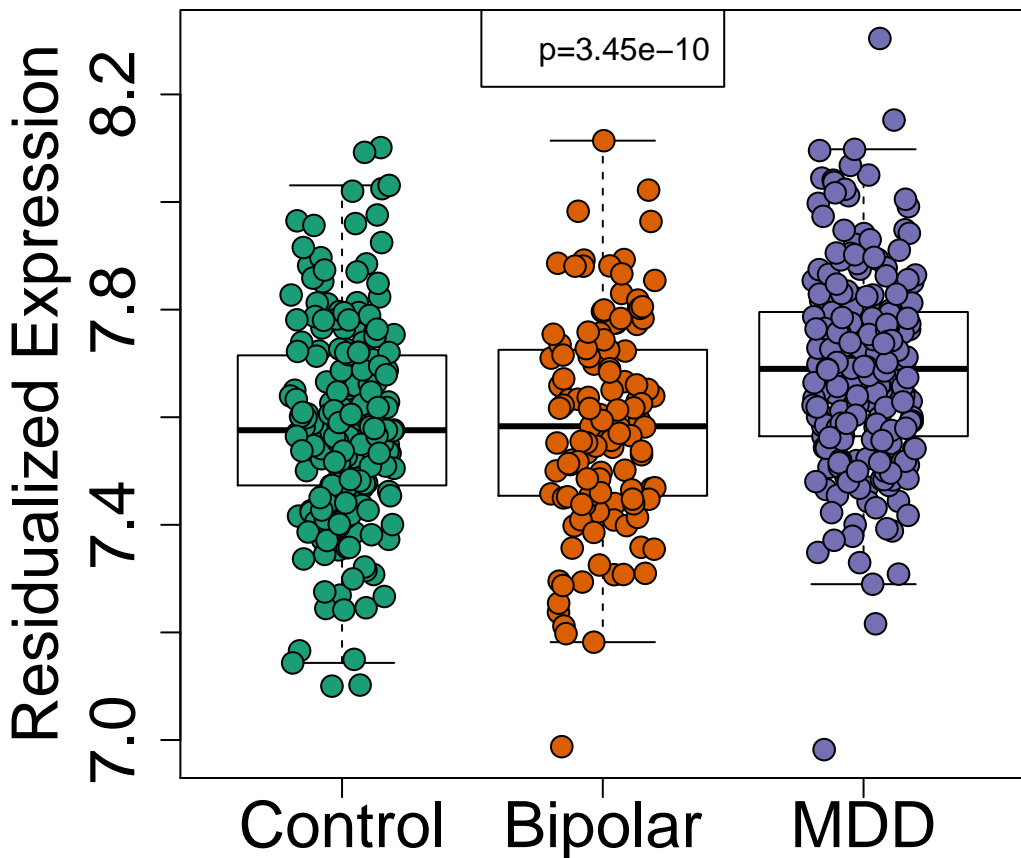
RPS21, InGen

chr20:62388365–62388456(+)



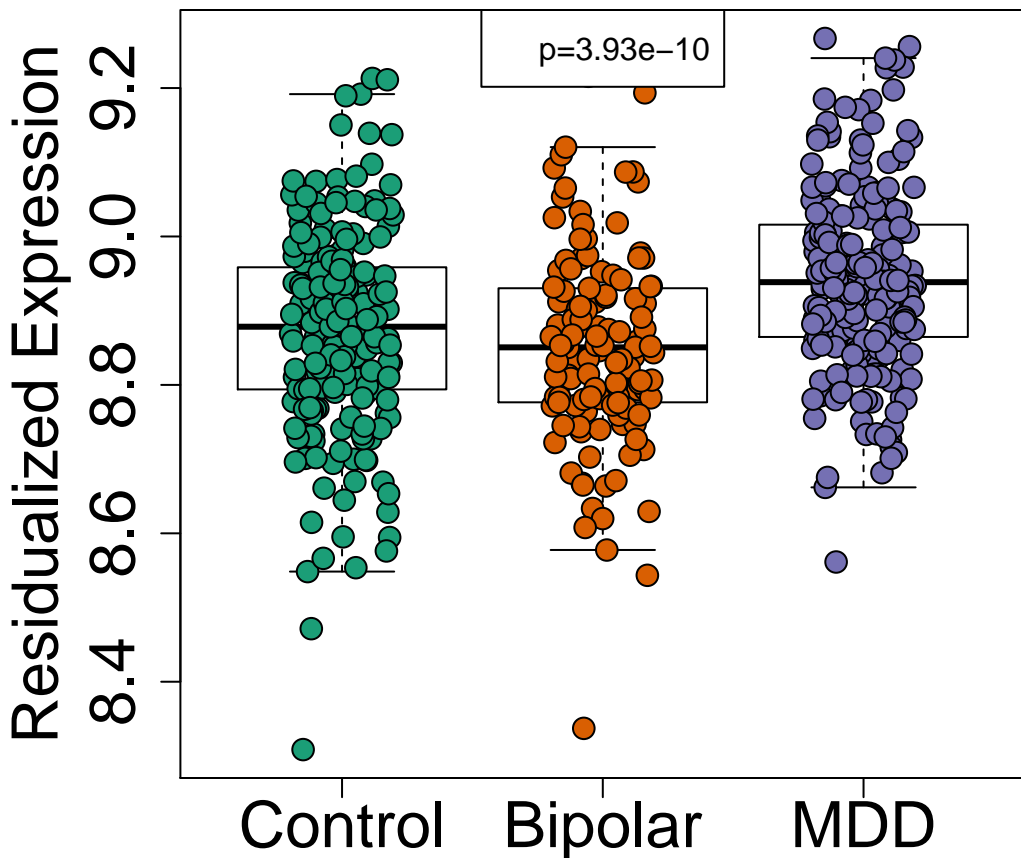
GPM6B, InGen

chrX:13785809–13816843(–)



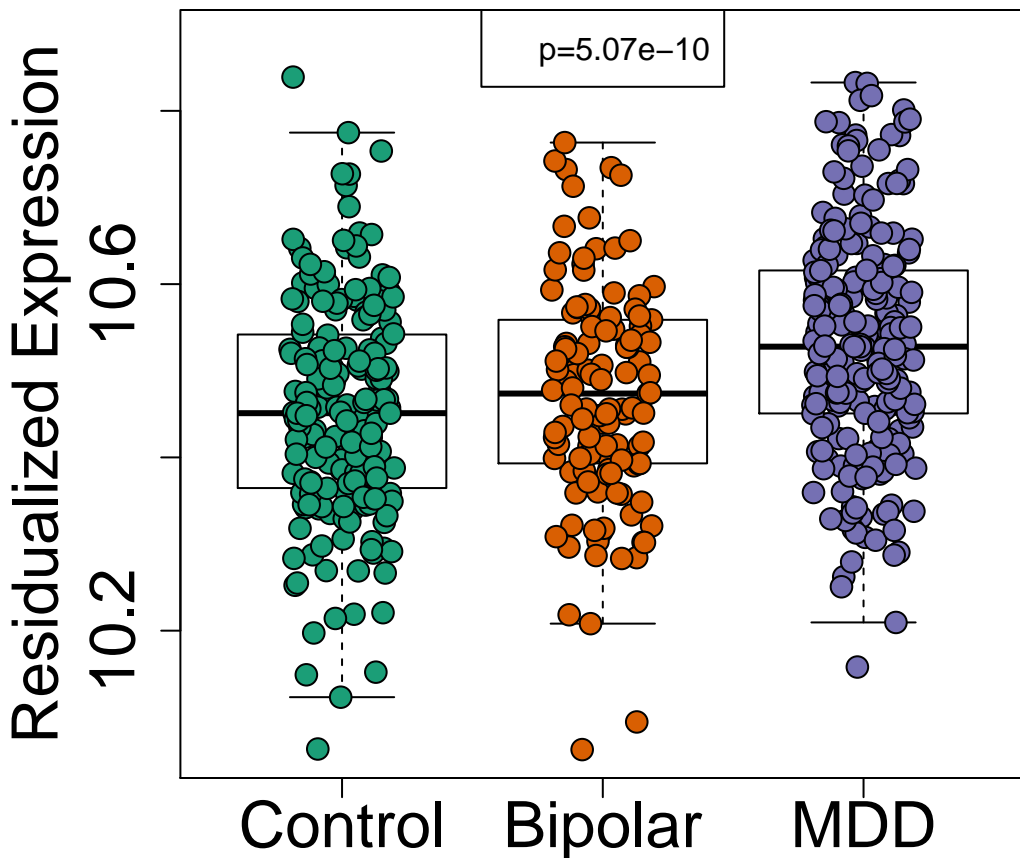
RPL18, InGen

chr19:48616203–48616725(-)



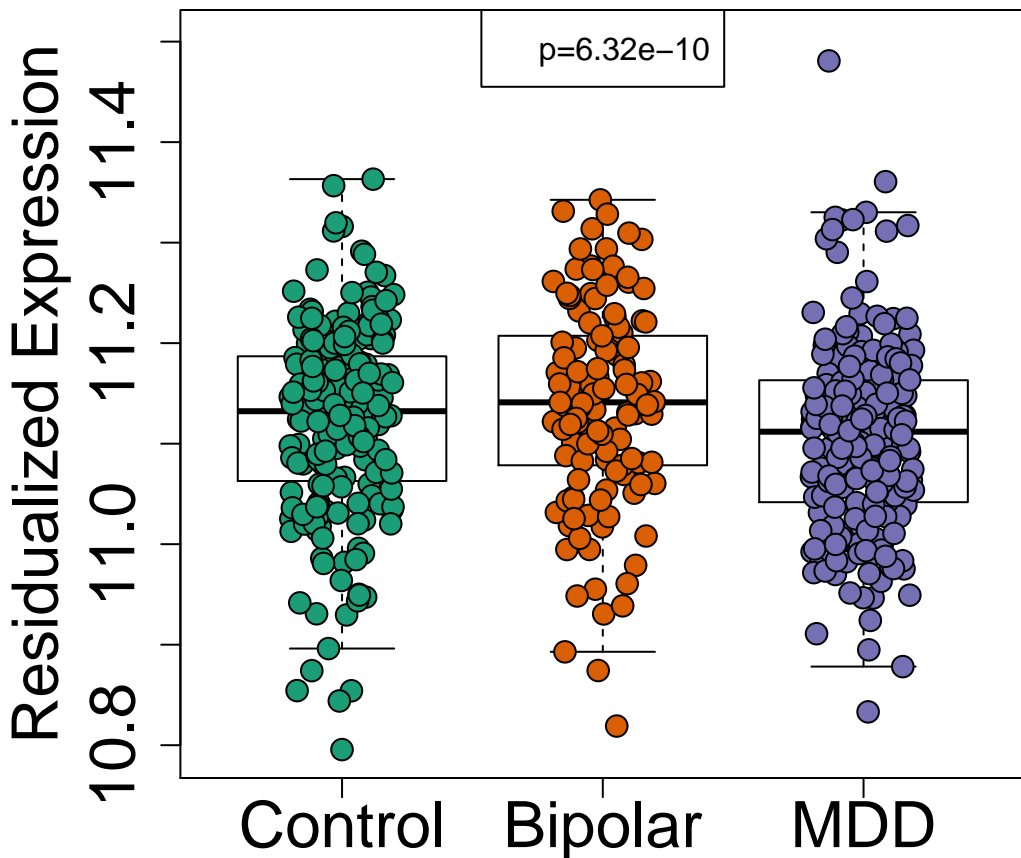
RPL22, InGen

chr1:6193055–6197651(-)



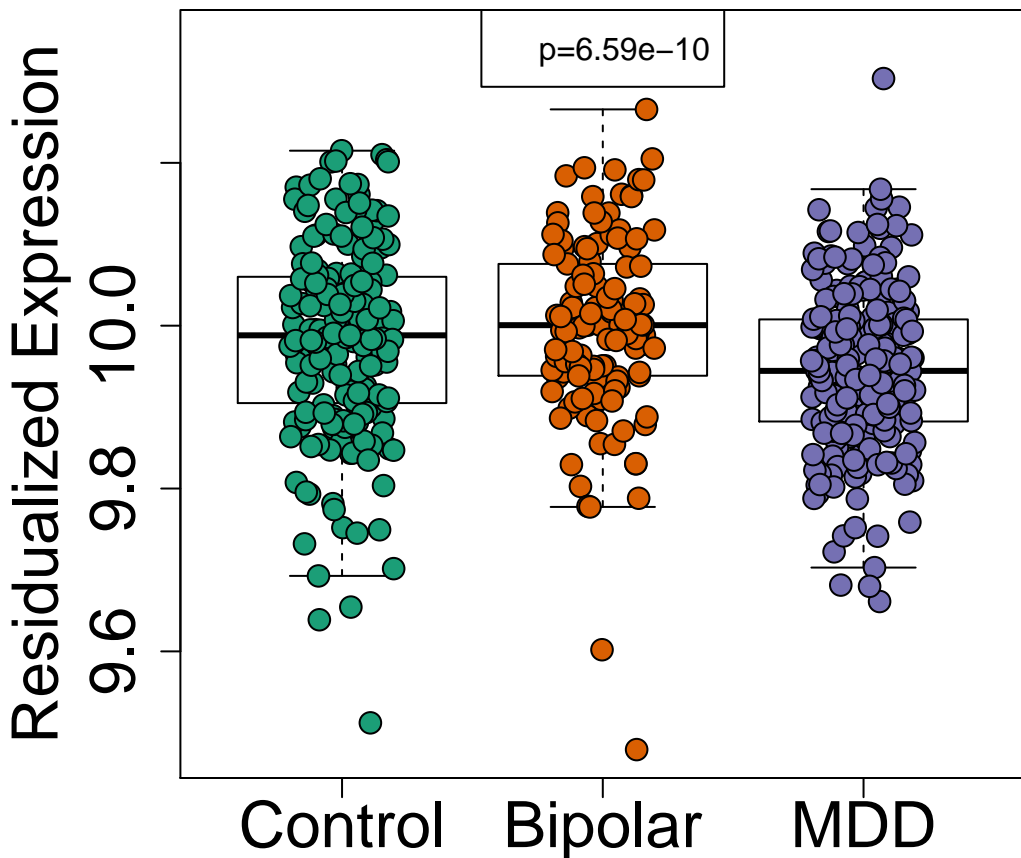
HSP90AB1, InGen

chr6:44248777–44249376(+)



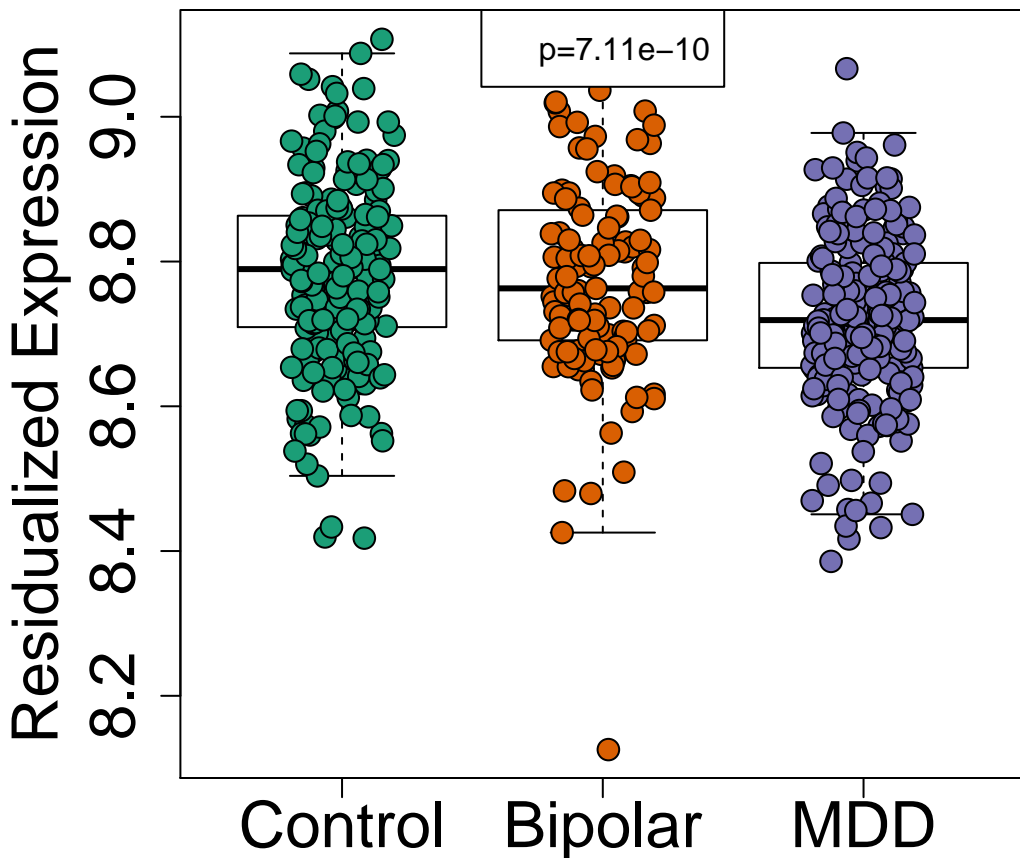
GNAS, InGen

chr20:58909424–58909520(+)



