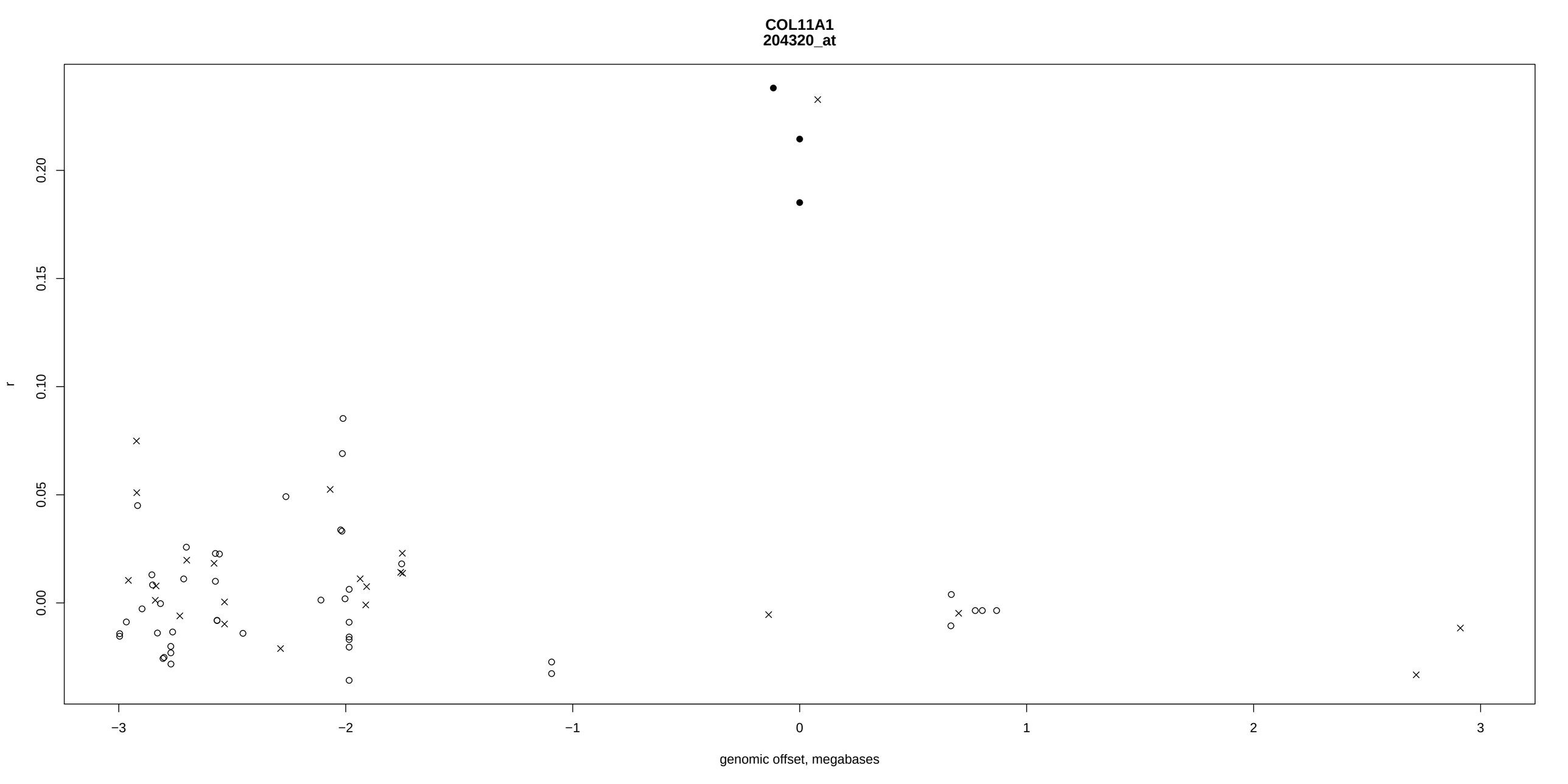
COL11A1
204320_at
r
0.00
