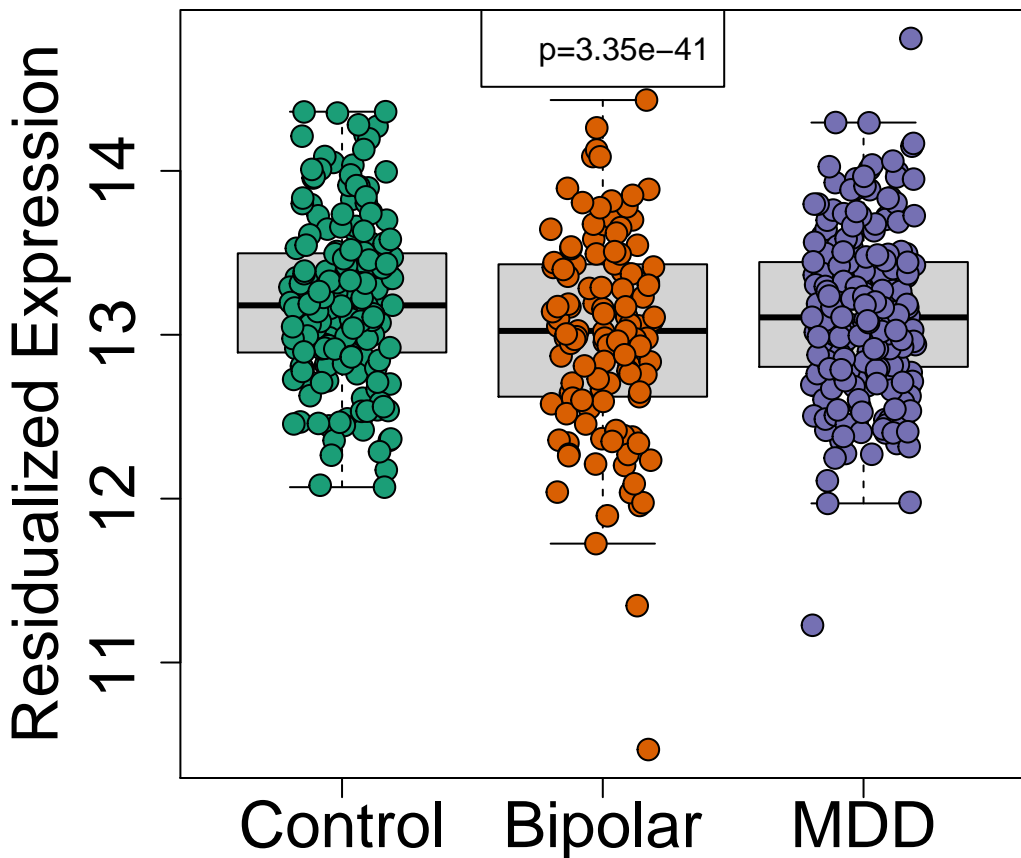


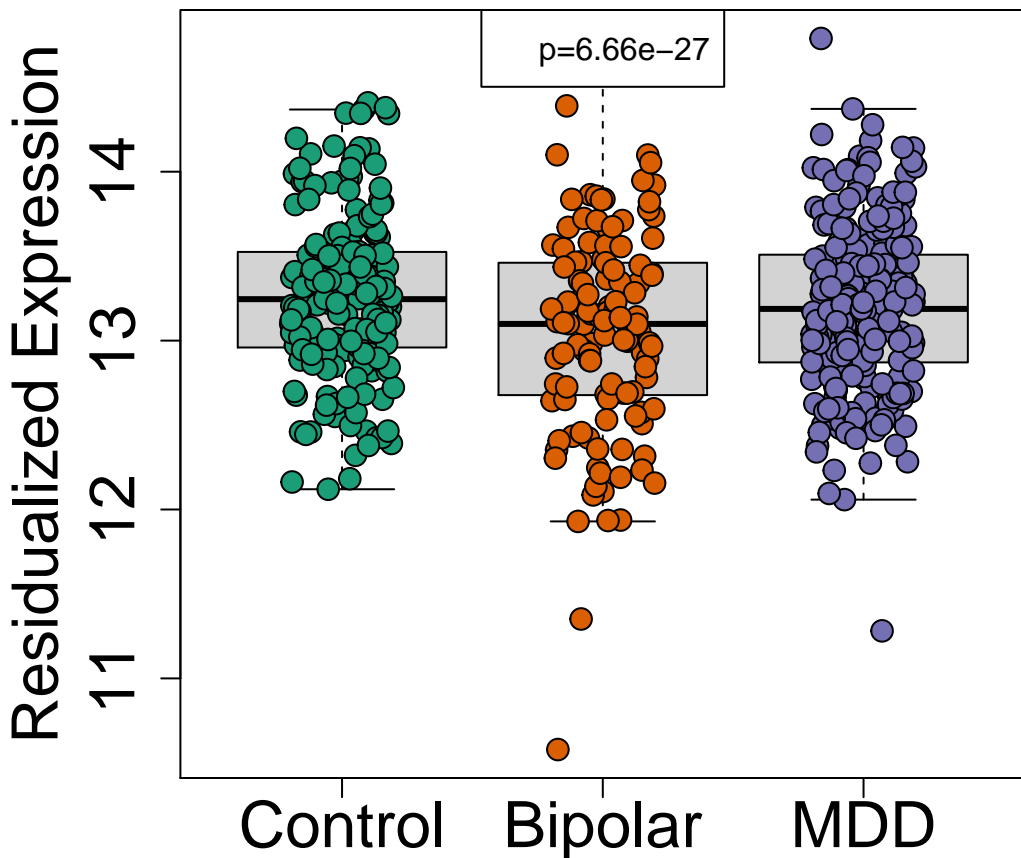
GFAP, InGen

chr17:44913431-44913727(-)



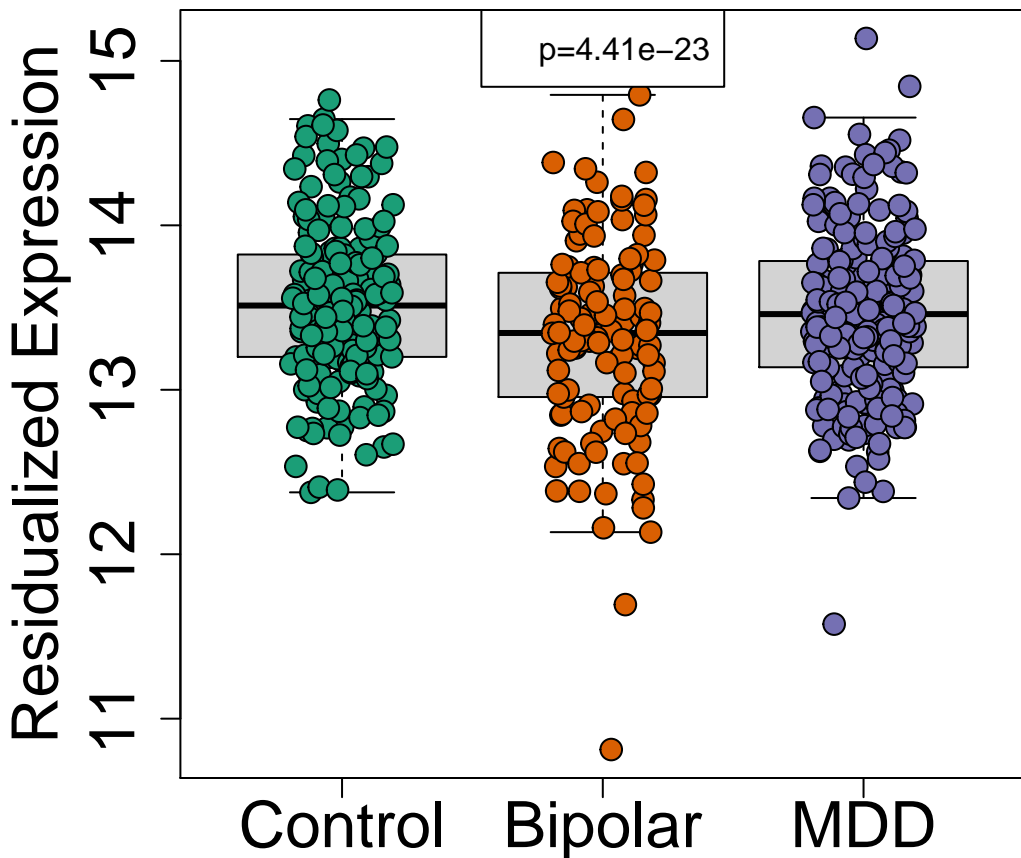
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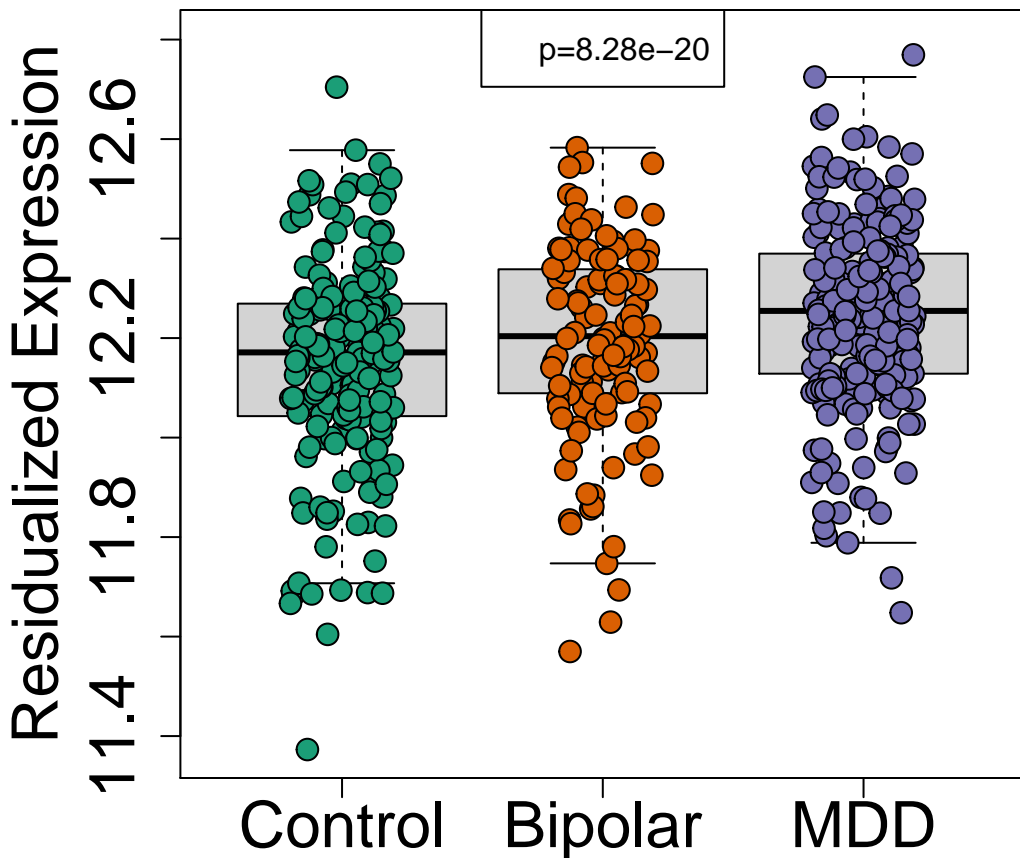
GFAP, InGen

chr17:44910659–44911235(-)



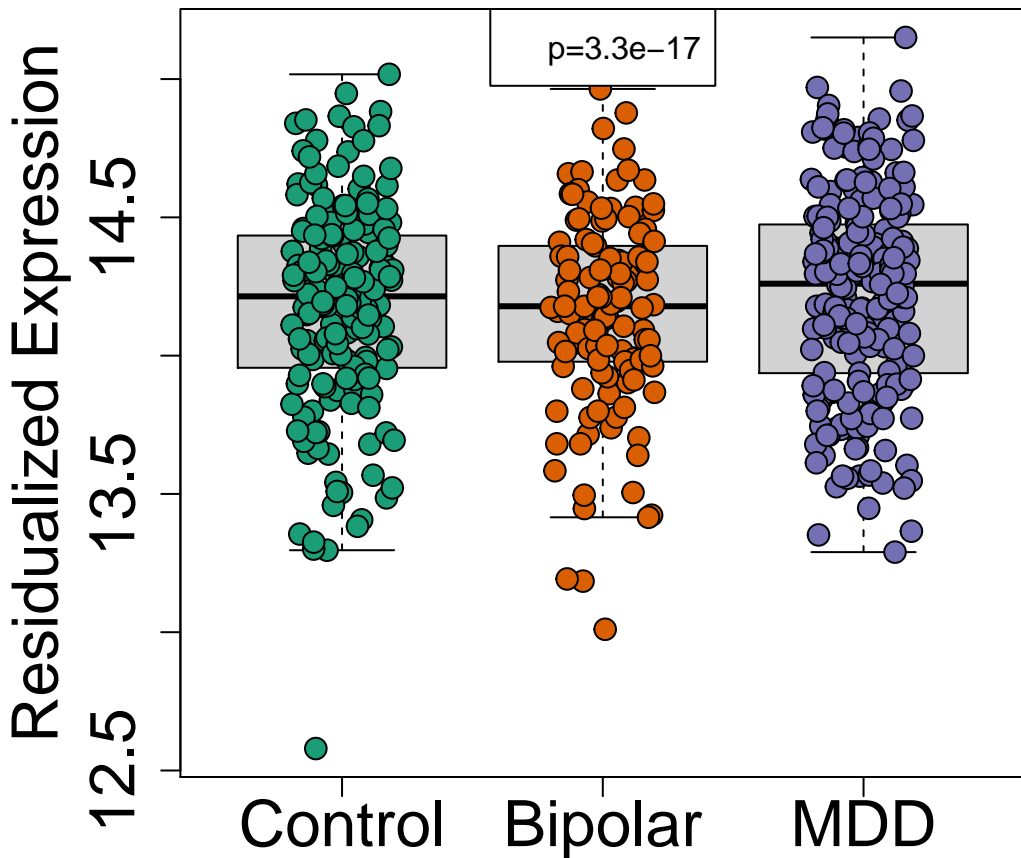
TMSB10, InGen

chr2:84905719–84906002(+)



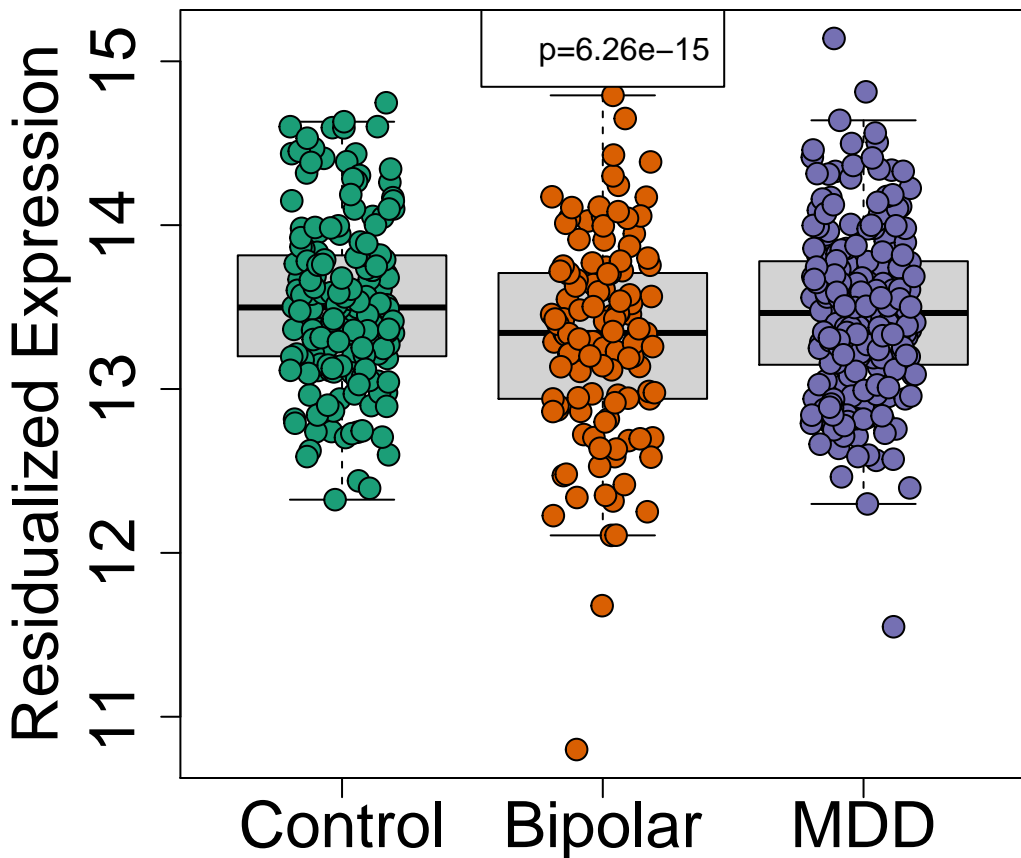
MBP, InGen

chr18:76988528–76988876(–)



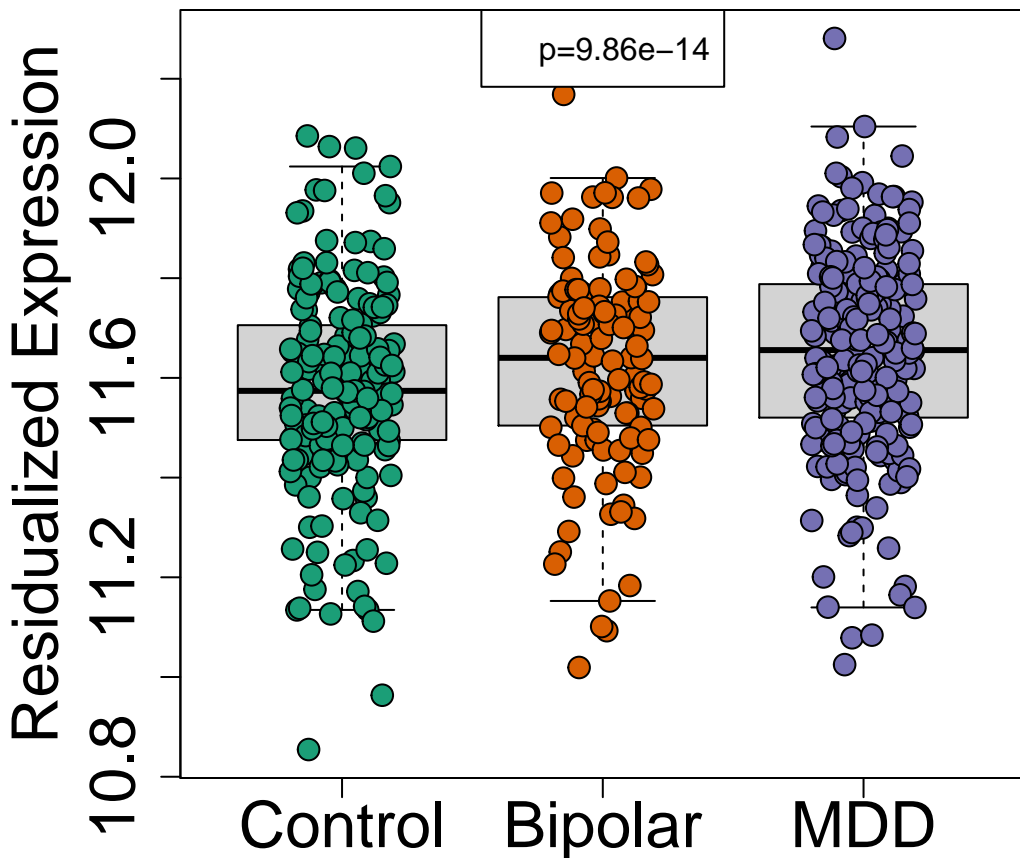
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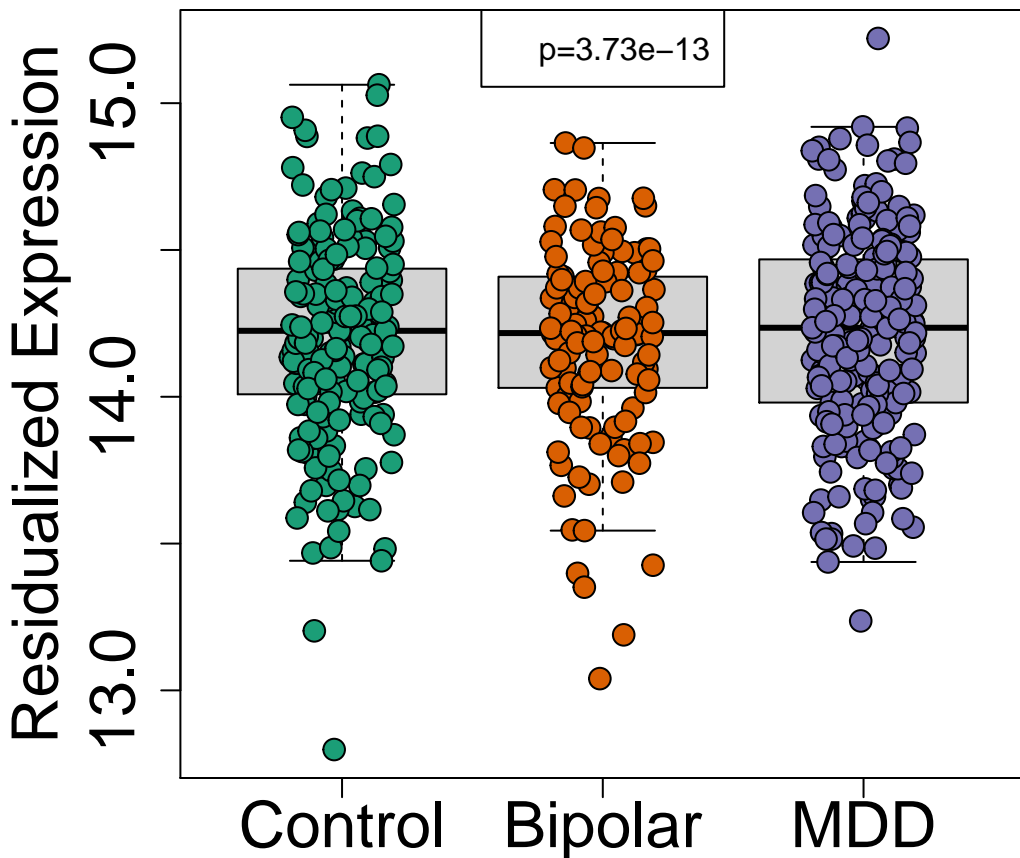
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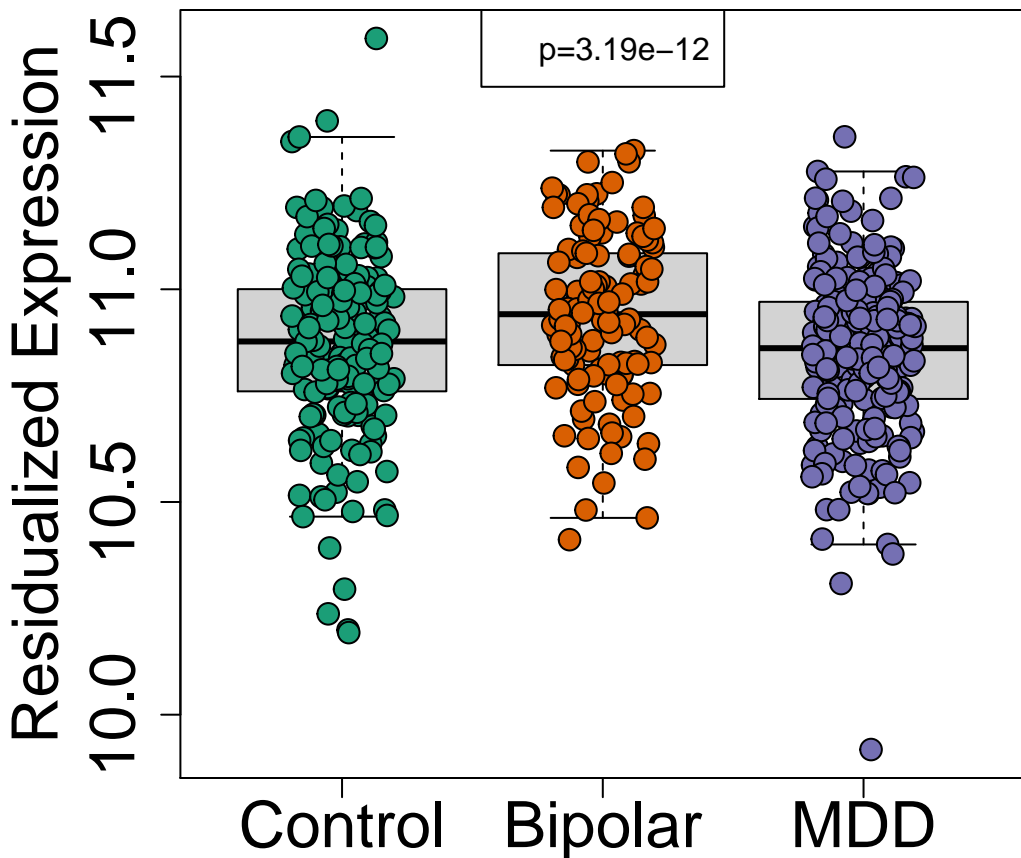
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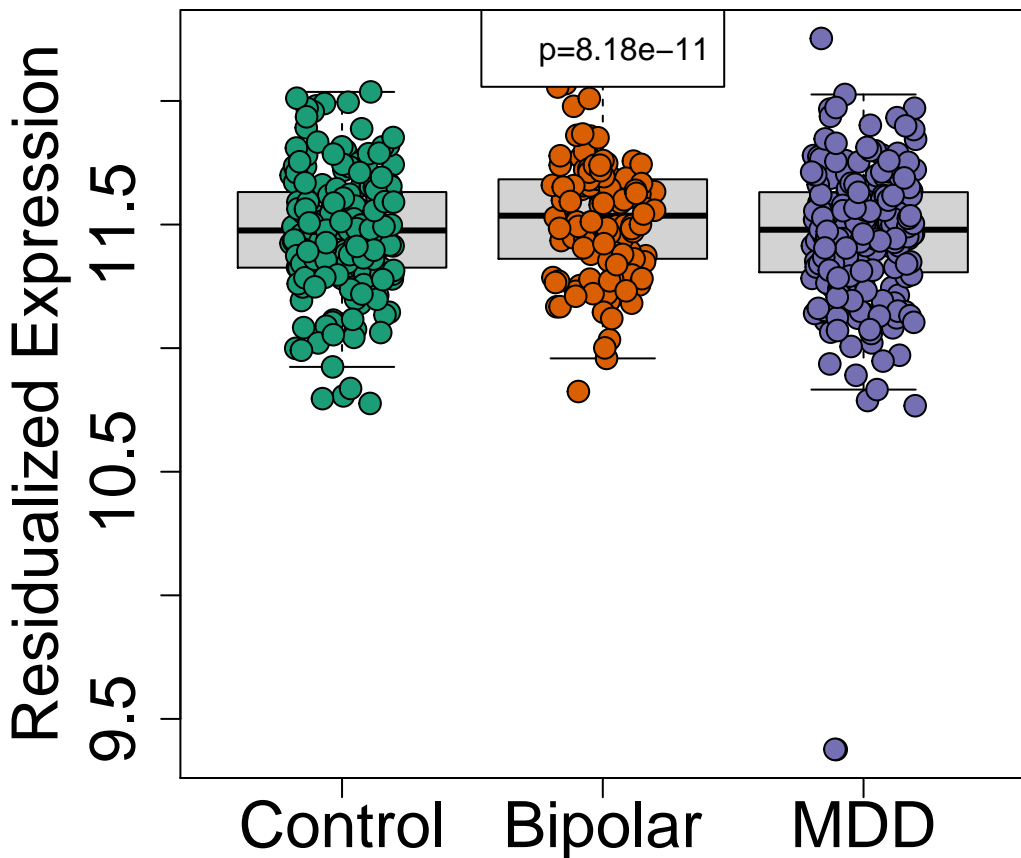
TUBA1B, InGen

chr12:49129723–49131297(-)



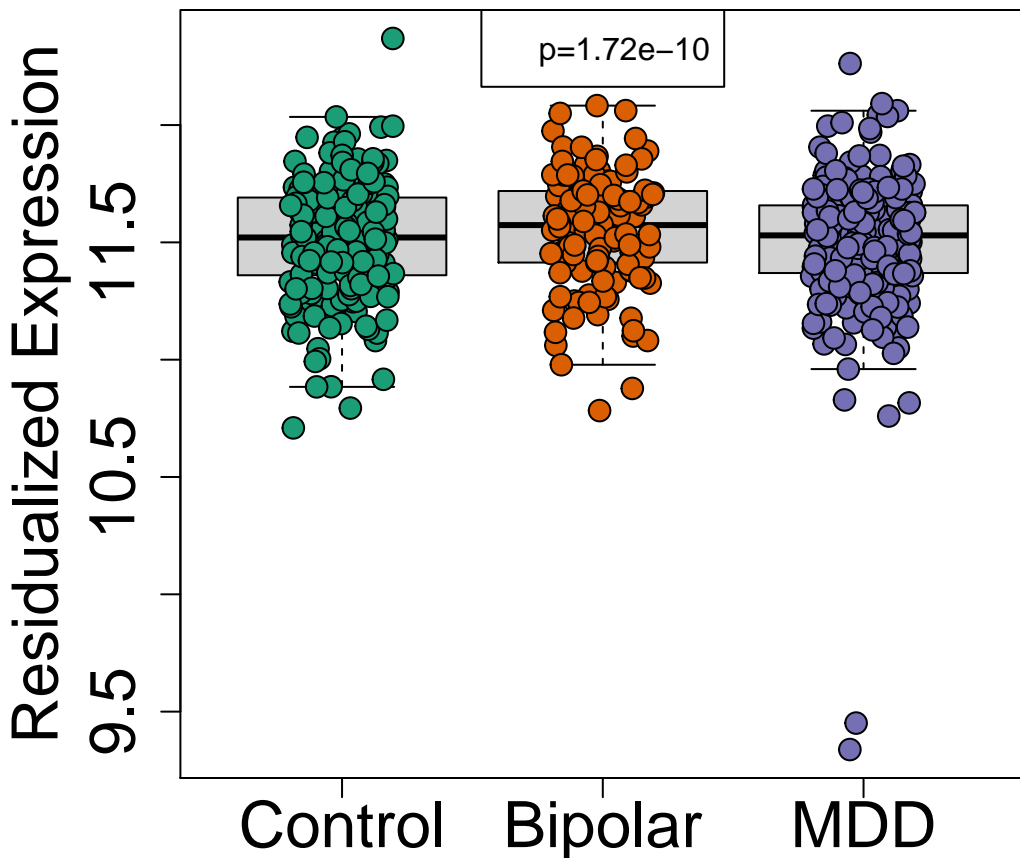
SPARCL1, InGen

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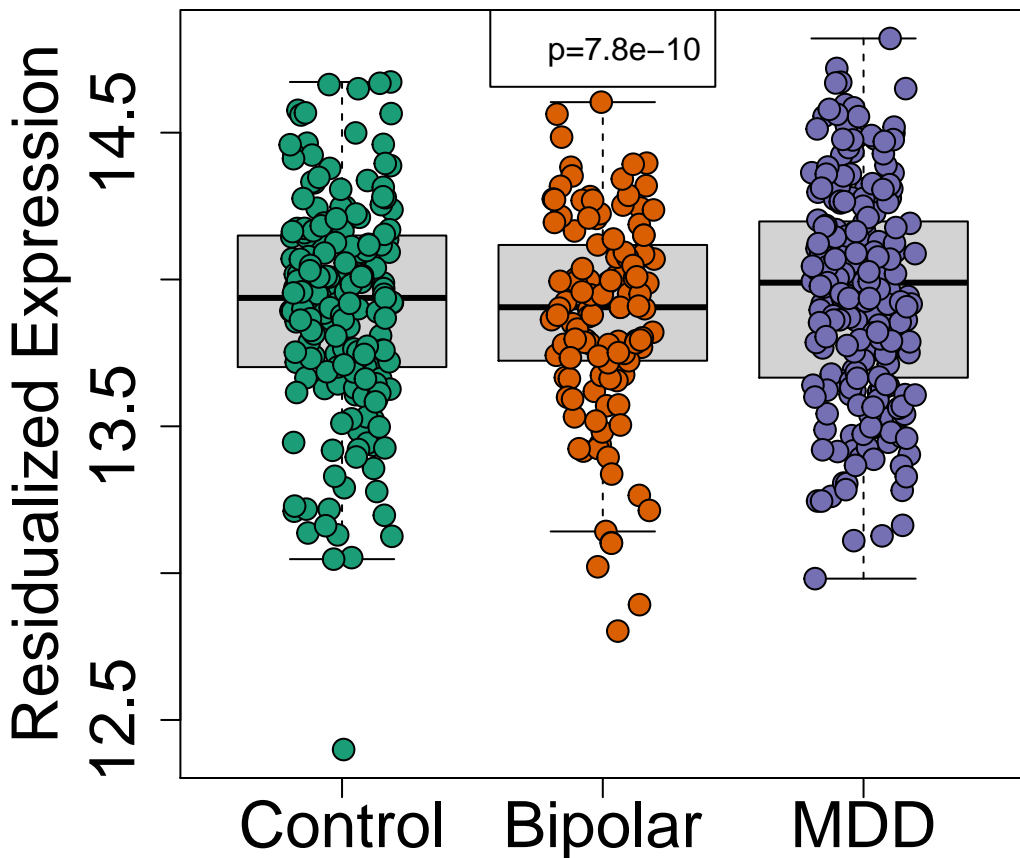
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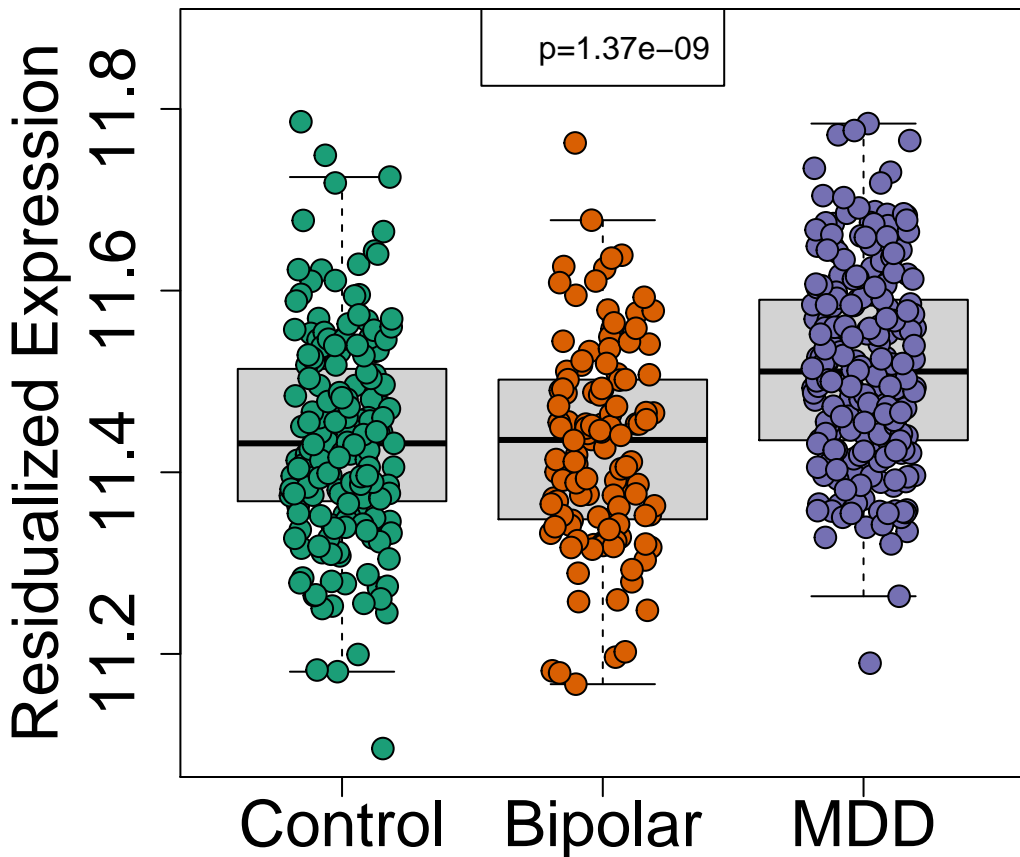
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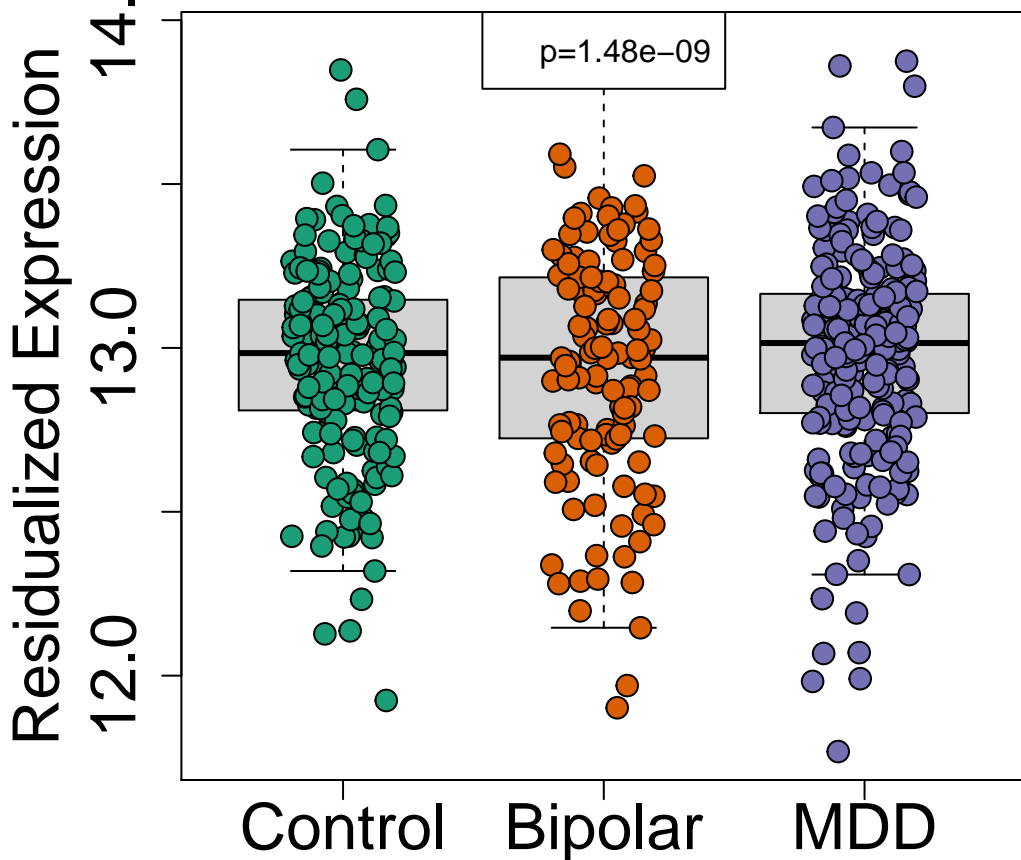
RPL31, InGen