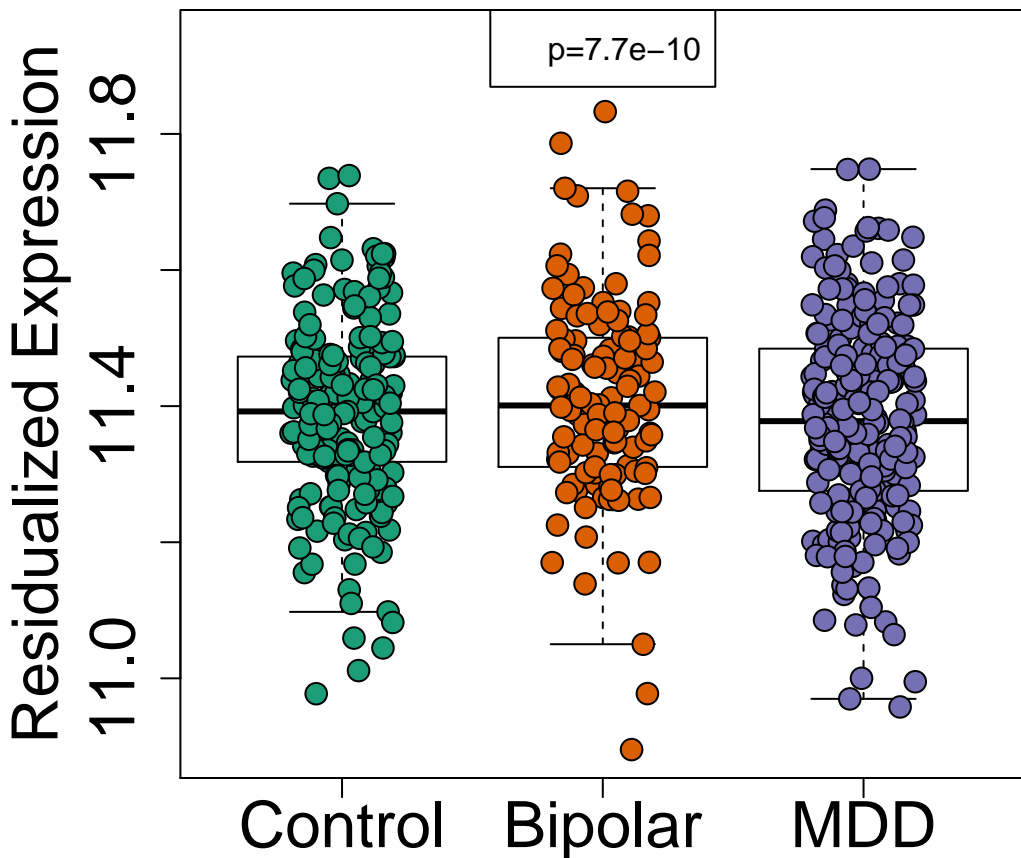
SPTBN1, InGen

chr2:54647262–54648985(+)



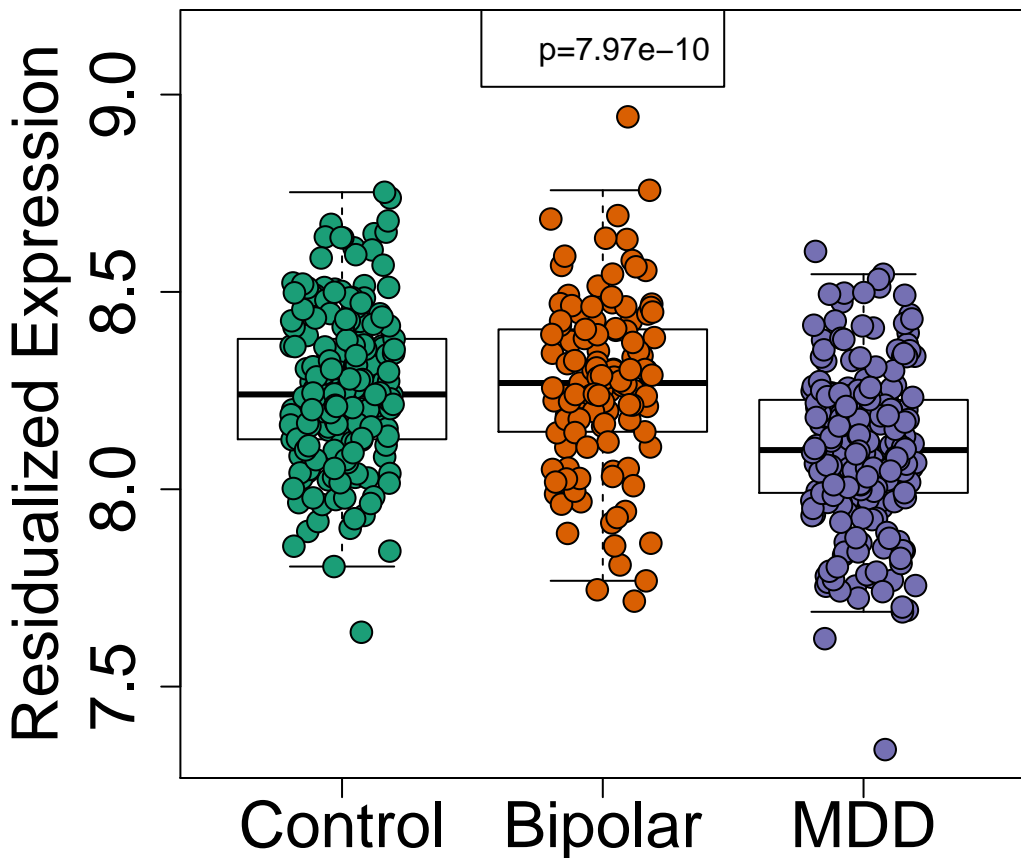
CPE, InGen

chr4:165495678–165497511(+)



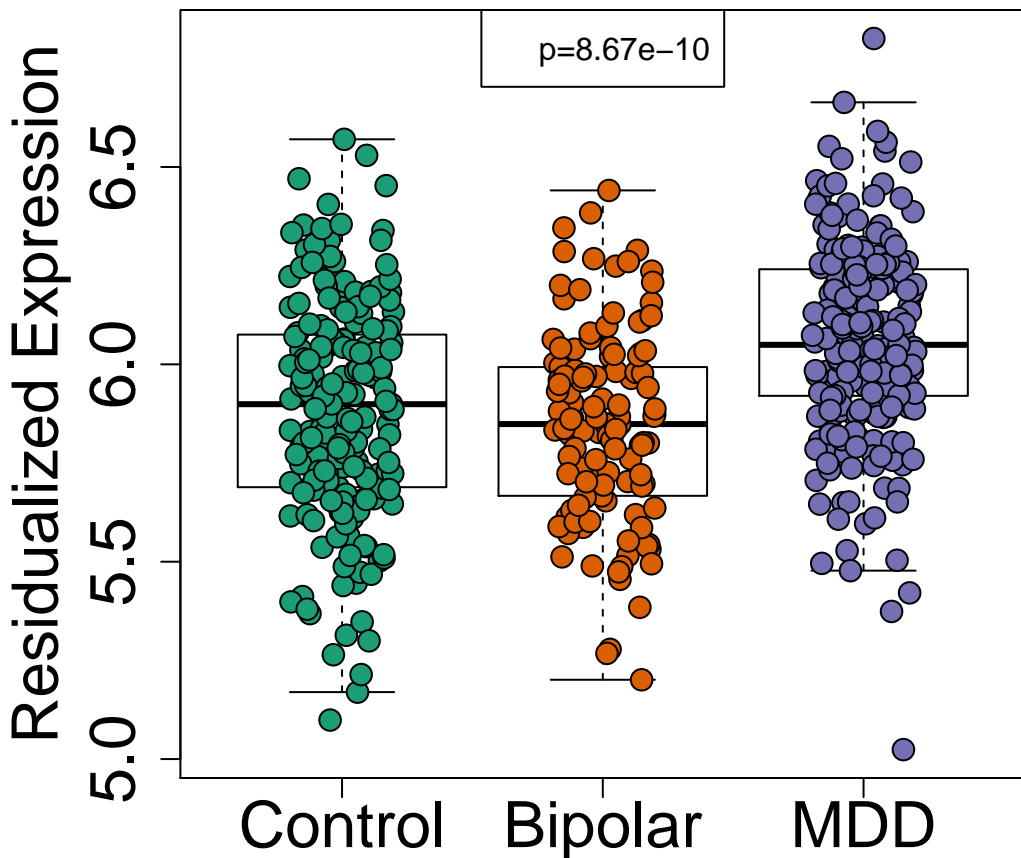
FUS, InGen

chr16:31182423–31182512(+)



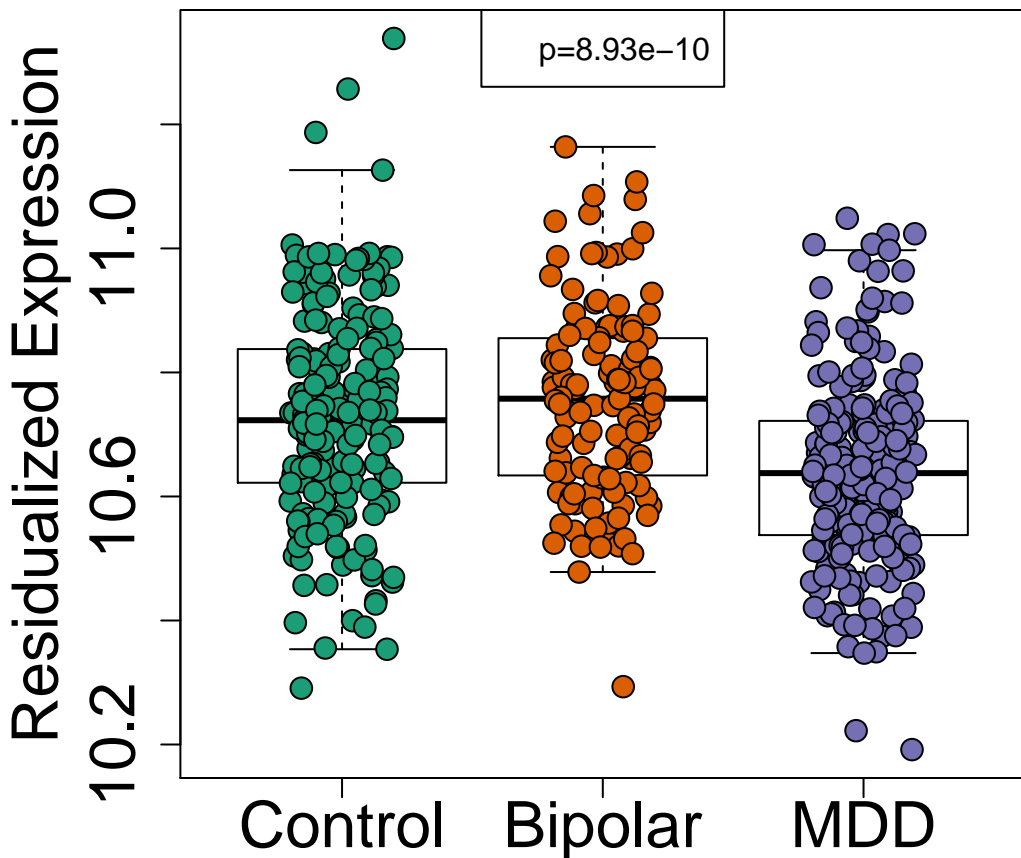
GLTSCR2, InGen

chr19:47750287–47750907(+)



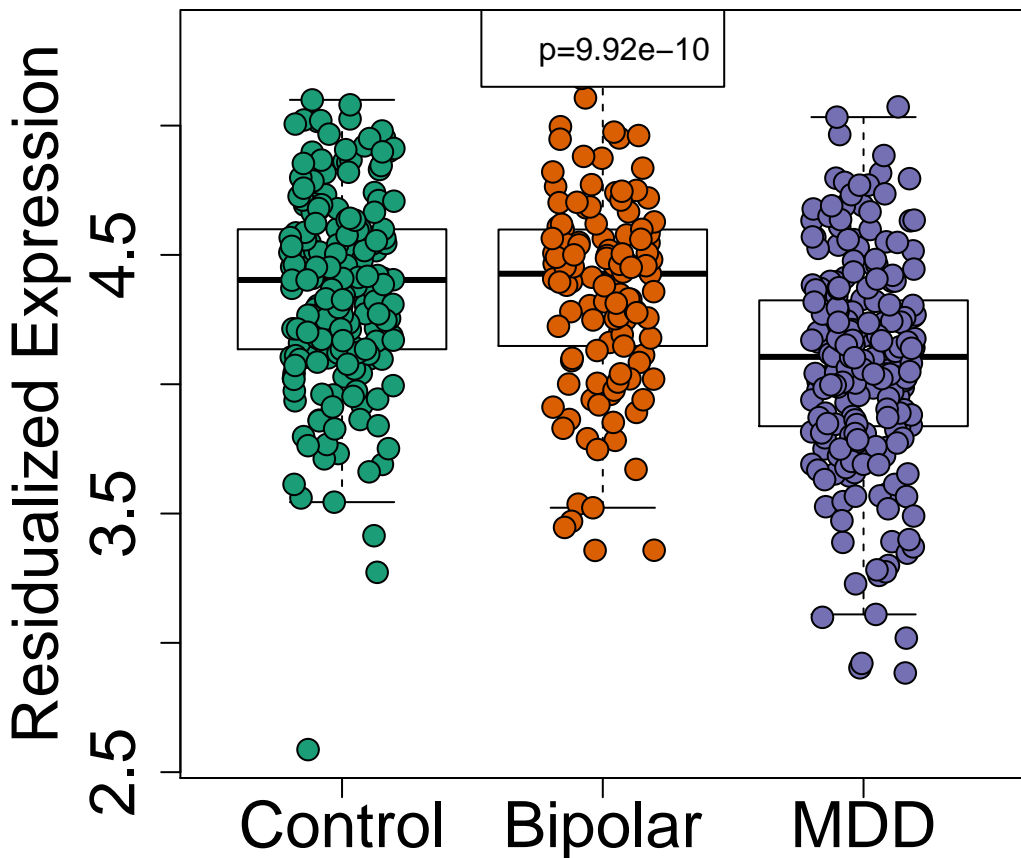
RPS21, InGen

chr20:62387915–62388308(+)



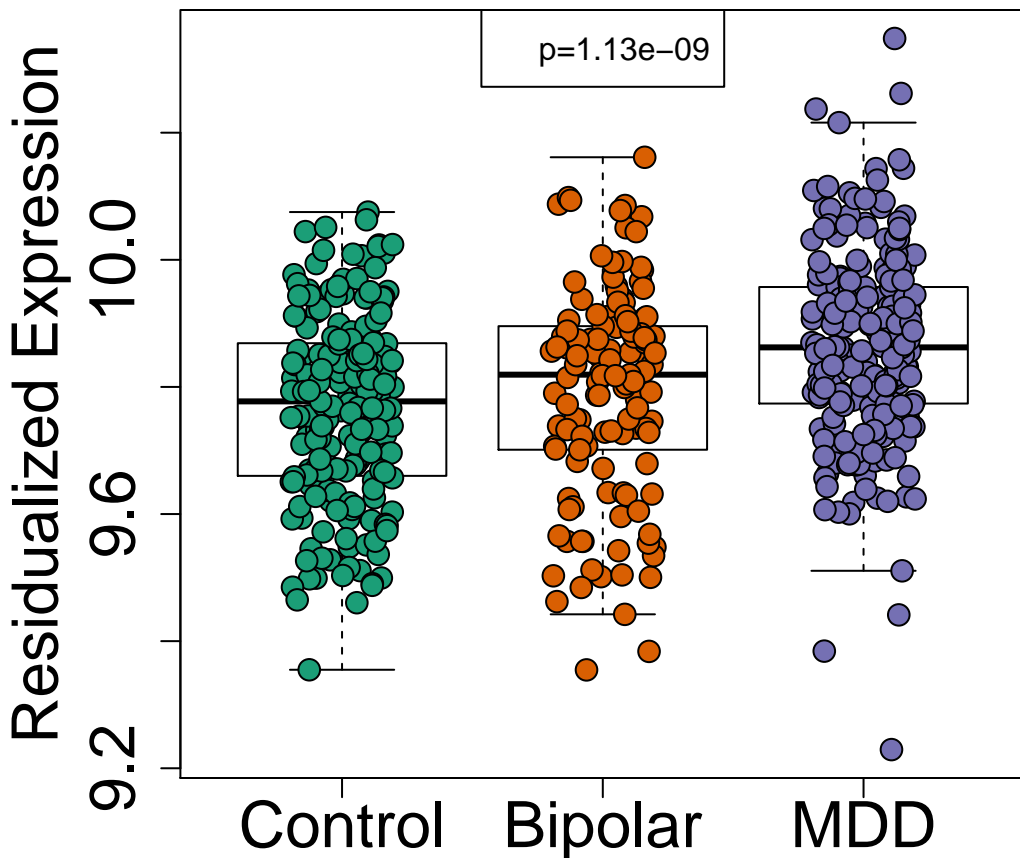
PPP4C, InGen

chr16:30076476–30081258(+)