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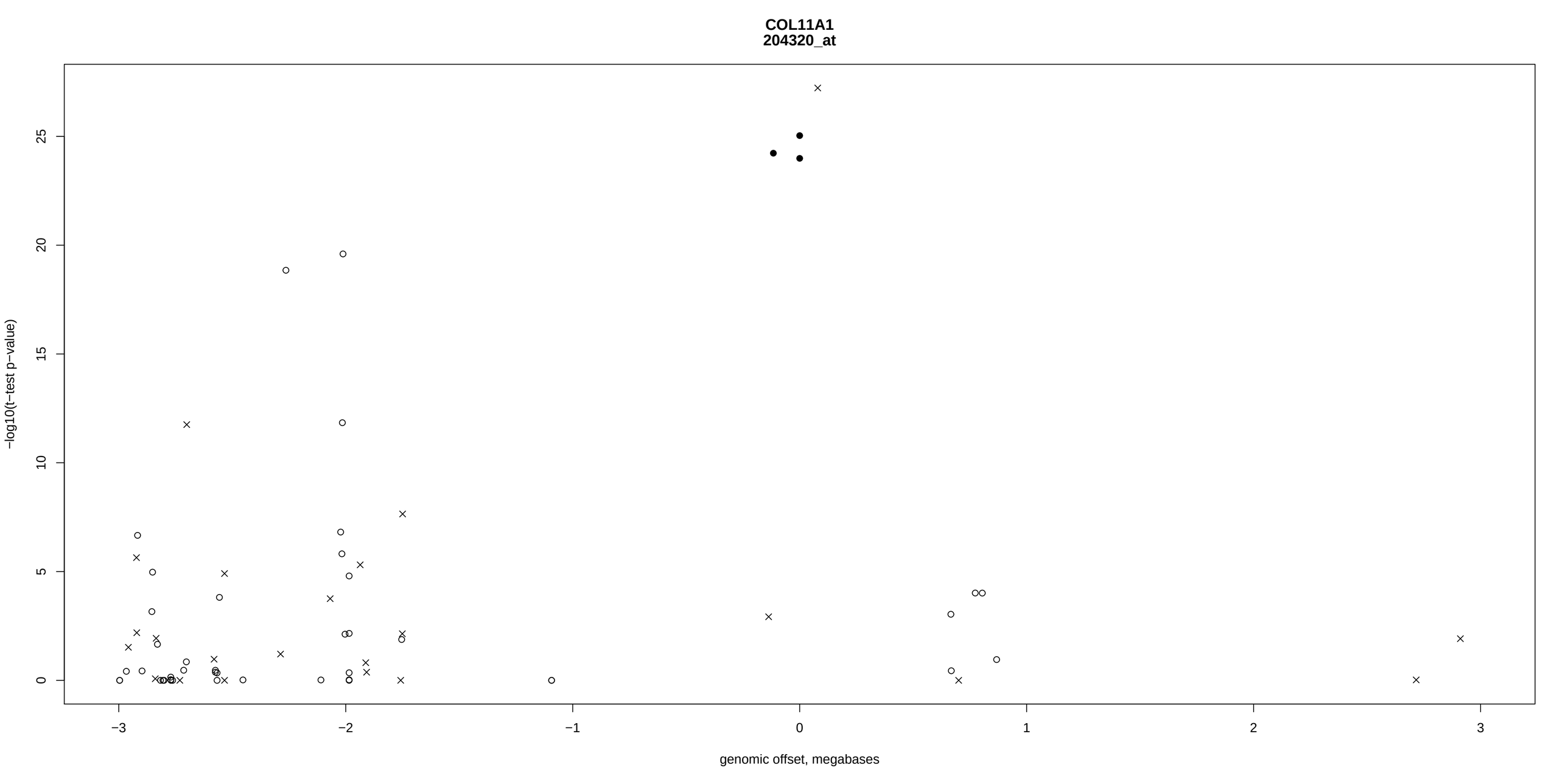