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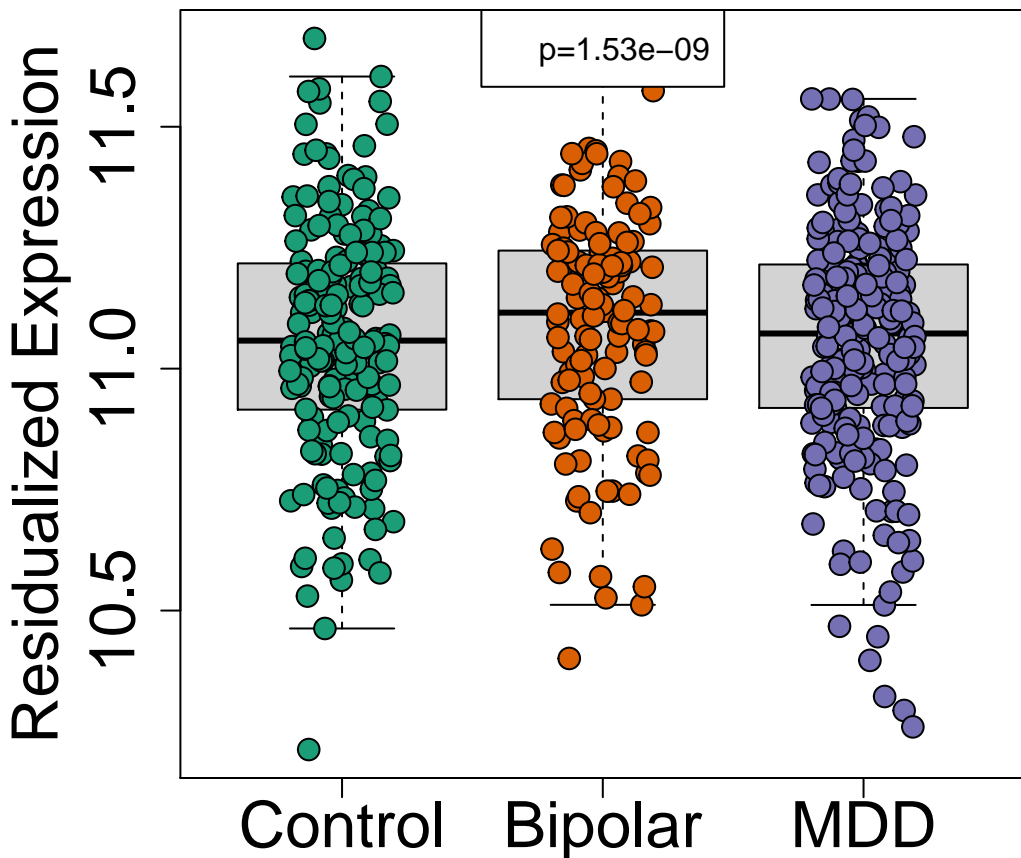
PLP1, InGen

chrX:103786727-103787797(+)



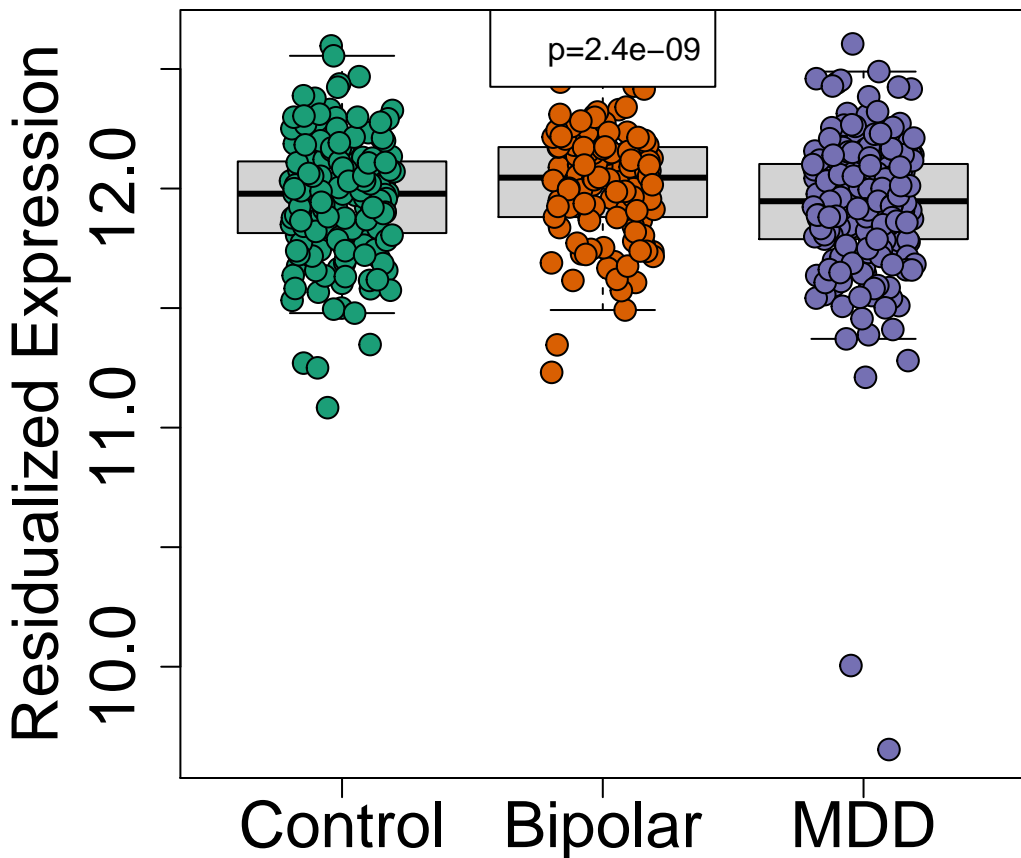
SLC1A2, InGen

chr11:35306243–35312197(-)



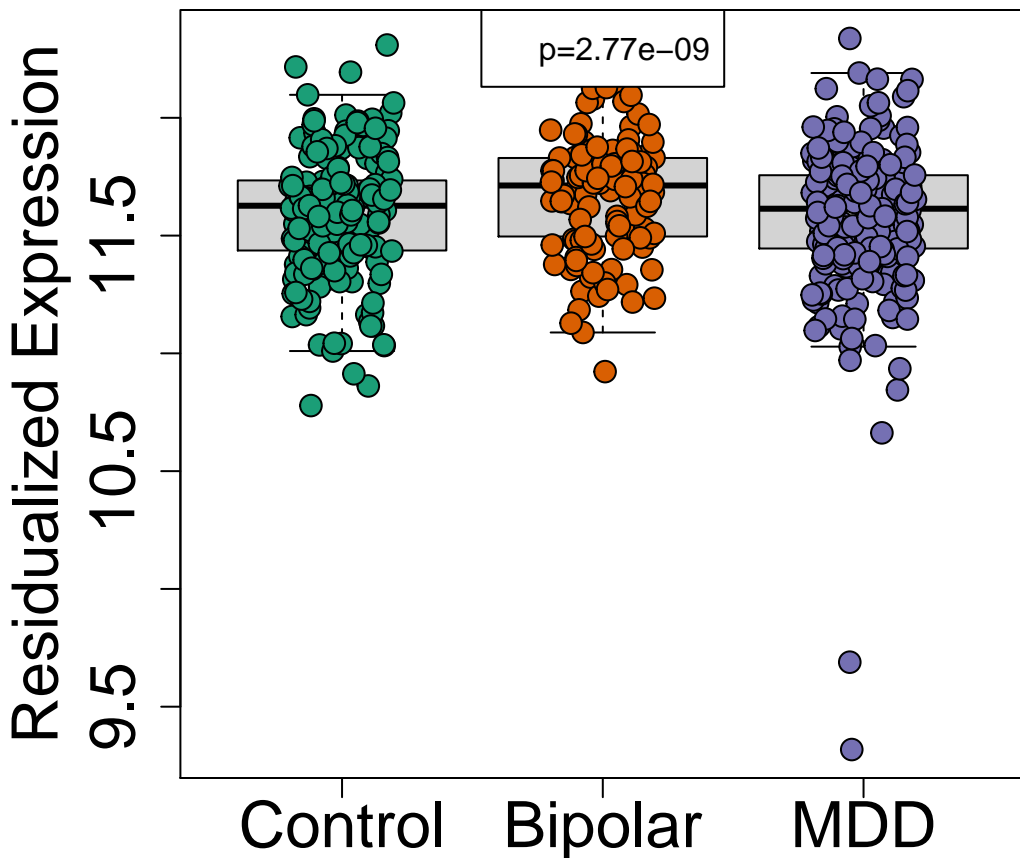
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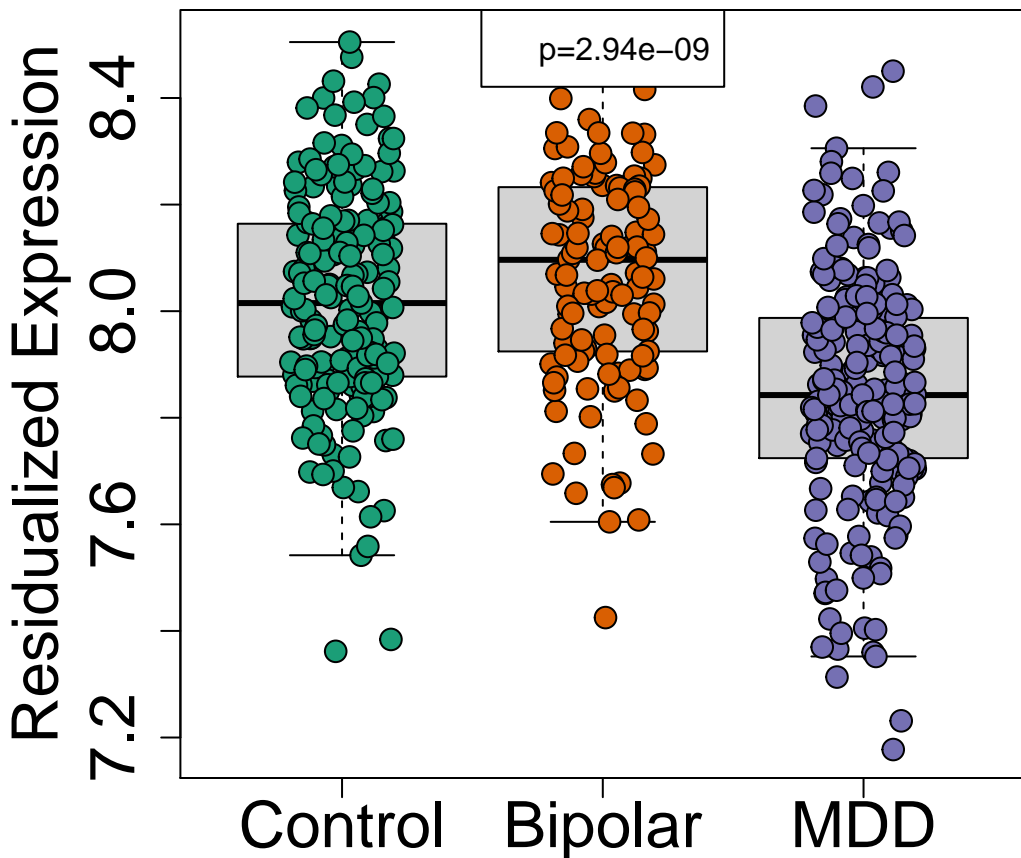
SPARCL1, InGen

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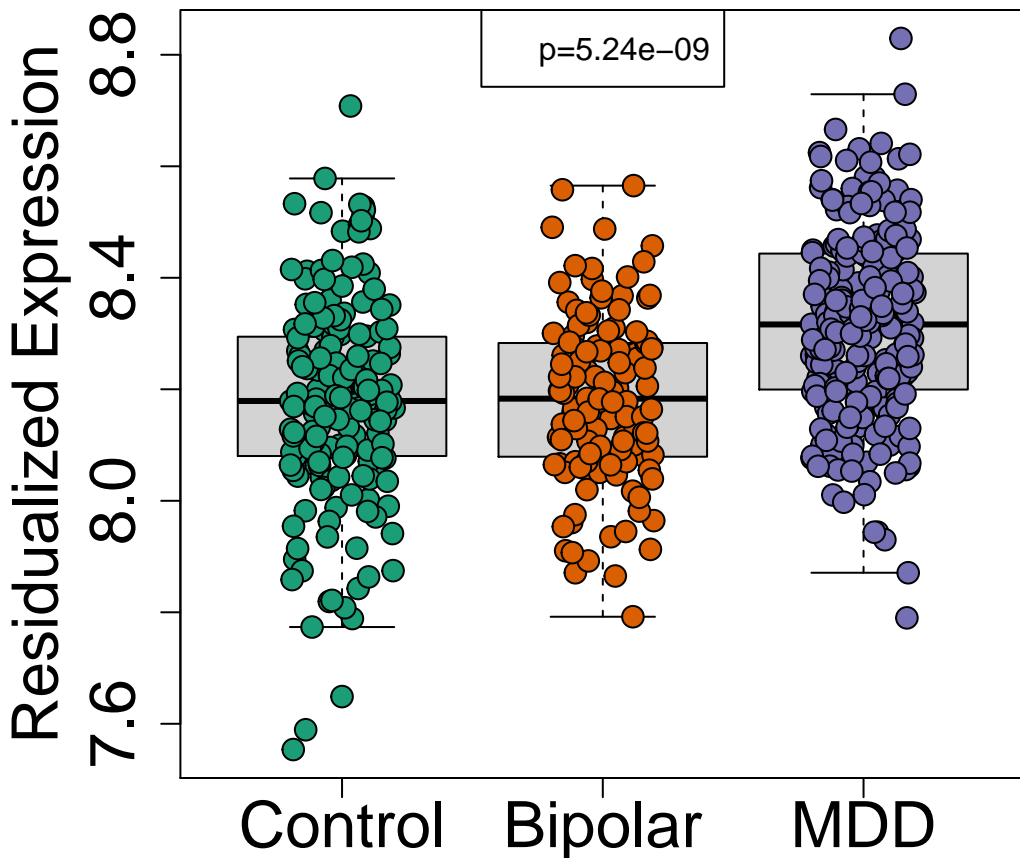
FUS, InGen

chr16:31184003–31184208(+)



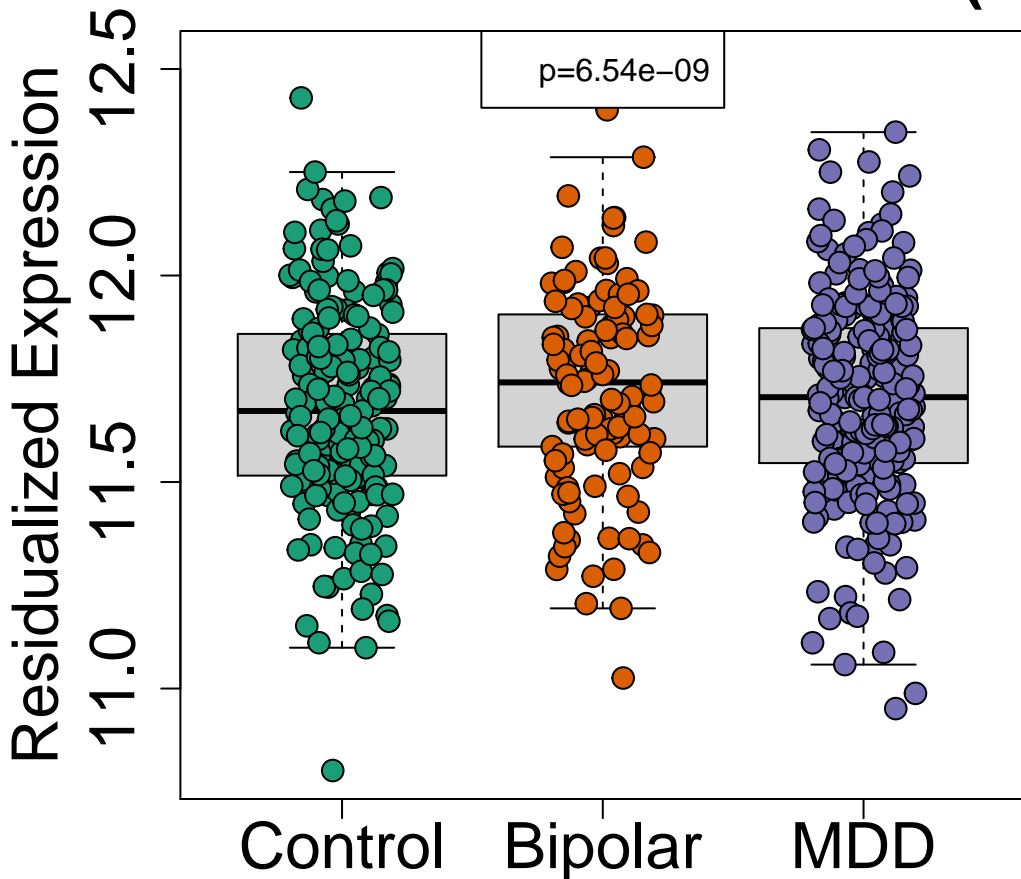
OAZ1, InGen

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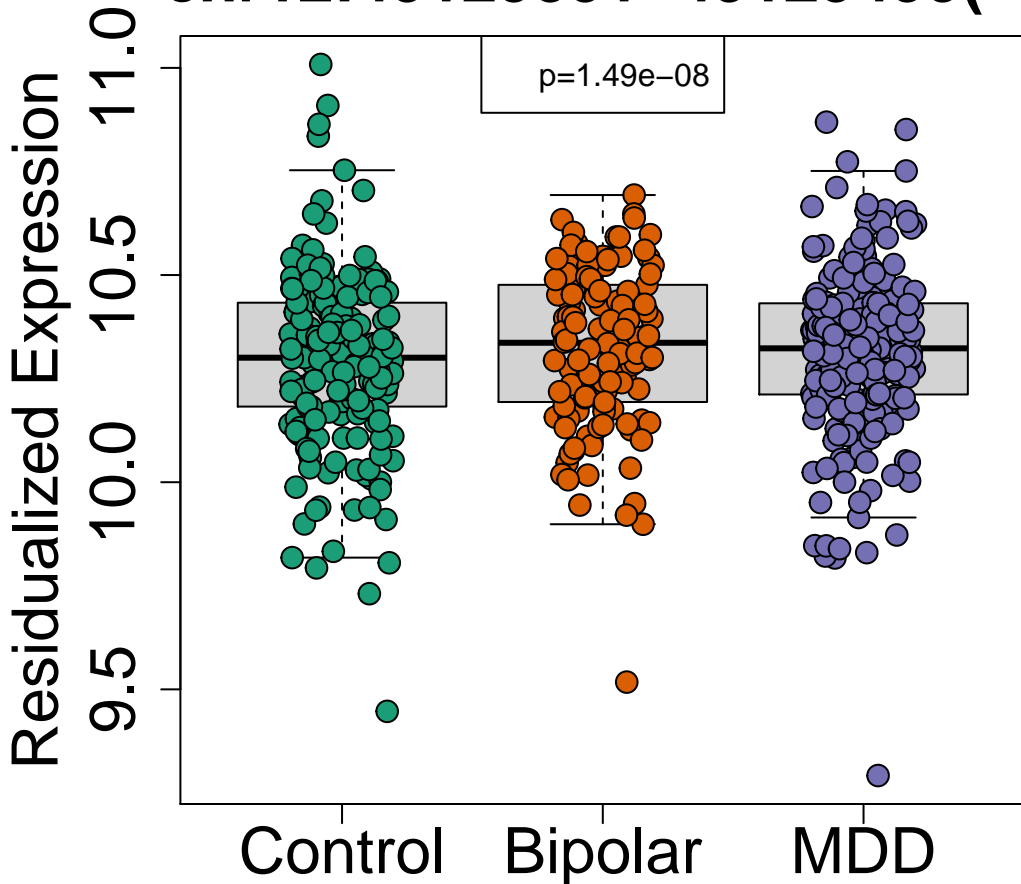
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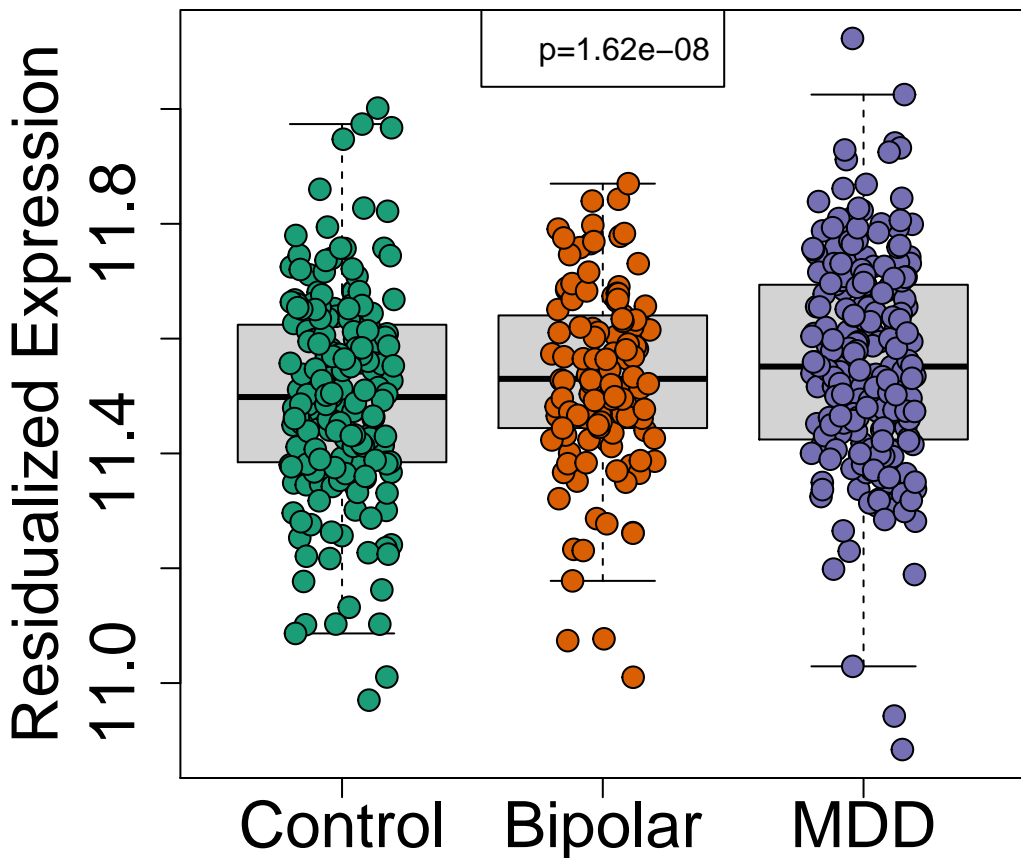
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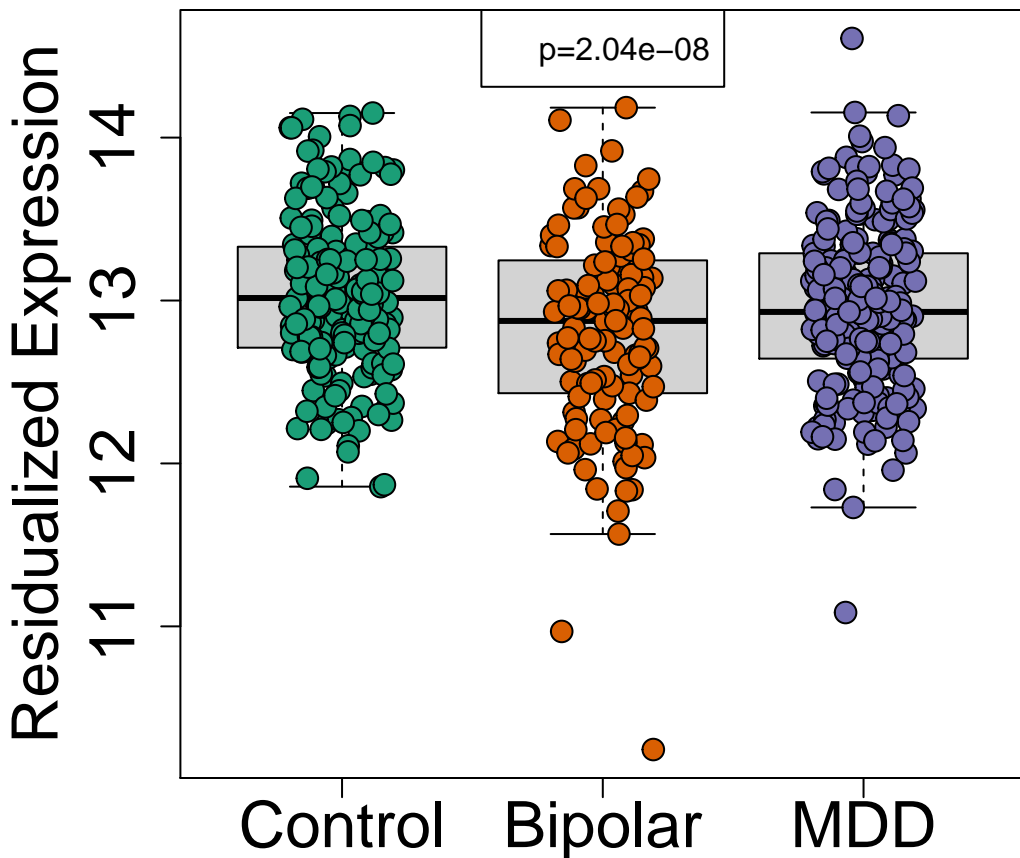
CALM1, InGen

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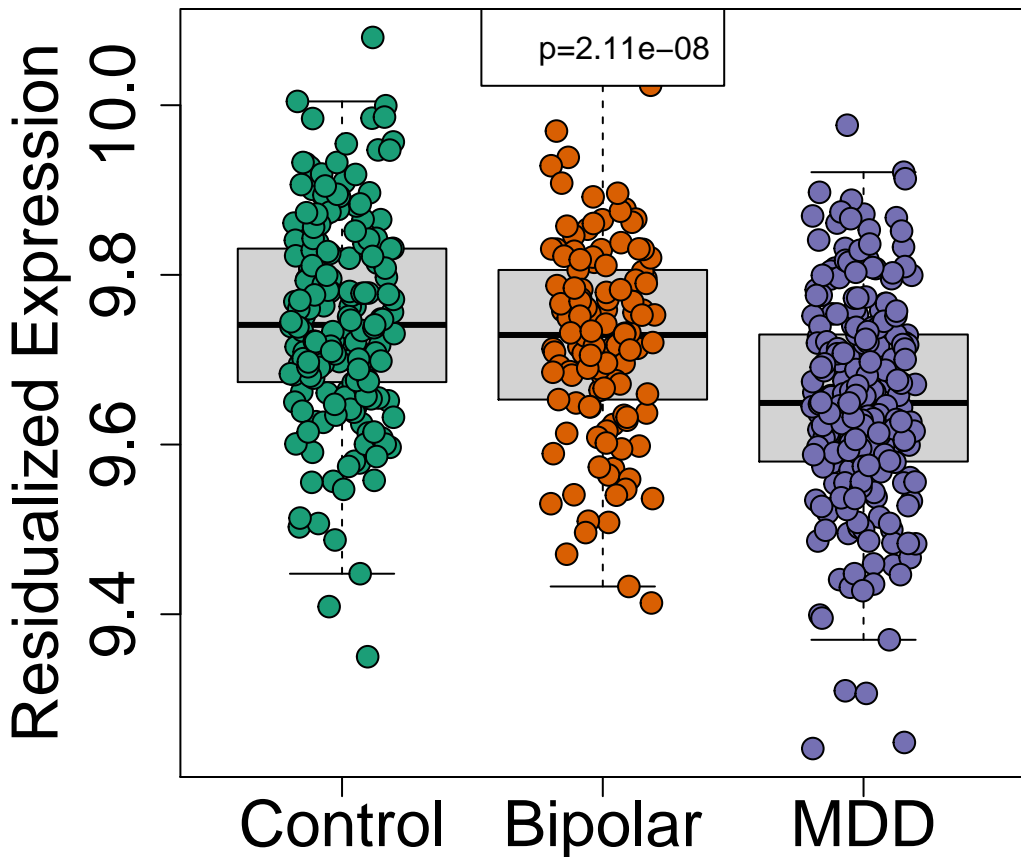
GFAP, InGen