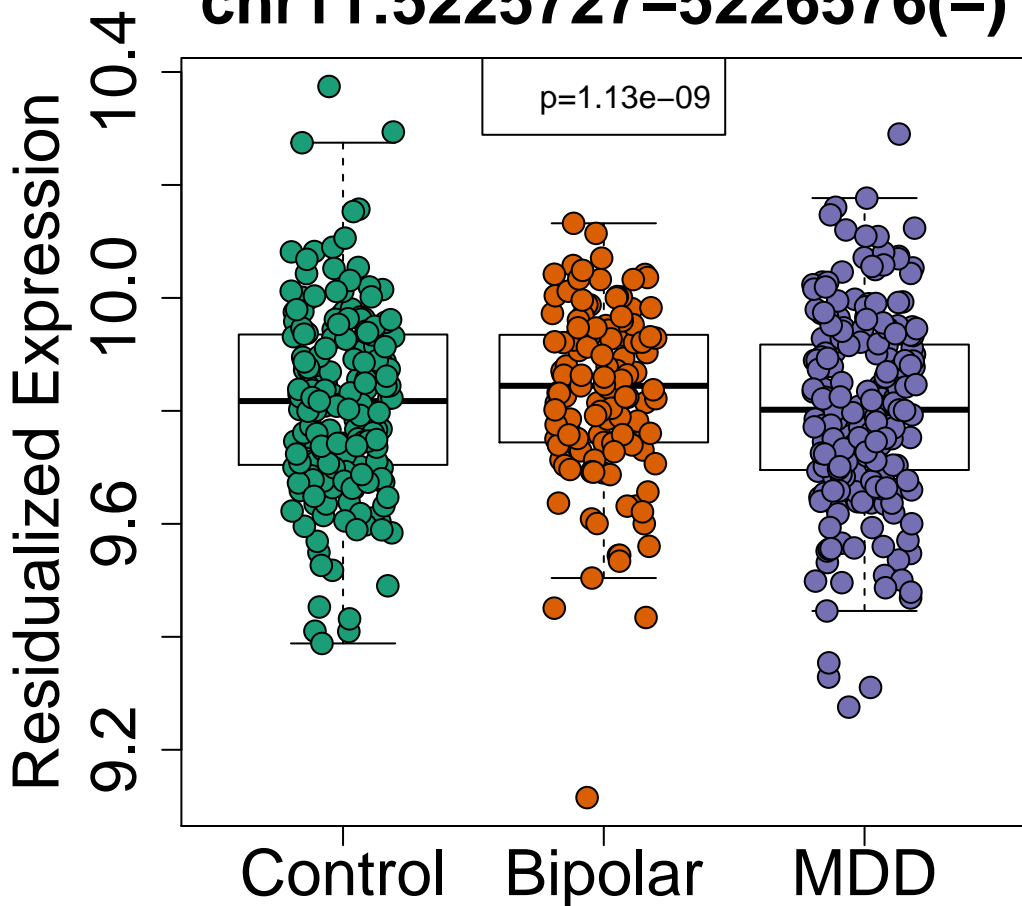
GLUD1, InGen

chr10:87060826–87060914(-)



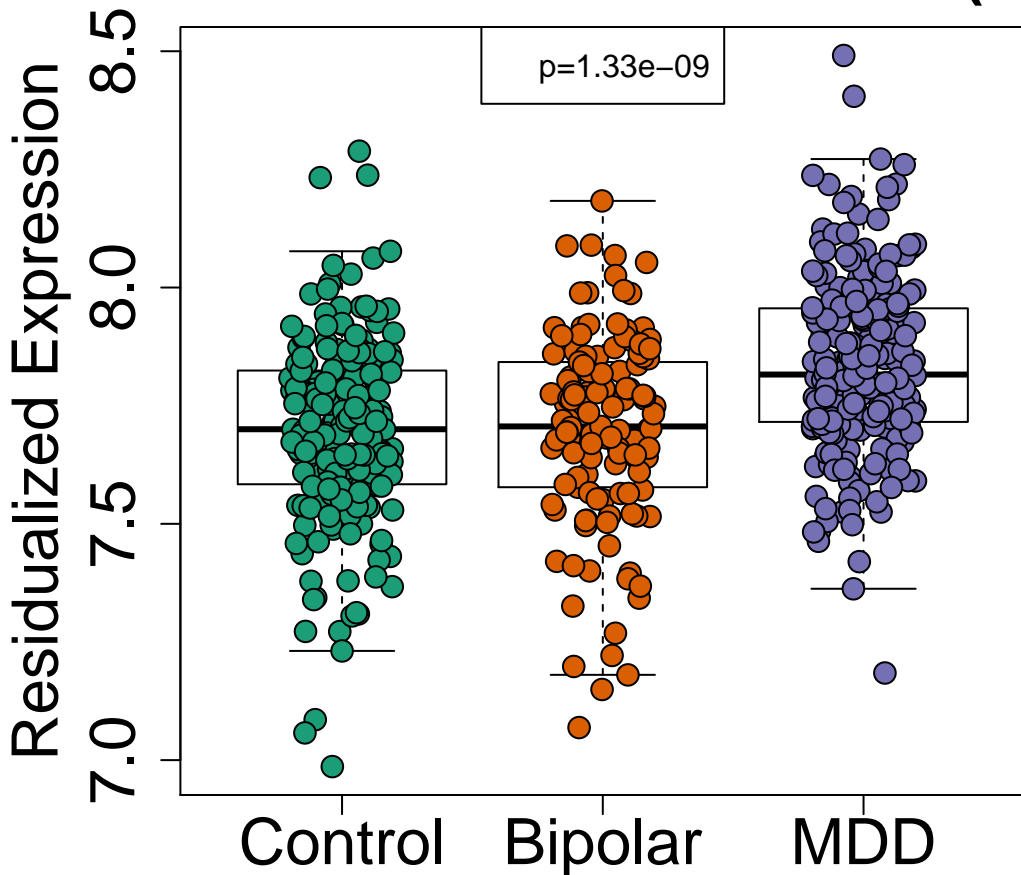
HBB, InGen

chr11:5225727-5226576(-)



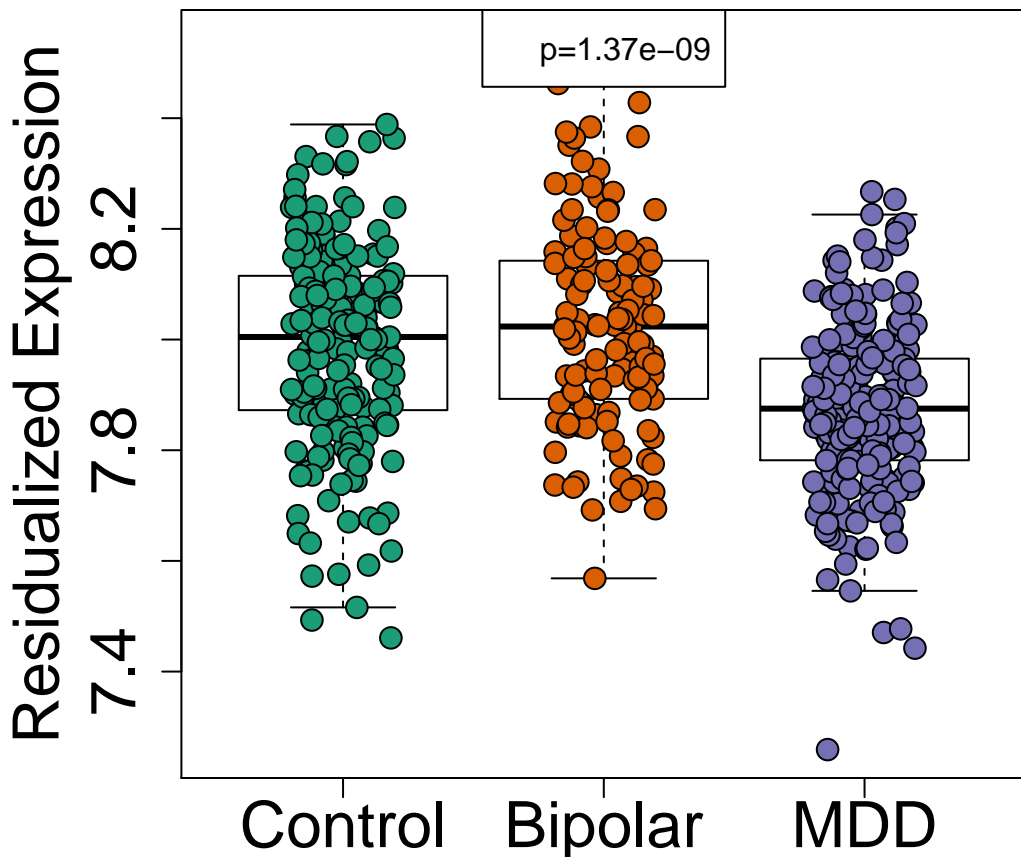
GNG3, InGen

chr11:62708395–62708677(+)



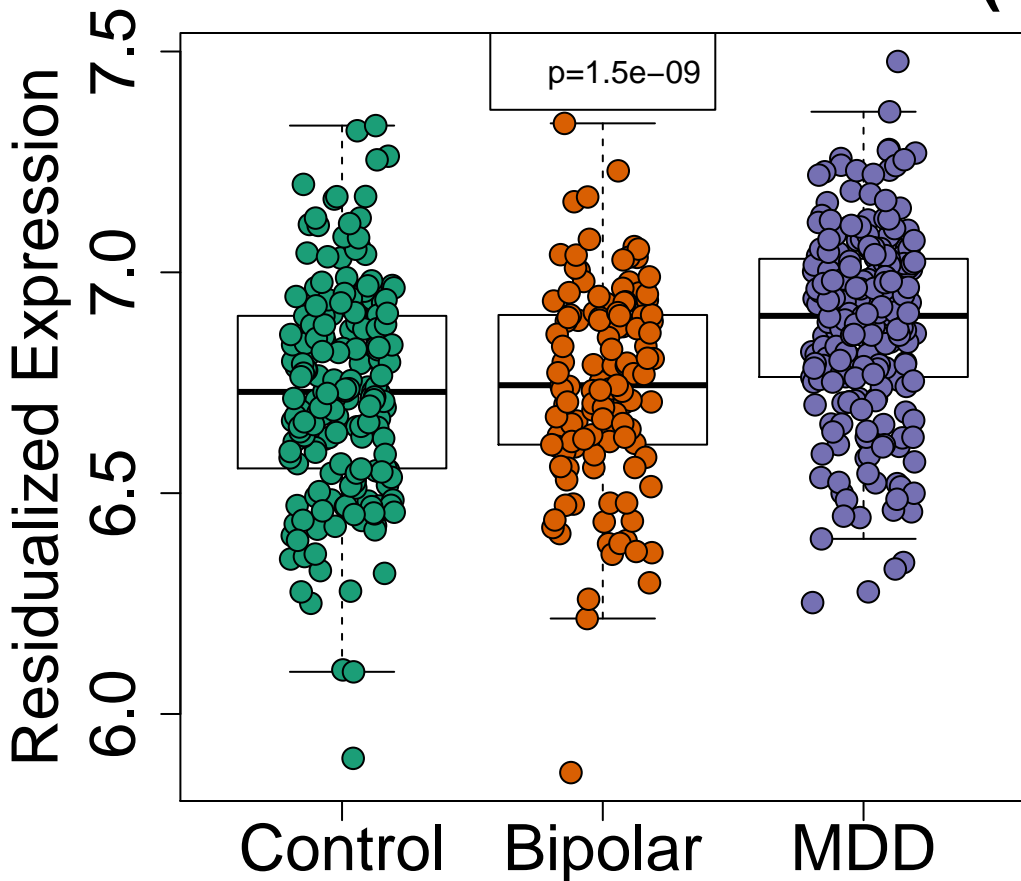
ATP2A2, InGen

chr12:110339722–110340658(+)



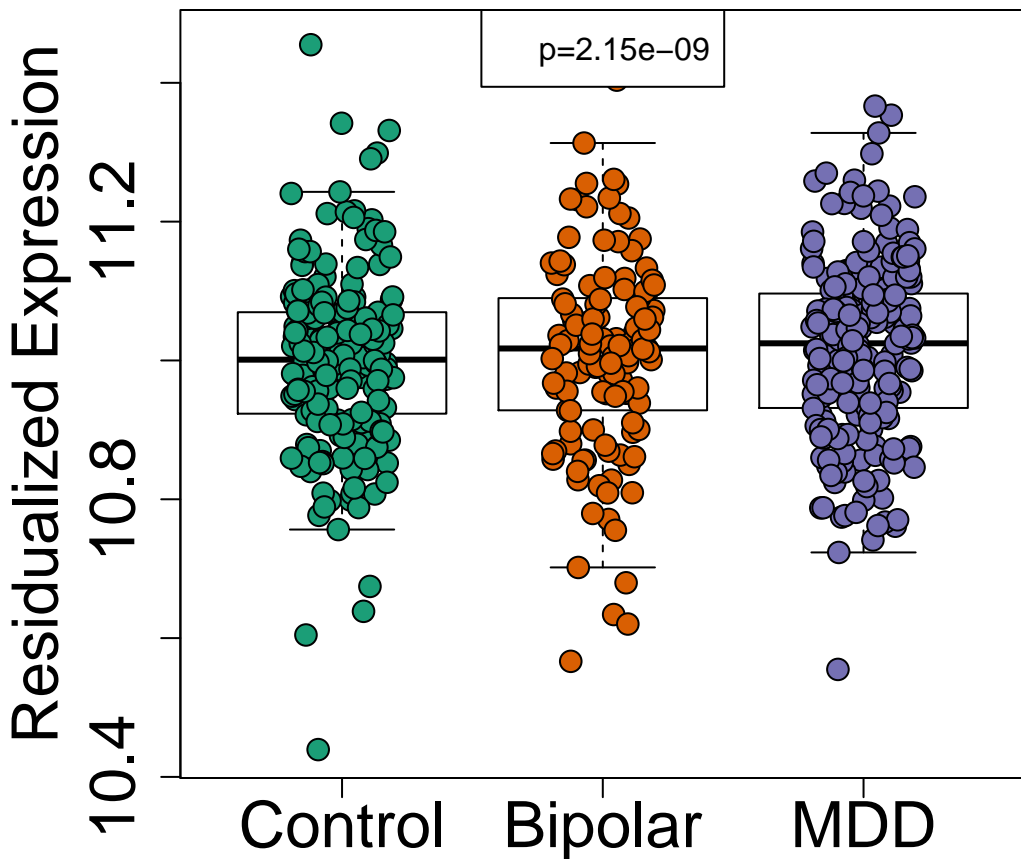
ATP1A1, InGen

chr1:116401007–116401129(+)



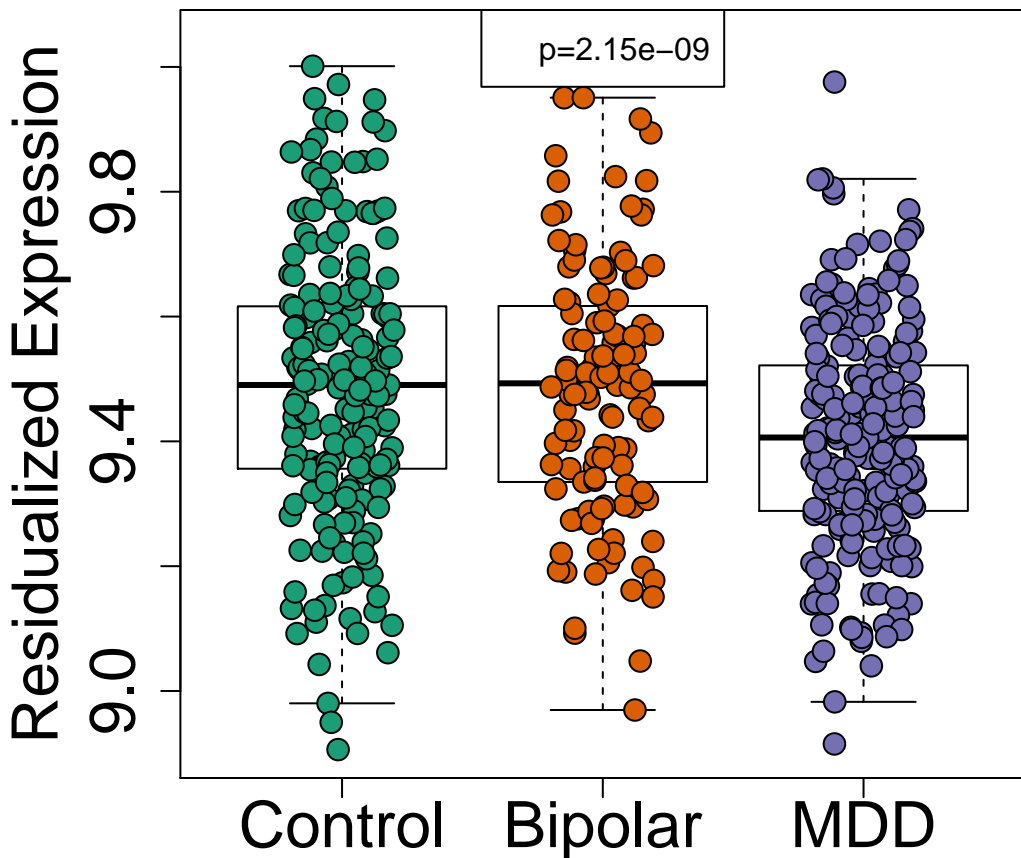
CPE, InGen

chr4:165487578–165493170(+)