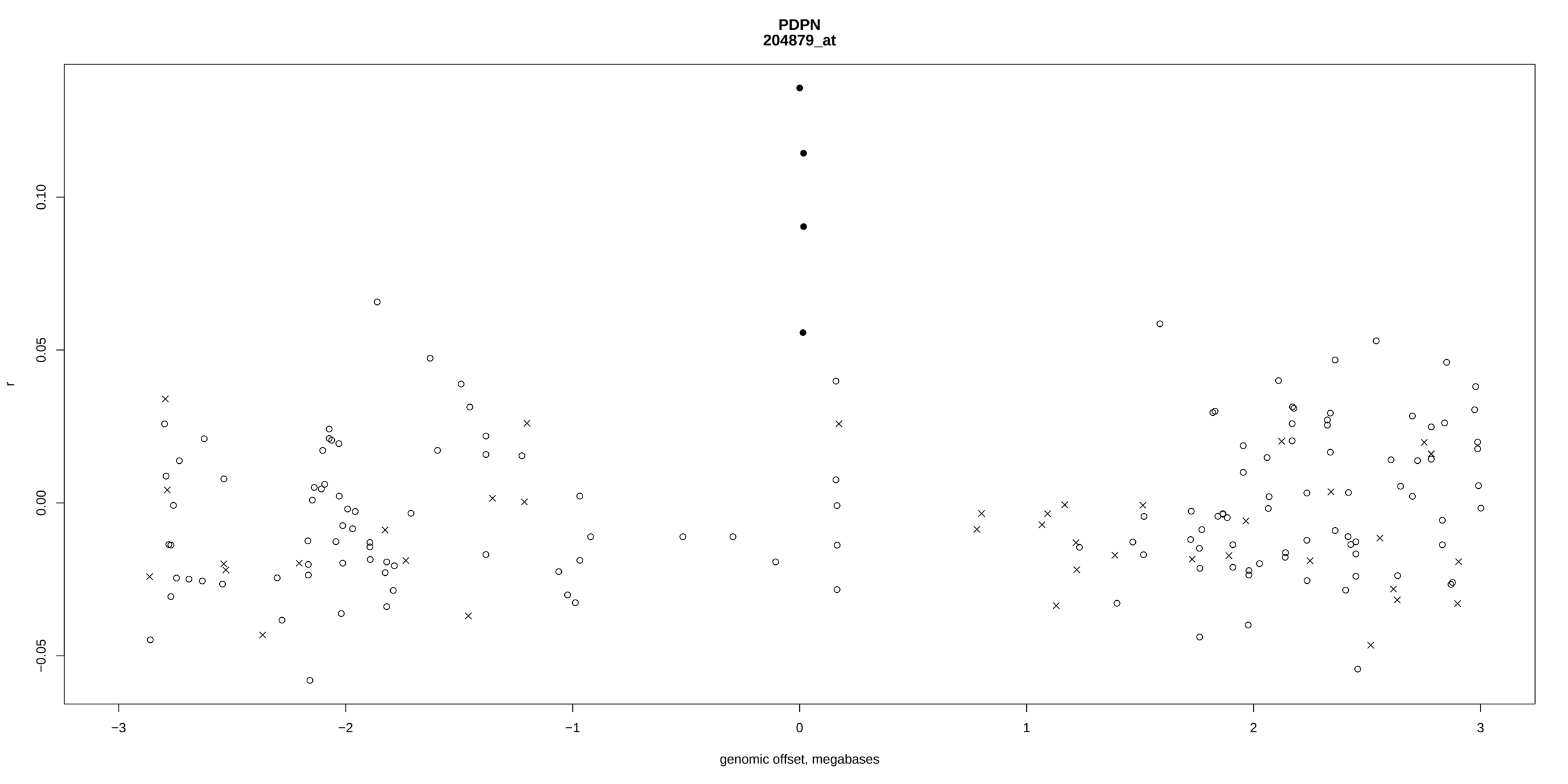