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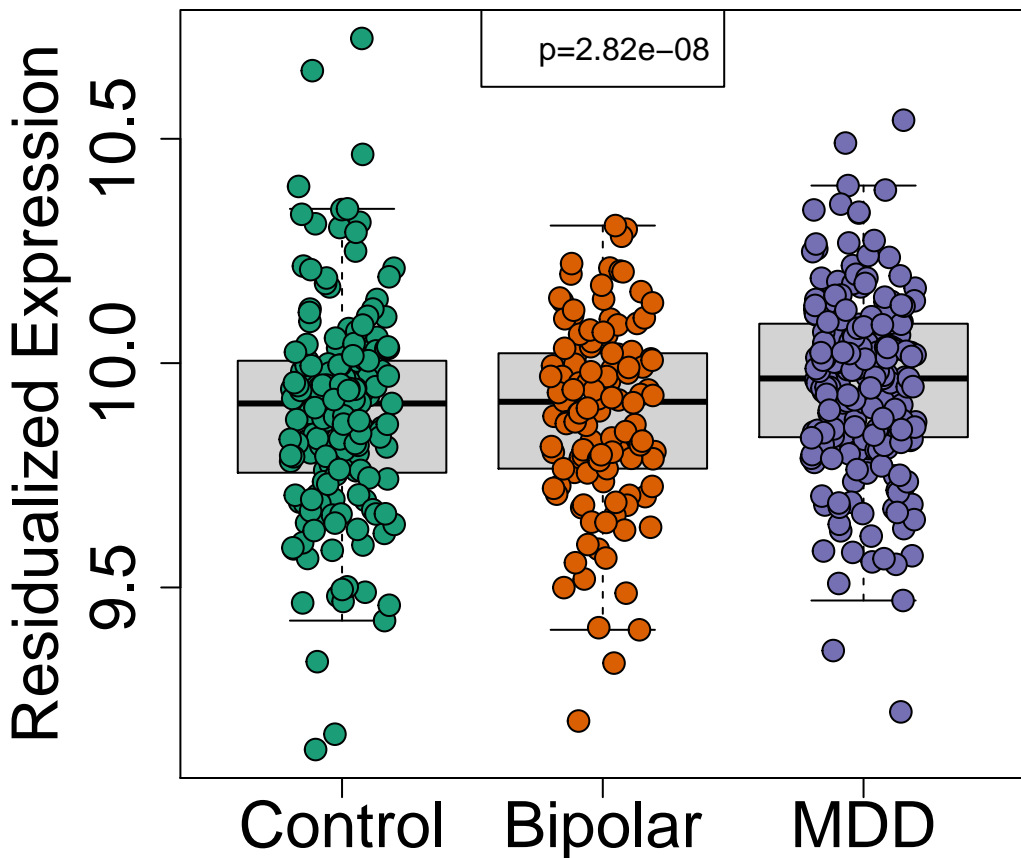
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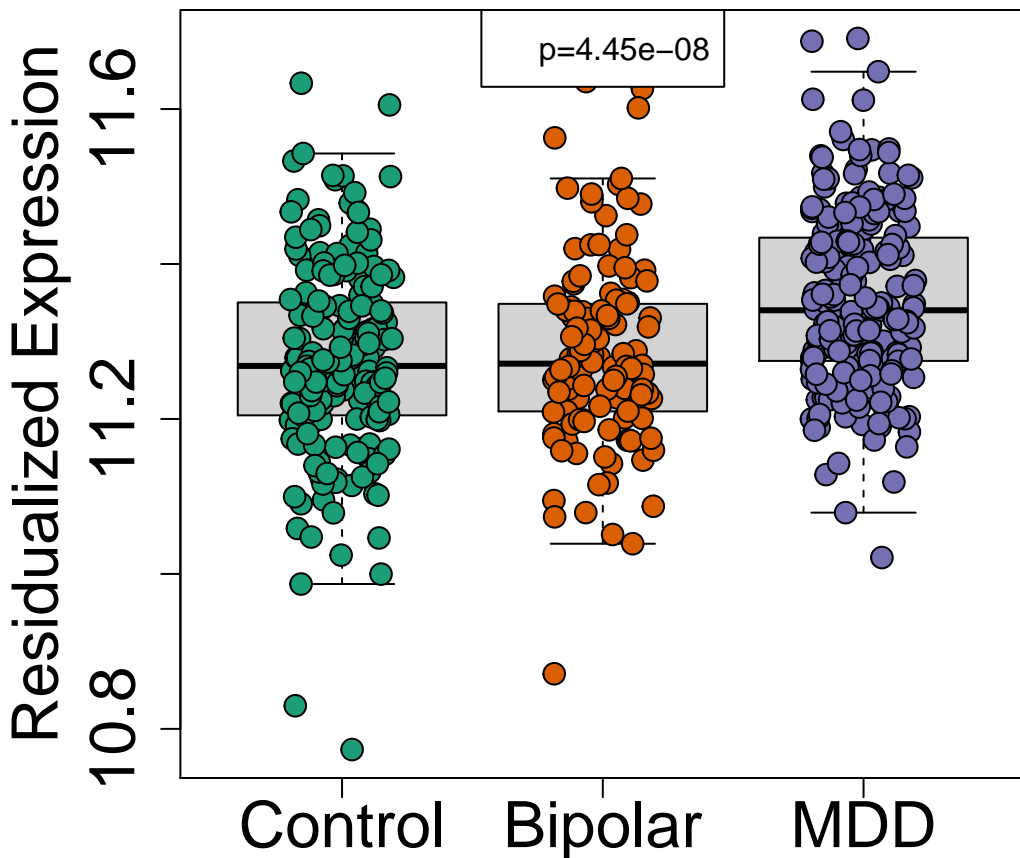
TUBA1B, InGen

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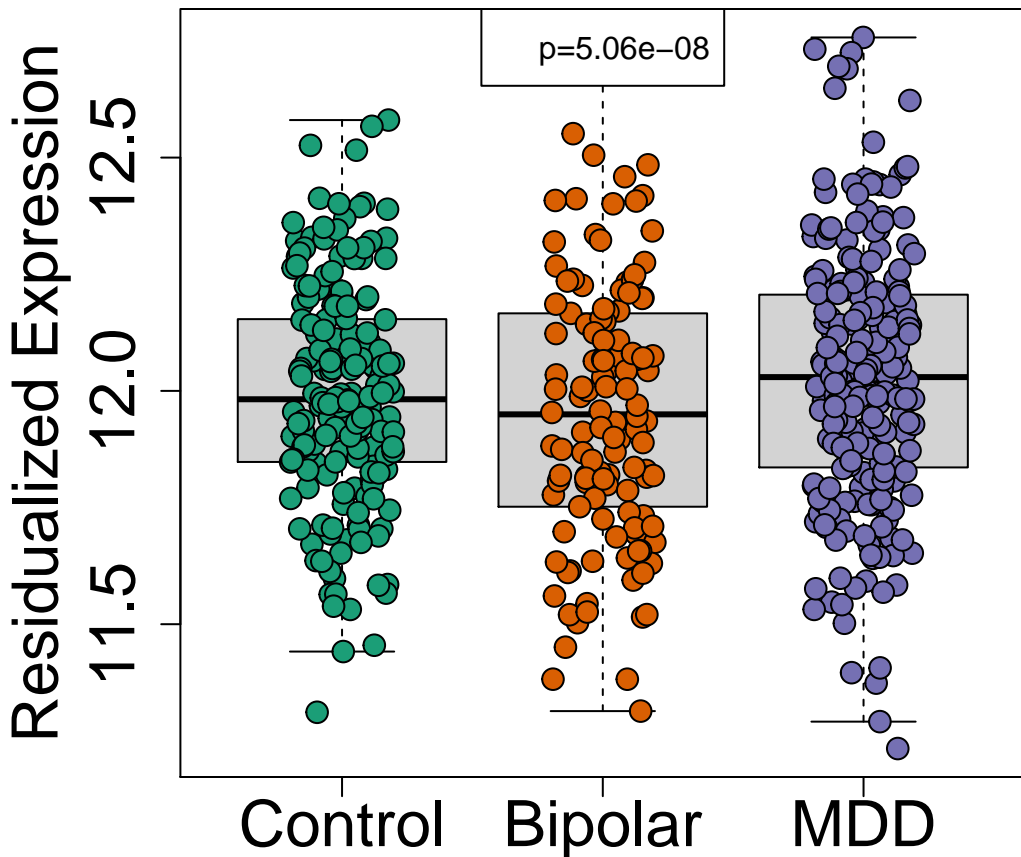
RPL31, InGen

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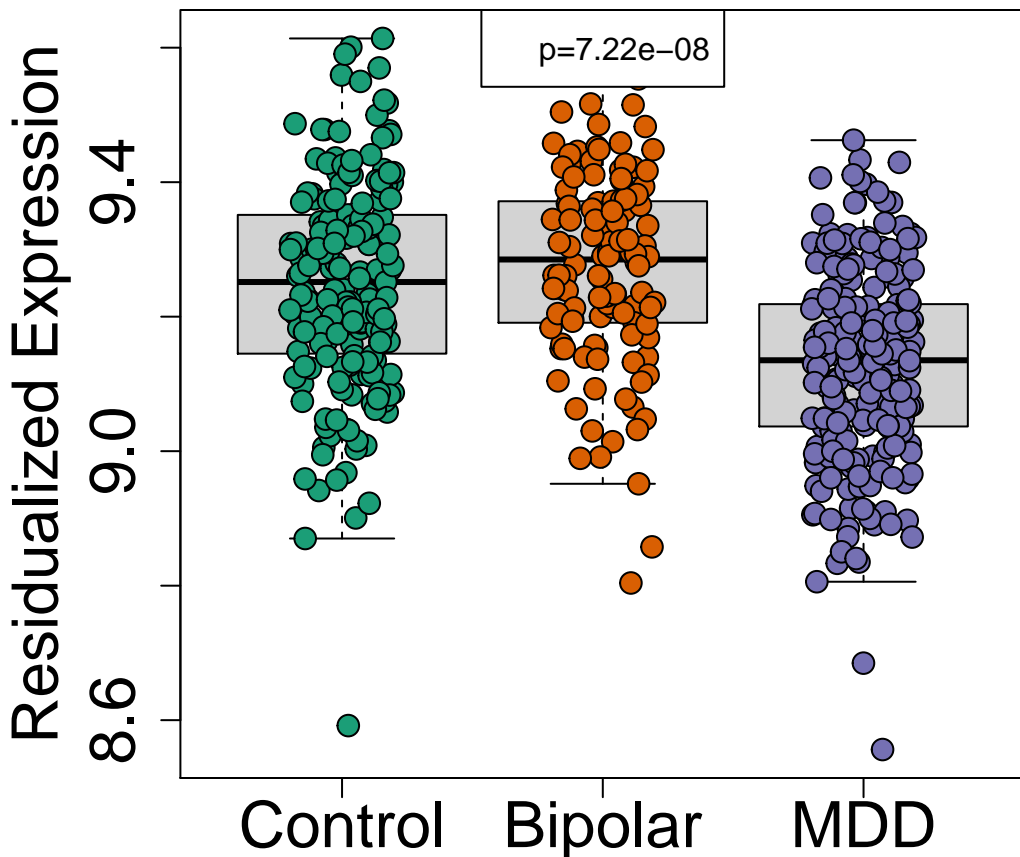
GLUL, InGen

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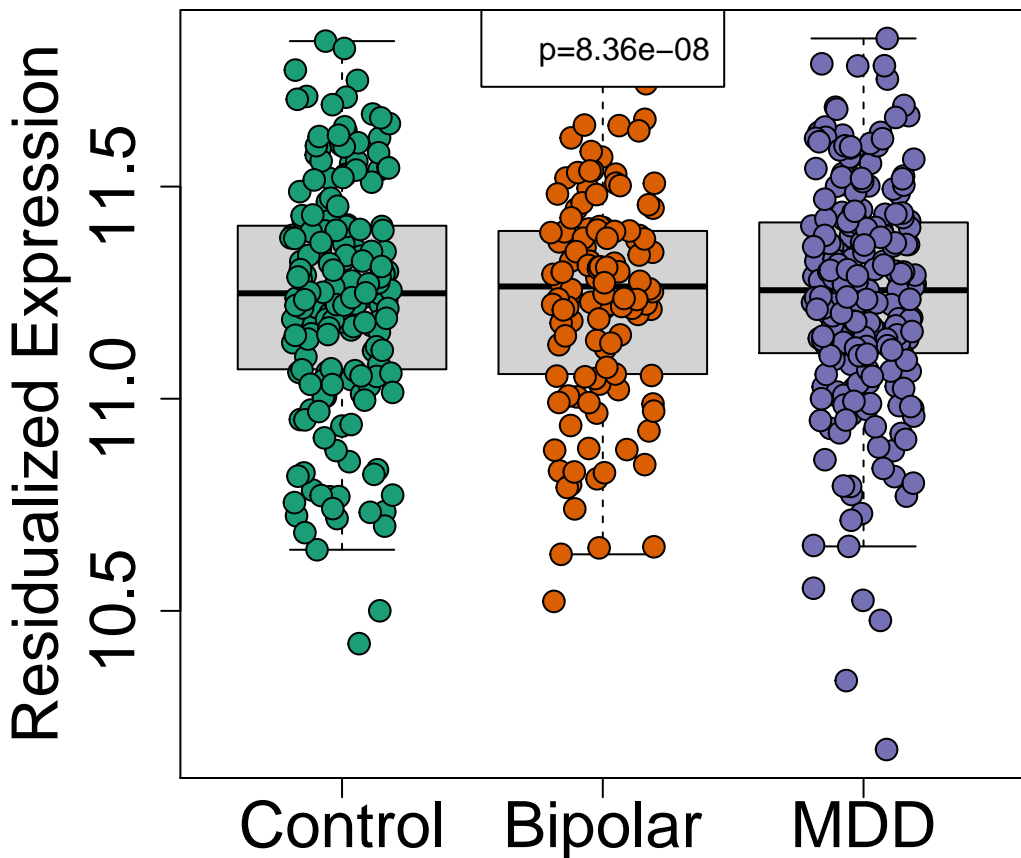
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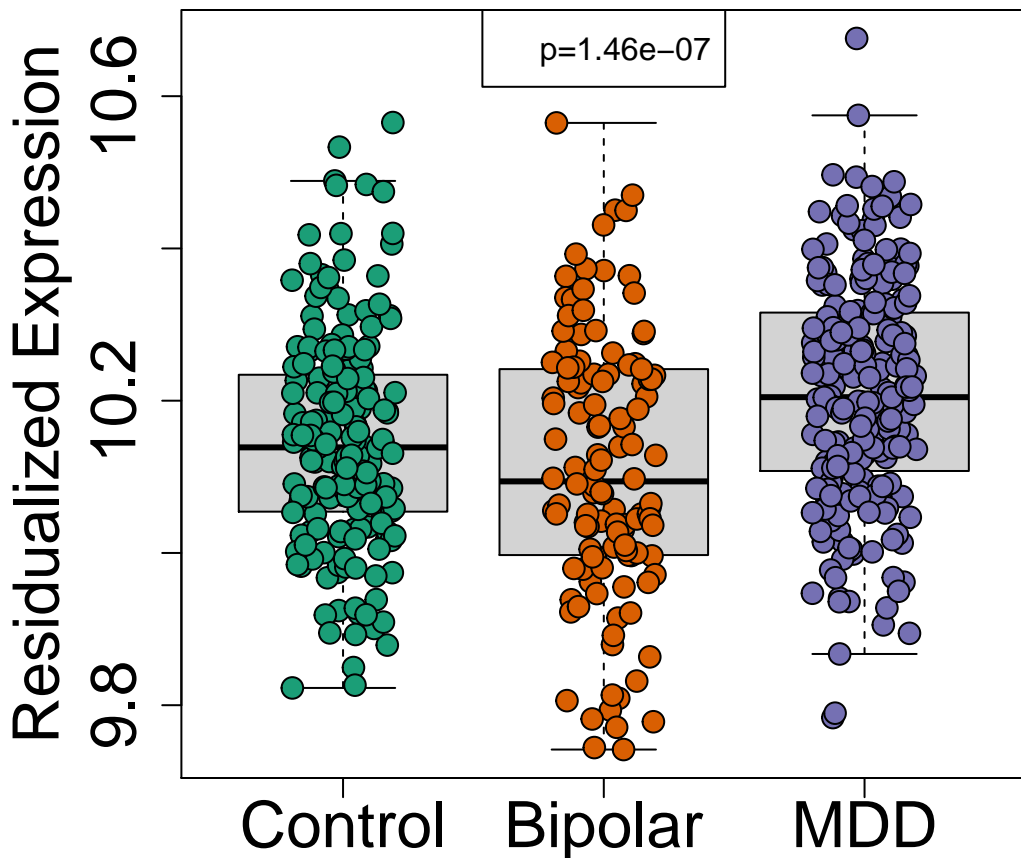
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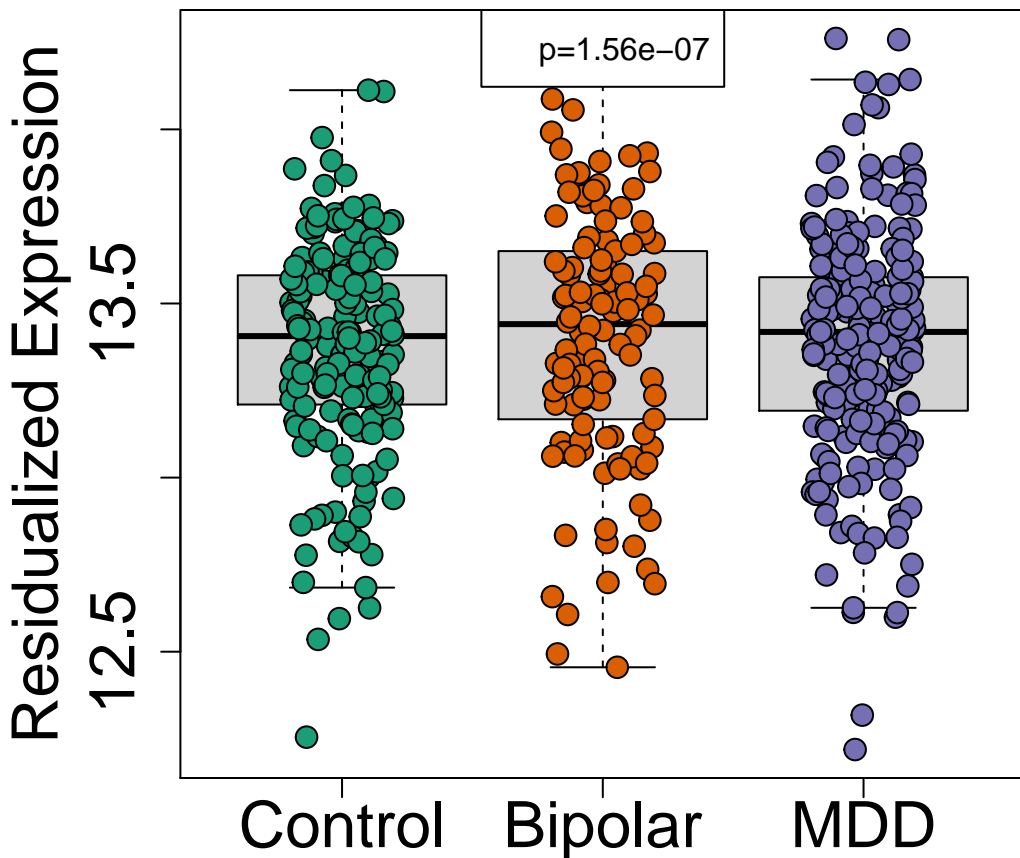
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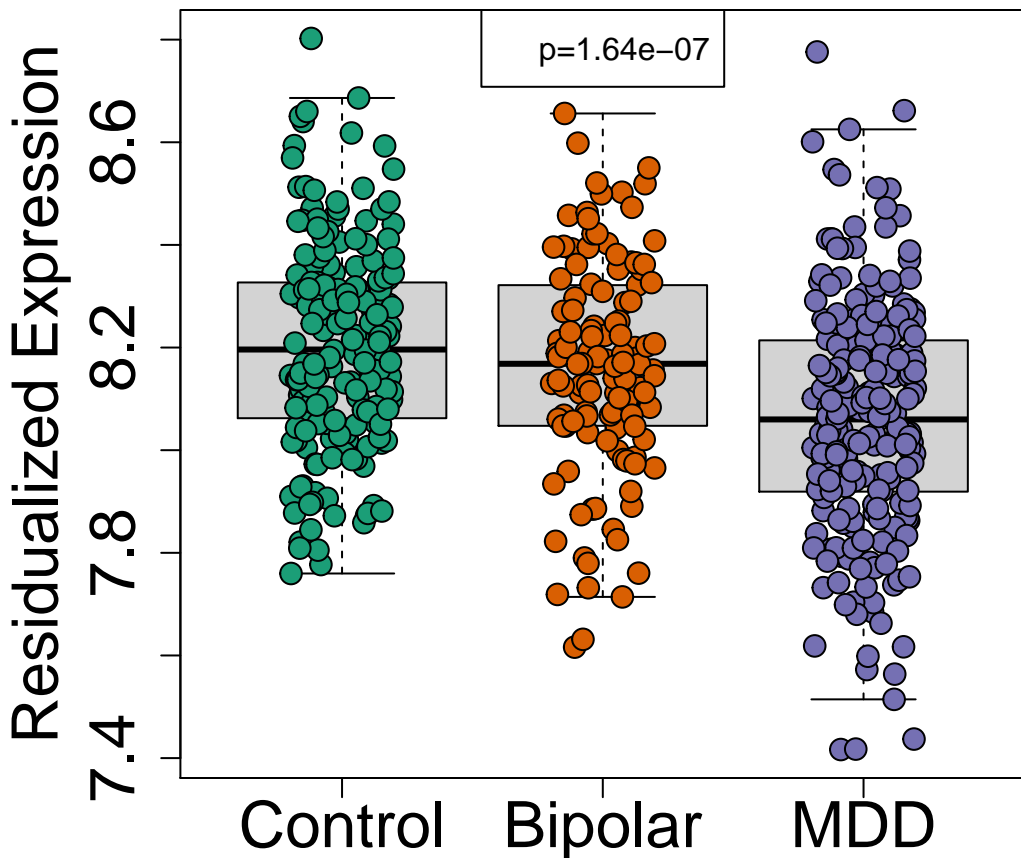
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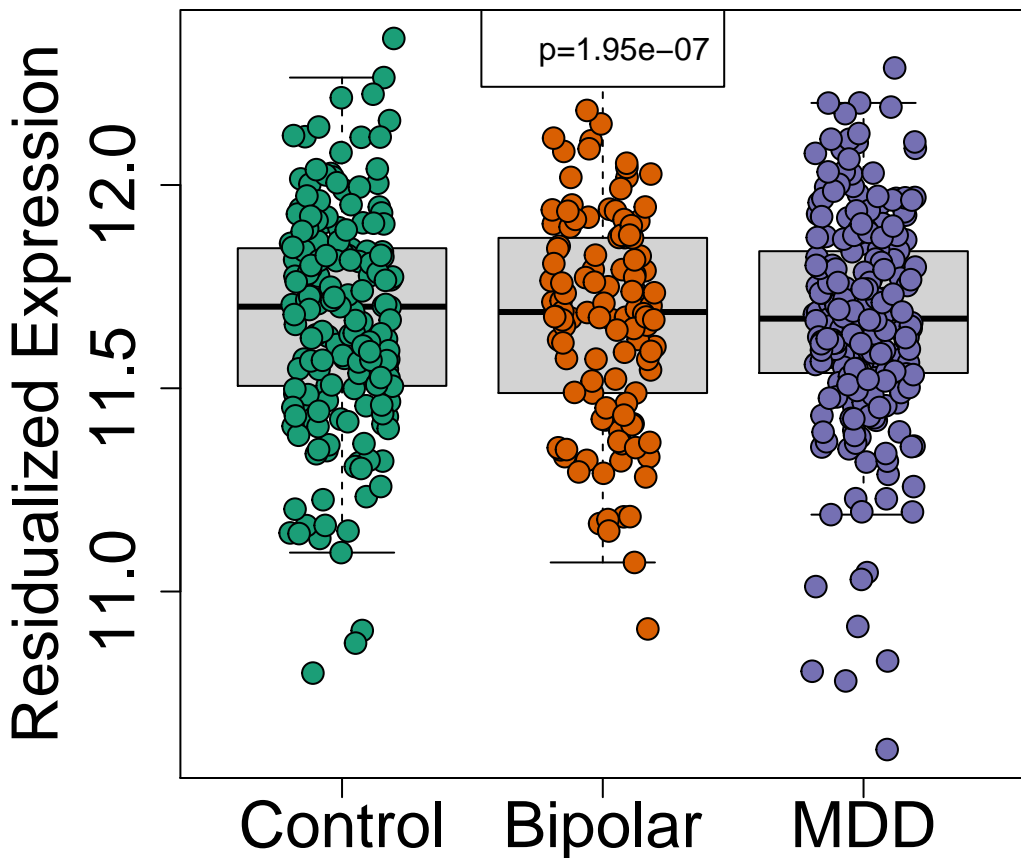
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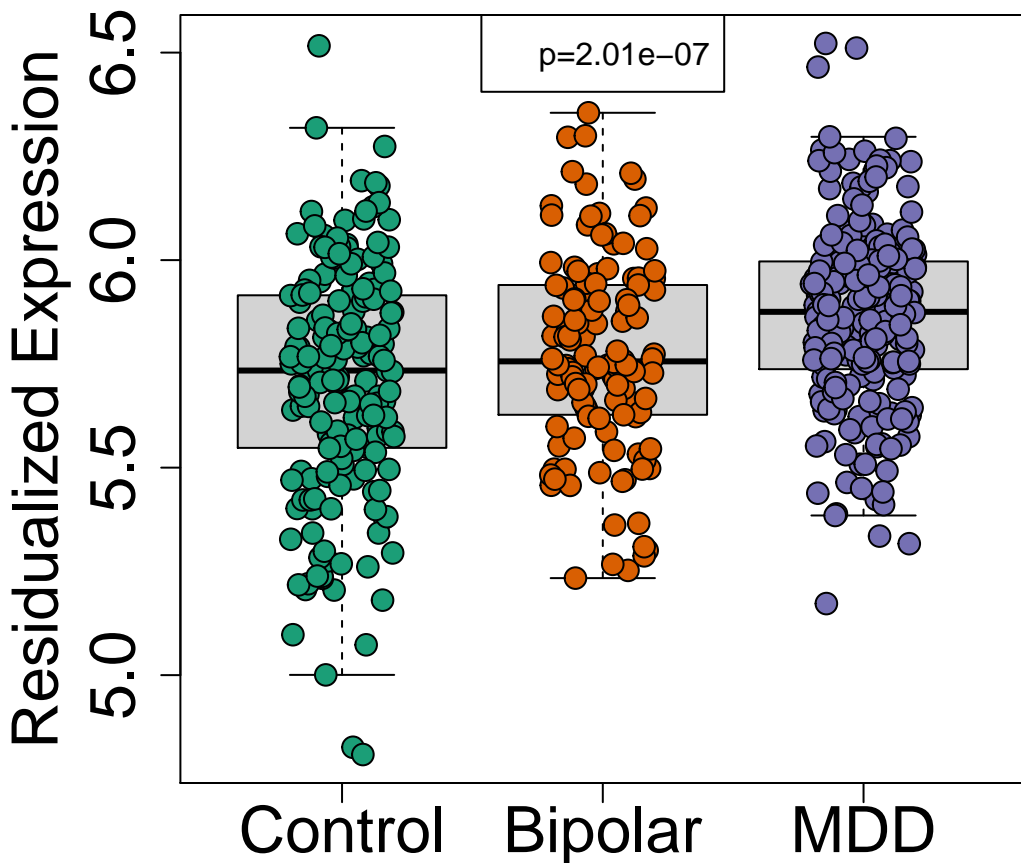
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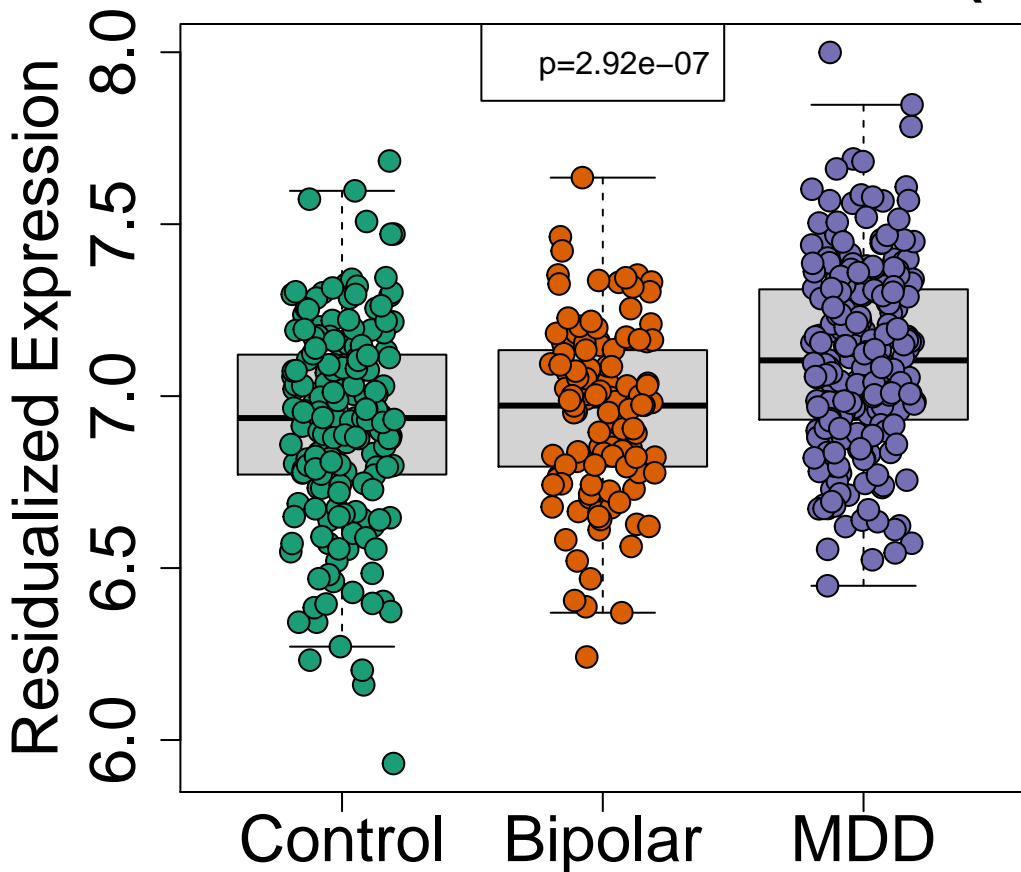
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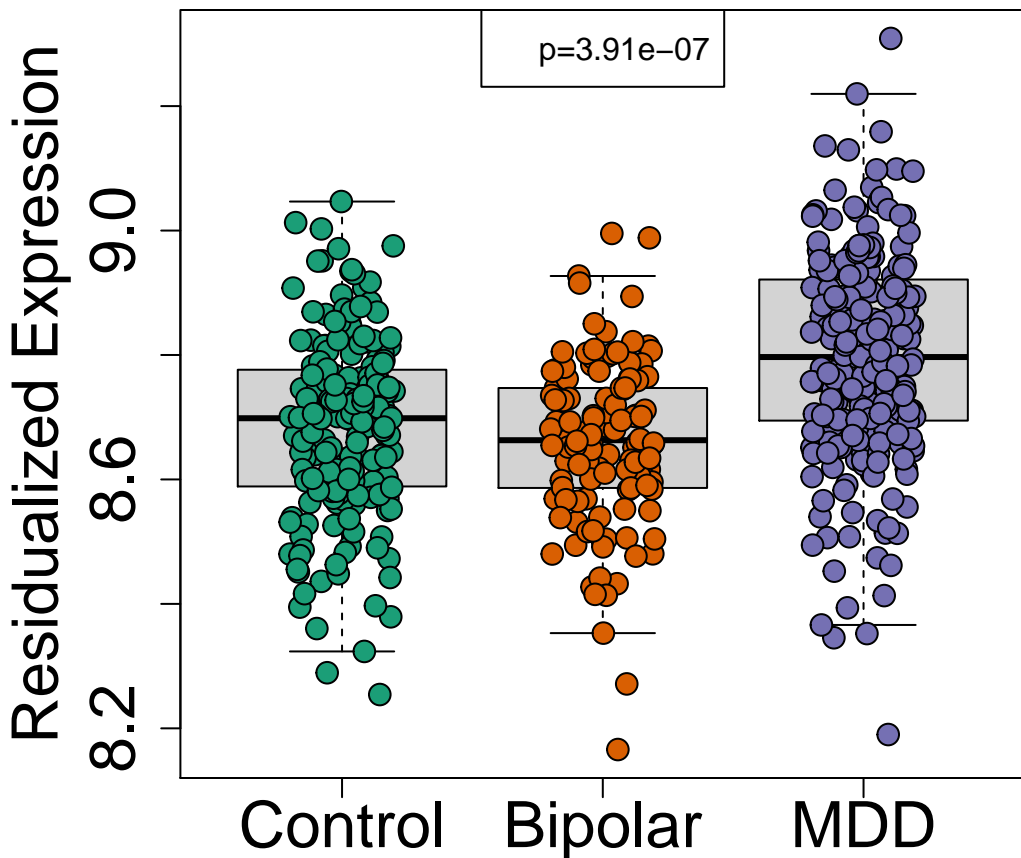
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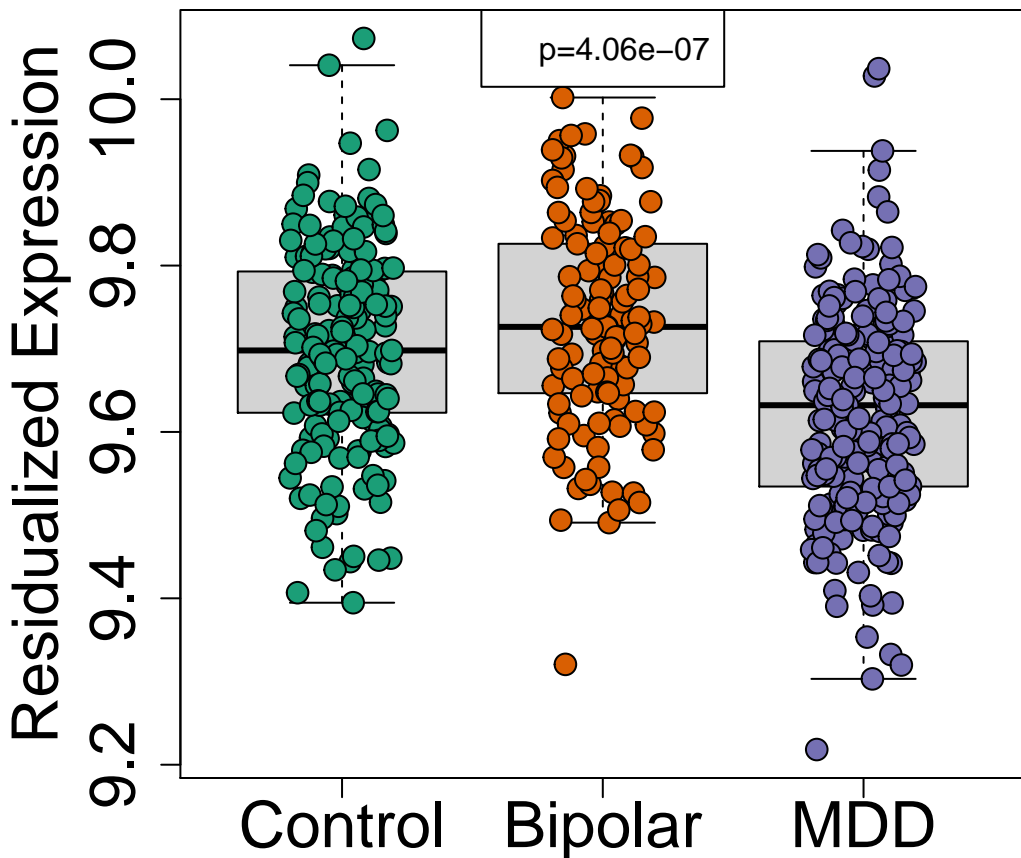
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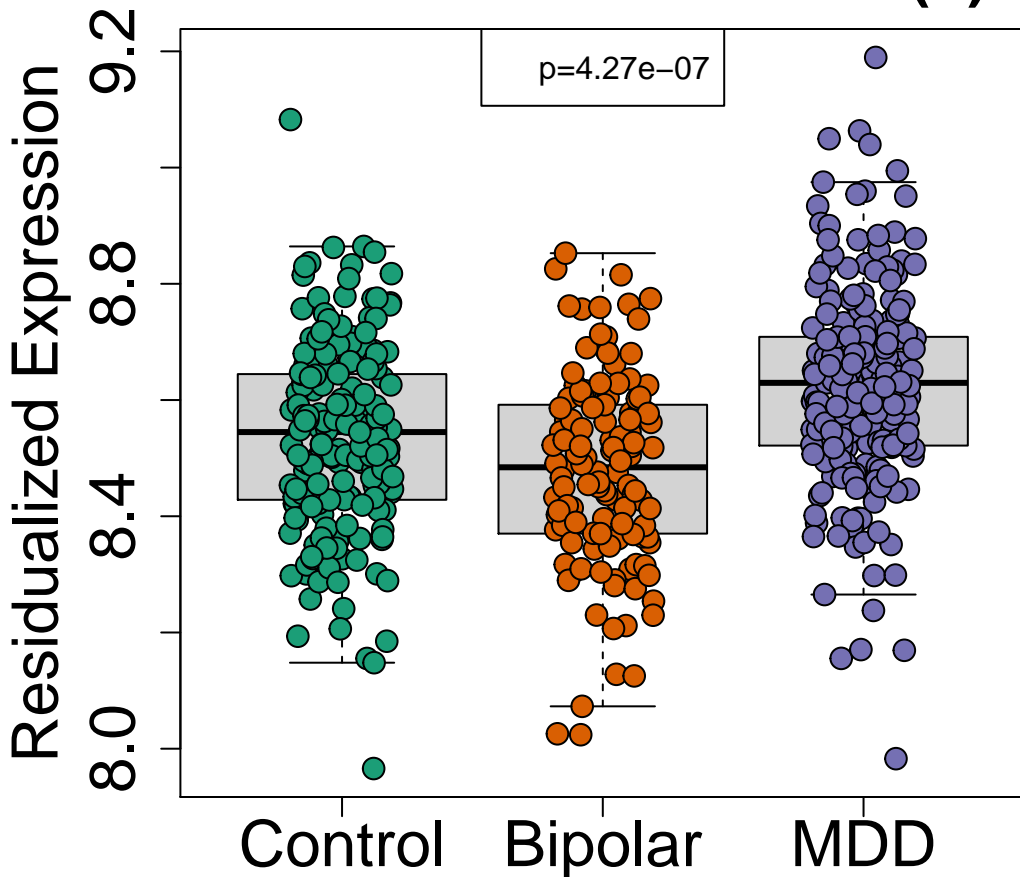
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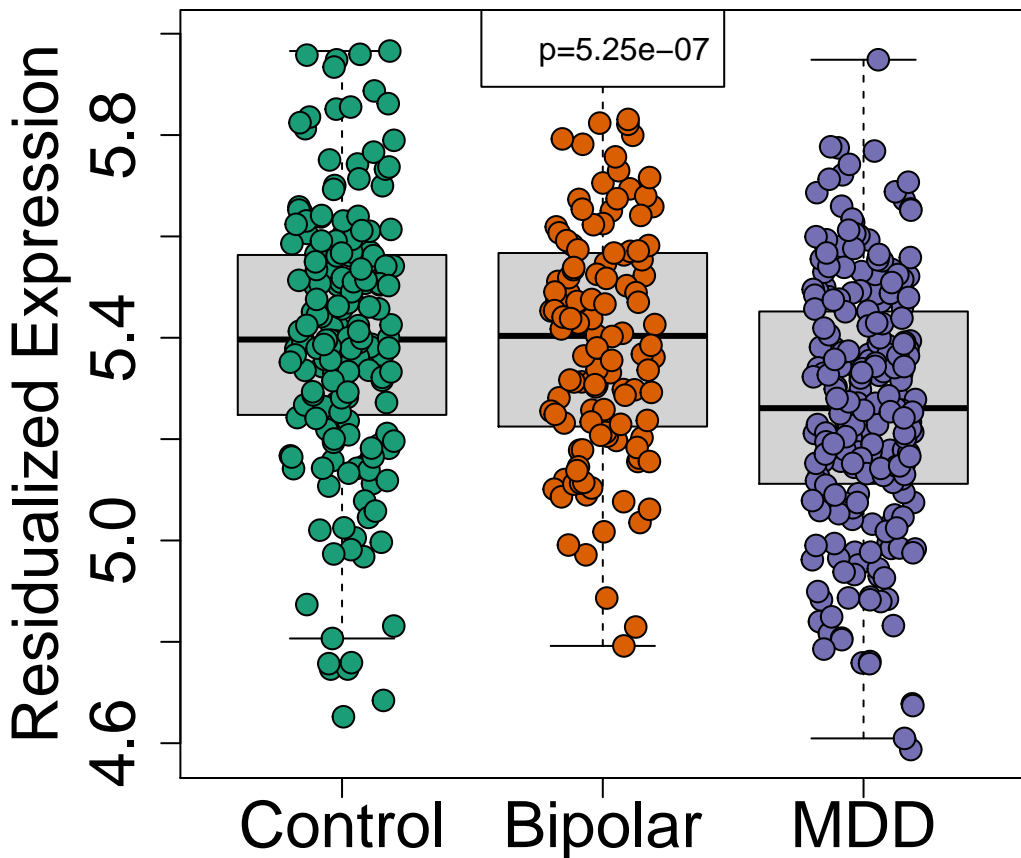
ENO1, InGen