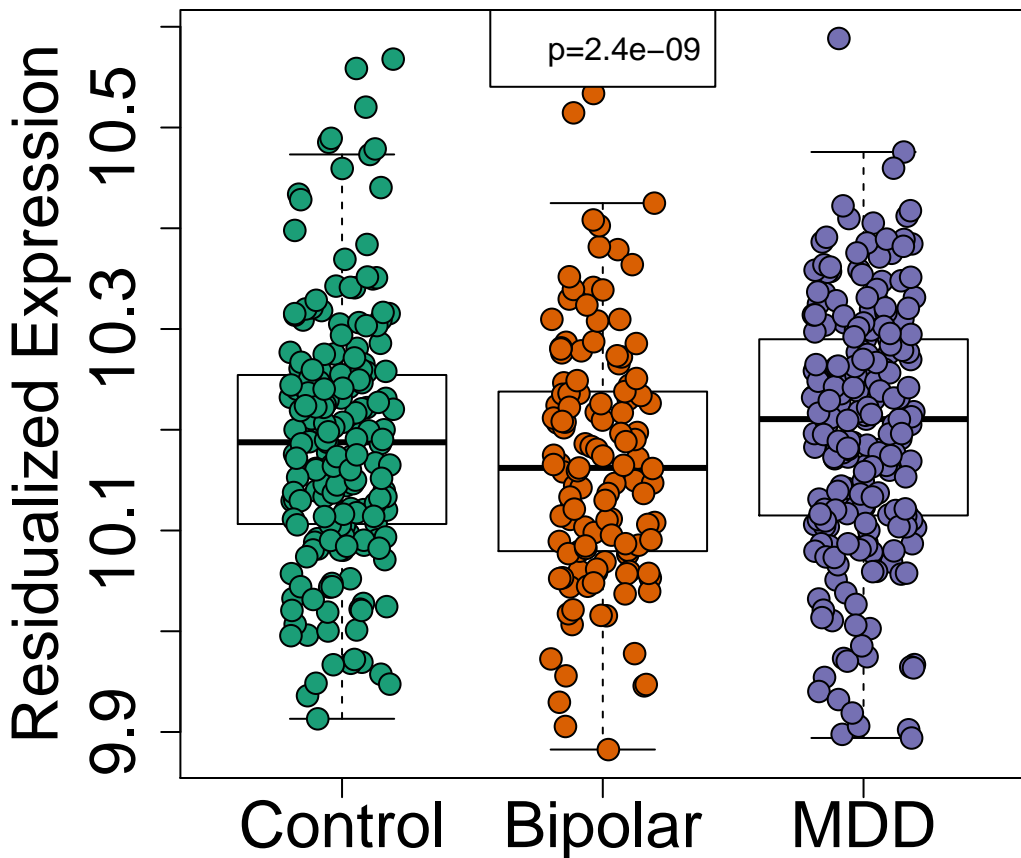
ATP1A3, InGen

chr19:41975798–41976415(-)



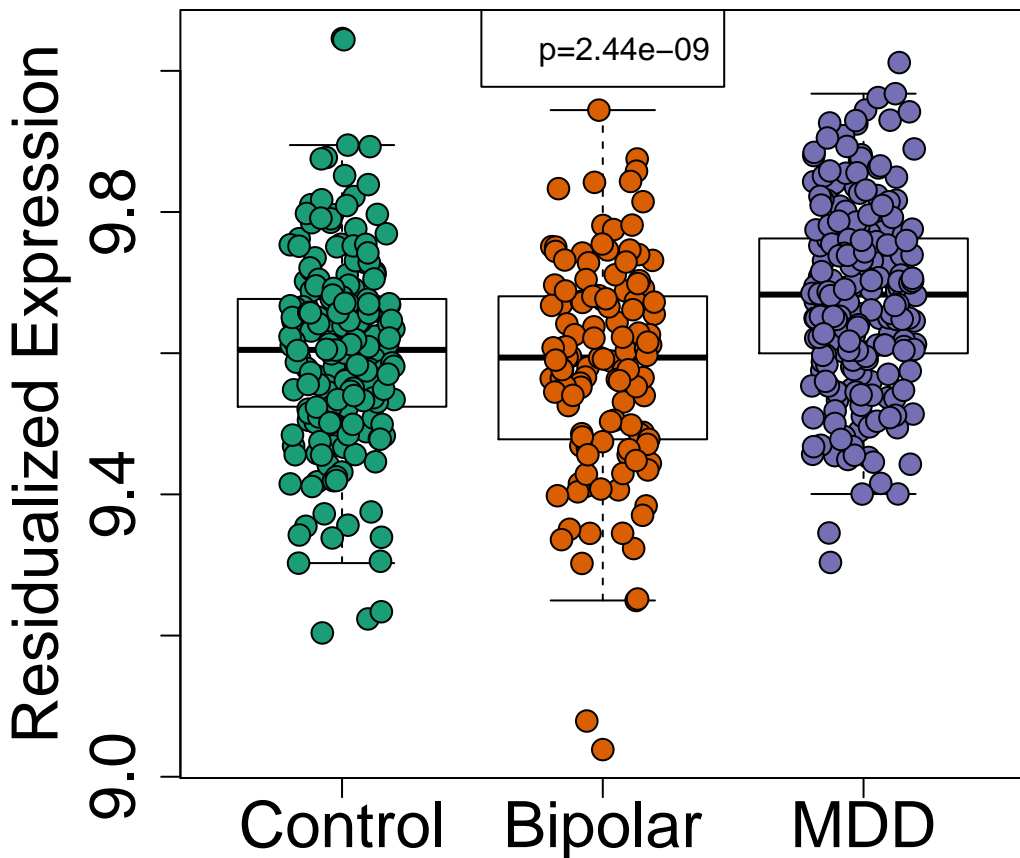
RPS6, InGen

chr9:19378919–19379486(-)



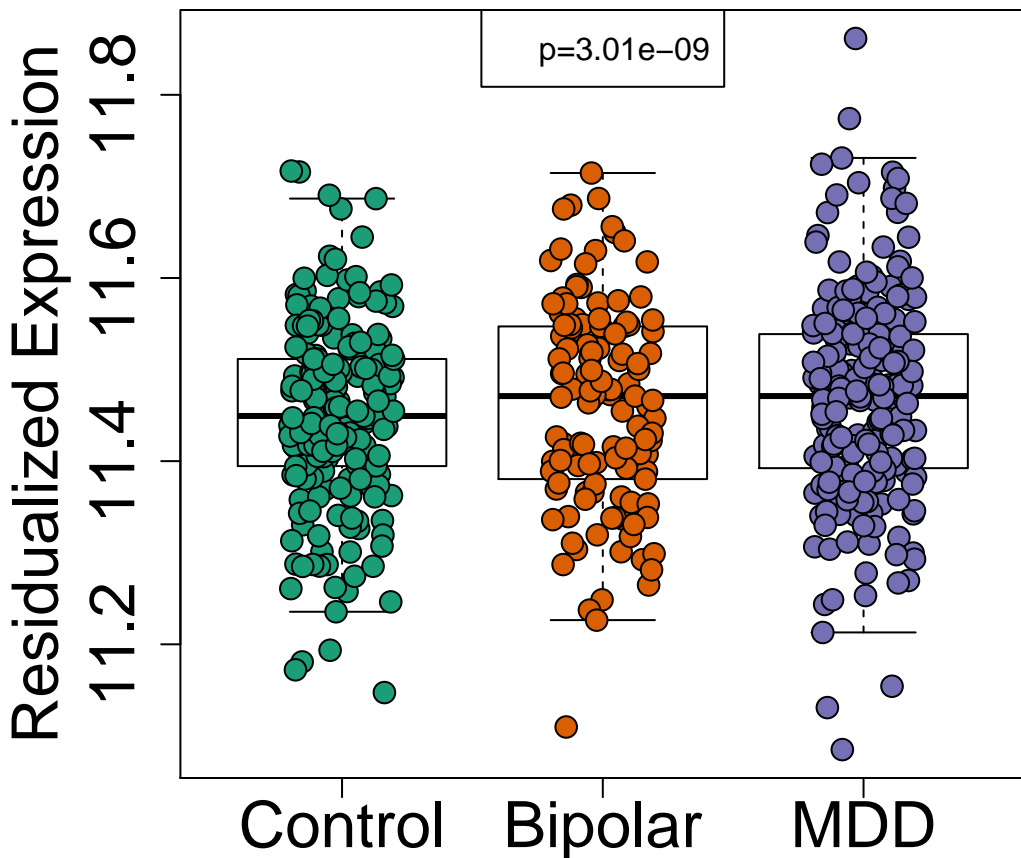
ATP5L, InGen

chr11:118407098–118408999(+)



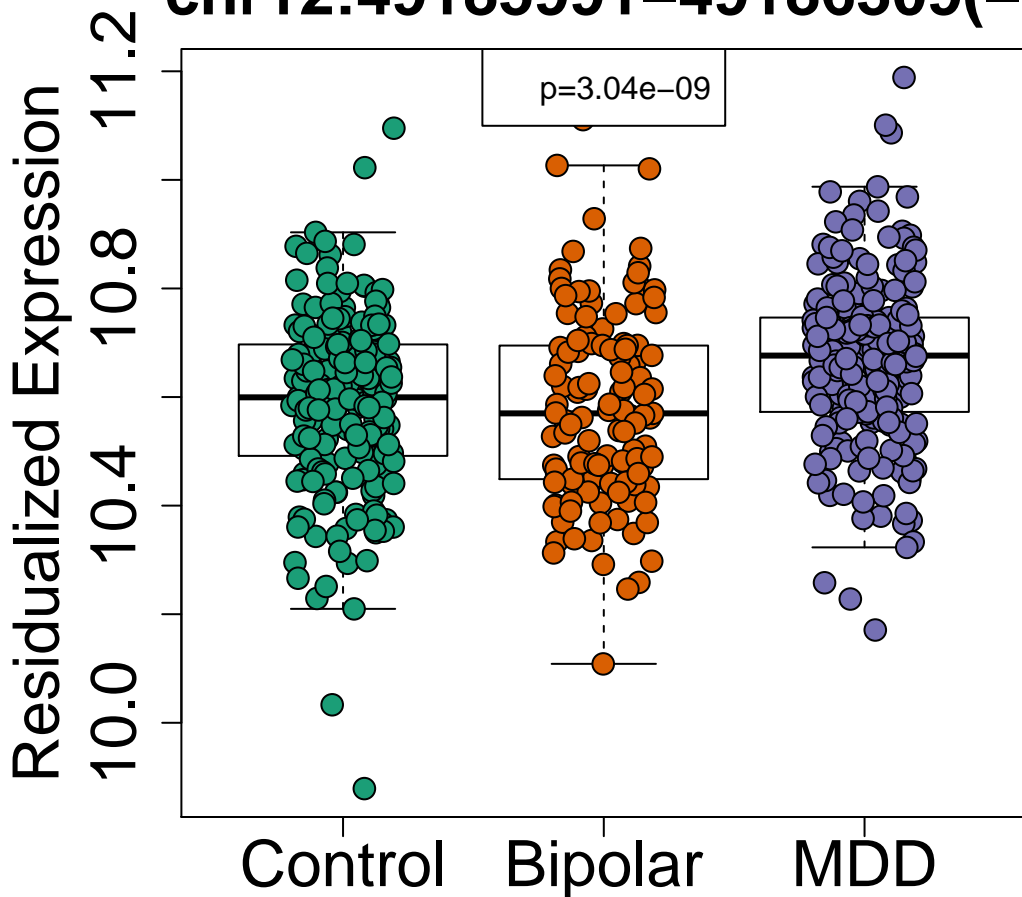
PPIA, InGen

chr7:44799875–44801286(+)



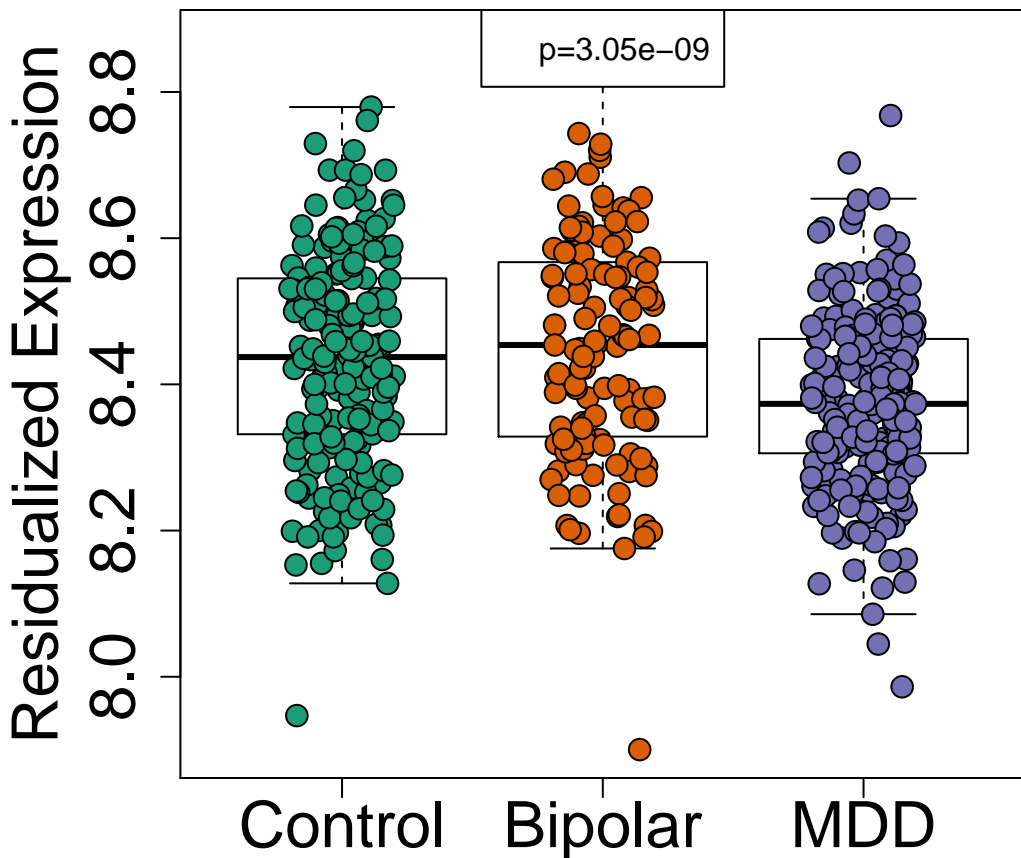
TUBA1A, InGen

chr12:49185991–49186309(-)



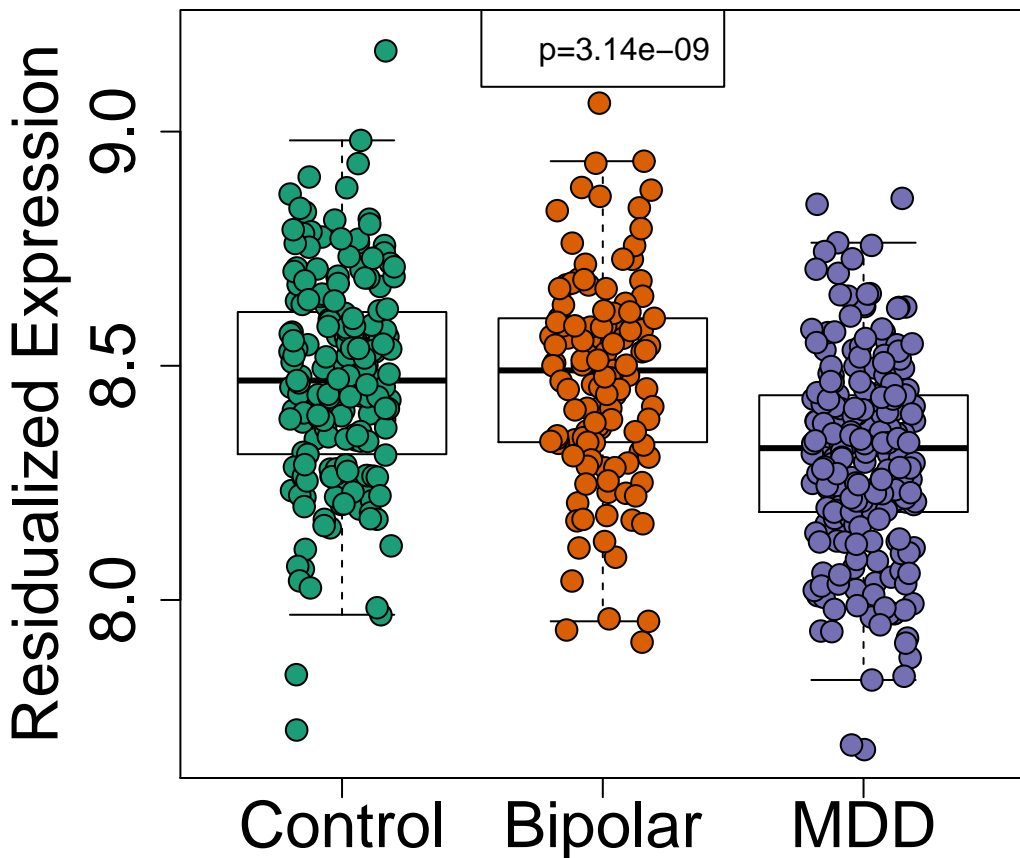
ATP9A, InGen

chr20:51629073–51639342(-)