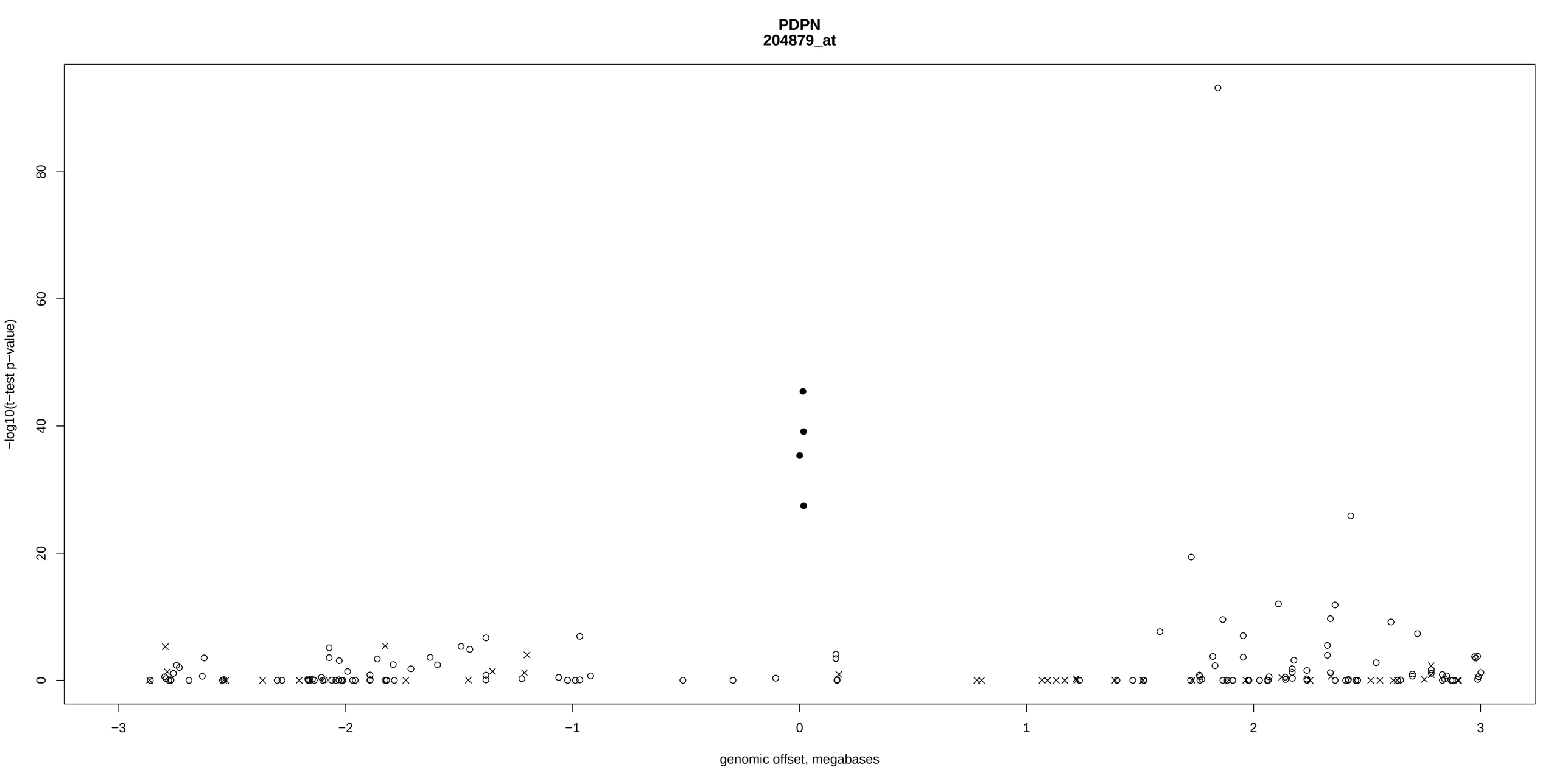
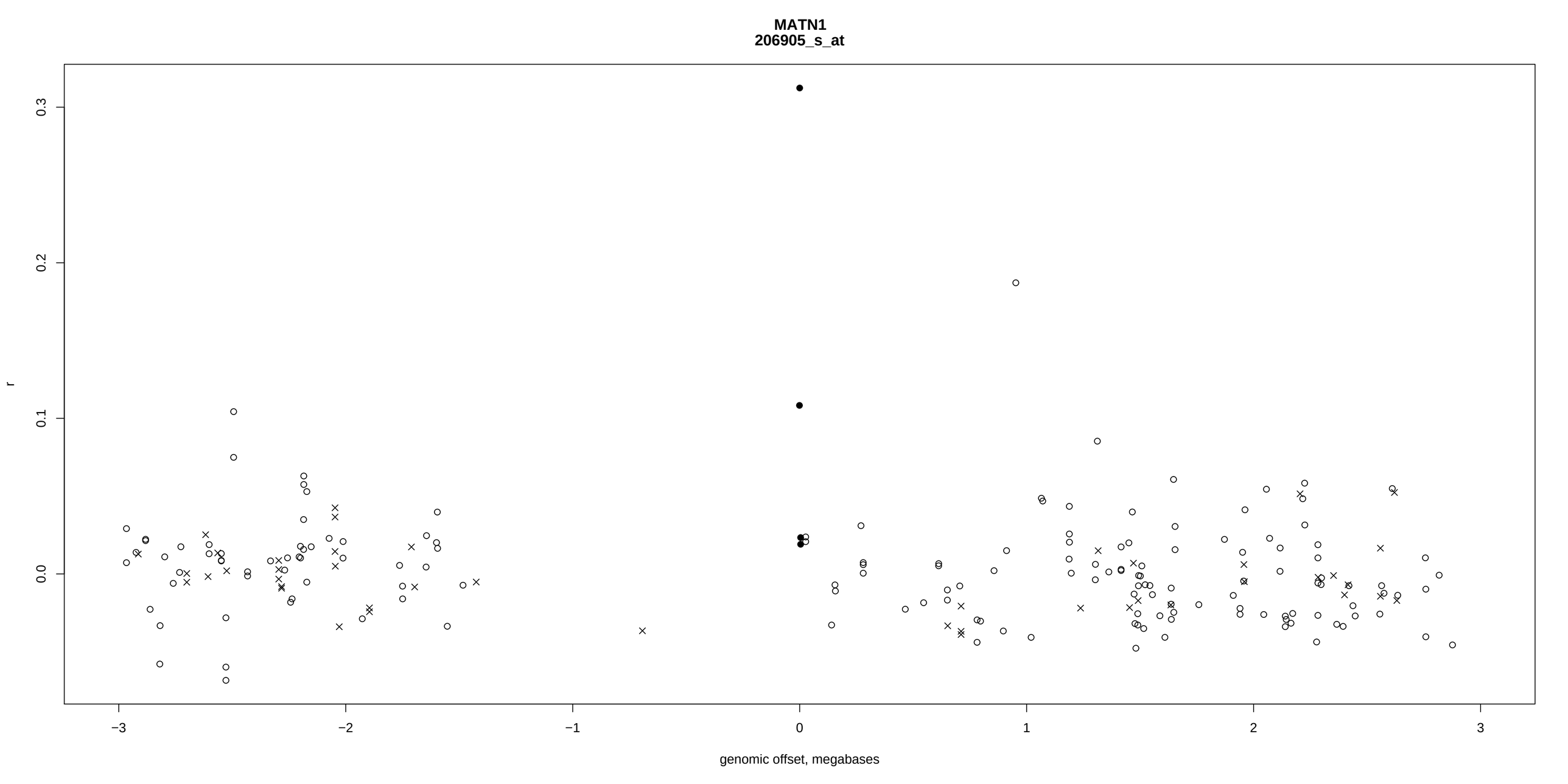