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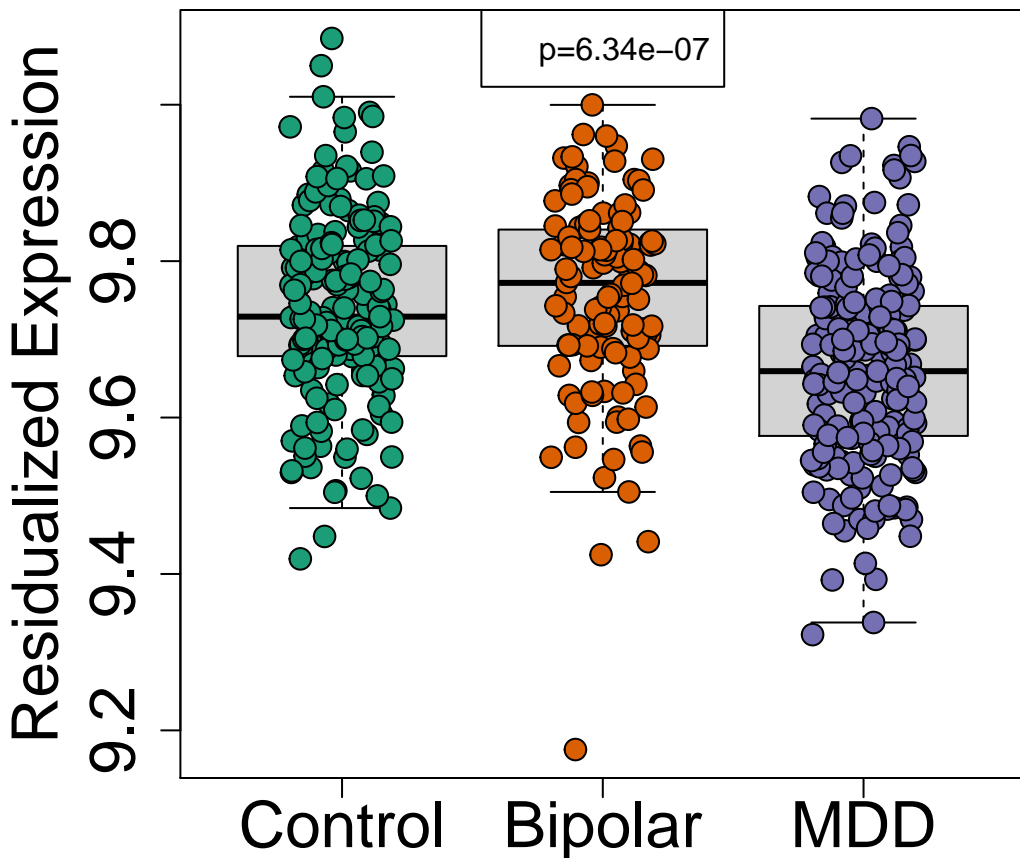
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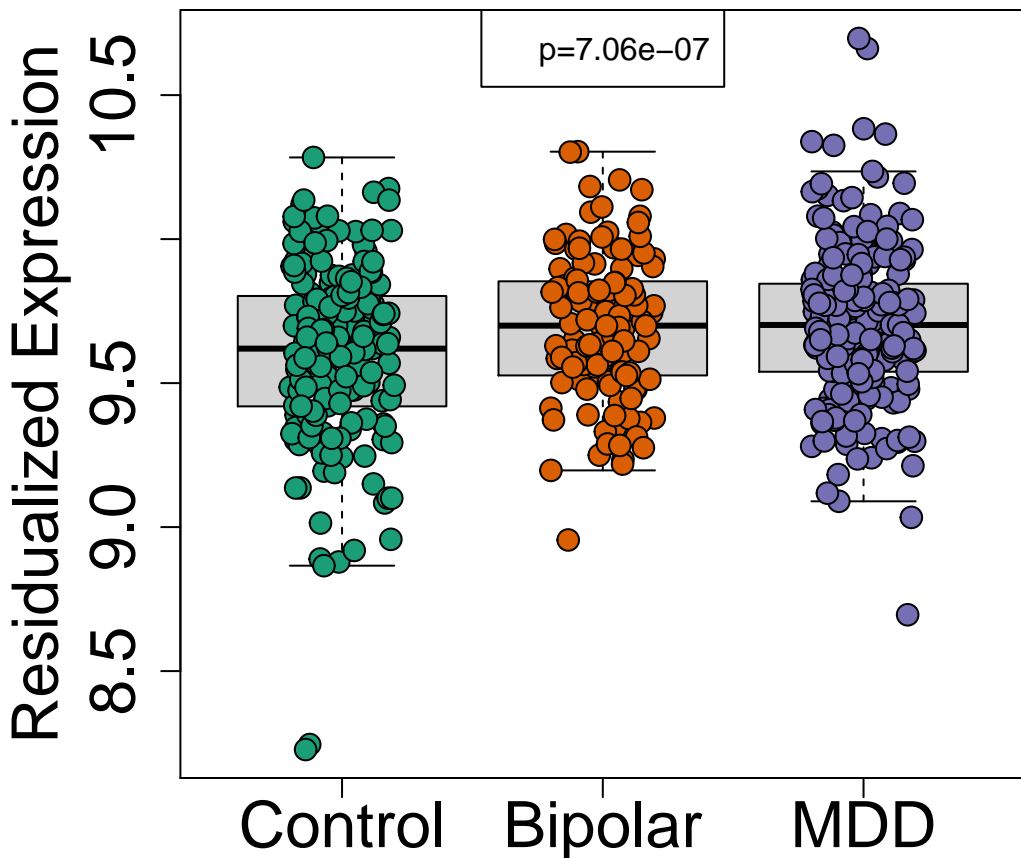
RPS10-NUDT3, InGen

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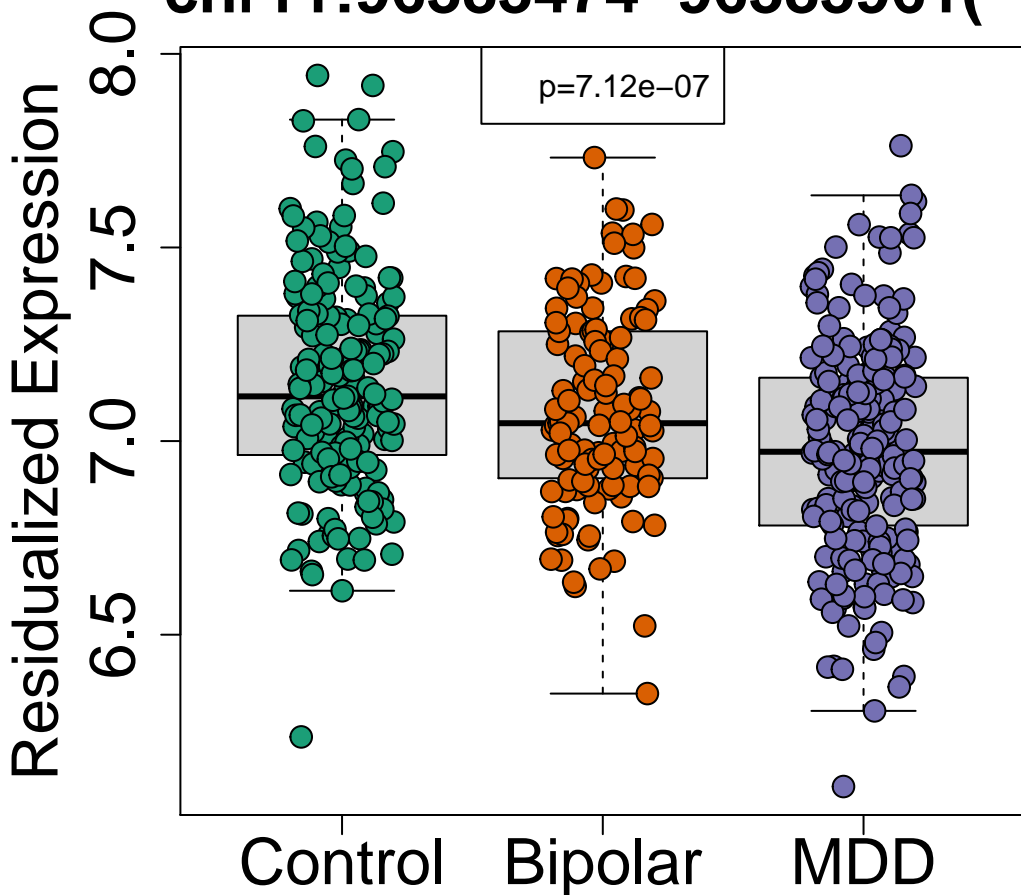
GAP43, InGen

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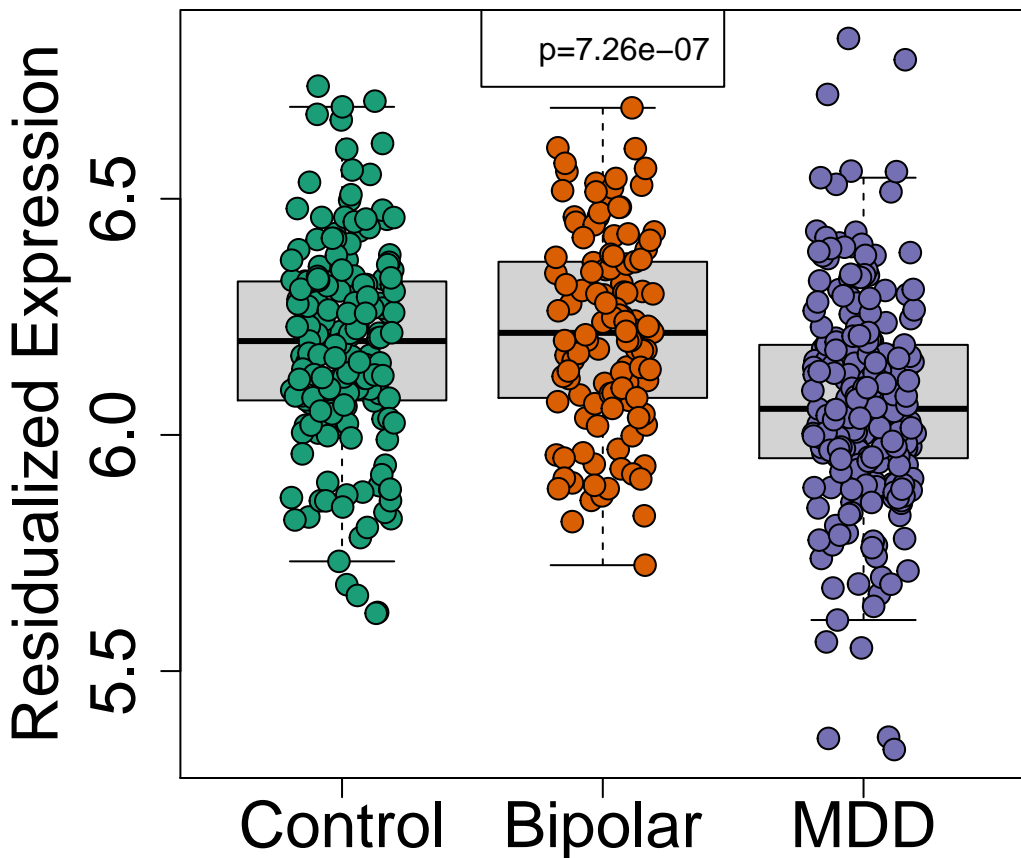
CCDC82, InGen

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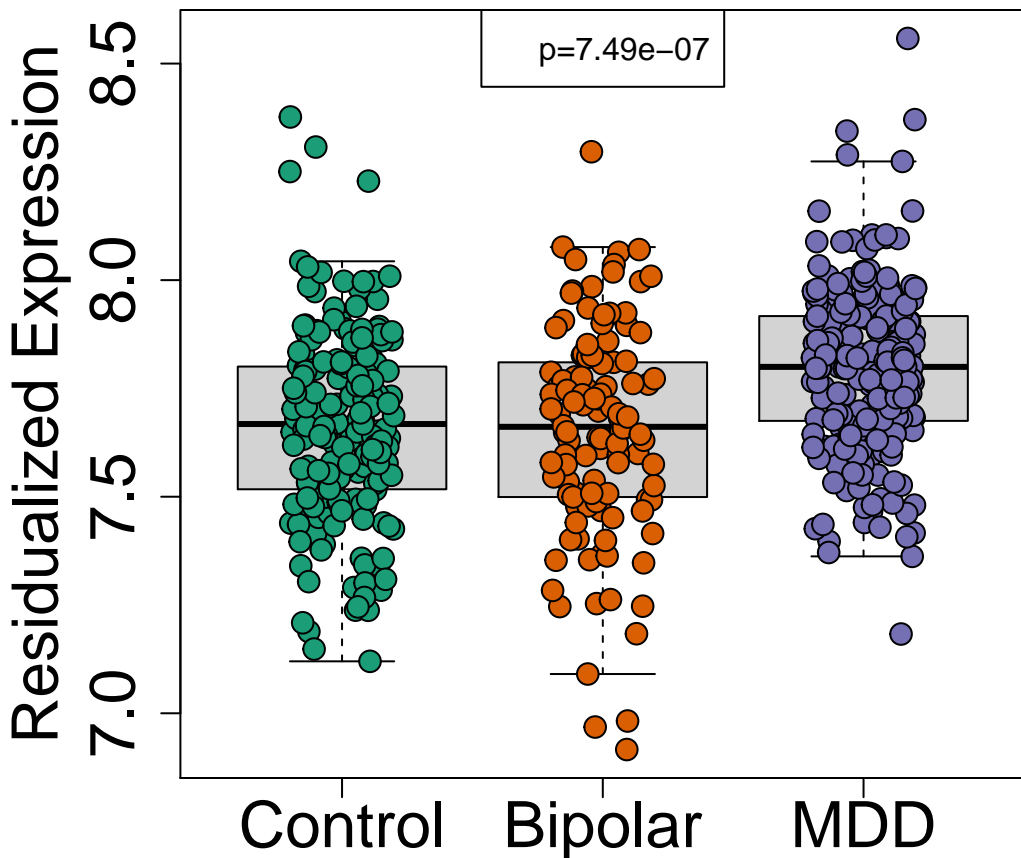
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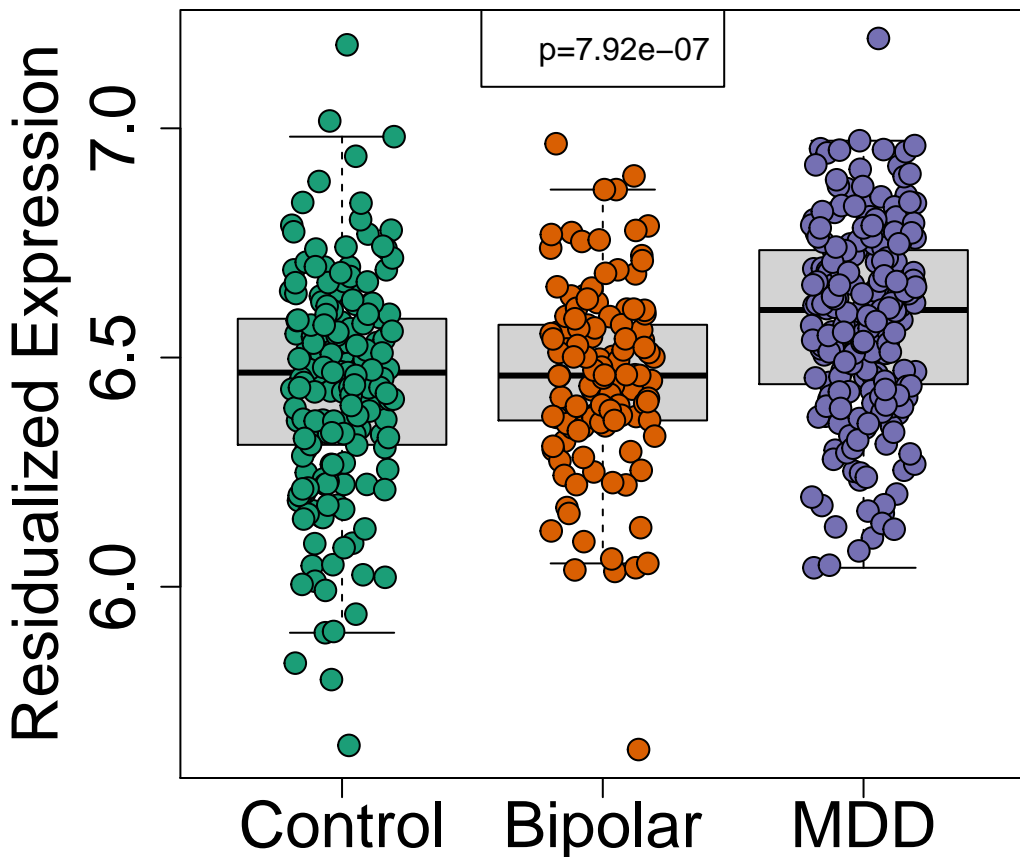
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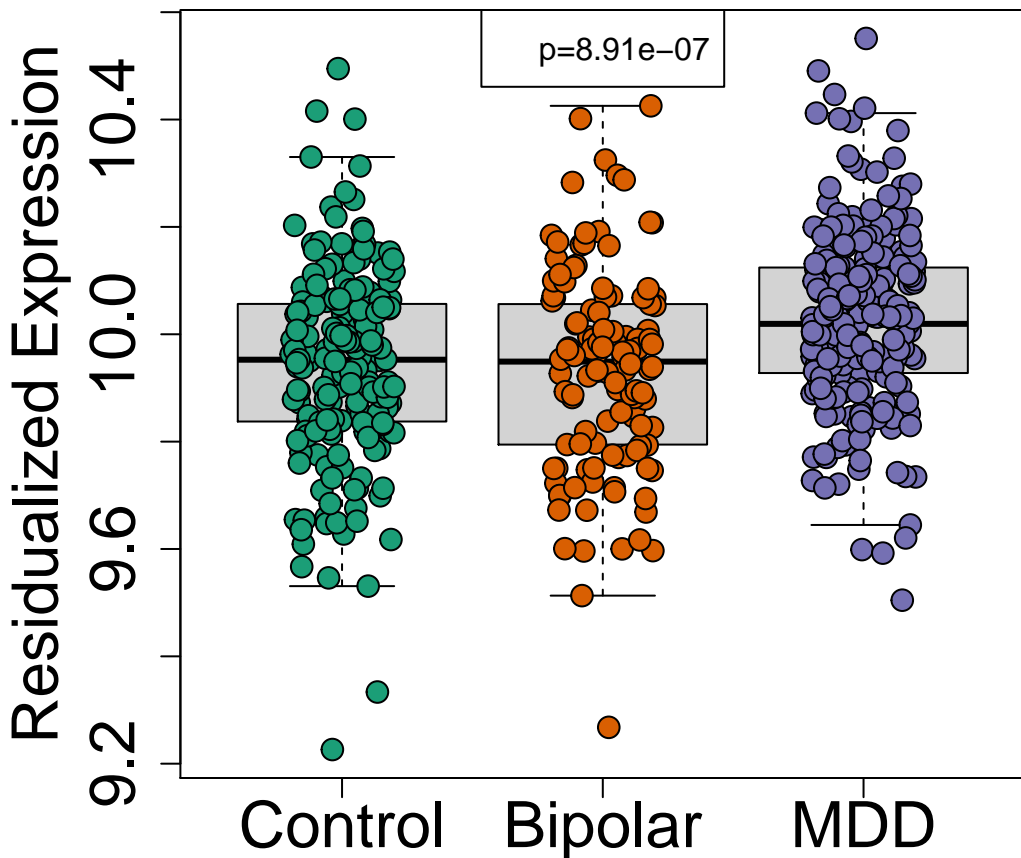
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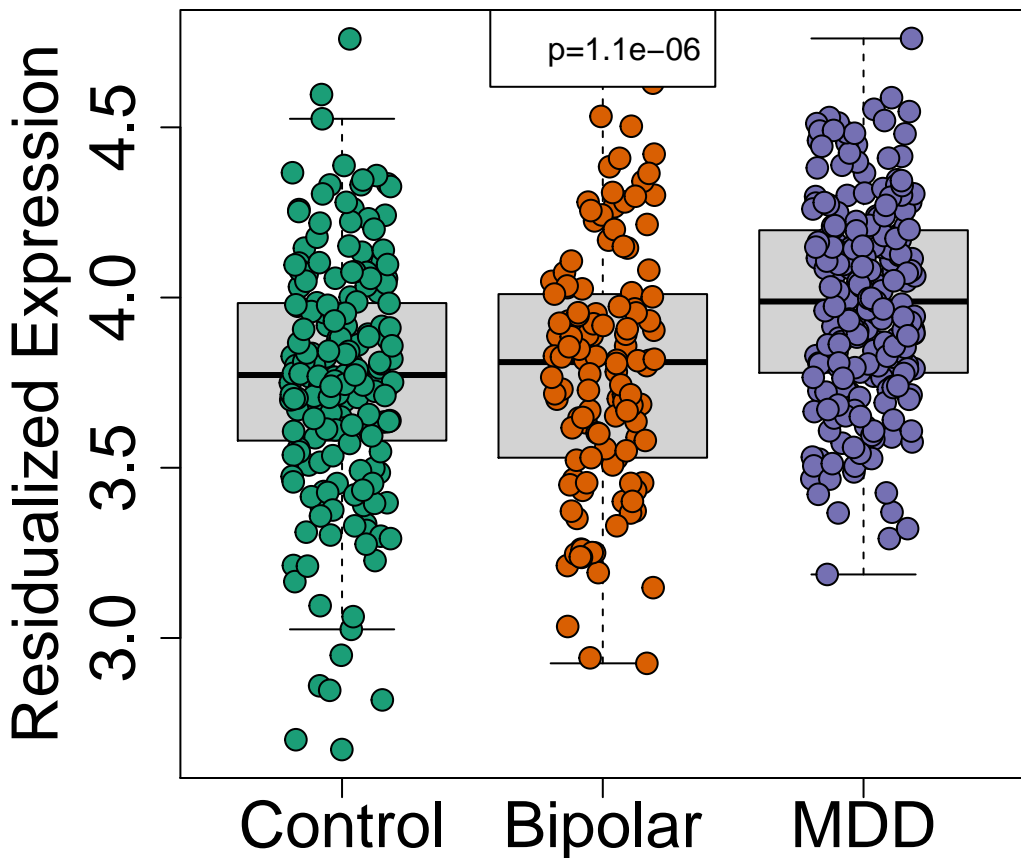
MDH1, InGen

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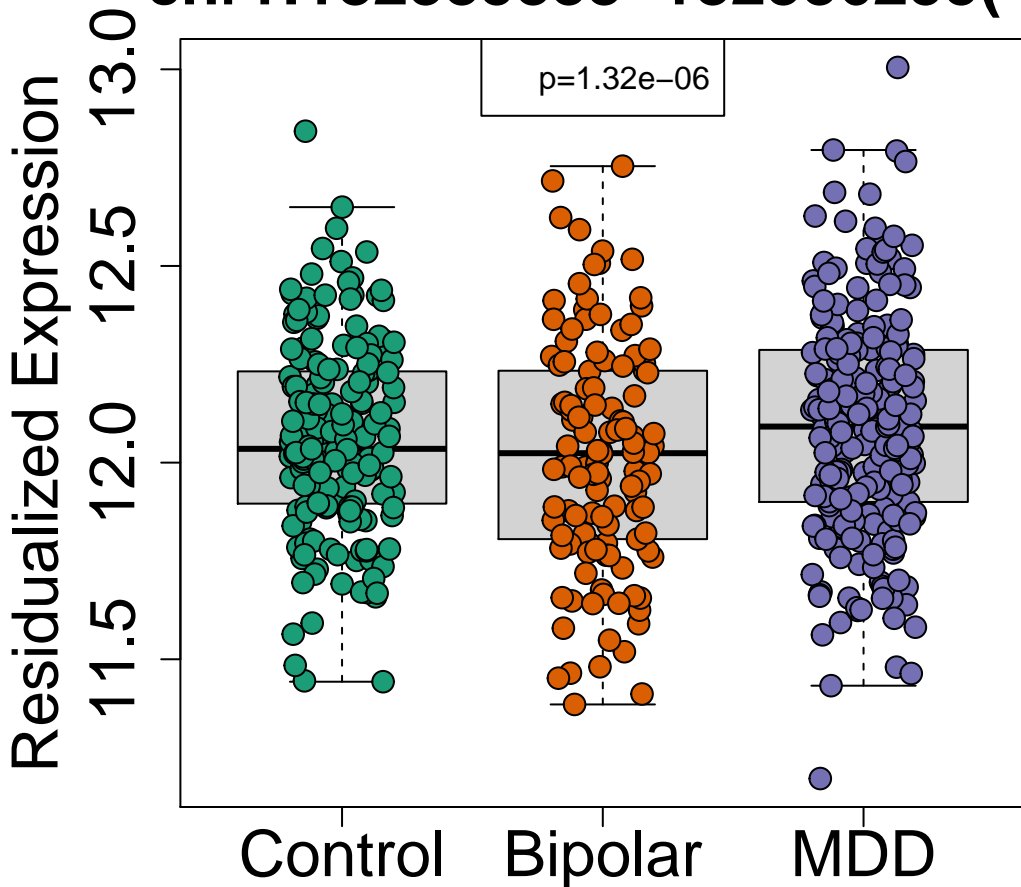
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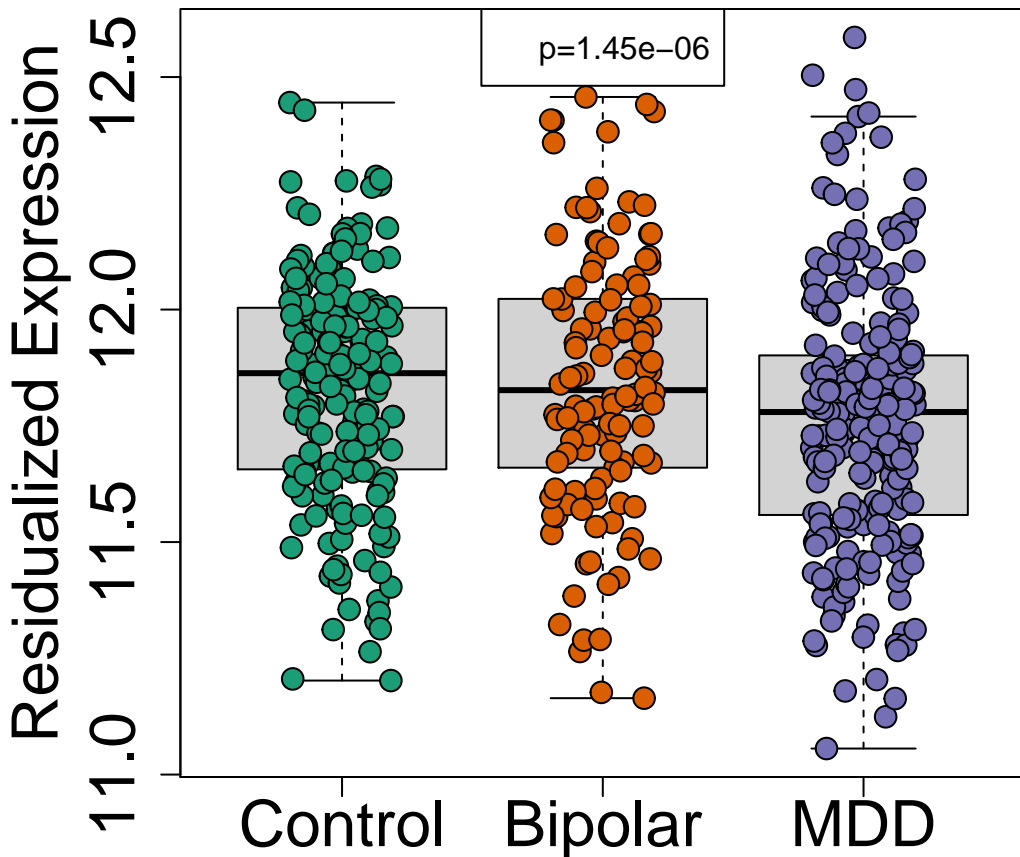
GLUL, InGen