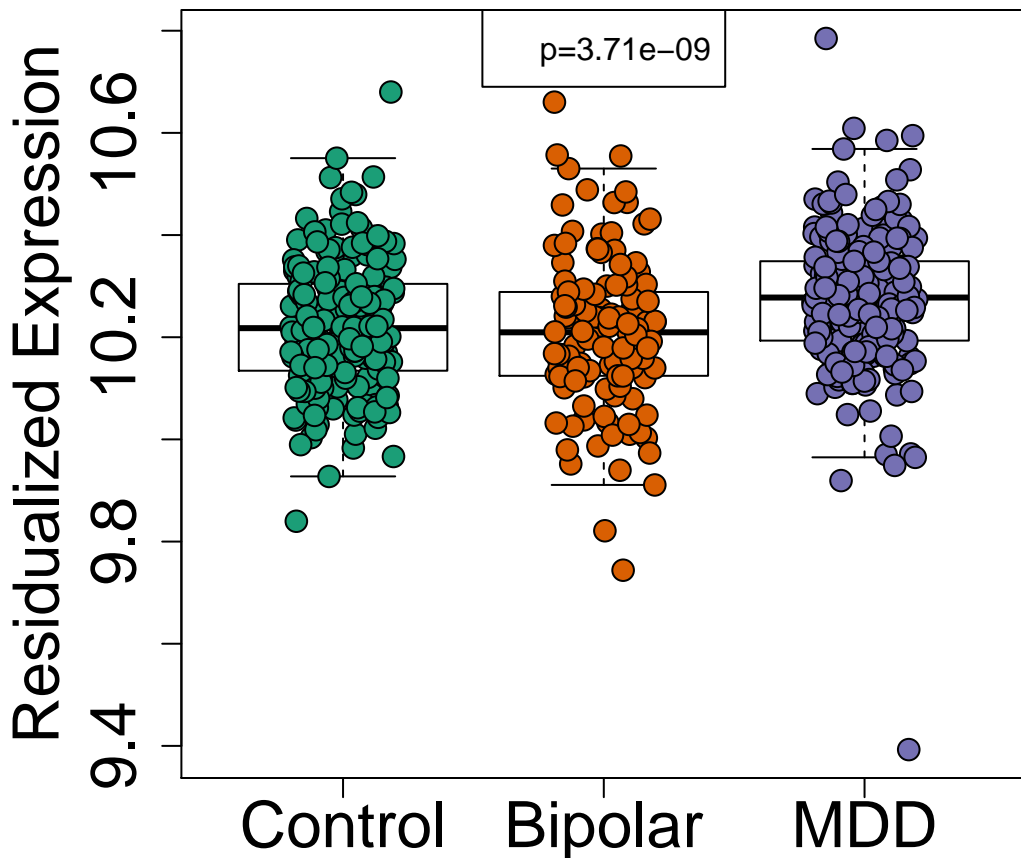
FUS, InGen

chr16:31180228–31182397(+)



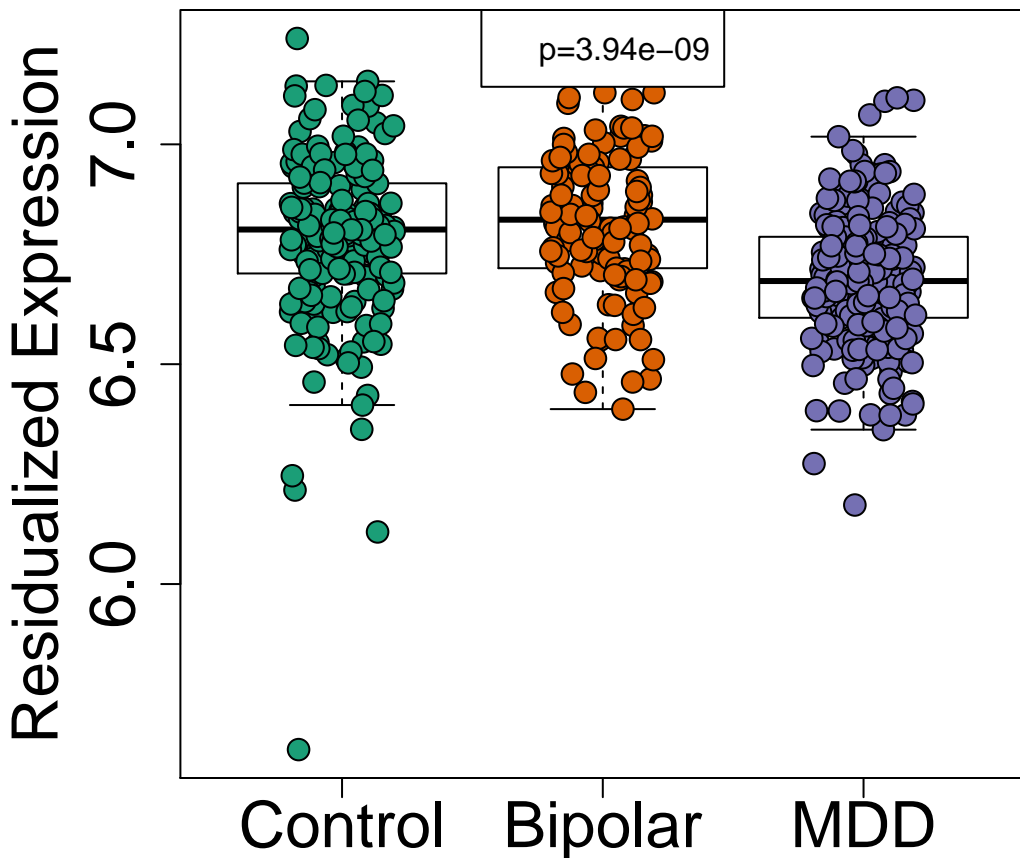
HSPA8, InGen

chr11:123059262–123059472(-)



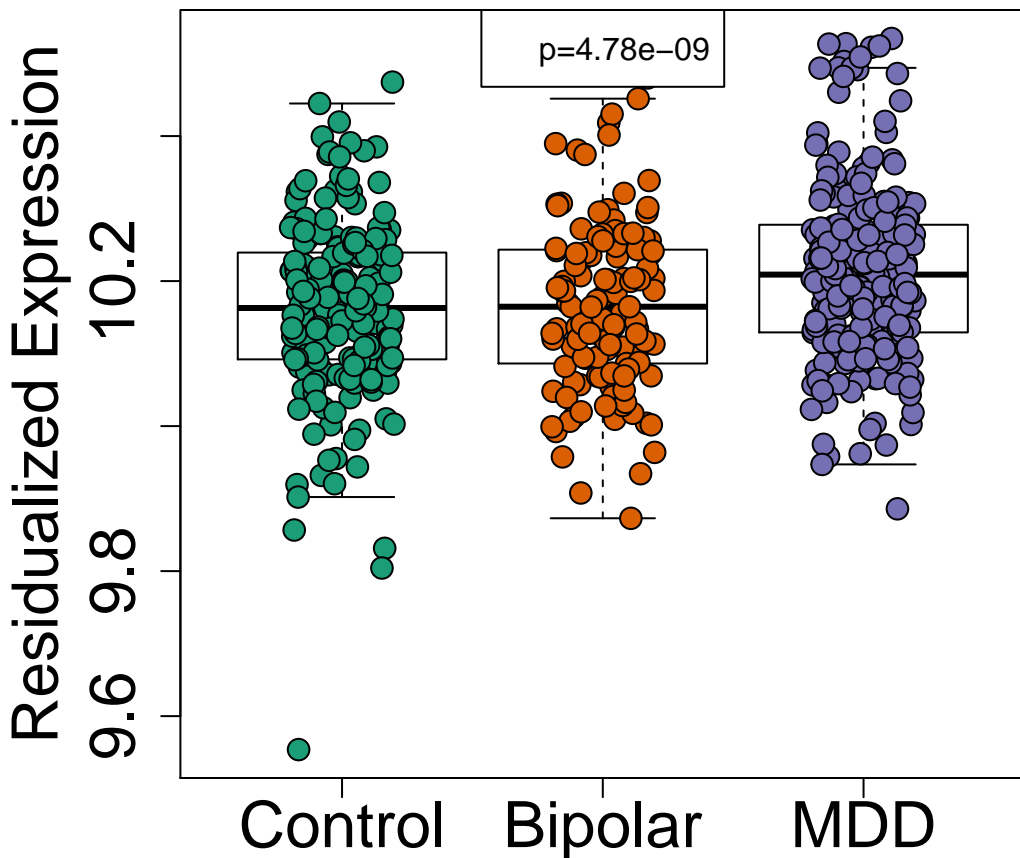
PI4KA, InGen

chr22:20798688–20799092(-)



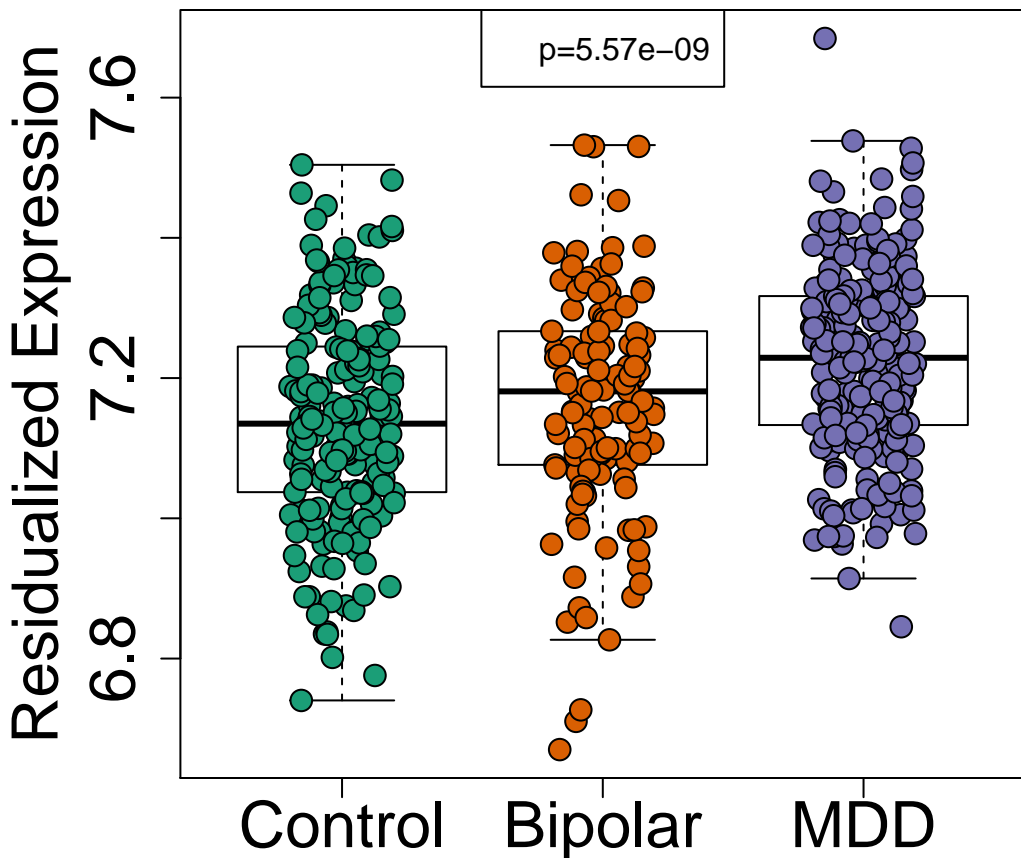
RPS4X, InGen

chrX:72275725-72276156(-)



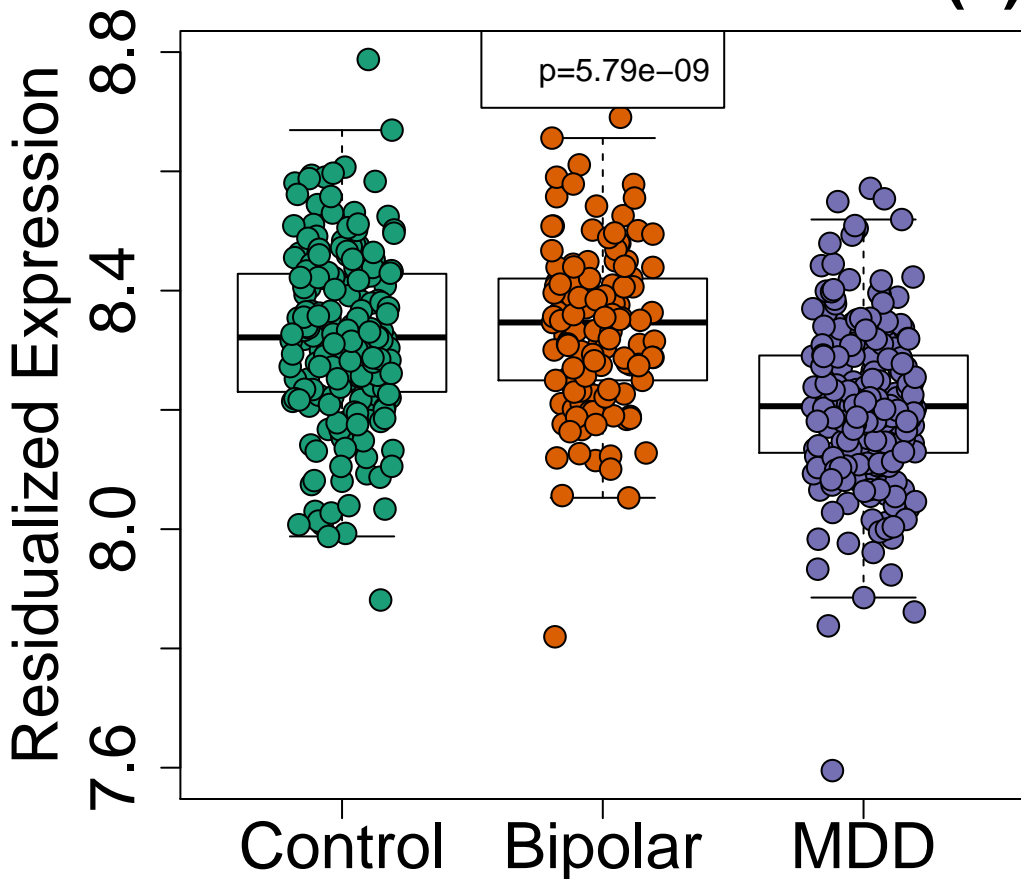
FKBP8, InGen

chr19:18532256–18532663(-)



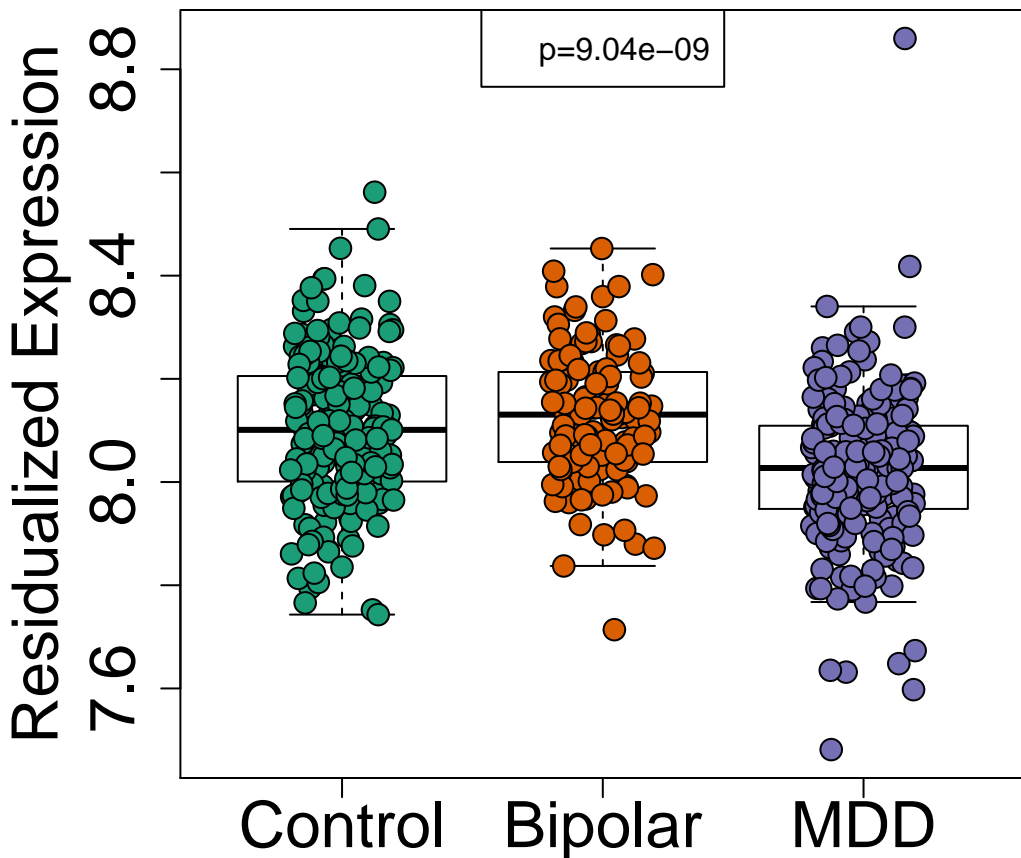
HNRNPA2B1, InGen

chr7:26197733–26200571(-)



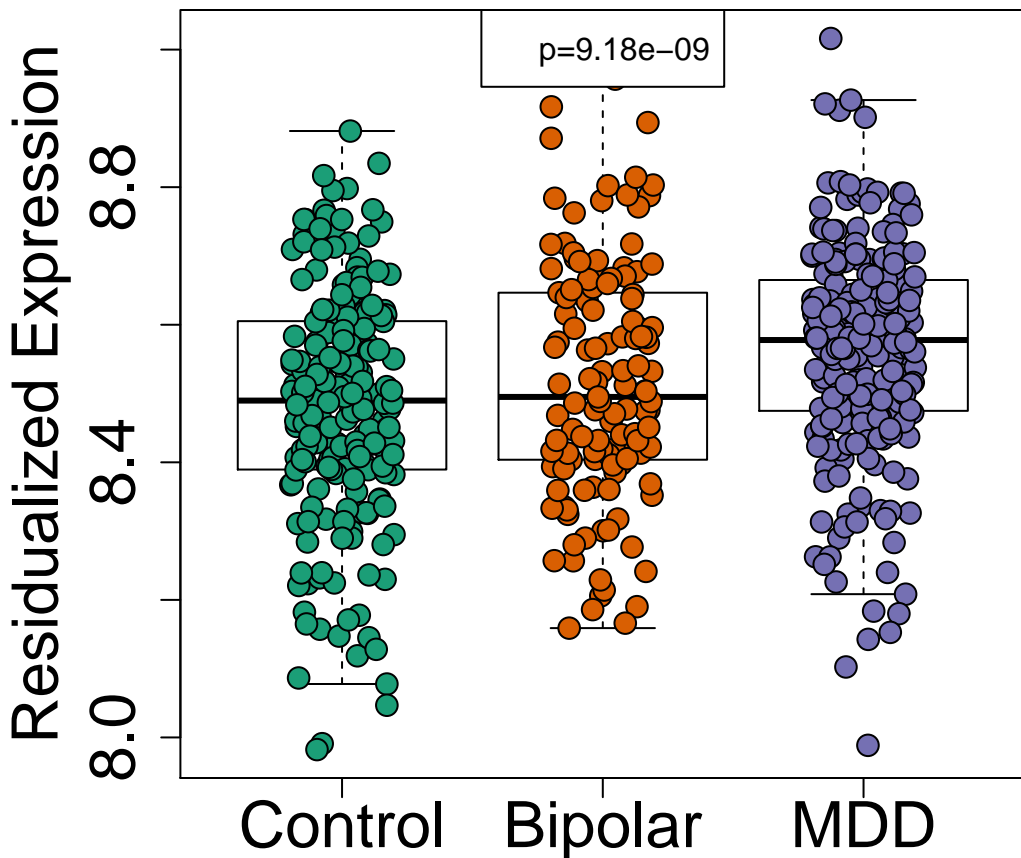
ATP6V0A1, InGen

chr17:42487368–42490486(+)