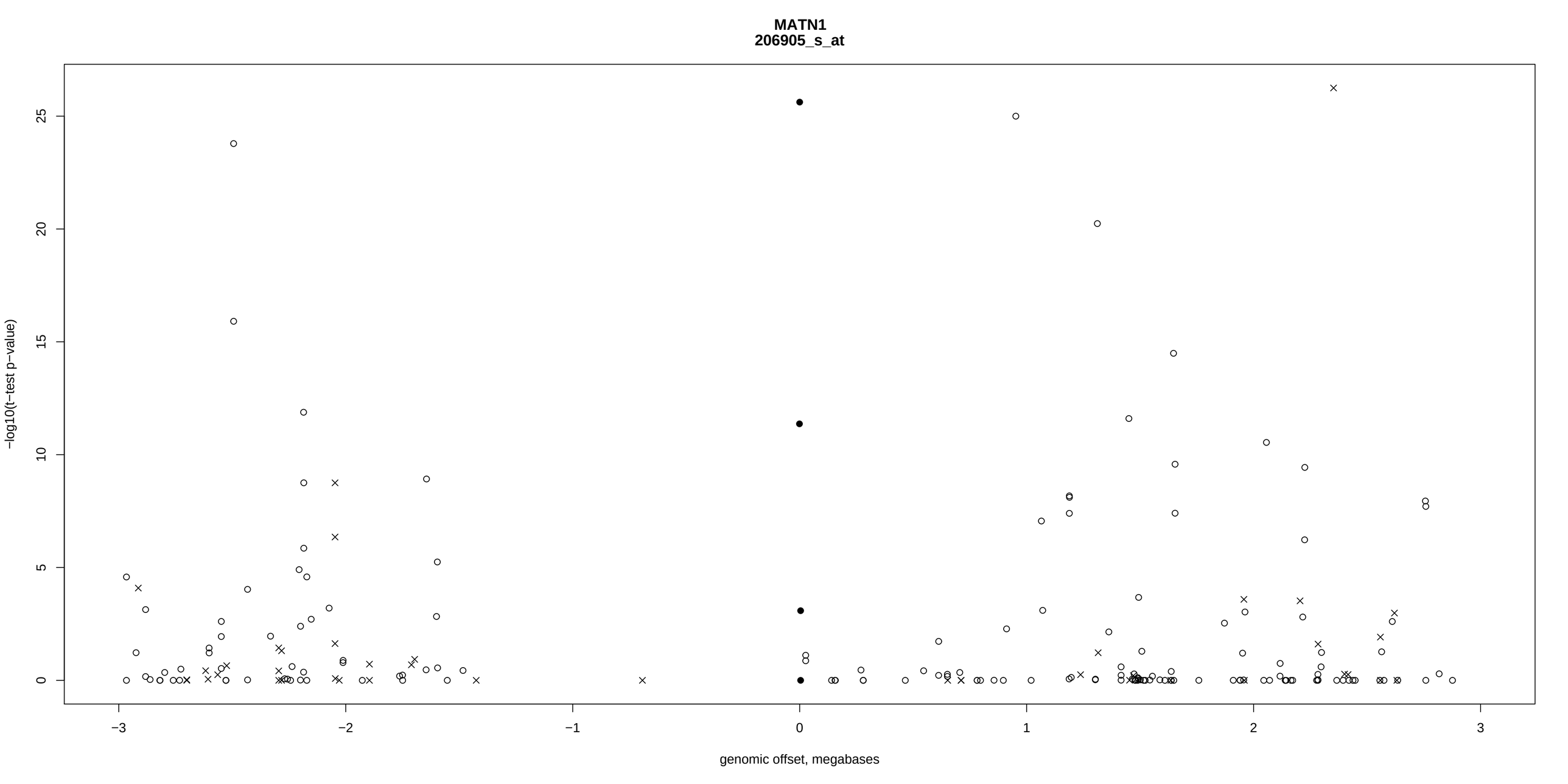
0.10
0.15
0.20
−3
−2
−1
0
1
2