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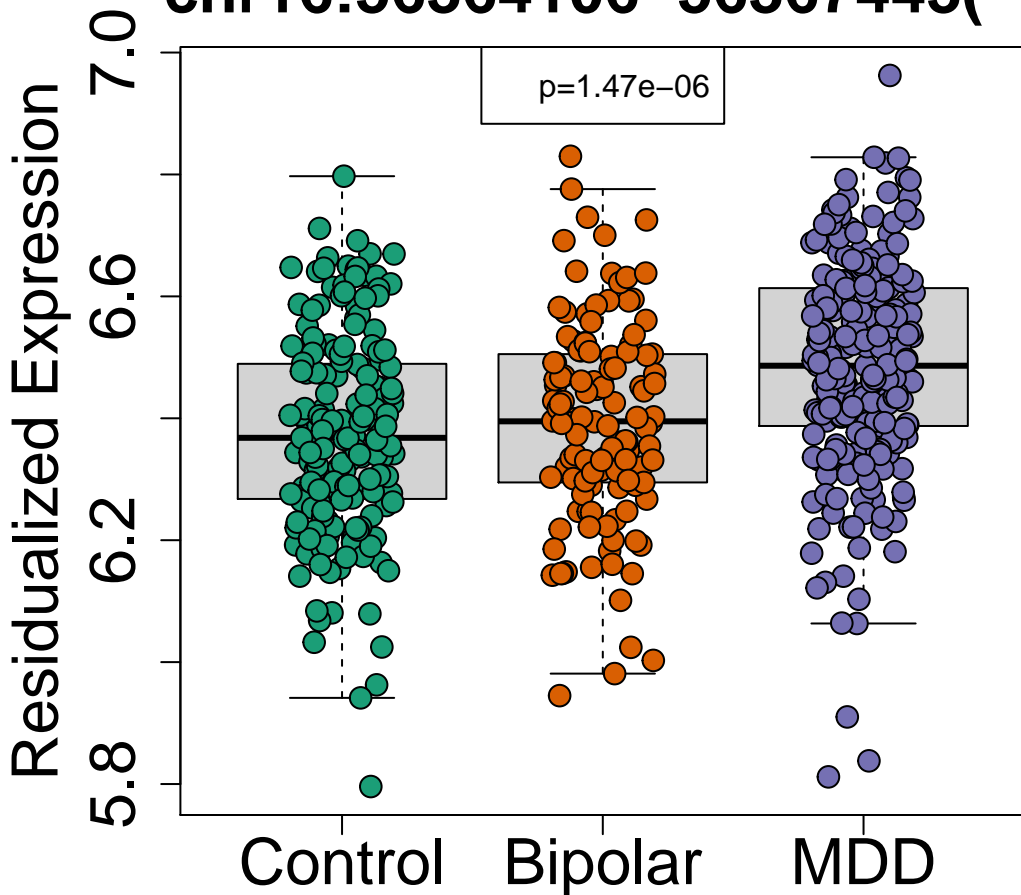
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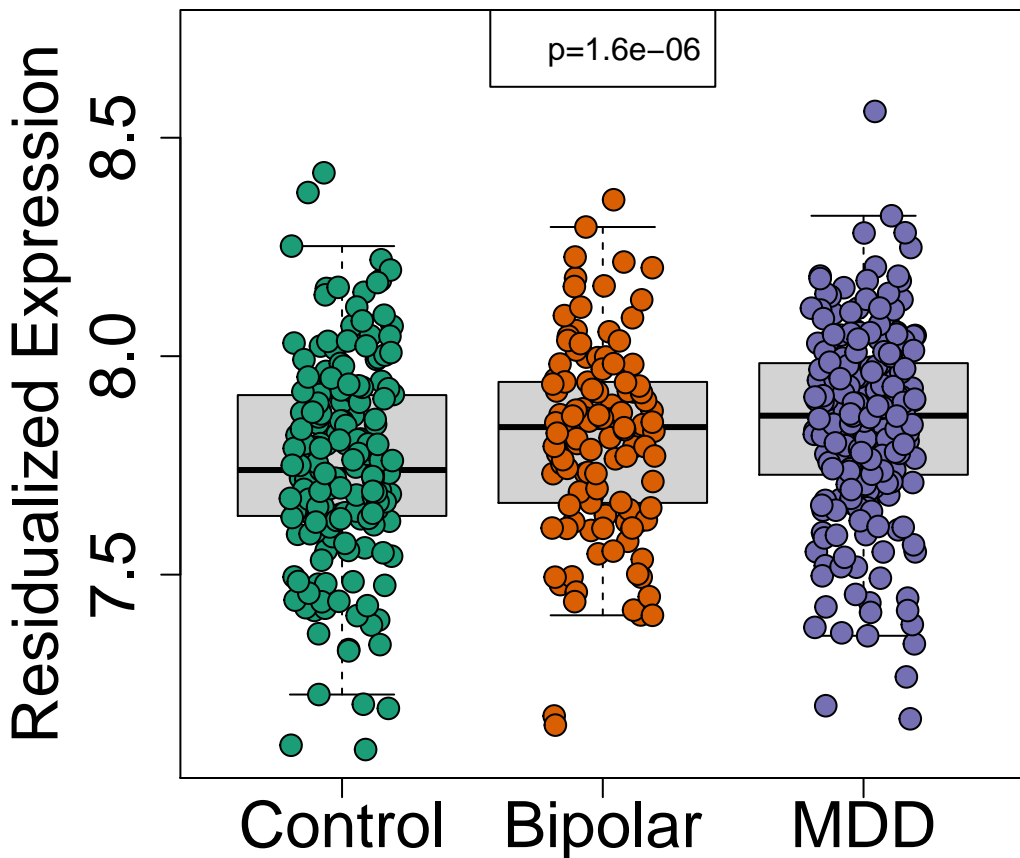
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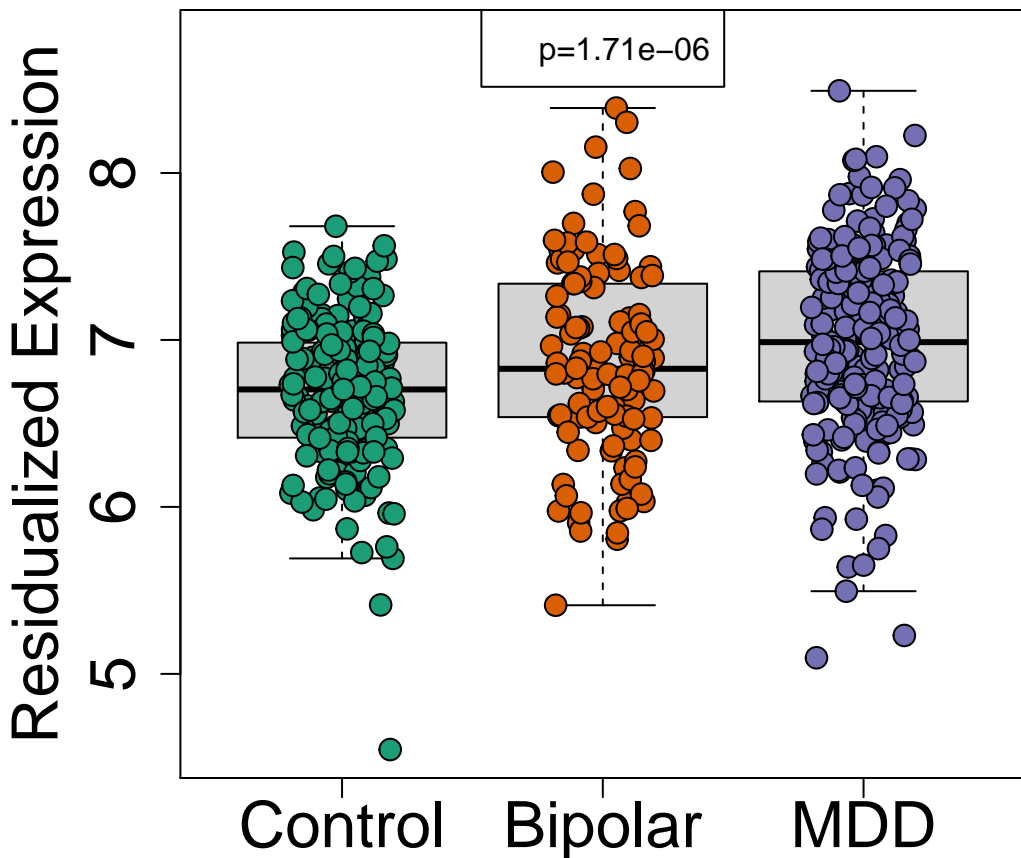
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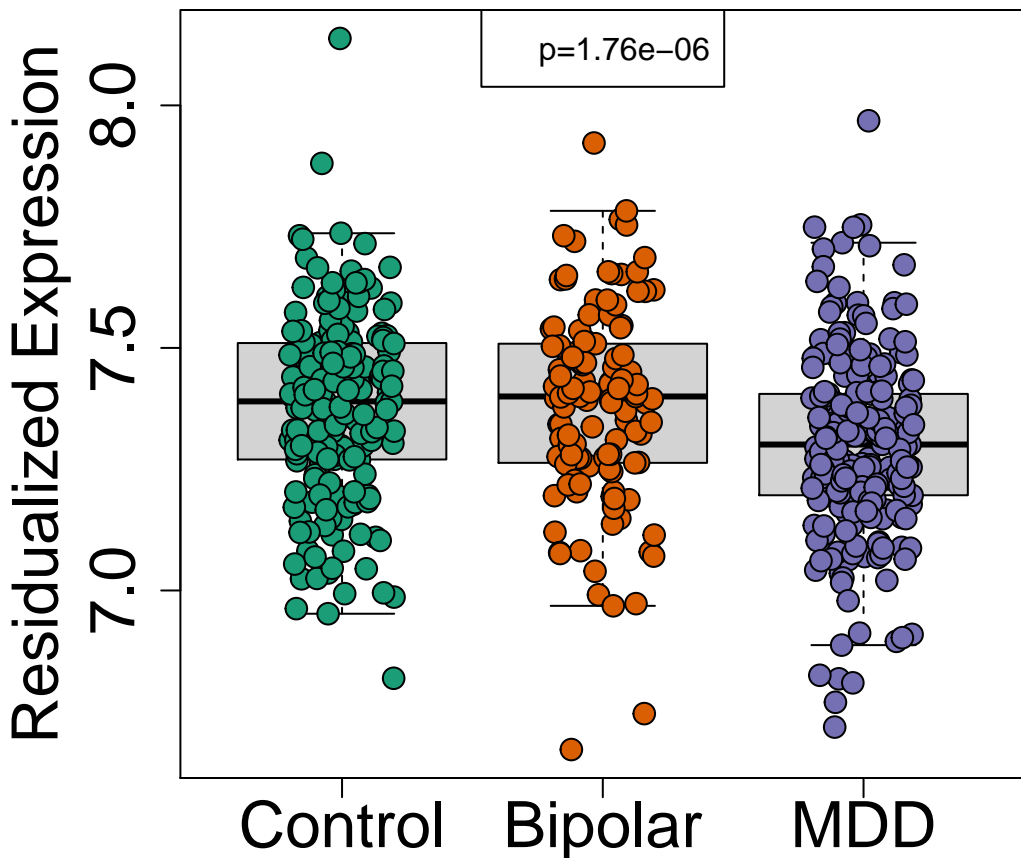
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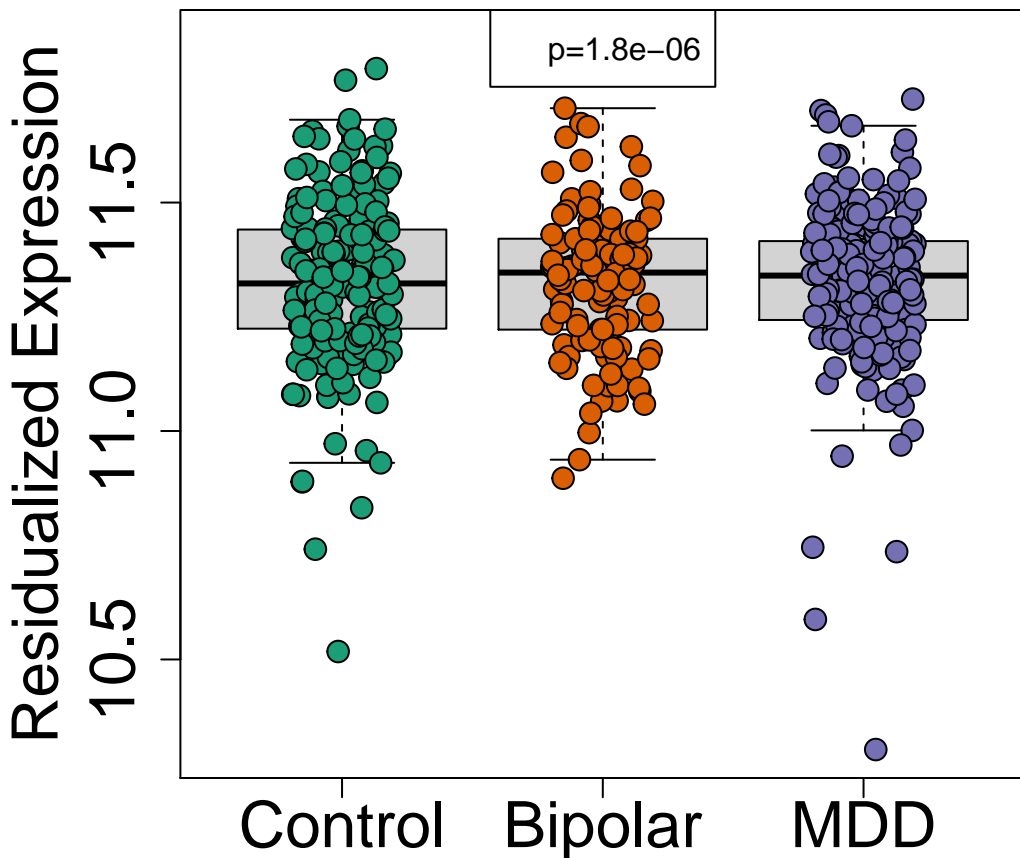
LRP1, InGen

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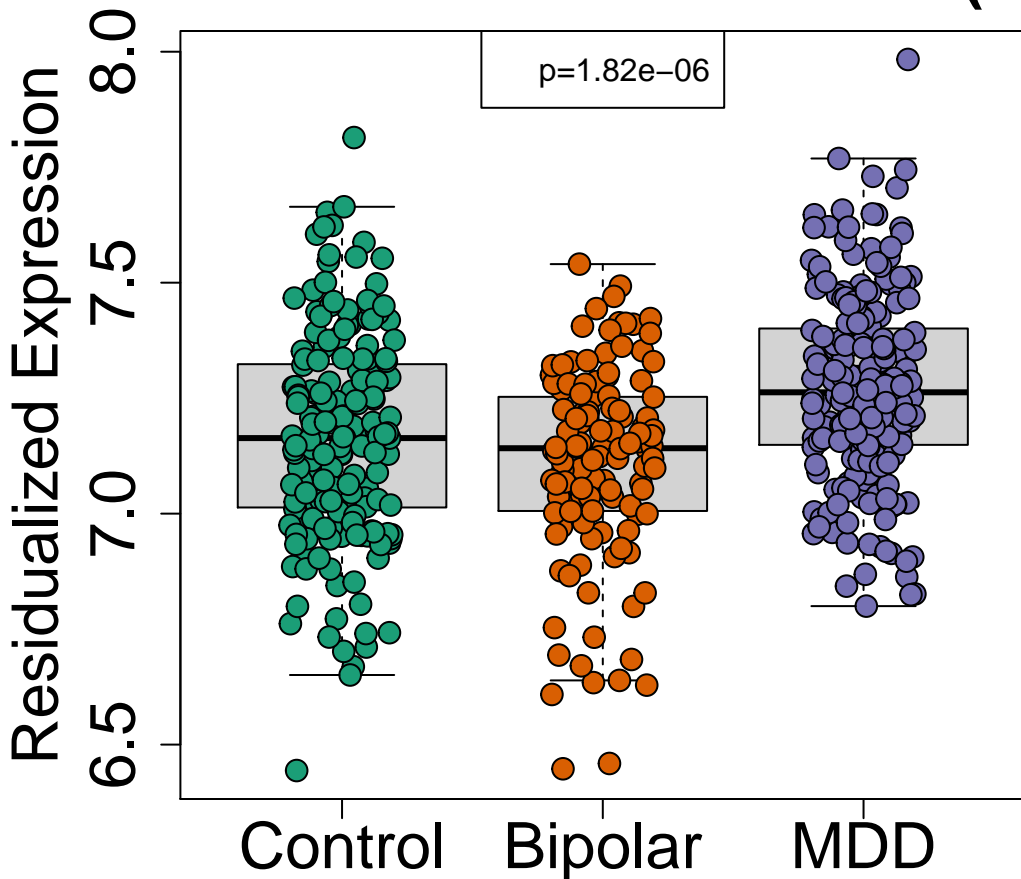
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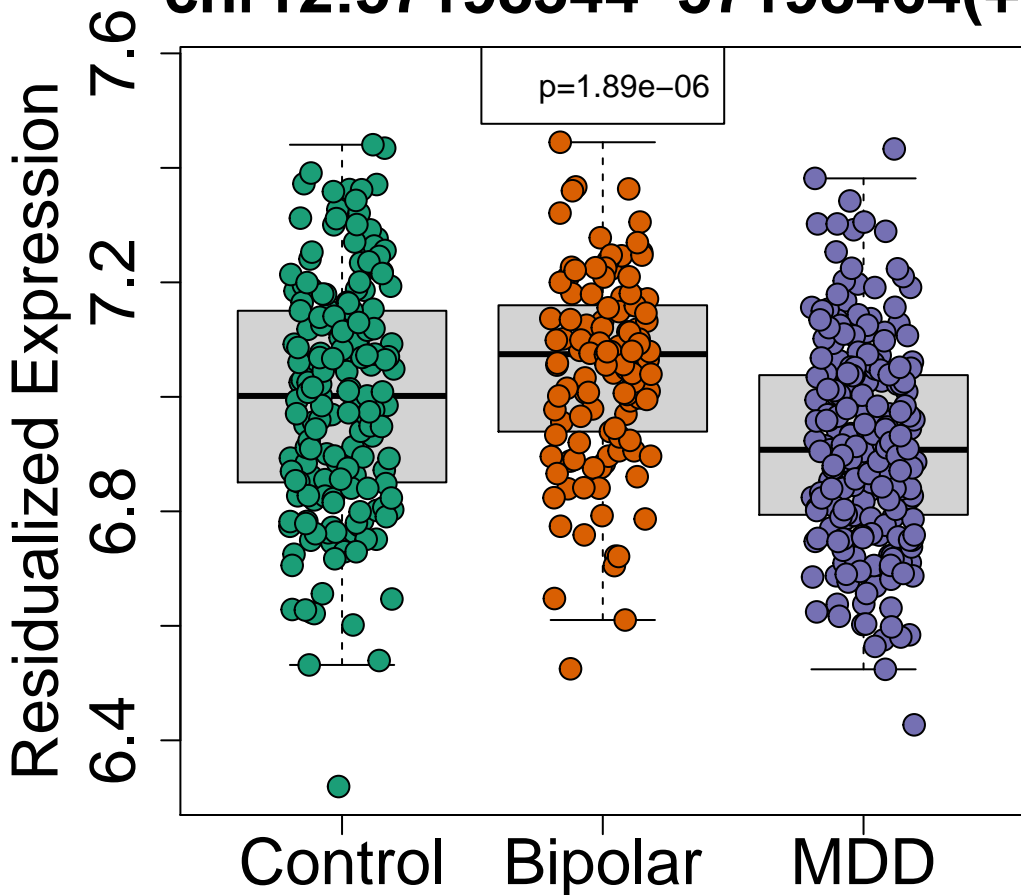
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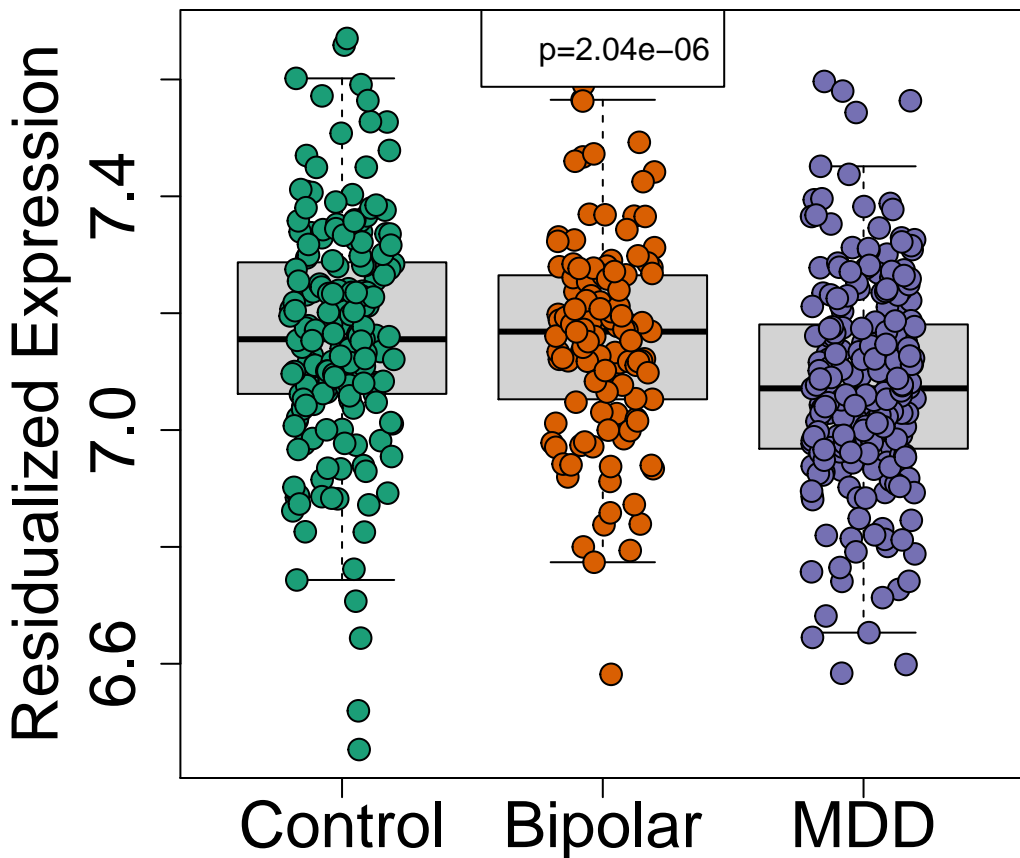
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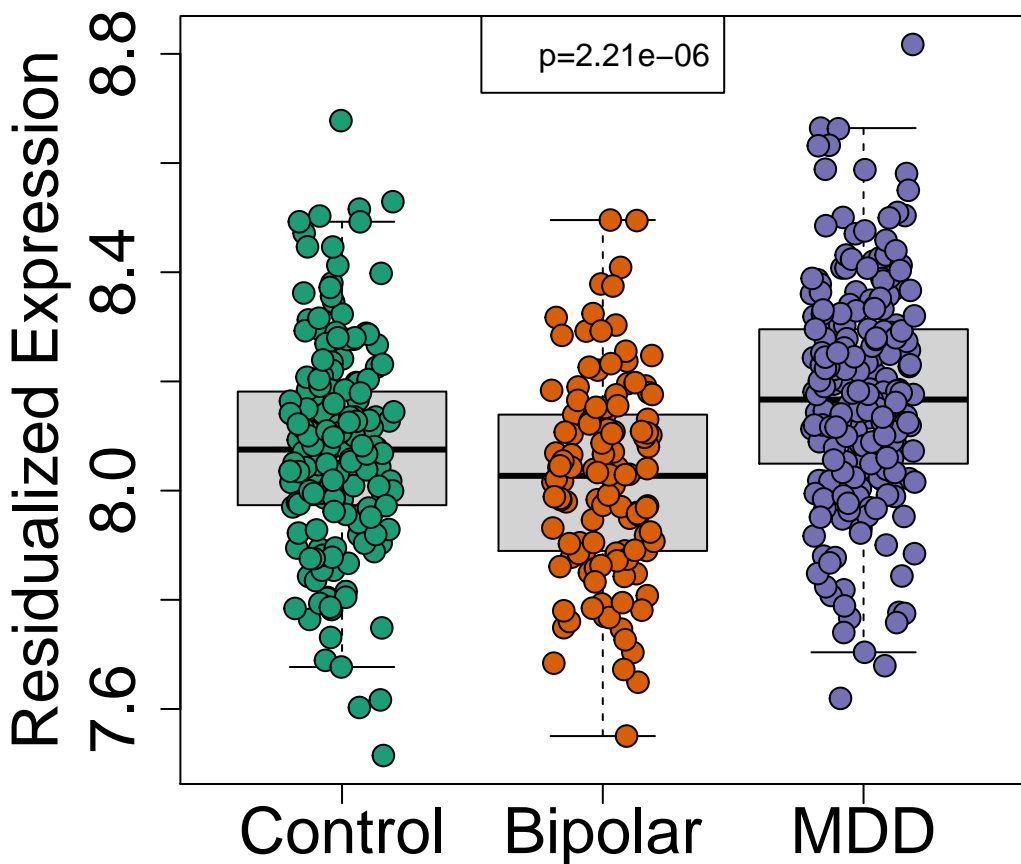
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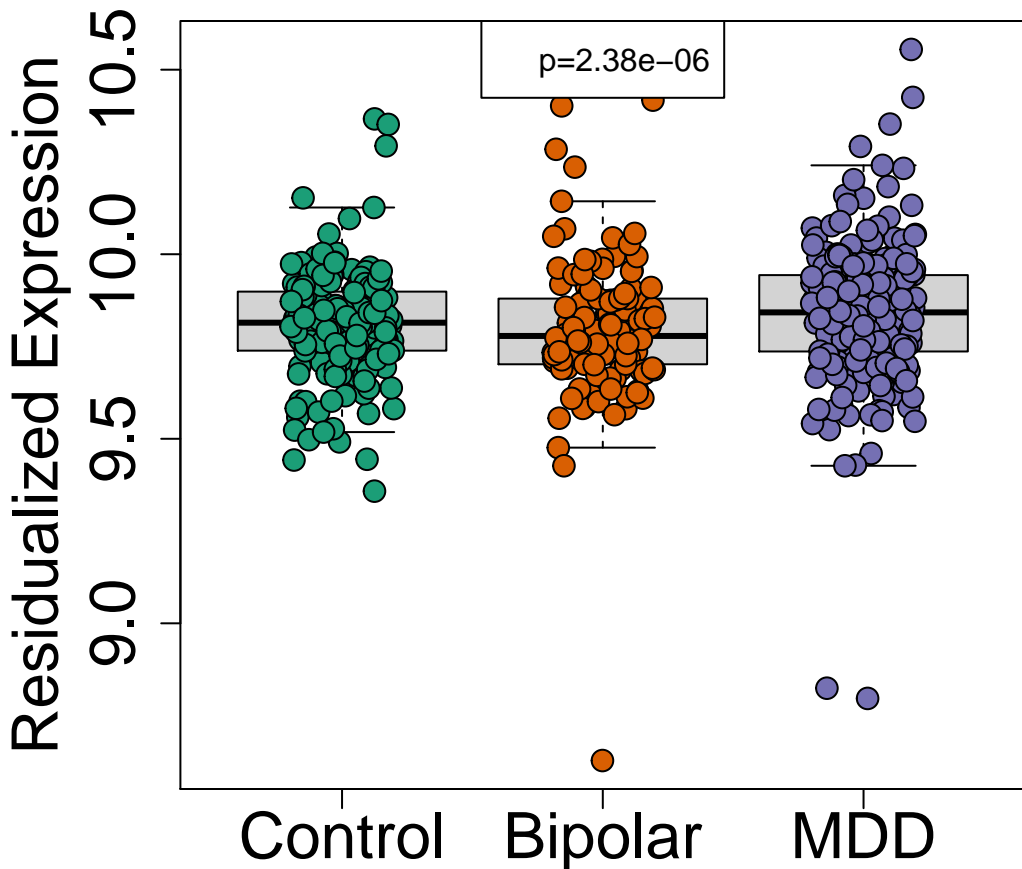
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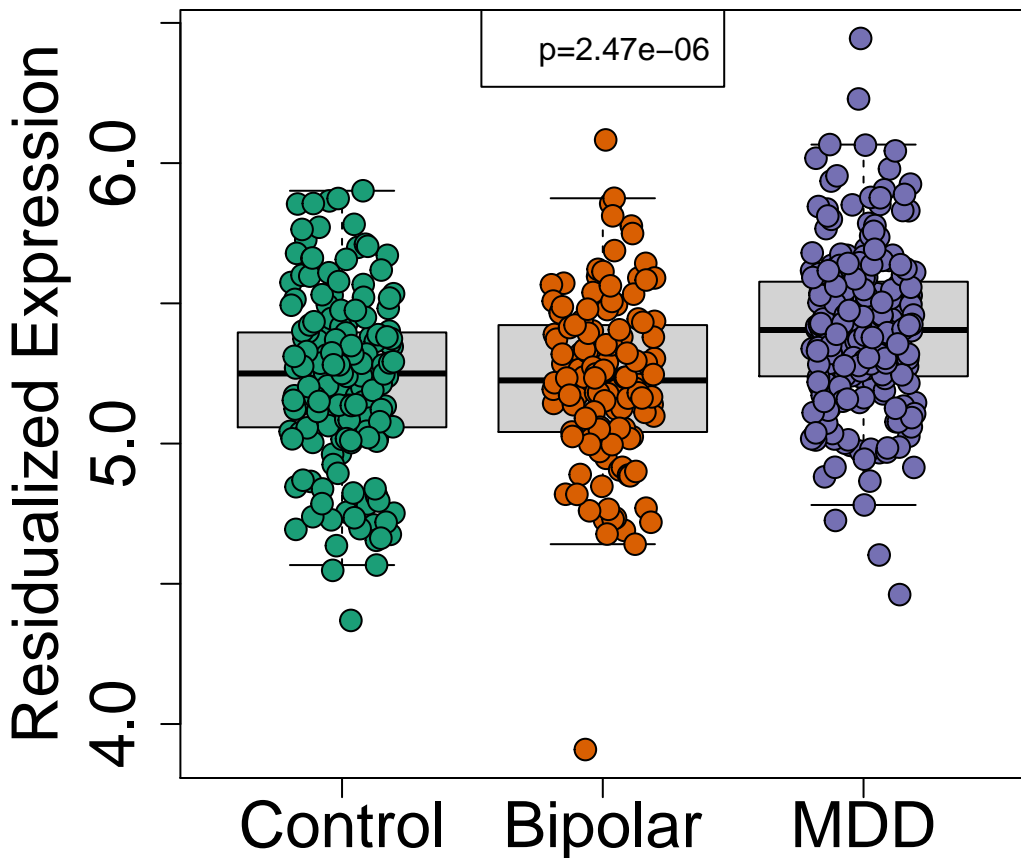
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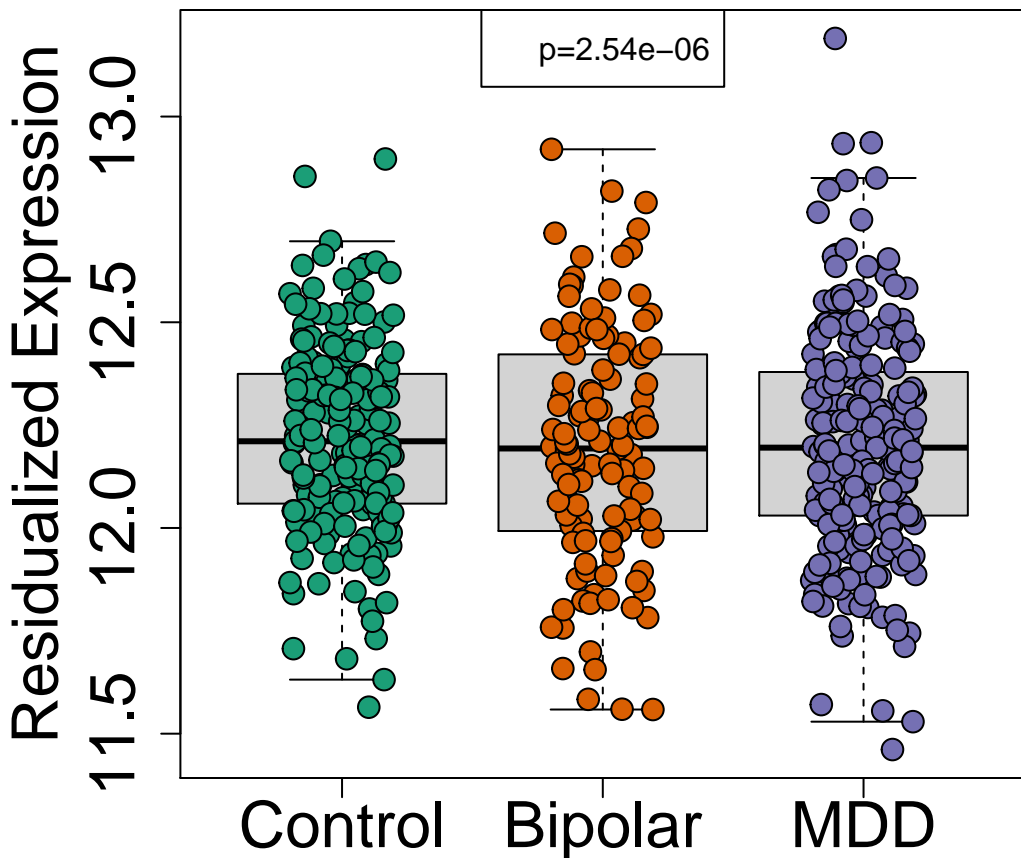
HDAC11, InGen

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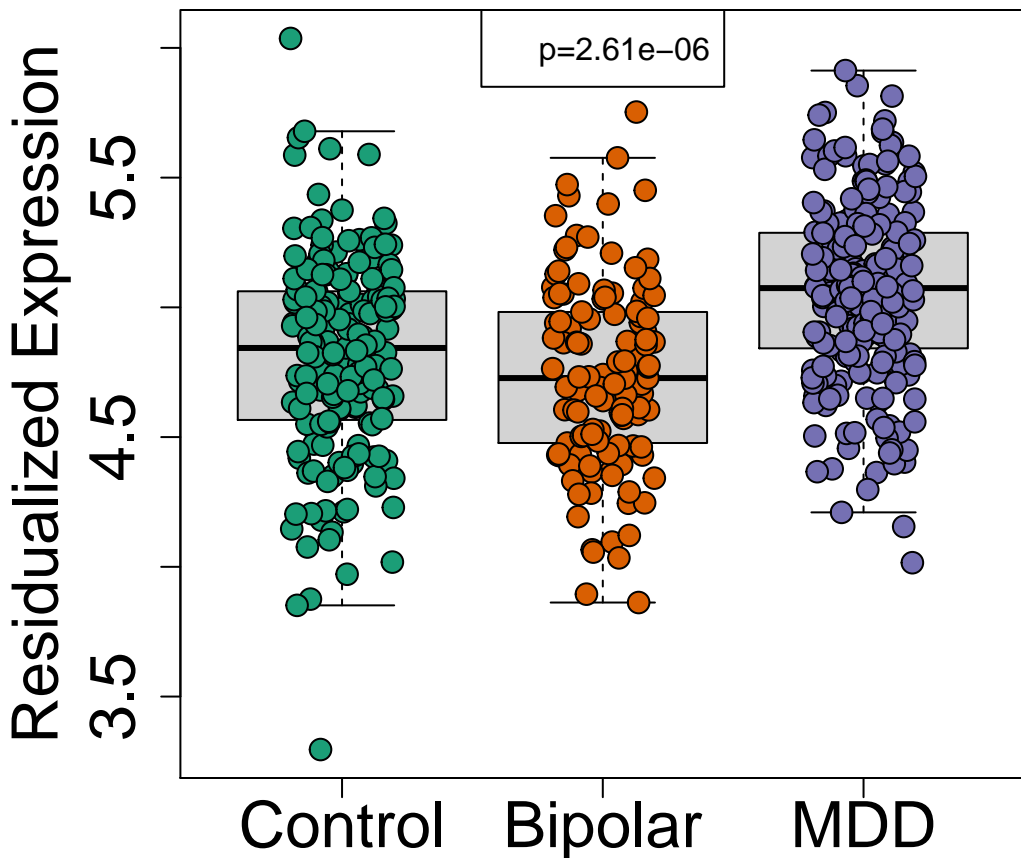
GLUL, InGen