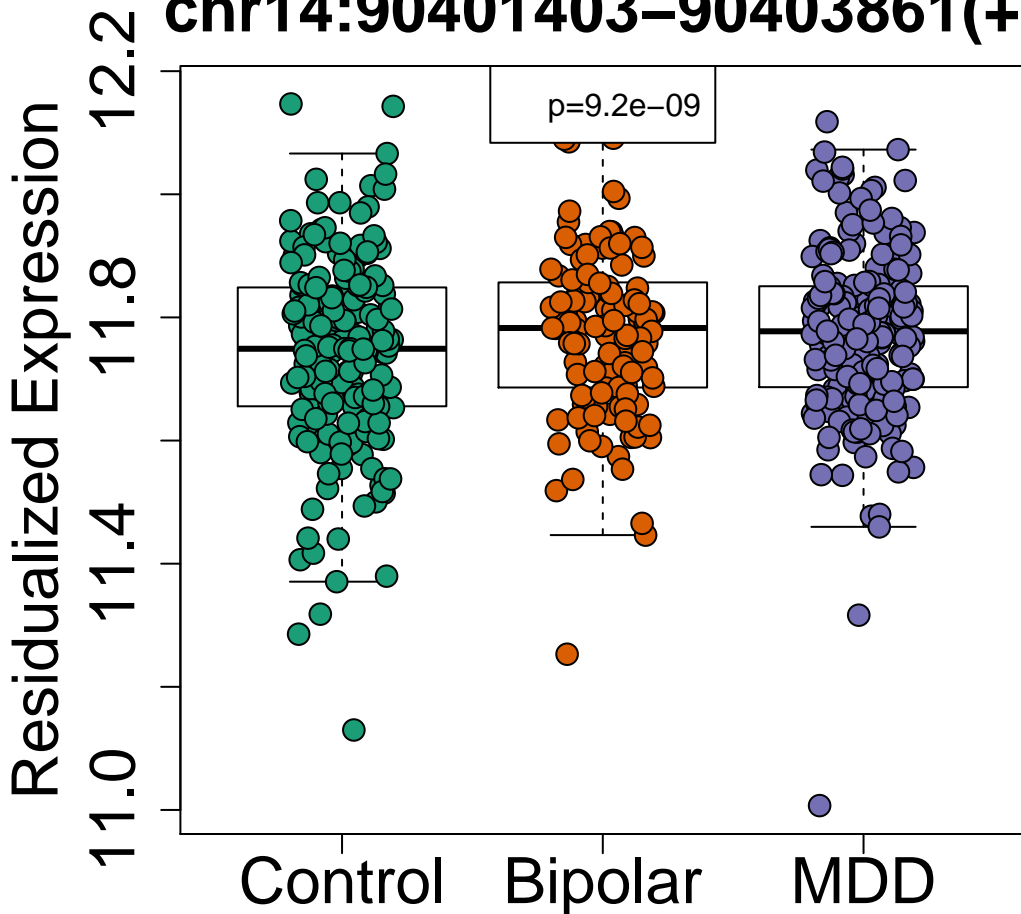
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chr12:54282487–54282572(+)



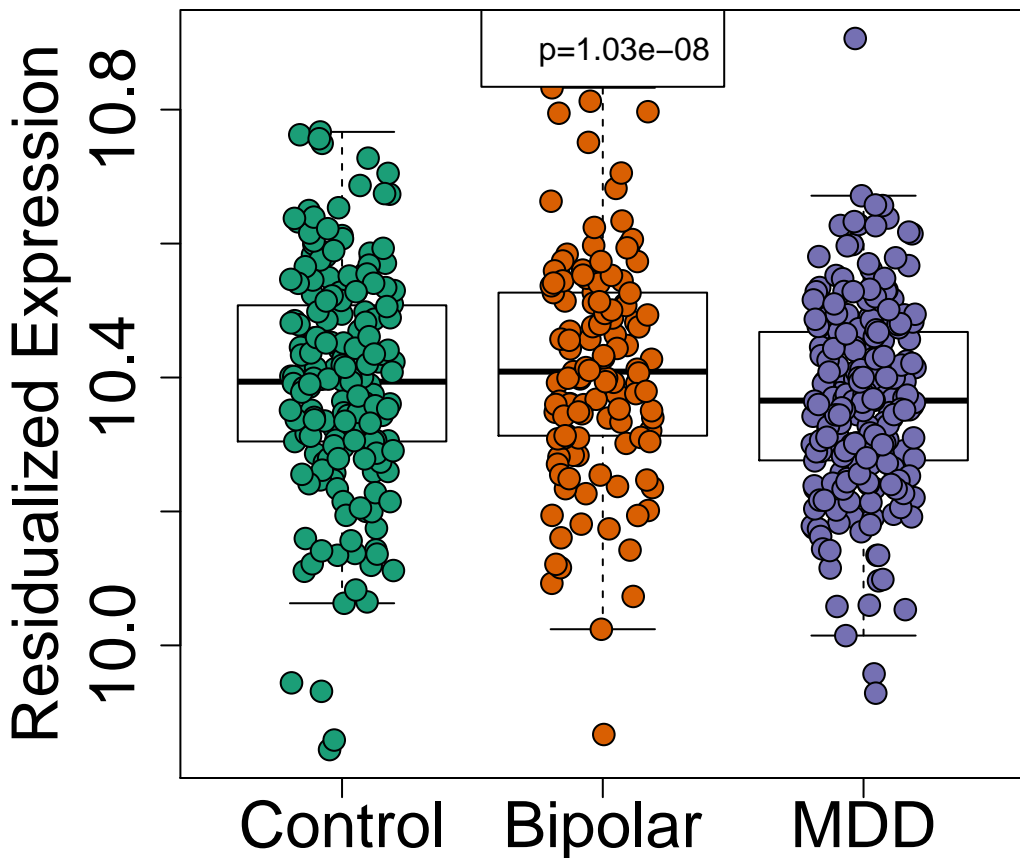
CALM1, InGen

chr14:90401403–90403861(+)



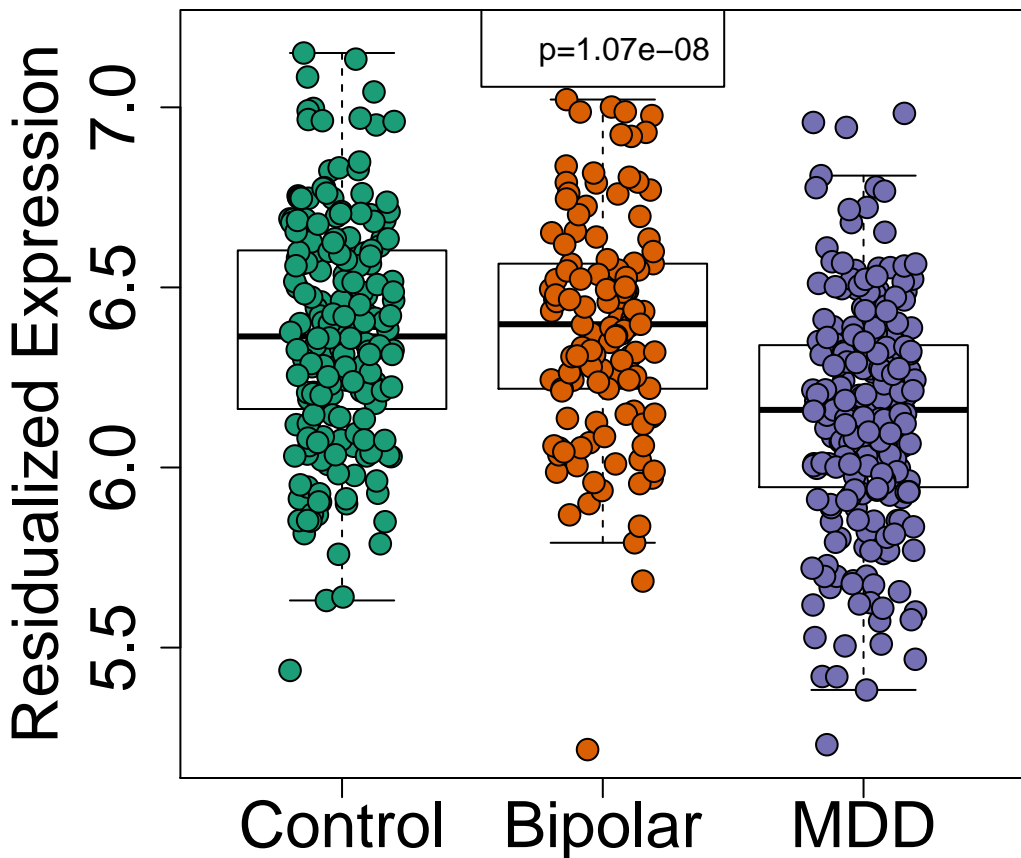
APP, InGen

chr21:26022043–26050999(–)



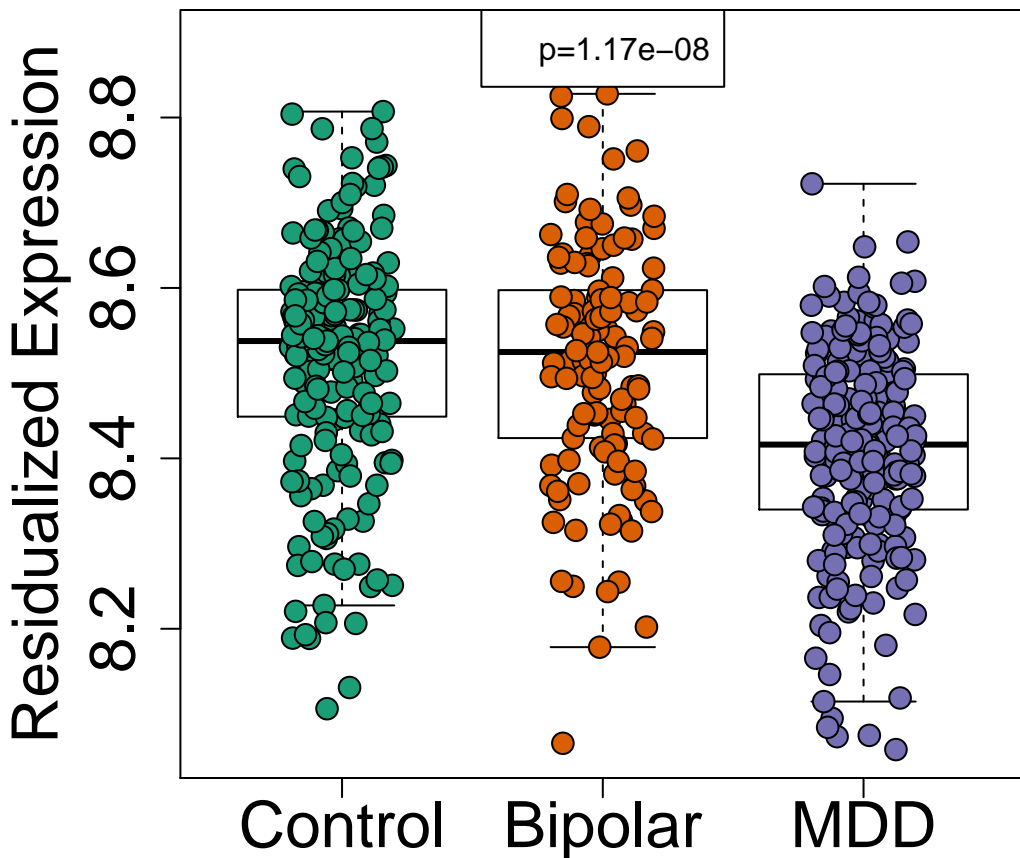
FUS, InGen

chr16:31191111–31191398(+)



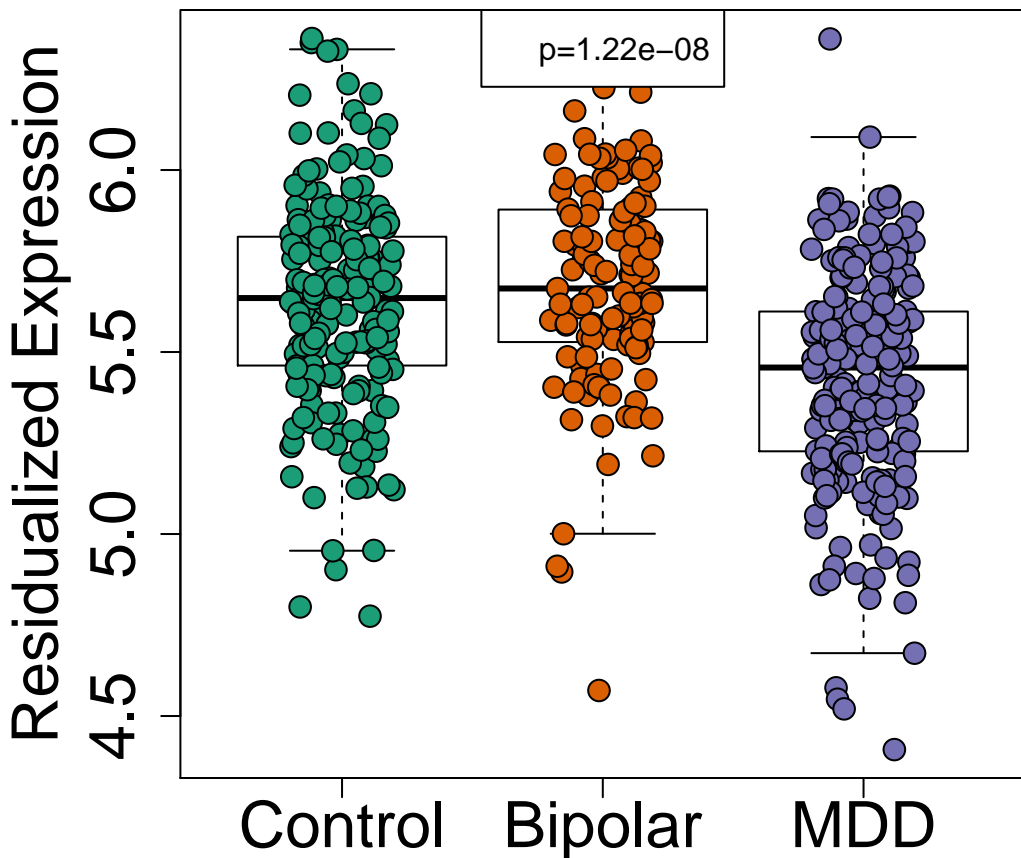
DYNC1H1, InGen

chr14:102005237-102005887(+)



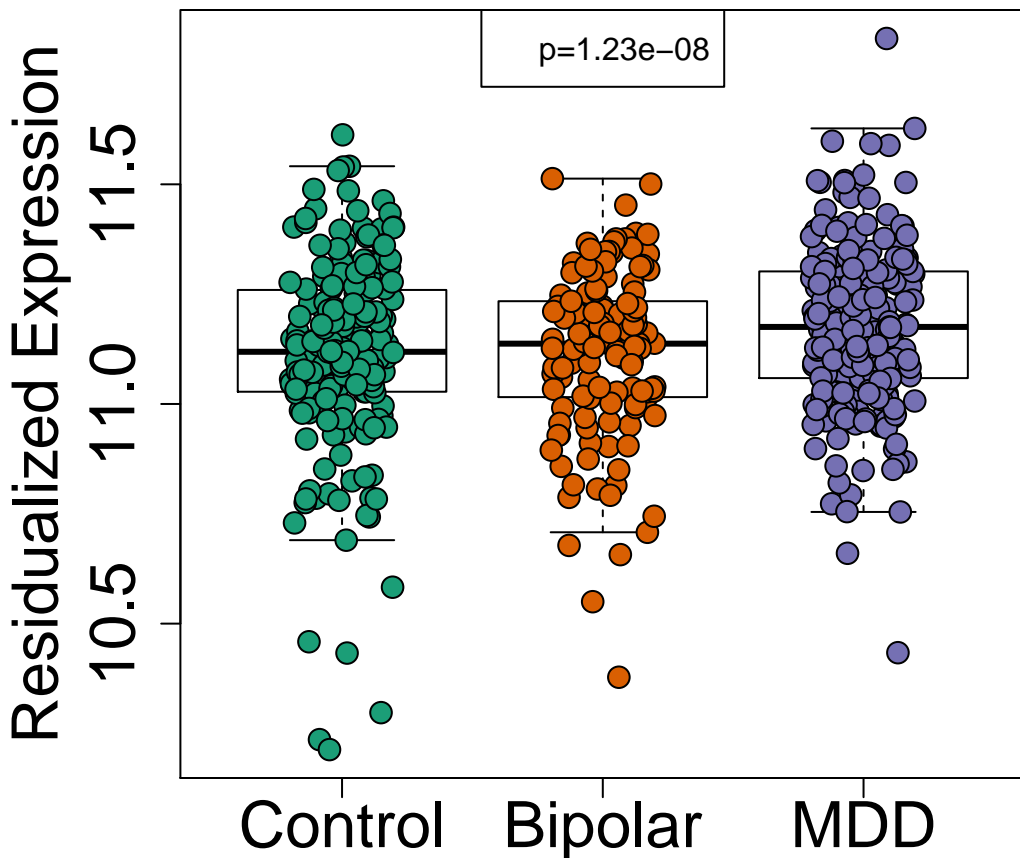
FUS, InGen

chr16:31186837–31188324(+)



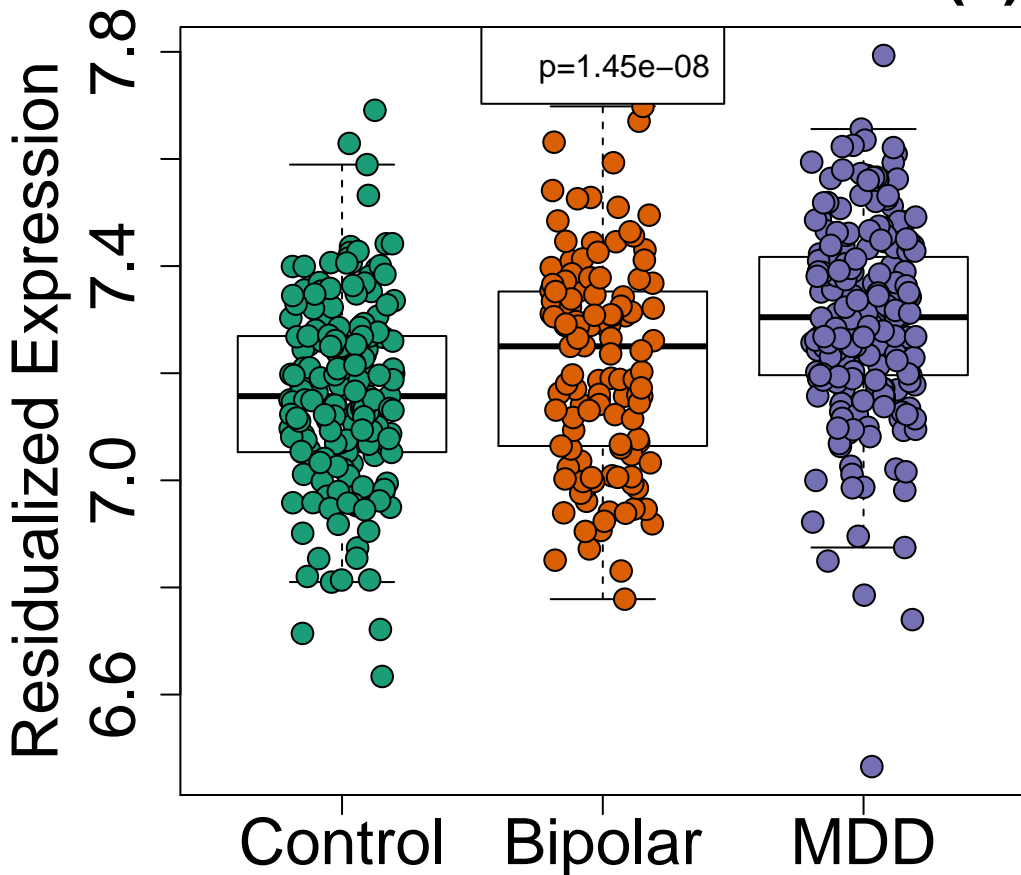
SPARCL1, InGen

chr4:87480521–87482423(–)



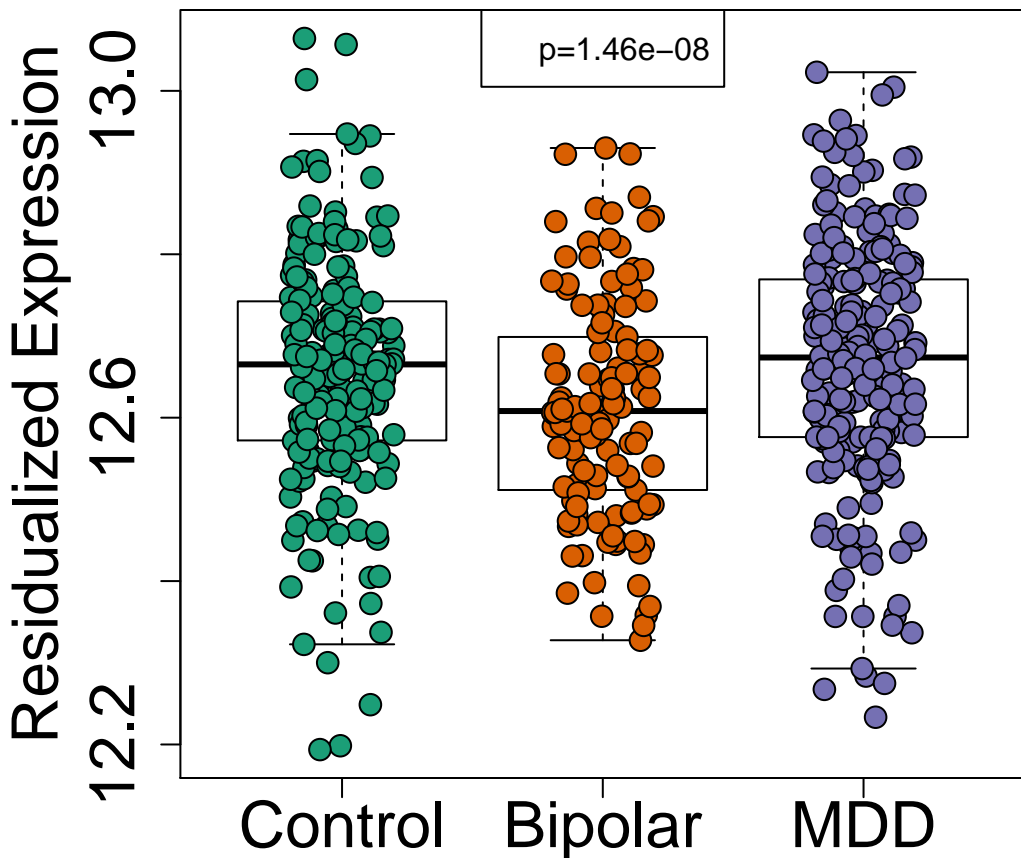
CTNNA2, InGen

chr2:79869936–79874075(+)



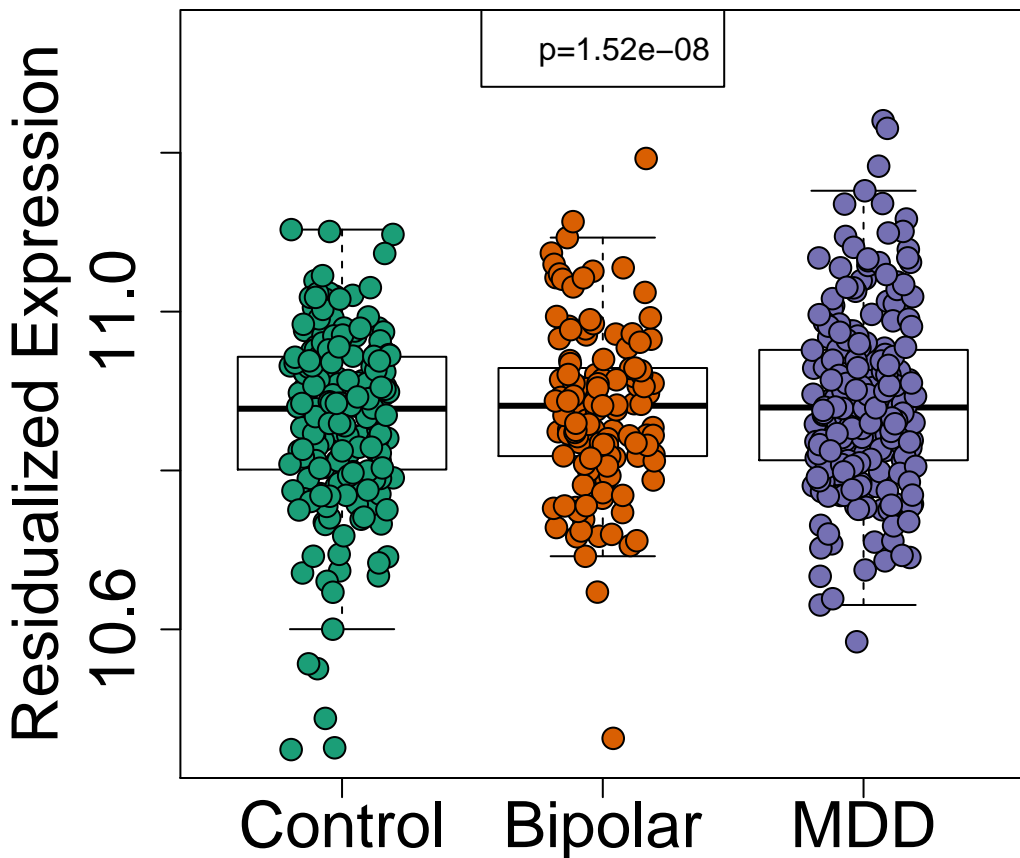
TMSB4X, InGen

chrX:12975169–12976245(+)



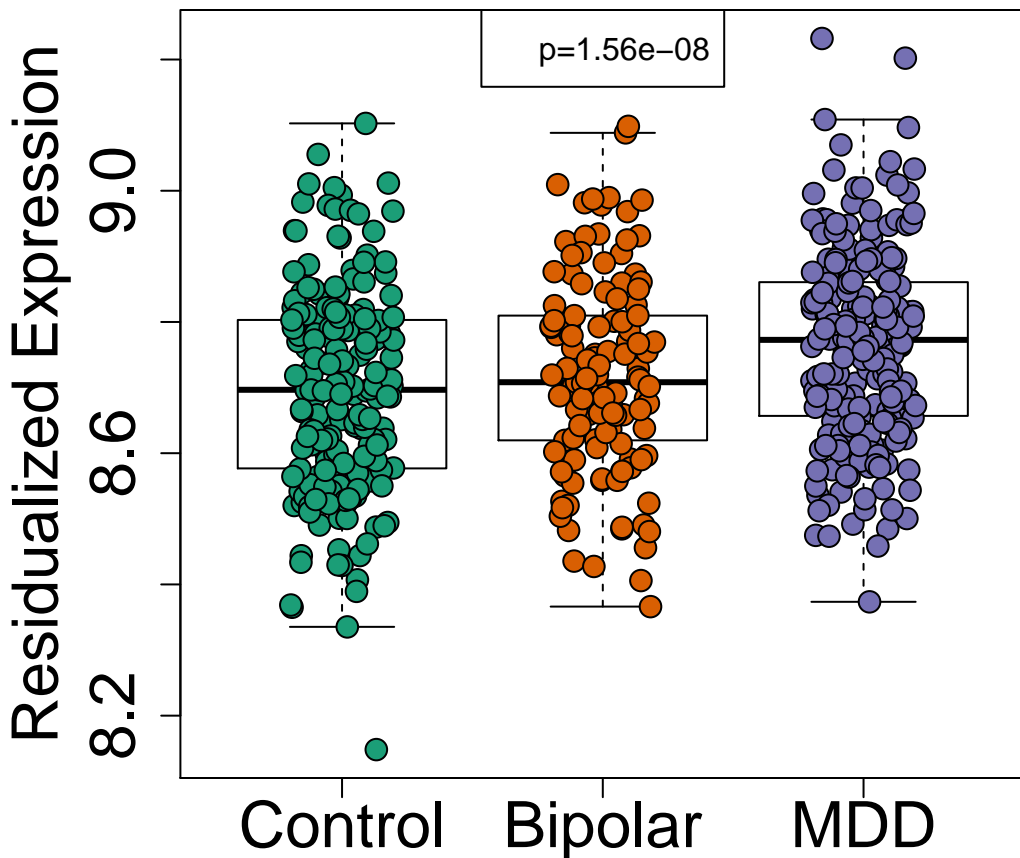
HSP90AB1, InGen

chr6:44249584–44249674(+)



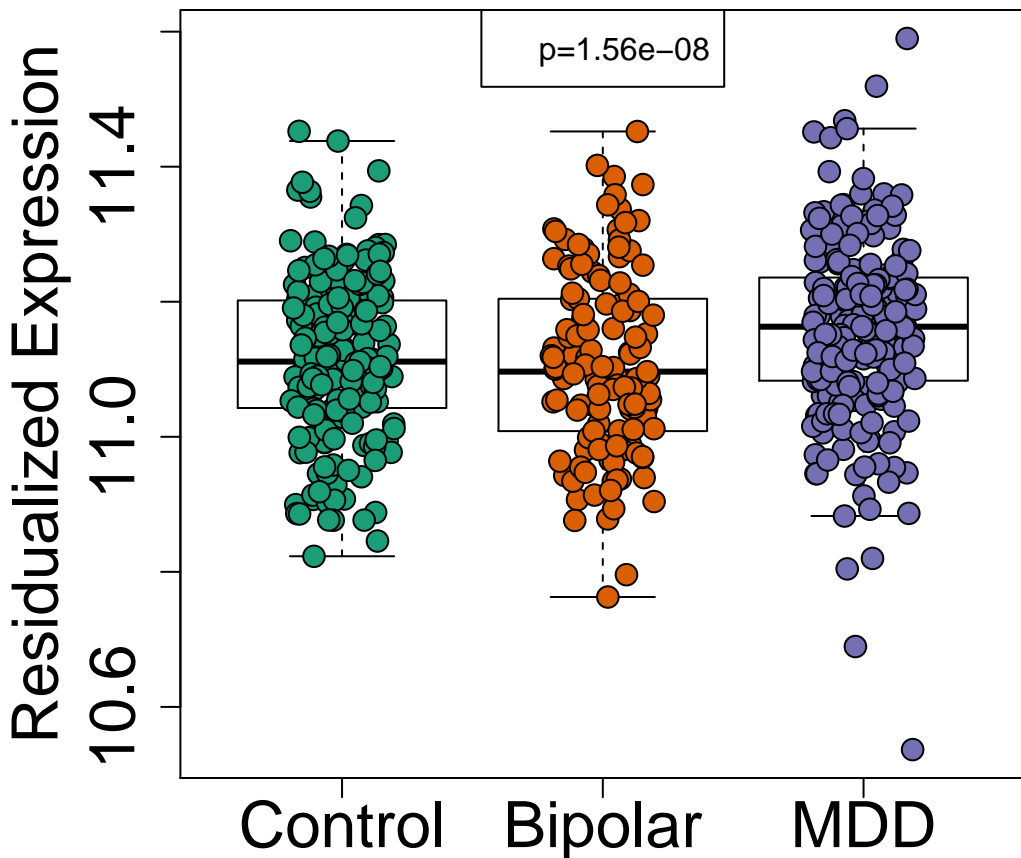
AZIN1, InGen

chr8:102828679–102829271(-)