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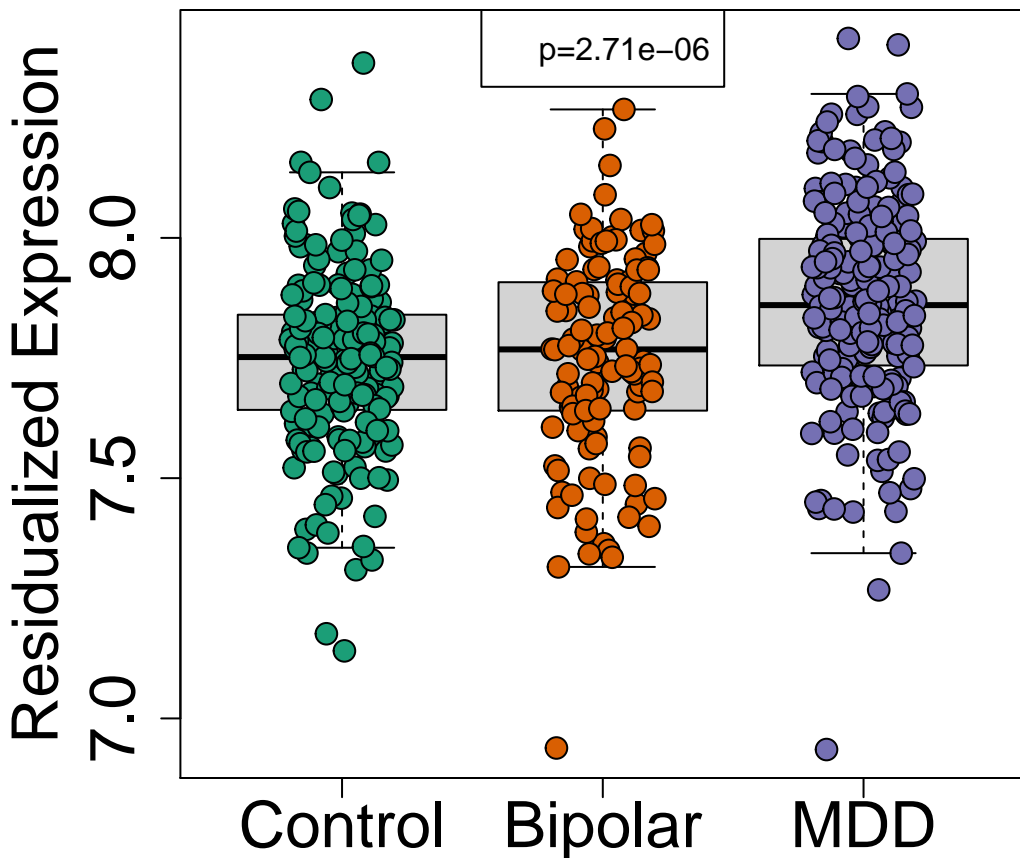
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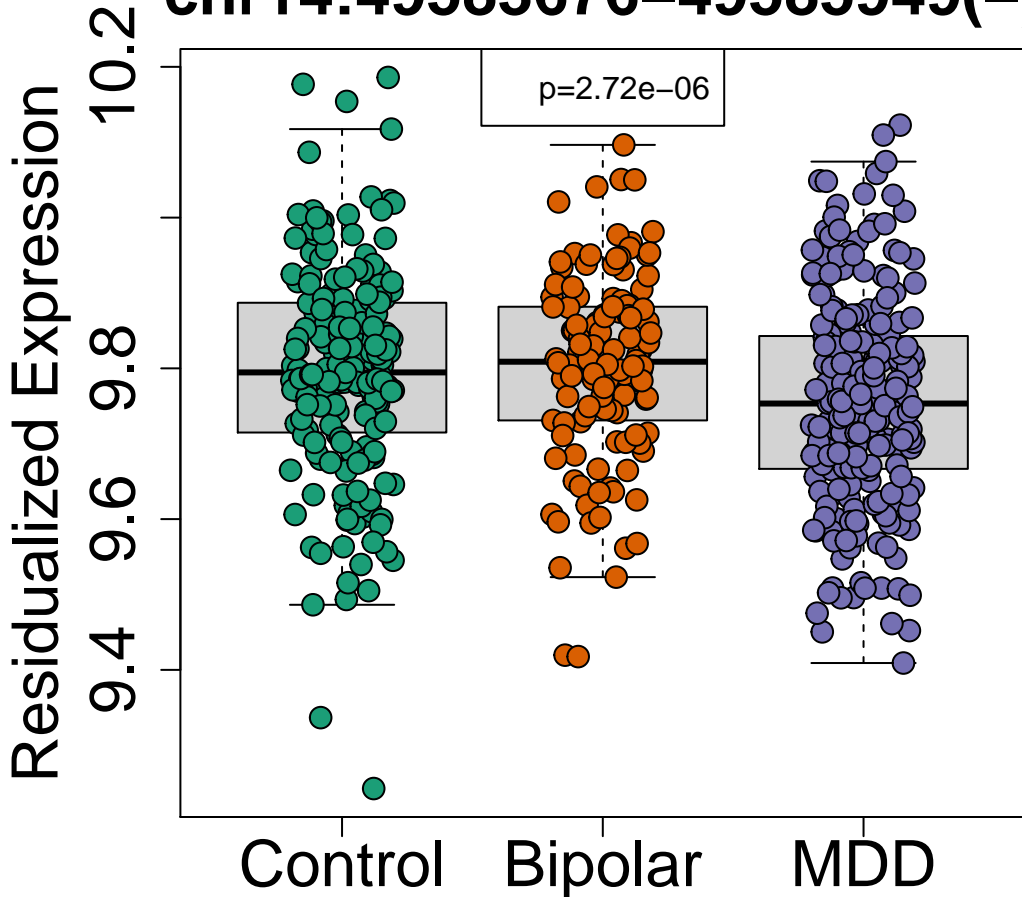
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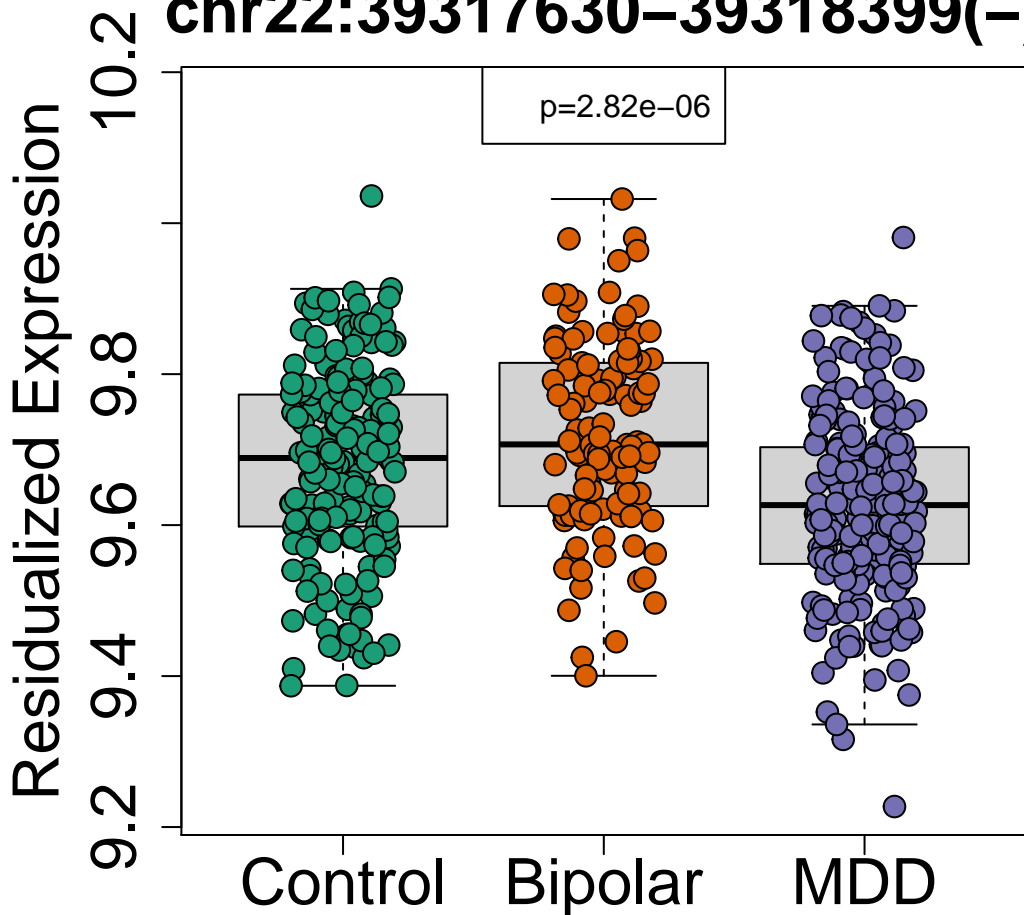
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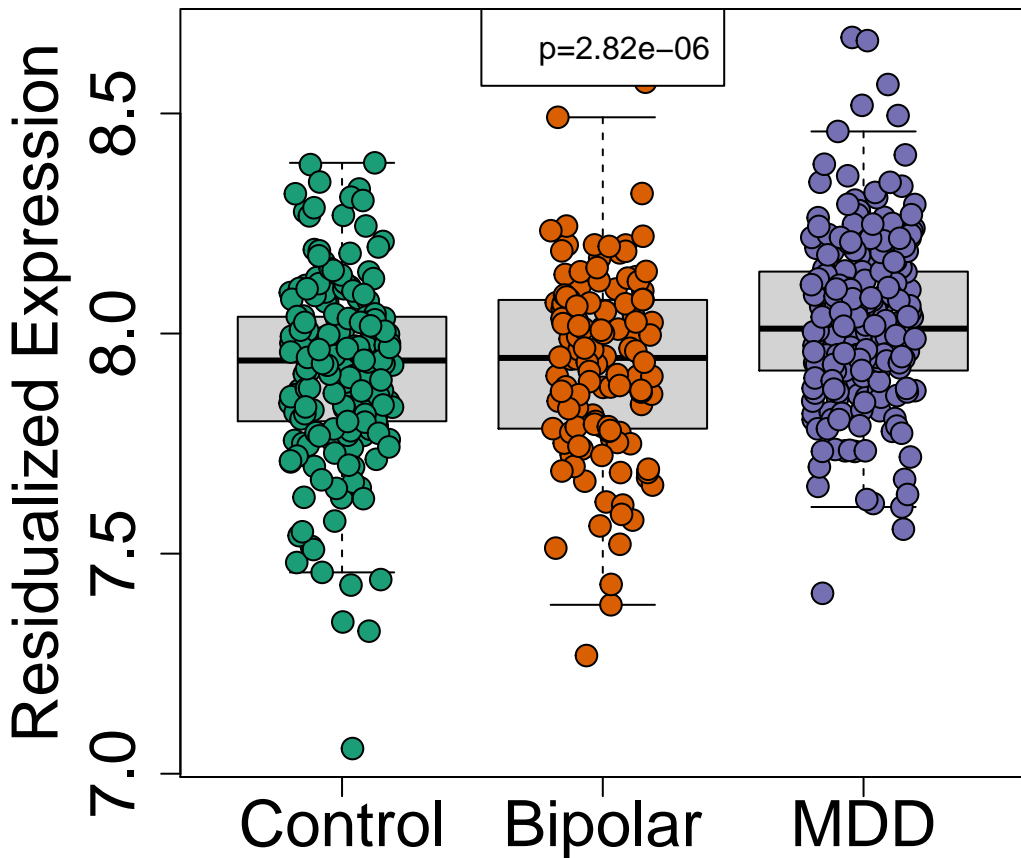
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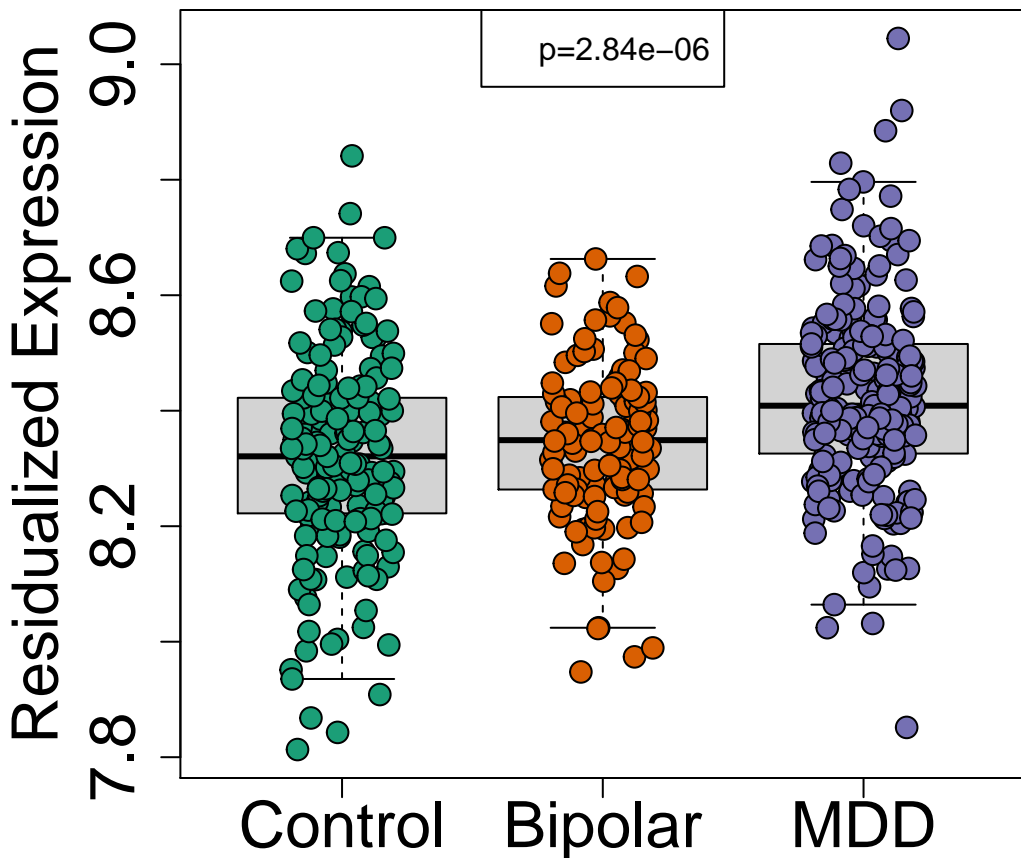
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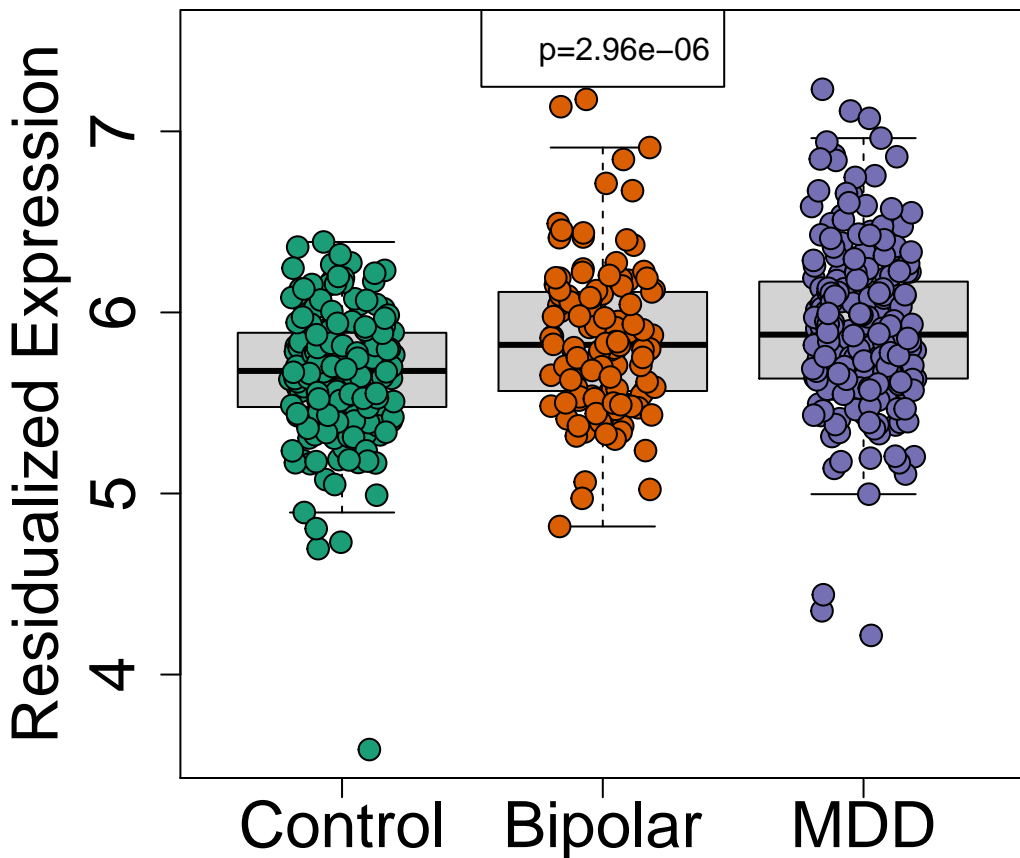
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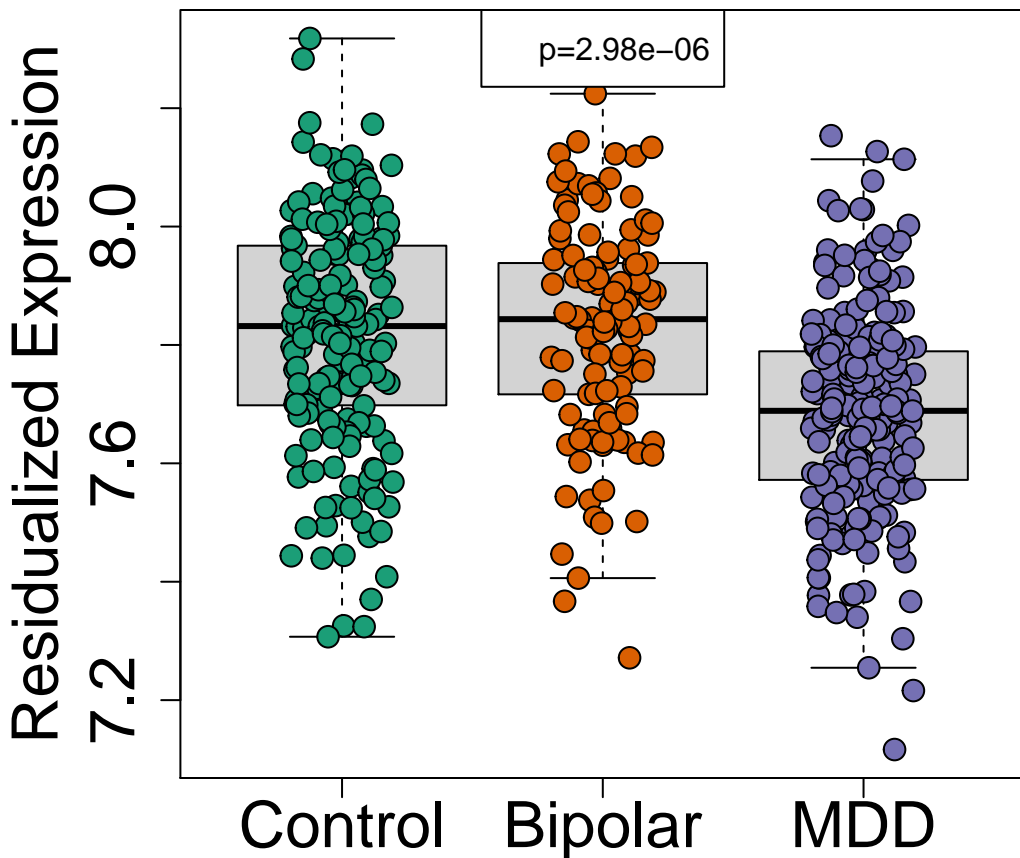
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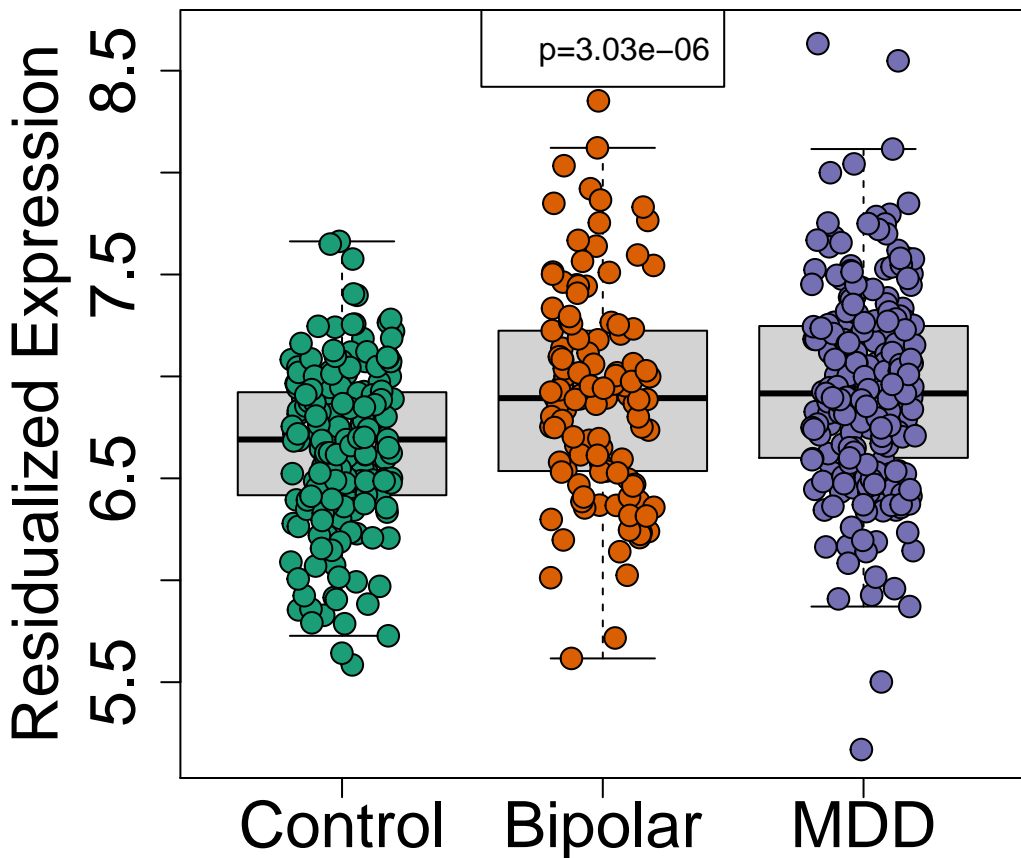
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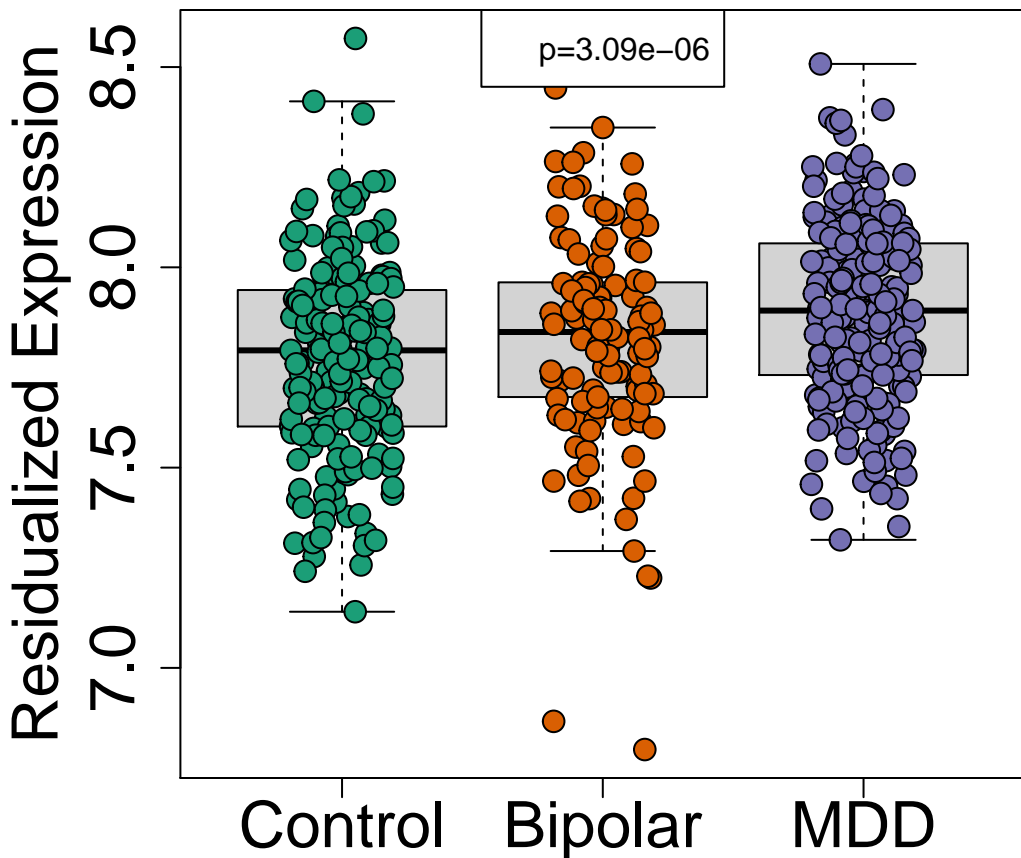
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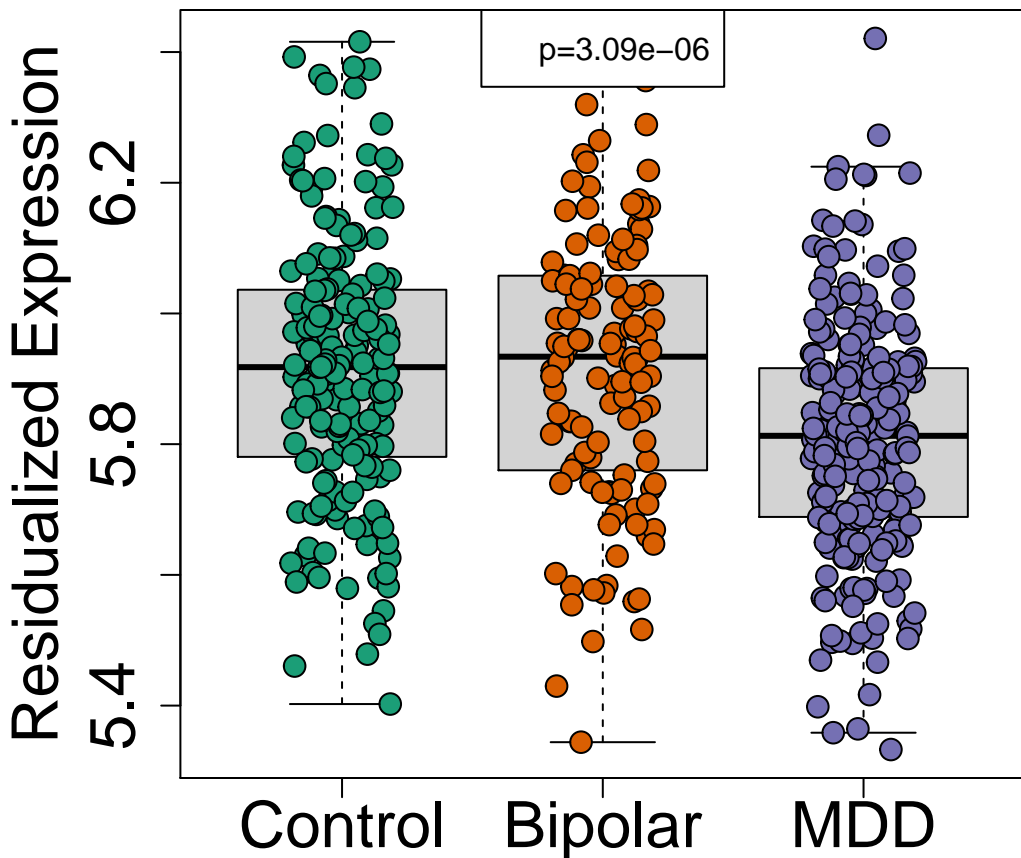
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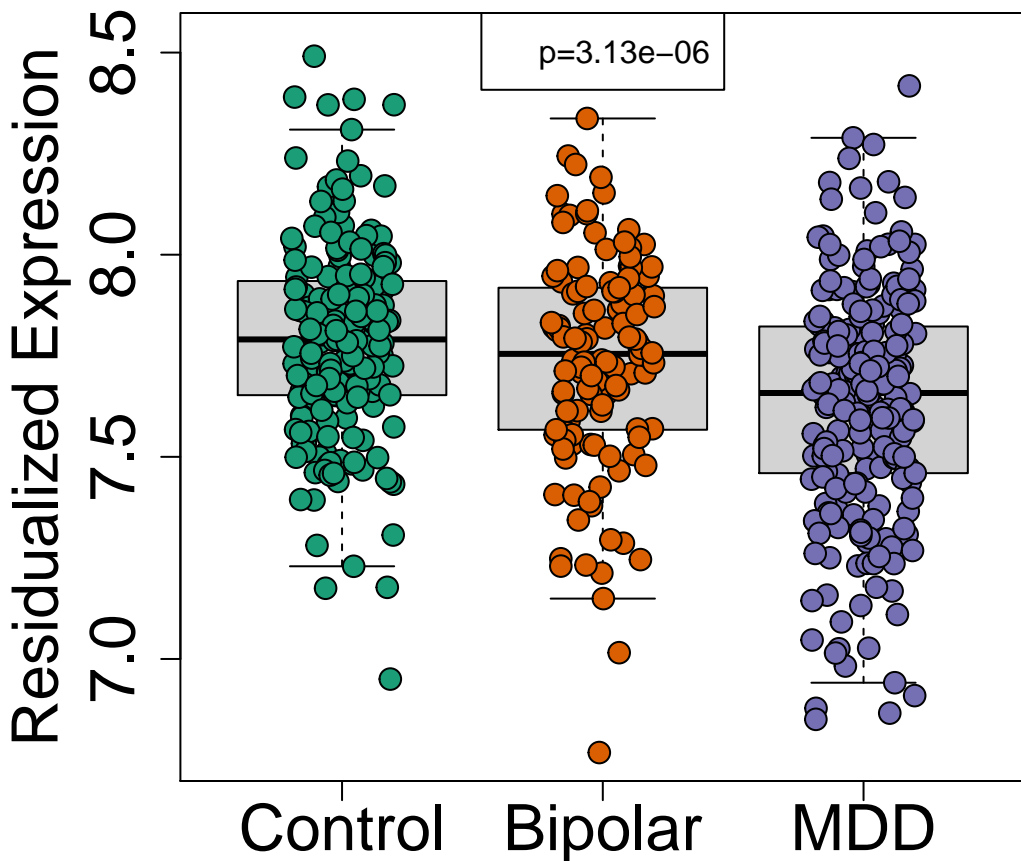
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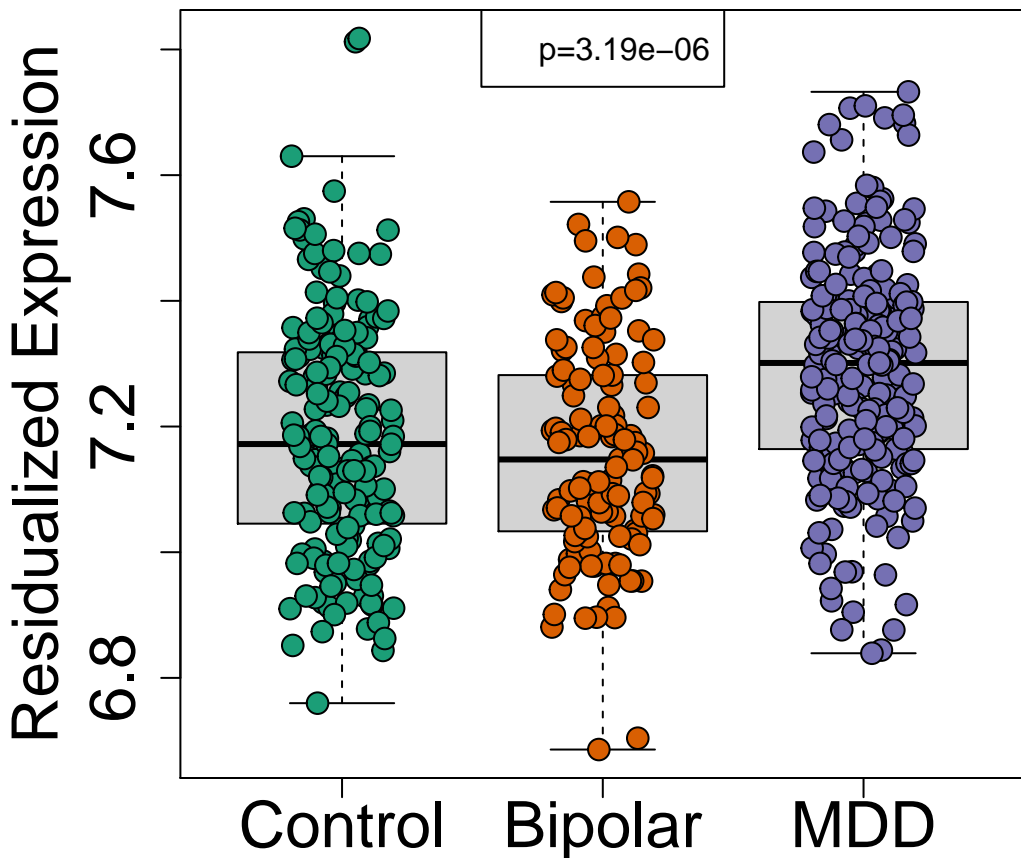
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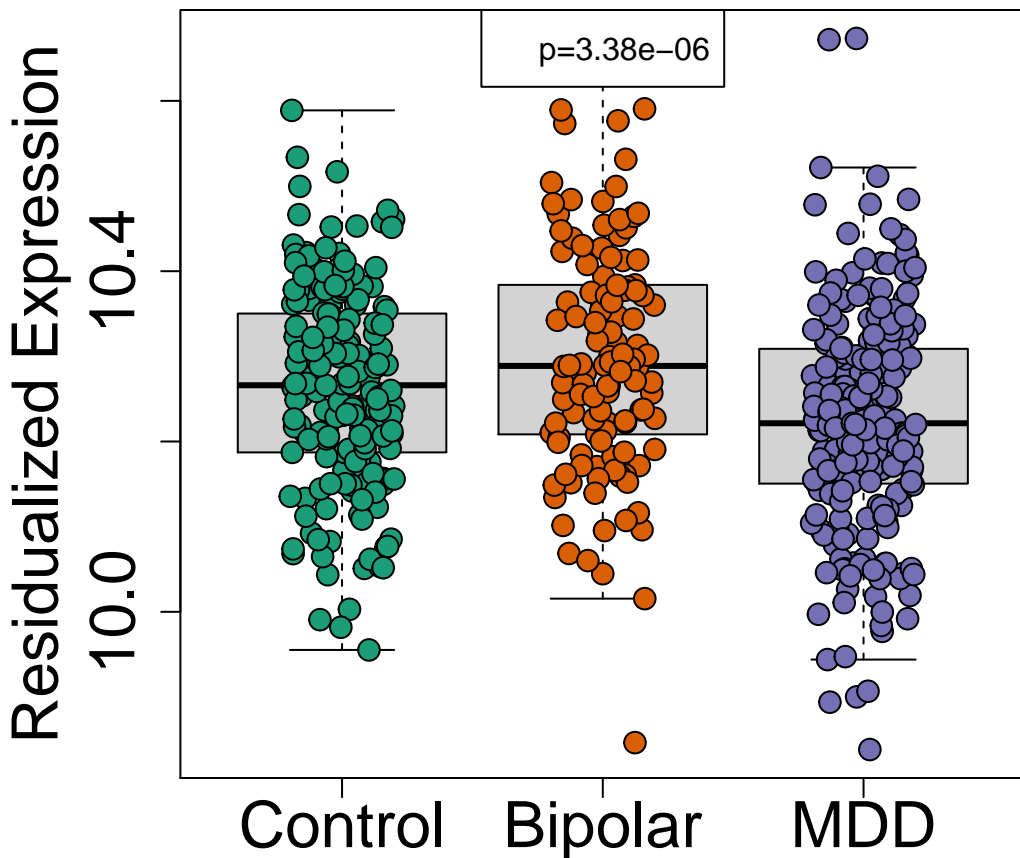
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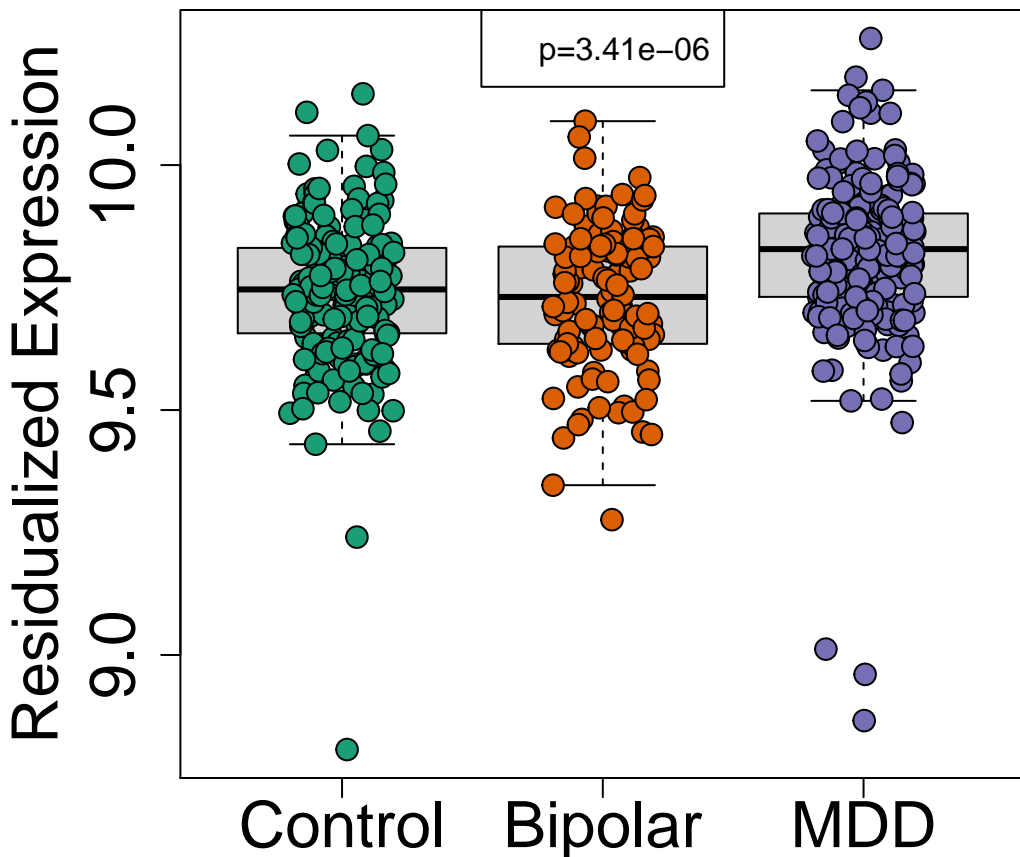
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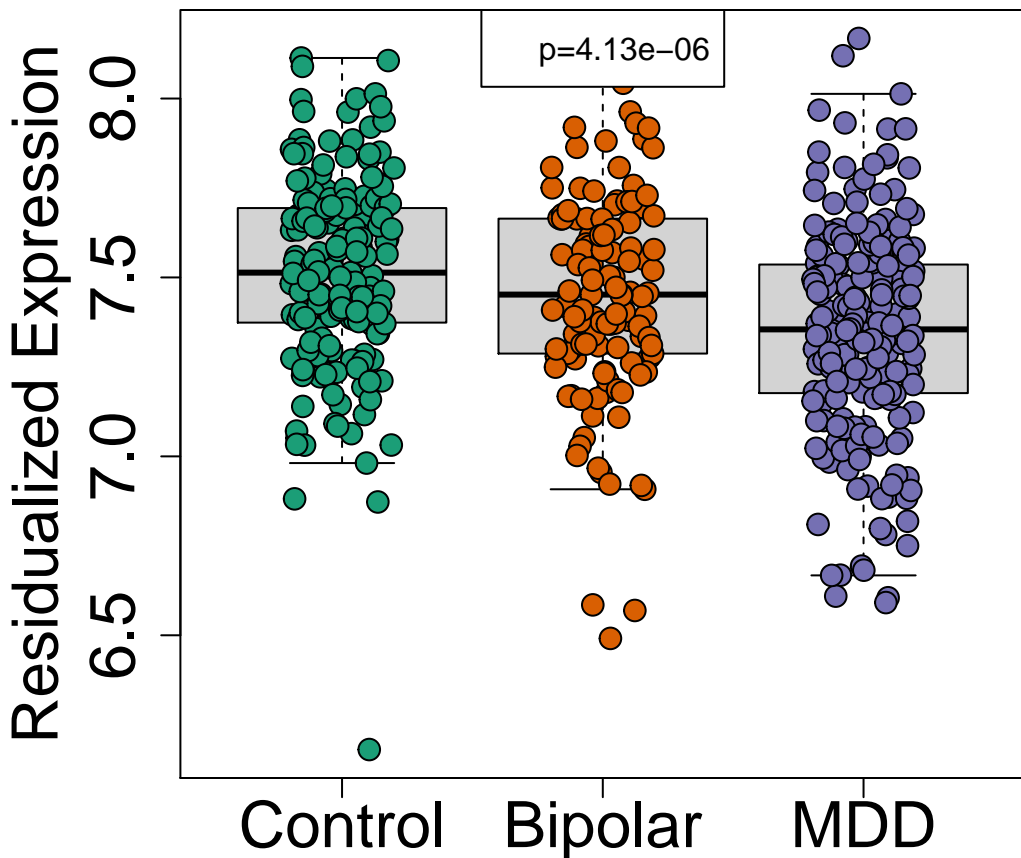
RPL7, InGen

chr8:73292798–73293598(–)



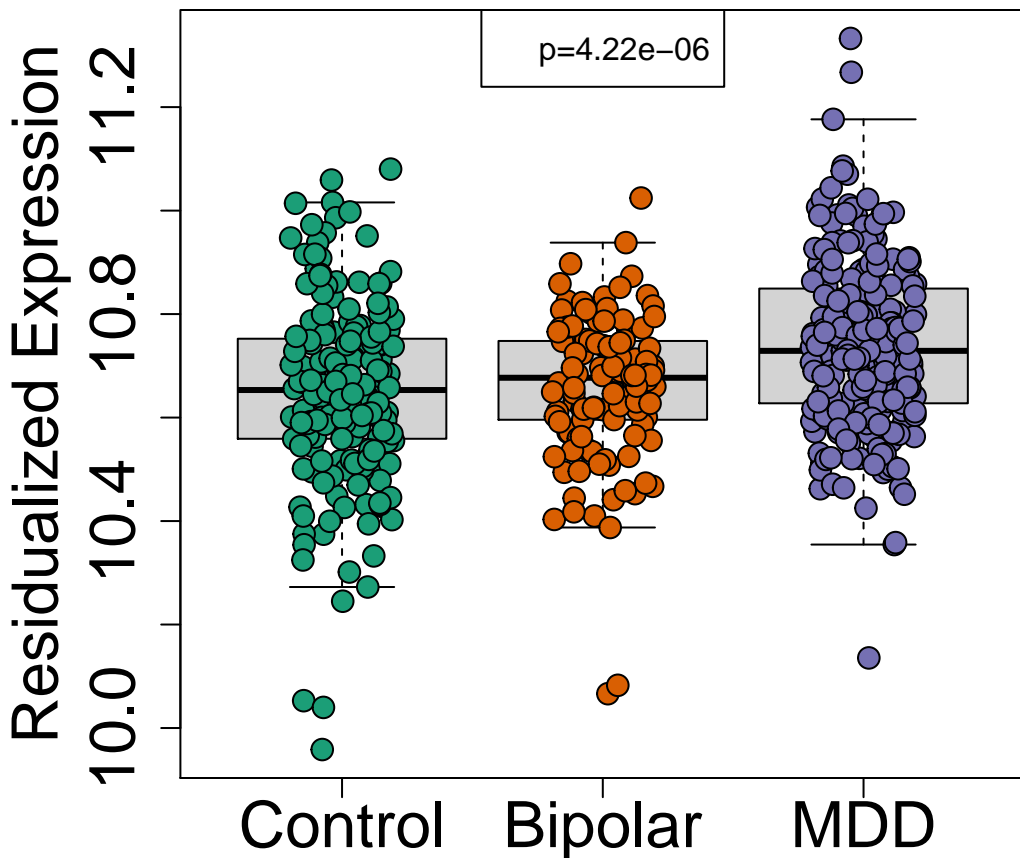
SRSF6, InGen

chr20:43460599–43460702(+)



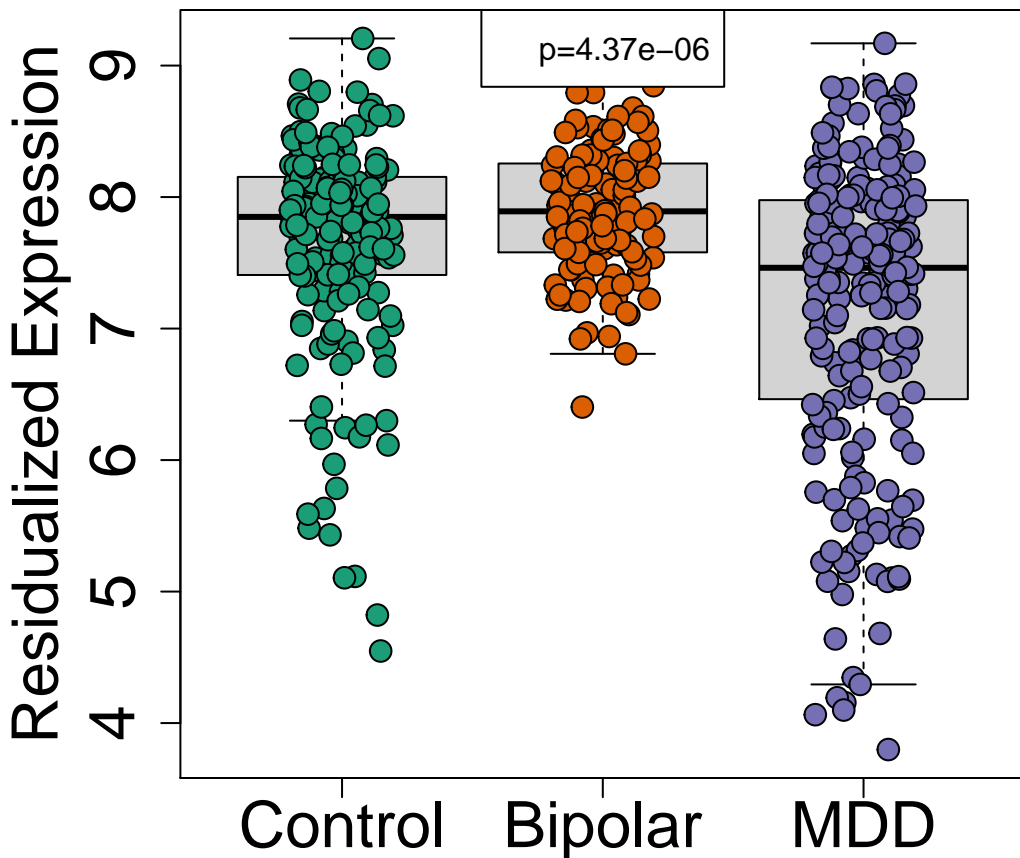
CALM3, InGen

chr19:46601438–46605826(+)



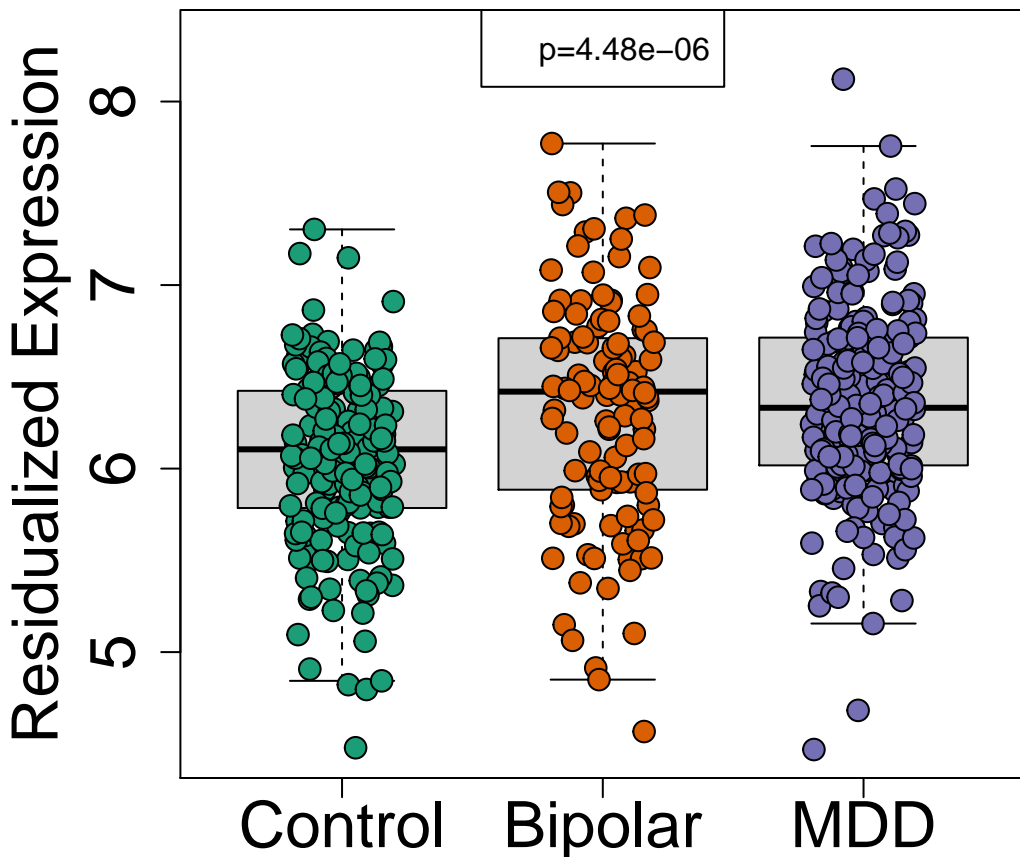
ITGB8, InGen

chr7:20380832–20381726(+)



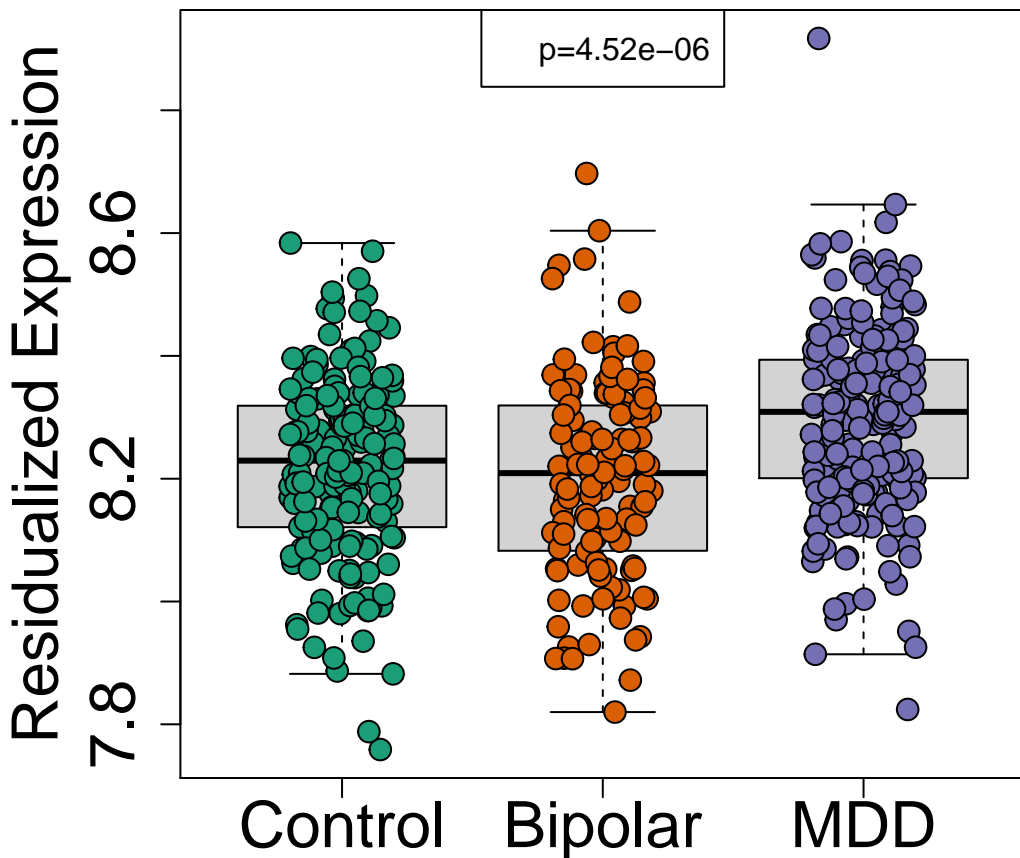
ARRDC3, InGen

chr5:91374277-91374921(-)



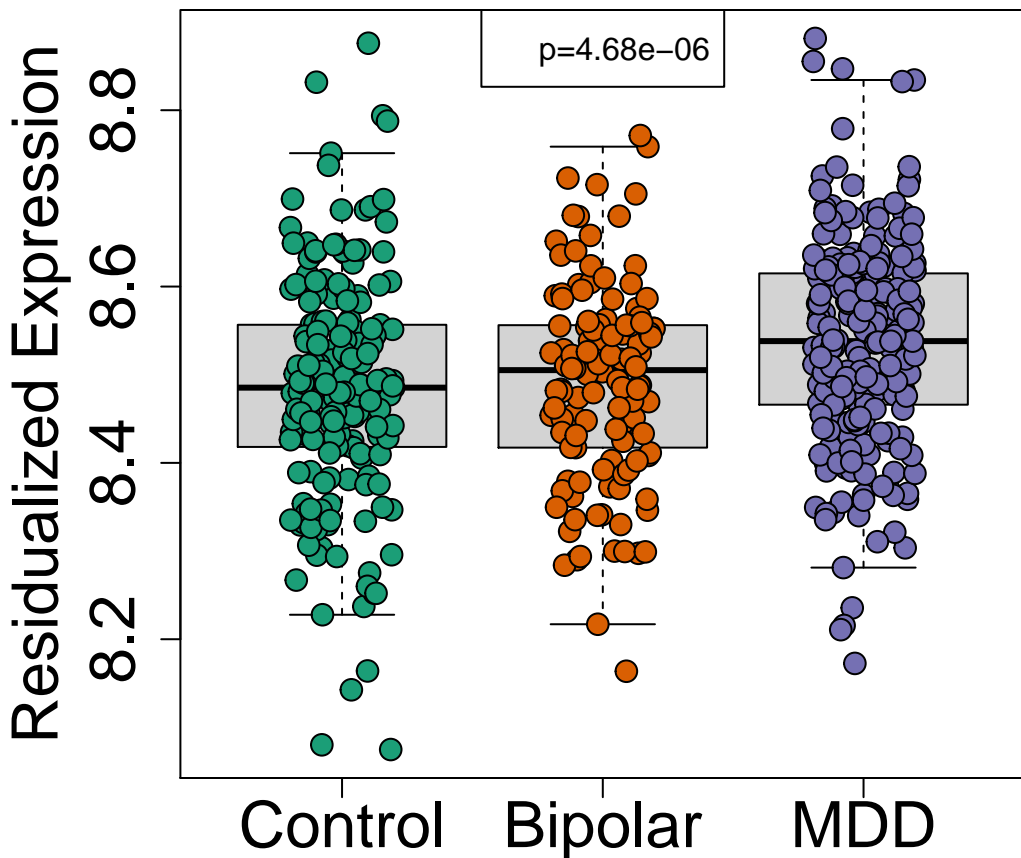
APBB1, InGen

chr11:6402726–6403144(-)



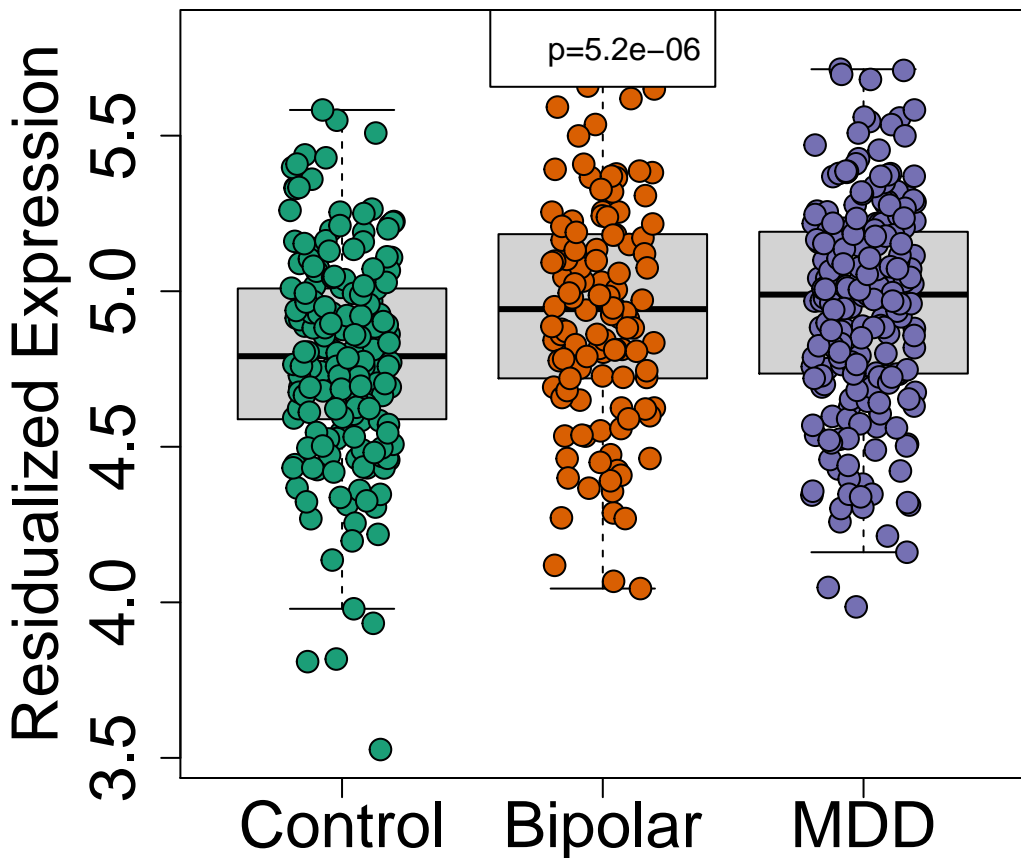
SPTBN1, InGen

chr2:54630030–54630854(+)



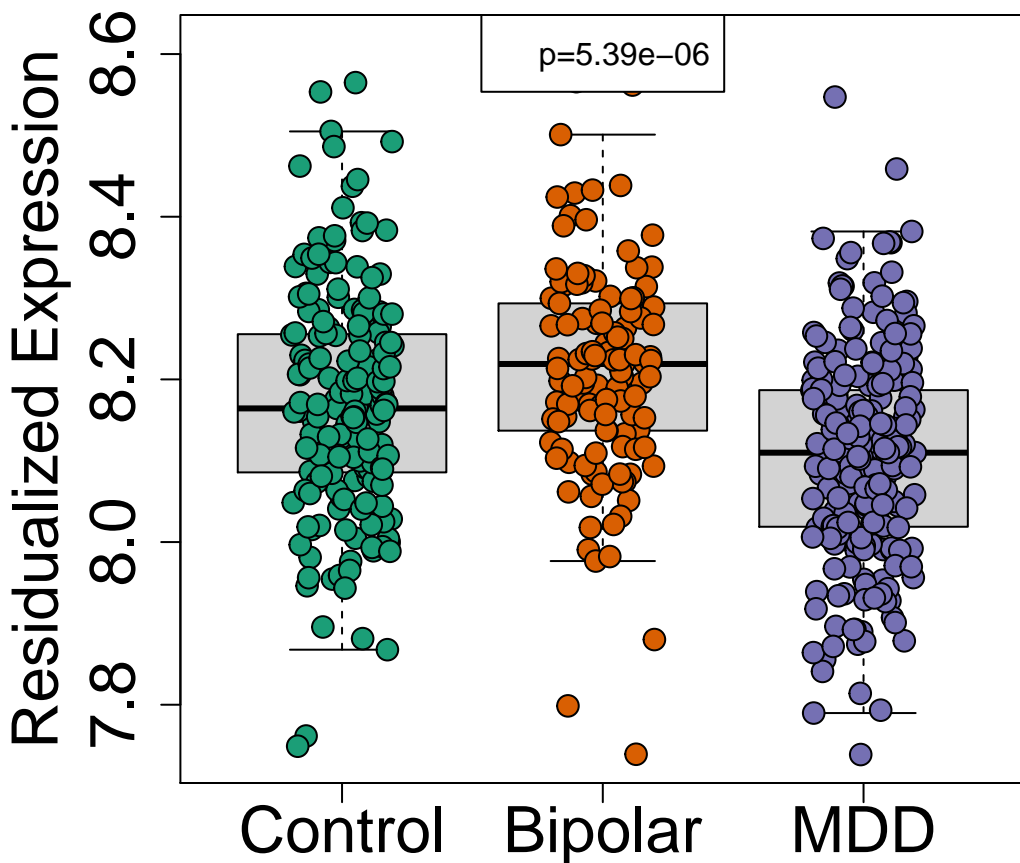
SLC2A1, InGen

chr1:42929315–42929592(-)



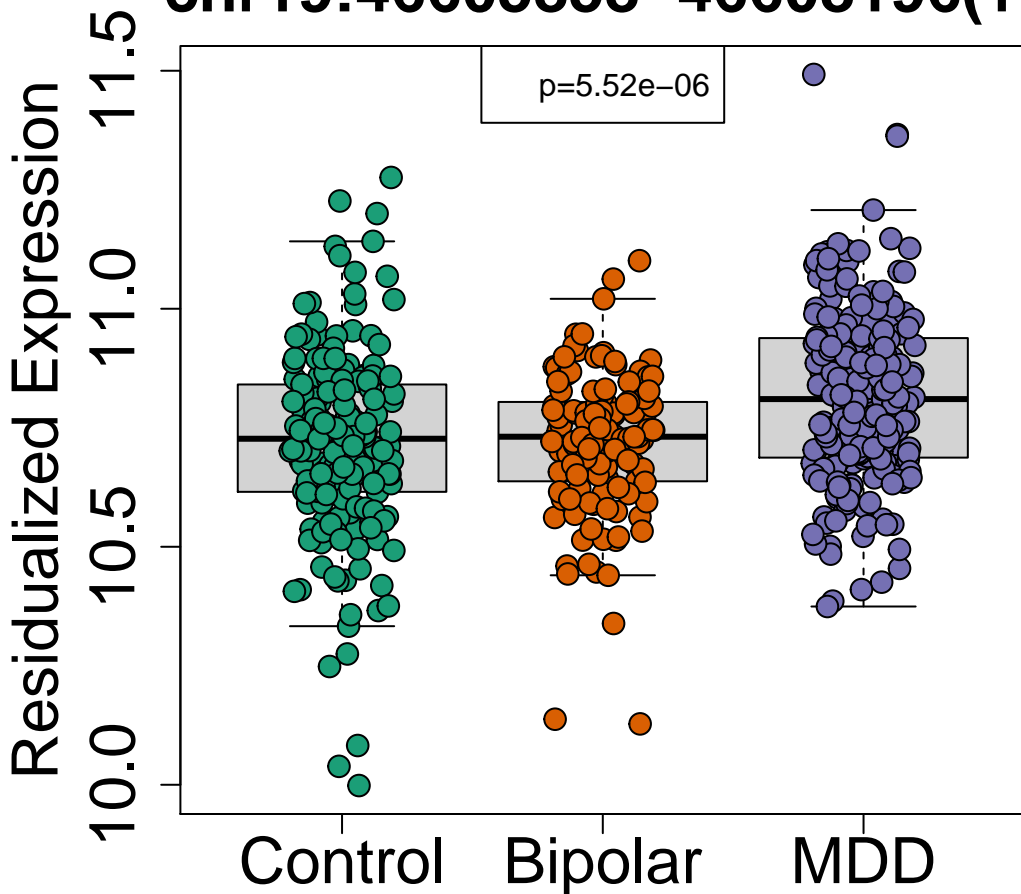
TCEB2, InGen

chr16:2772103–2775450(-)



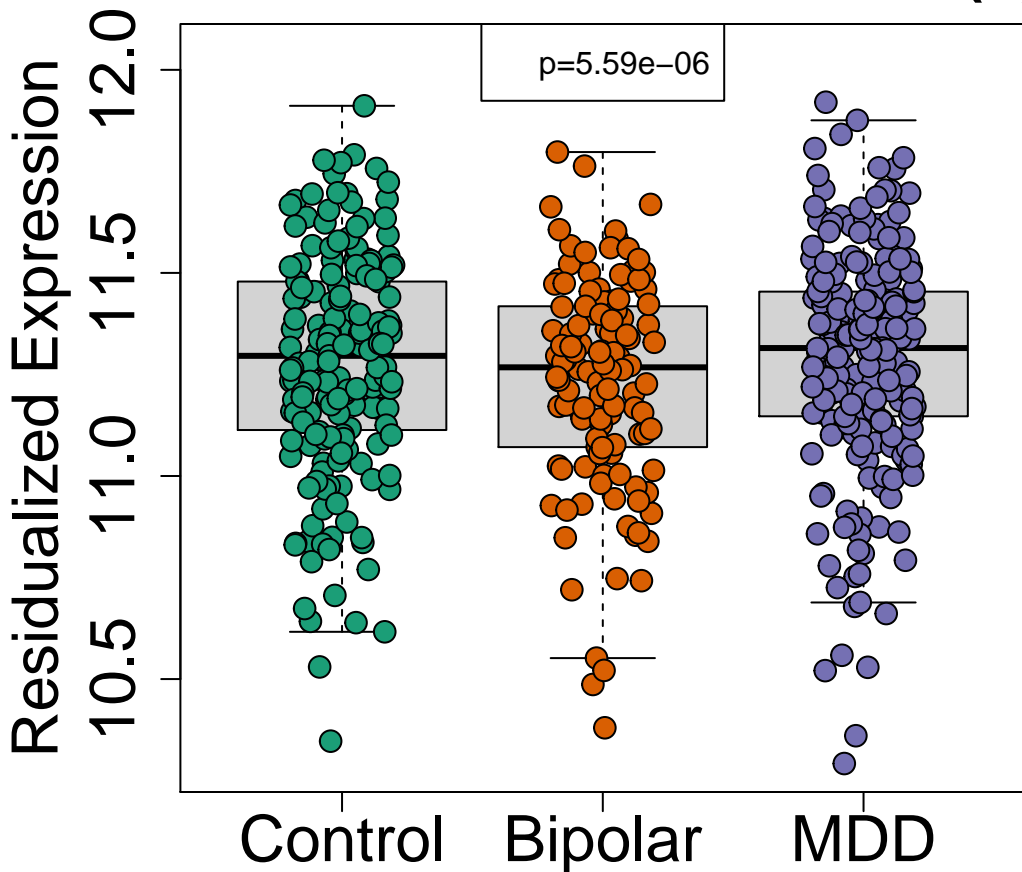
CALM3, InGen

chr19:46605858–46608196(+)



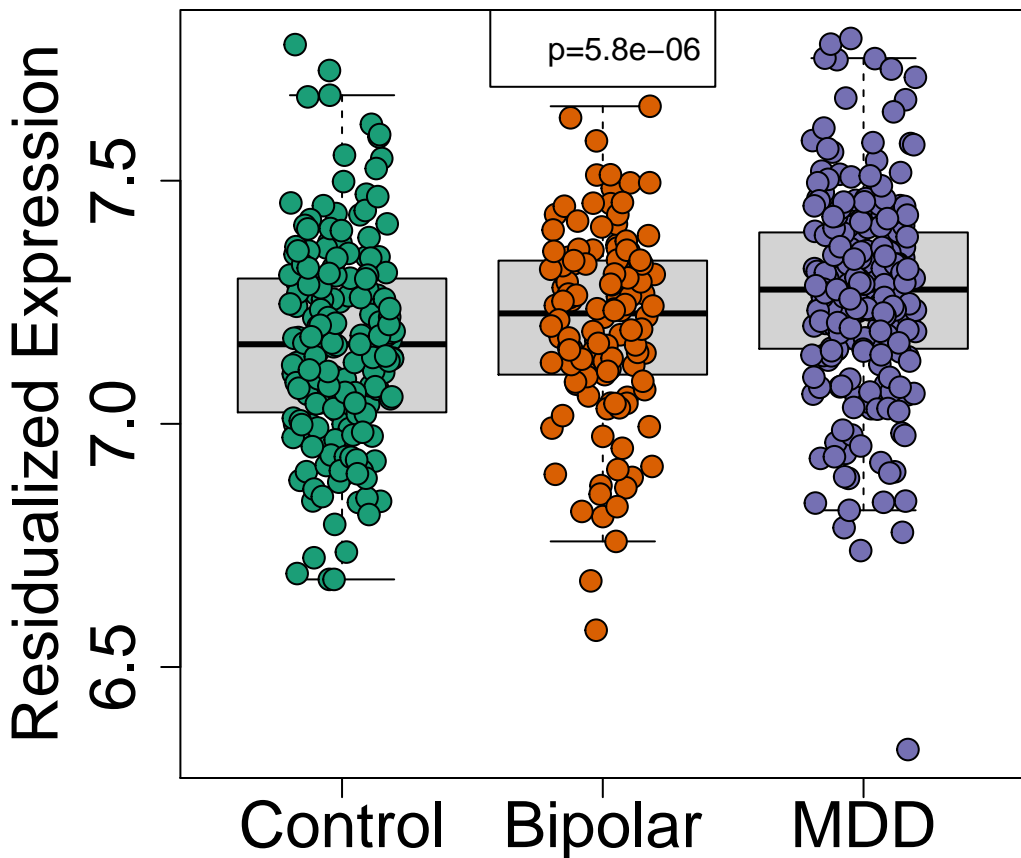
SLC1A2, InGen

chr11:35292521–35301518(-)