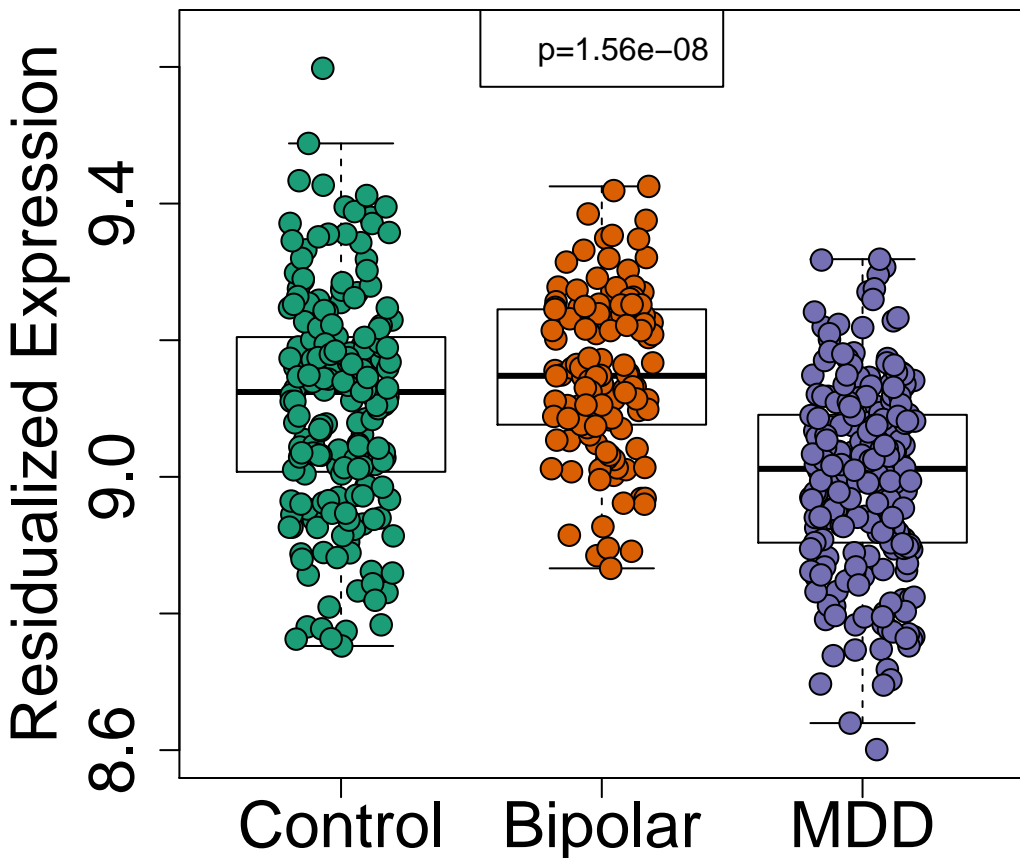
RPL13A, InGen

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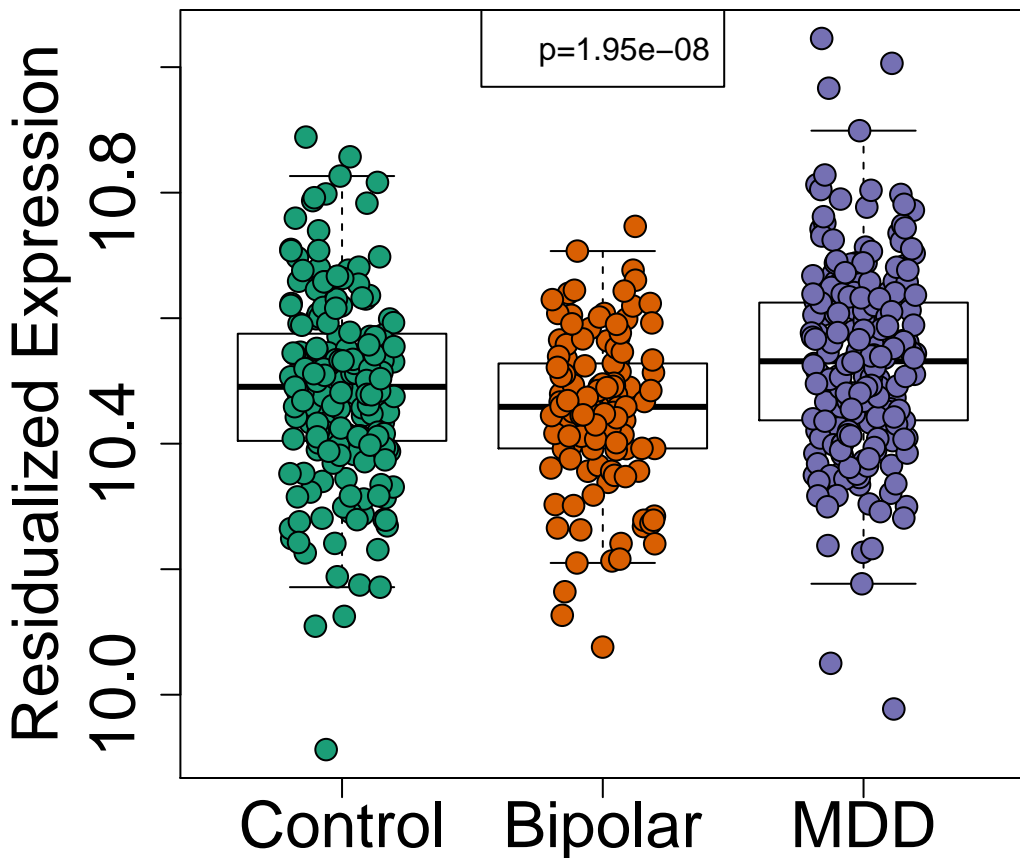
RPS21, InGen

chr20:62387675–62387842(+)



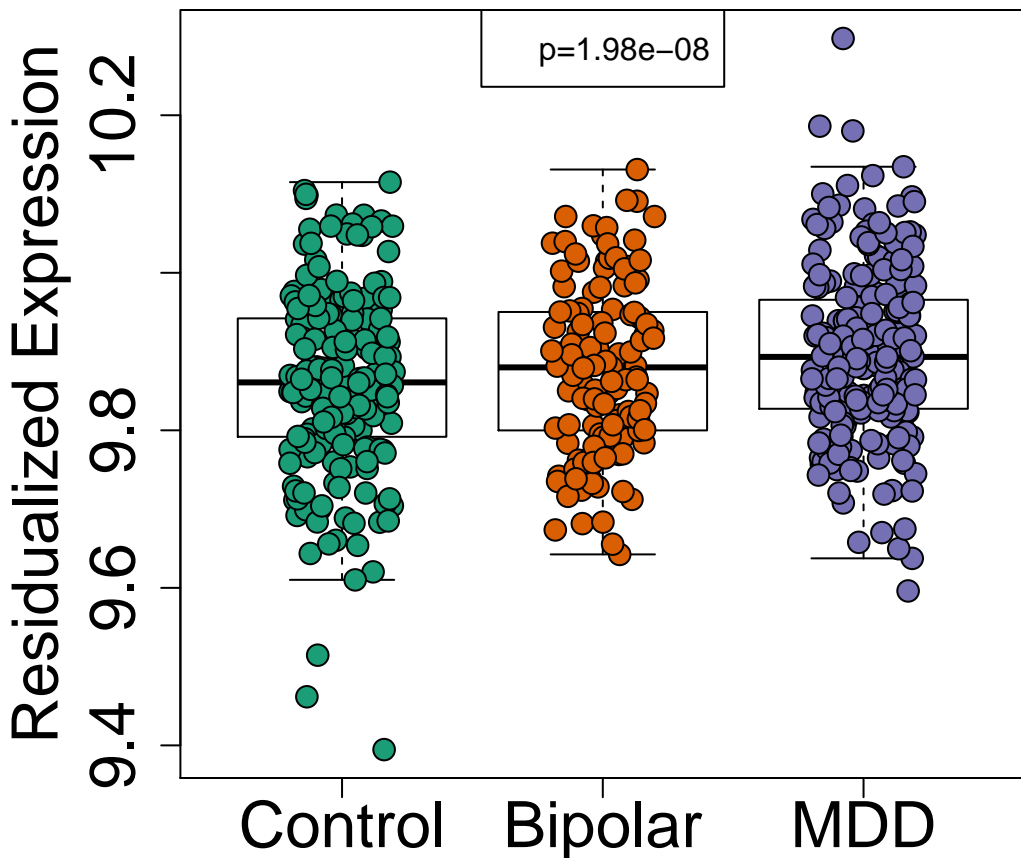
MDH1, InGen

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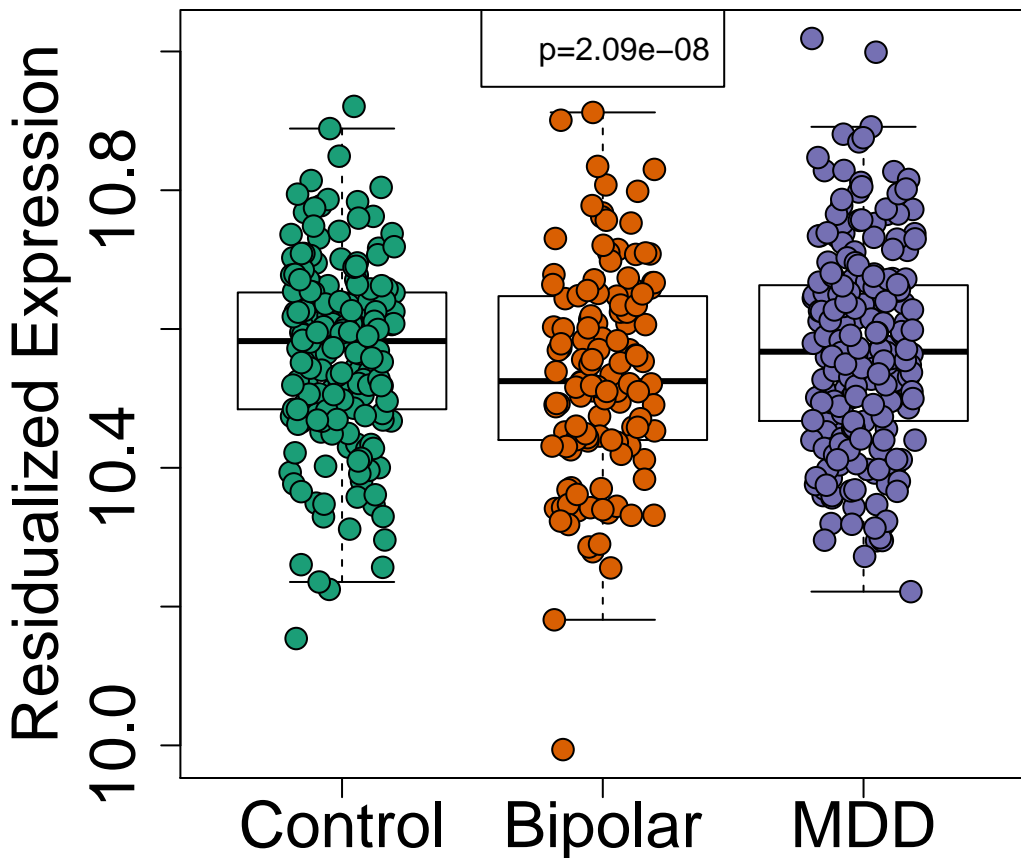
NACA, InGen

chr12:56712909–56713061(–)



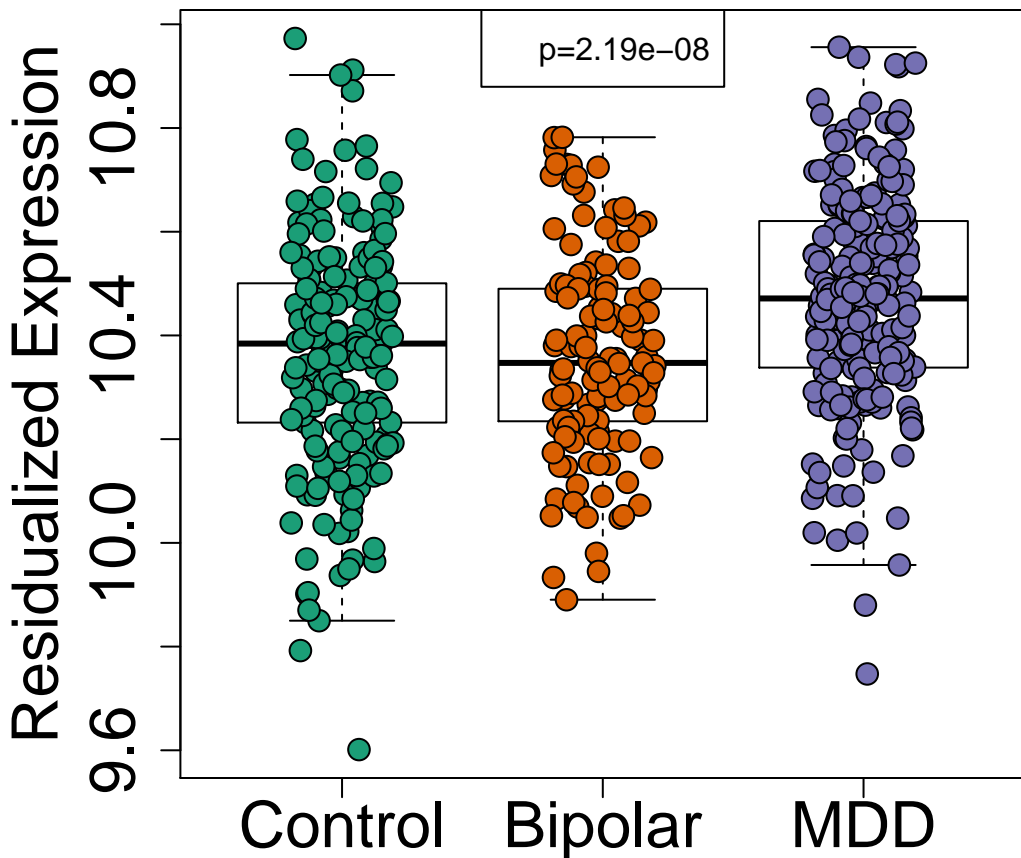
EIF4A2, InGen

chr3:186787265–186787494(+)



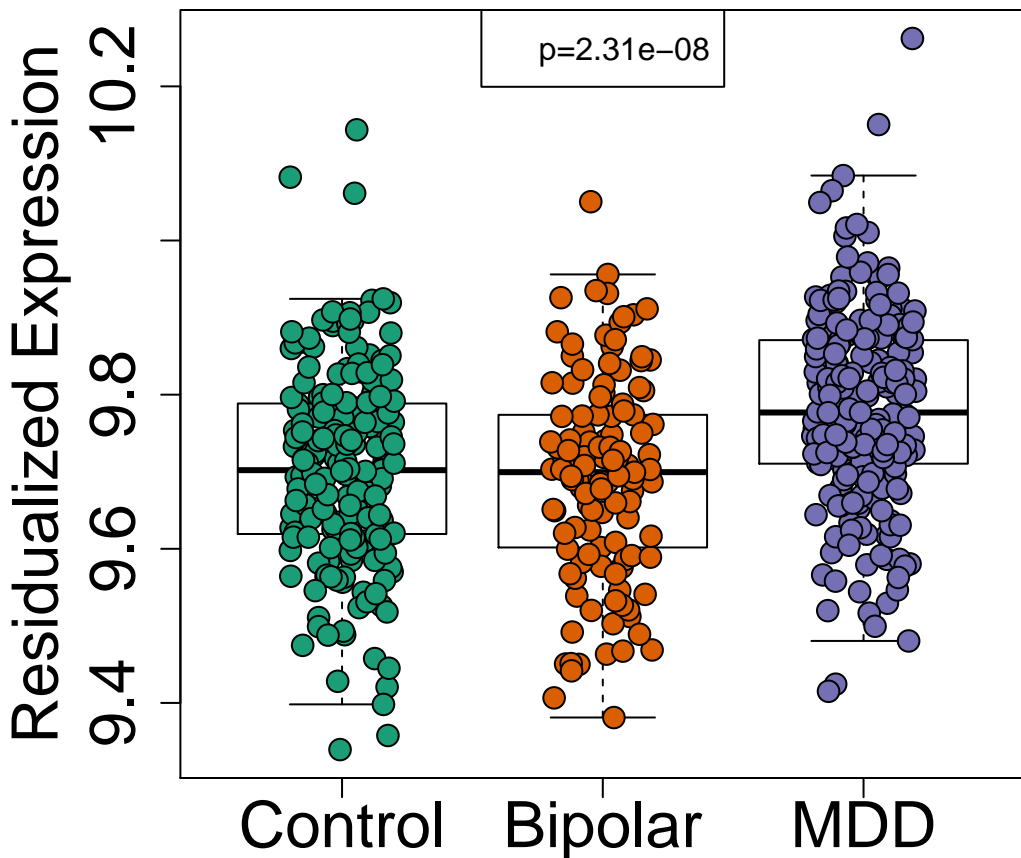
UCHL1, InGen

chr4:41257127-41257608(+)



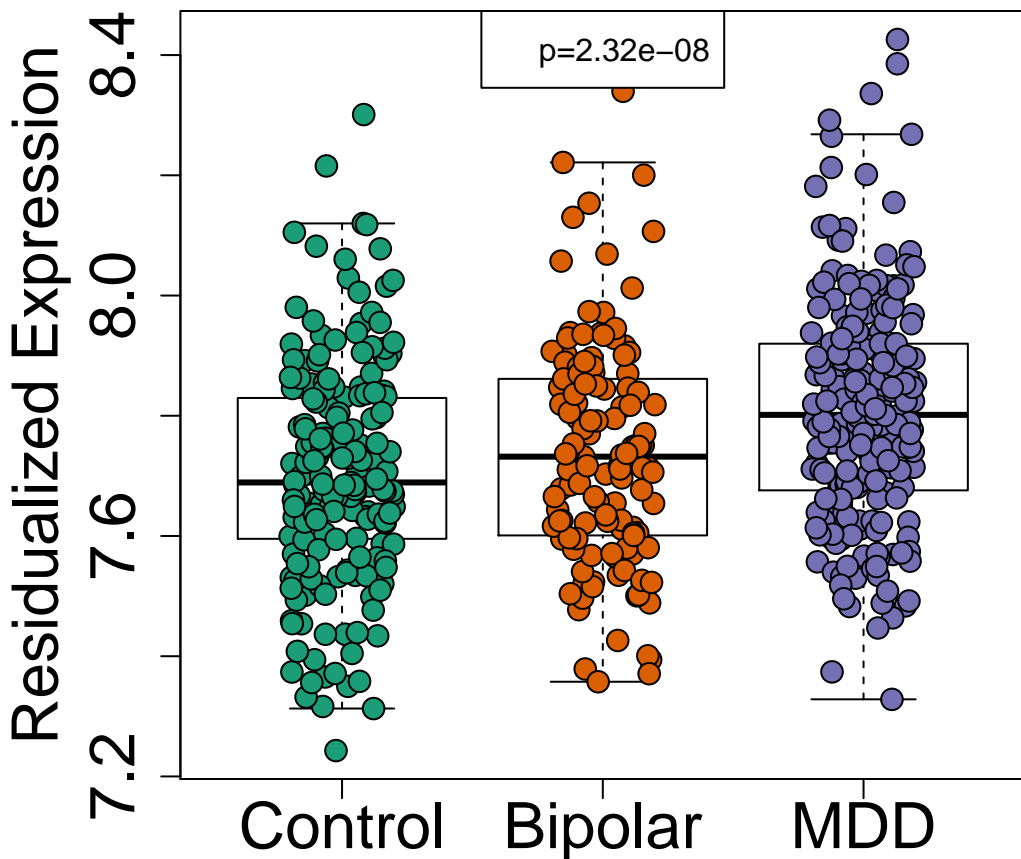
ENO1, InGen

chr1:8866502–8867116(–)



XPO1, InGen

chr2:61492749–61492914(–)



MEG3, InGen

chr14:100831535–100836166(+)