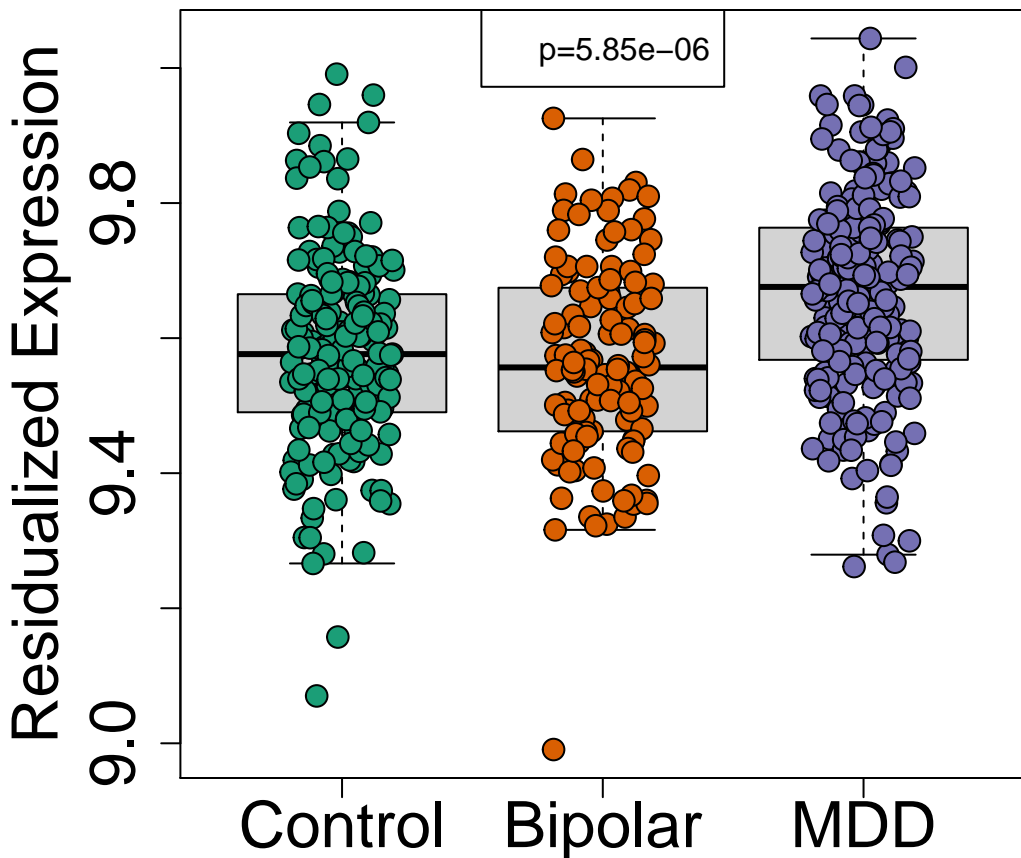
TPM3, InGen

chr1:154176249–154182987(-)



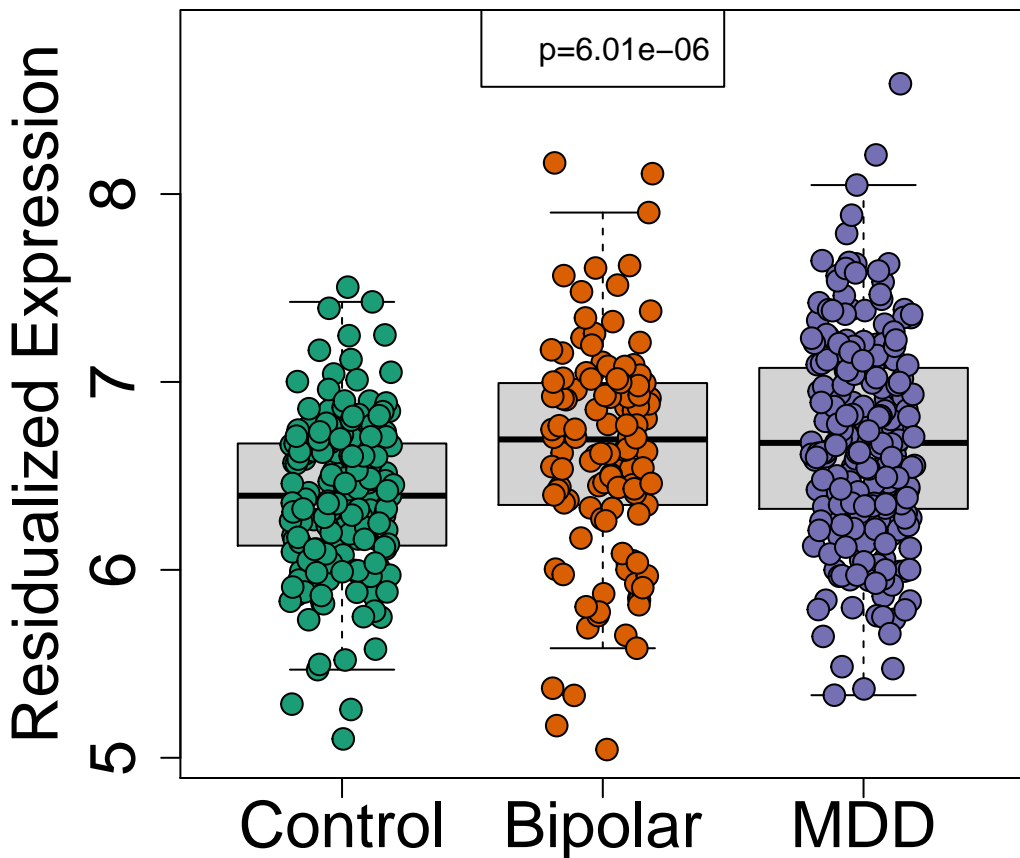
ENO1, InGen

chr1:8865483–8866278(-)



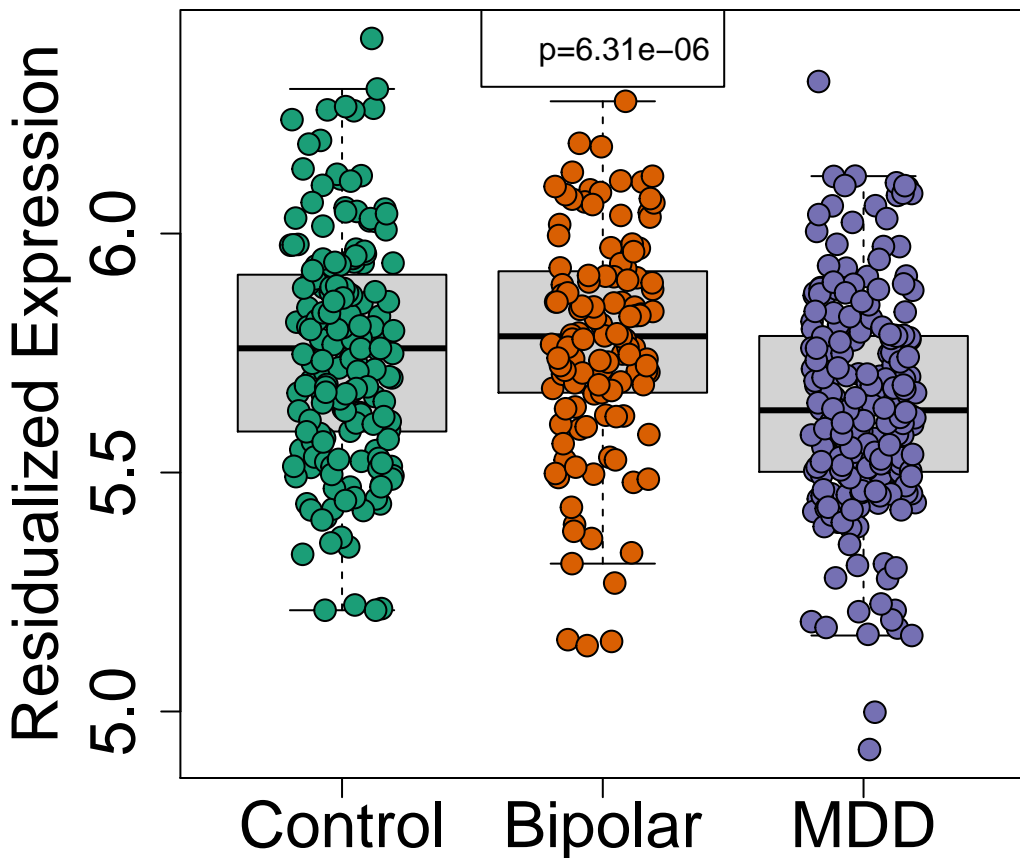
ARRDC3, InGen

chr5:91376769–91378693(–)



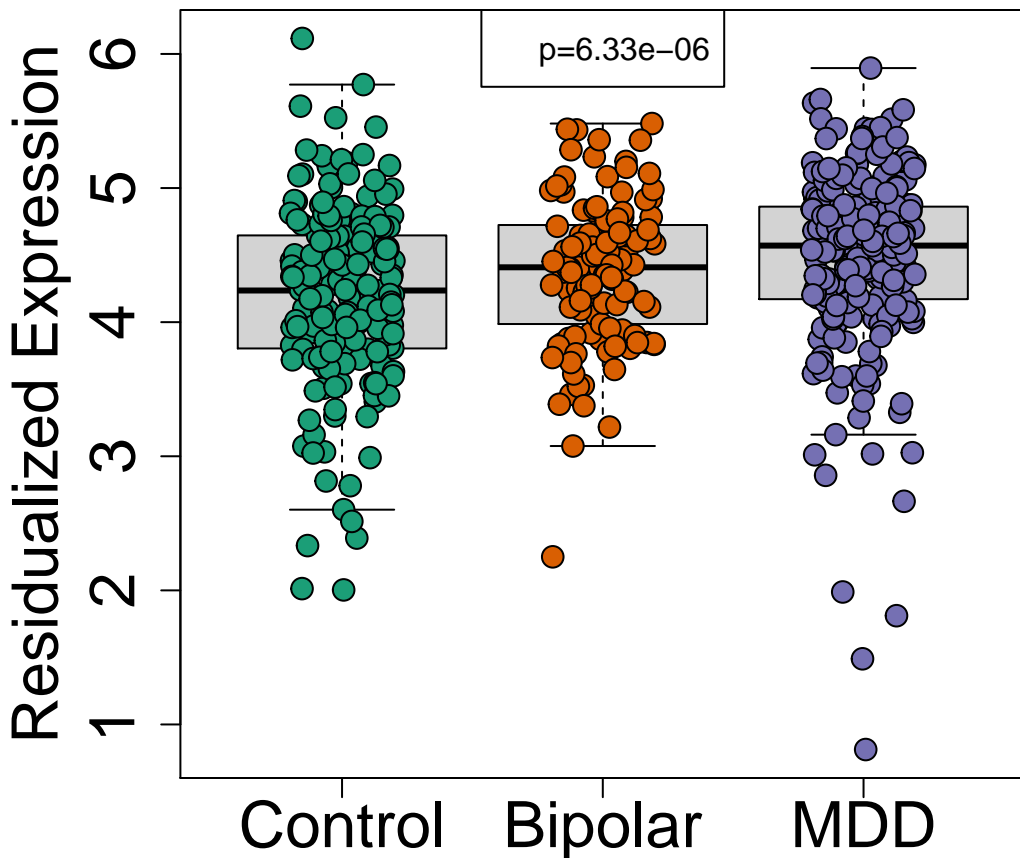
LRP1, InGen

chr12:57196278–57196981(+)



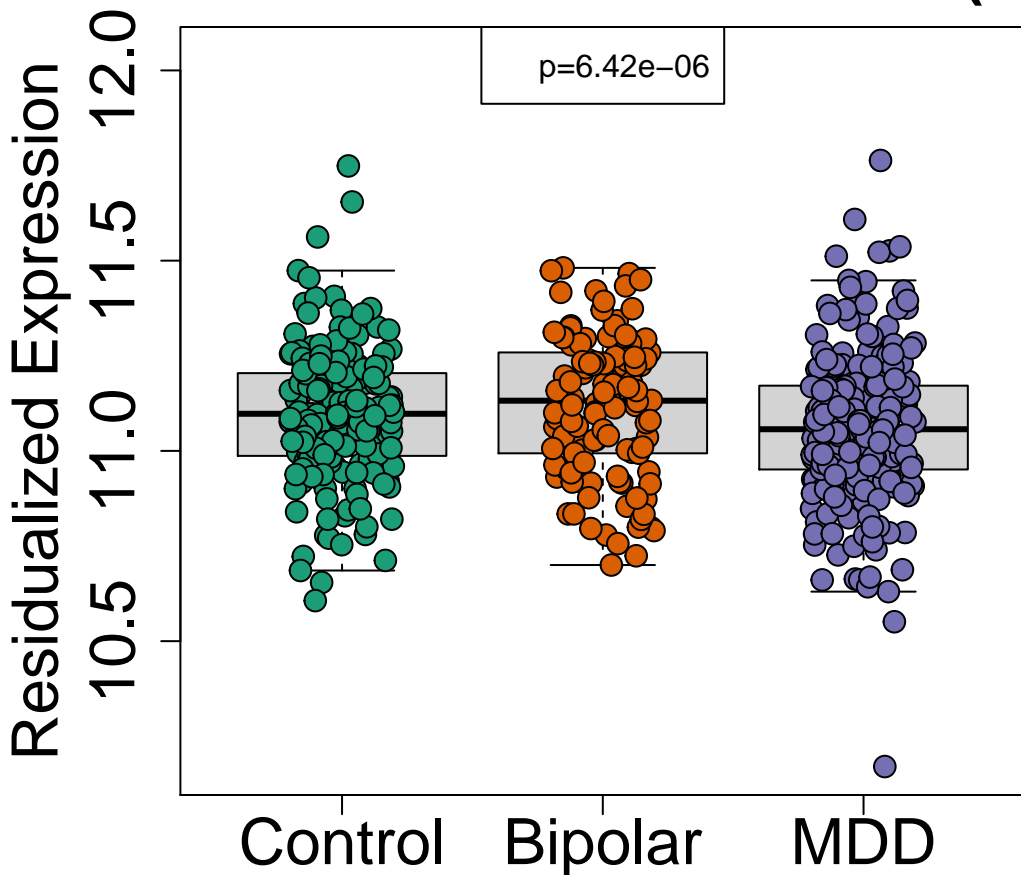
KRBOX4, ExonSkip

chrX:46471186–46472749(+)



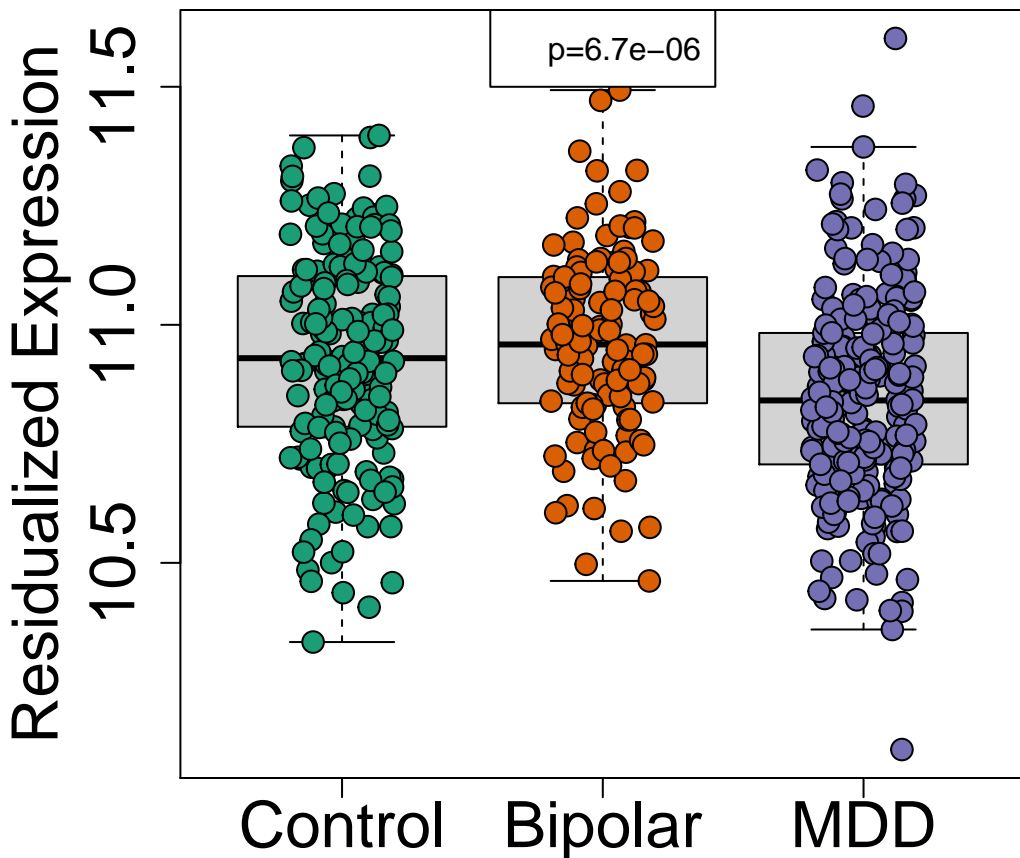
GNAS, InGen

chr20:58905481–58909161(+)



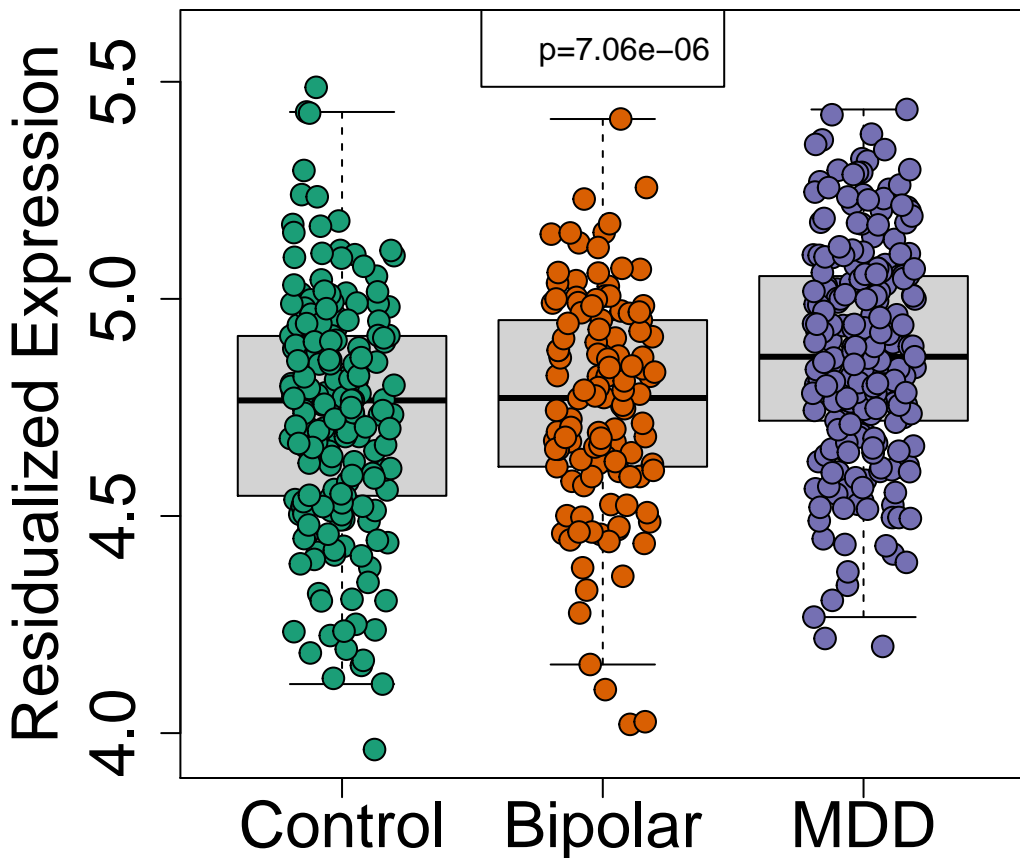
MTURN, InGen

chr7:30146300–30157437(+)



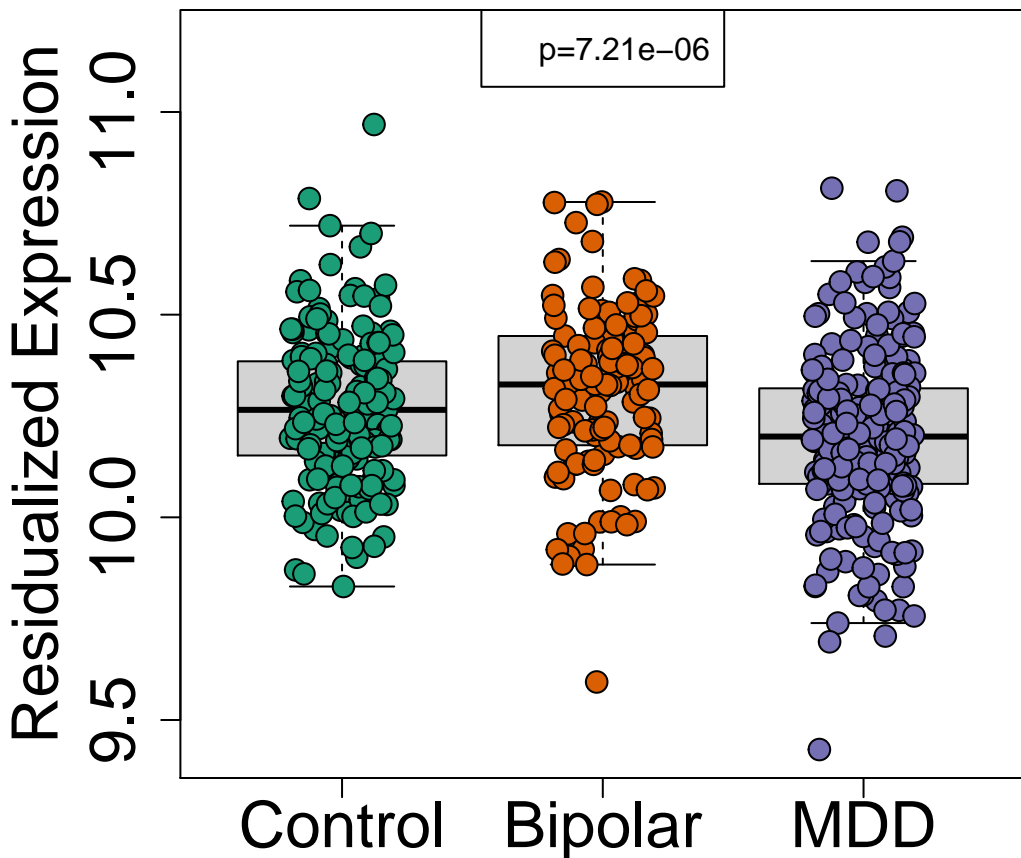
CSNK1G2, InGen

chr19:1978751–1978858(+)



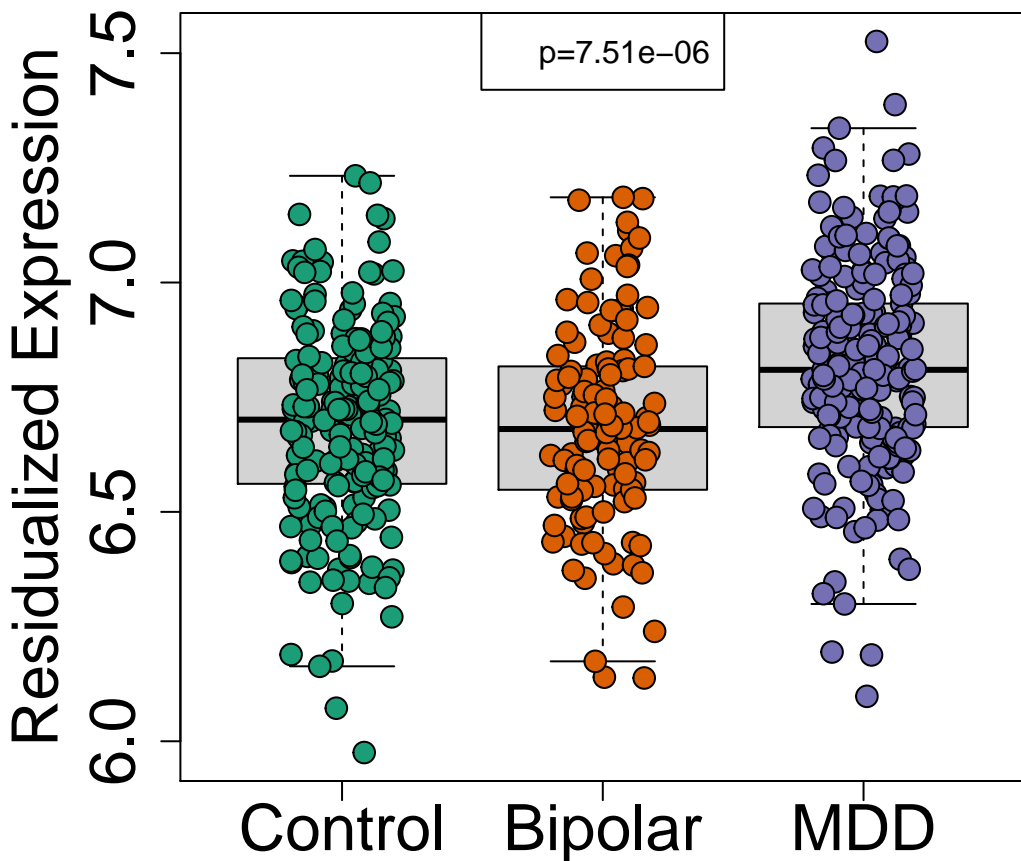
GNAS, InGen

chr20:58909217–58909349(+)



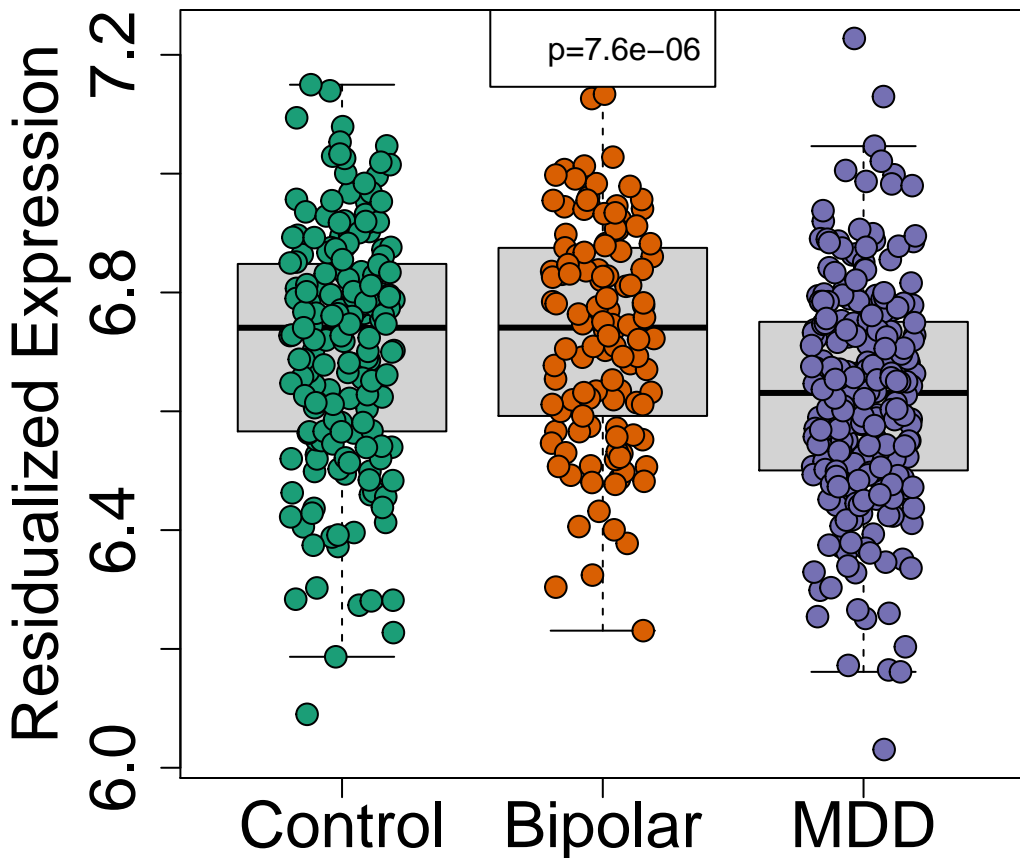
GDE1, InGen

chr16:19503618–19504880(-)



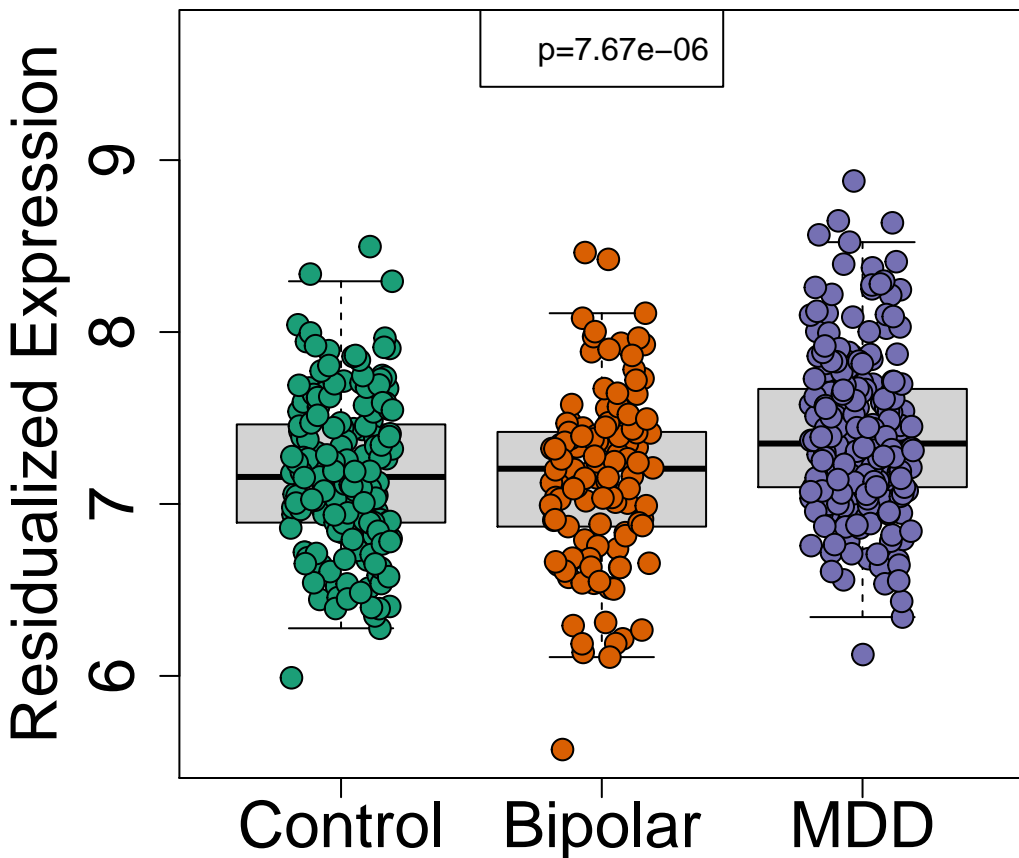
LRP1, InGen

chr12:57198671–57199211(+)



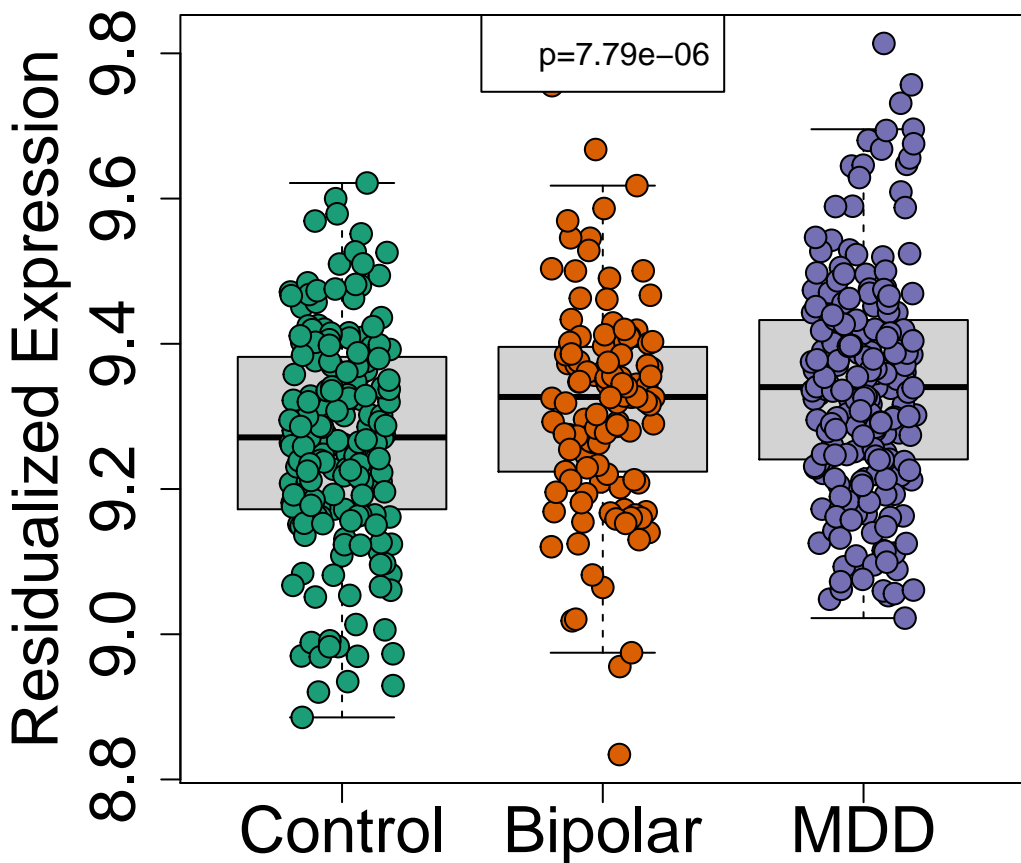
TXNIP, InGen

chr1:145993887–145994015(-)



NEDD8-MDP1, InGen

chr14:24217224-24218132(-)



LRP1, InGen

chr12:57197665–57198155(+)