3
genomic offset, megabases

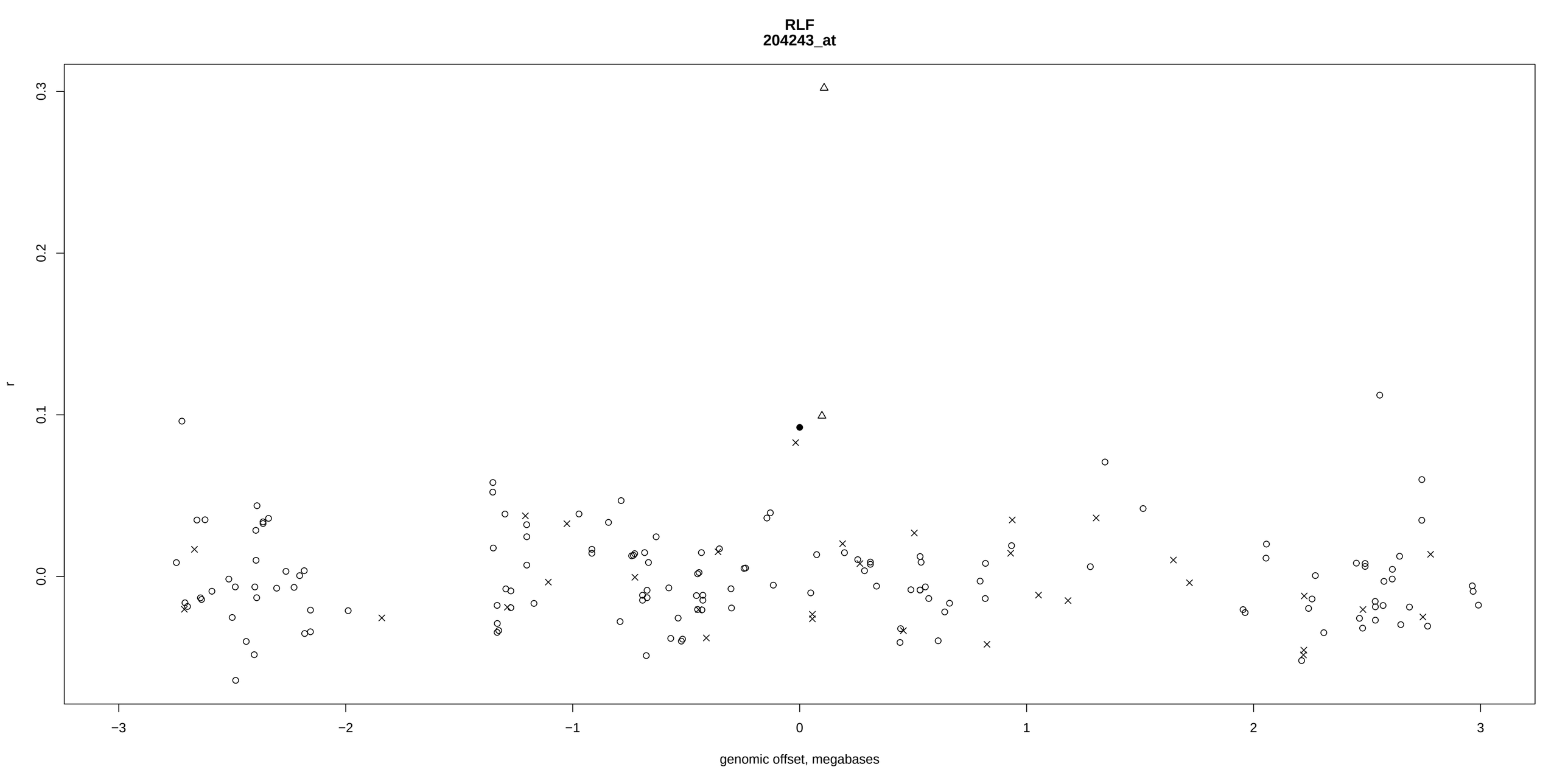

COL11A1
204320_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

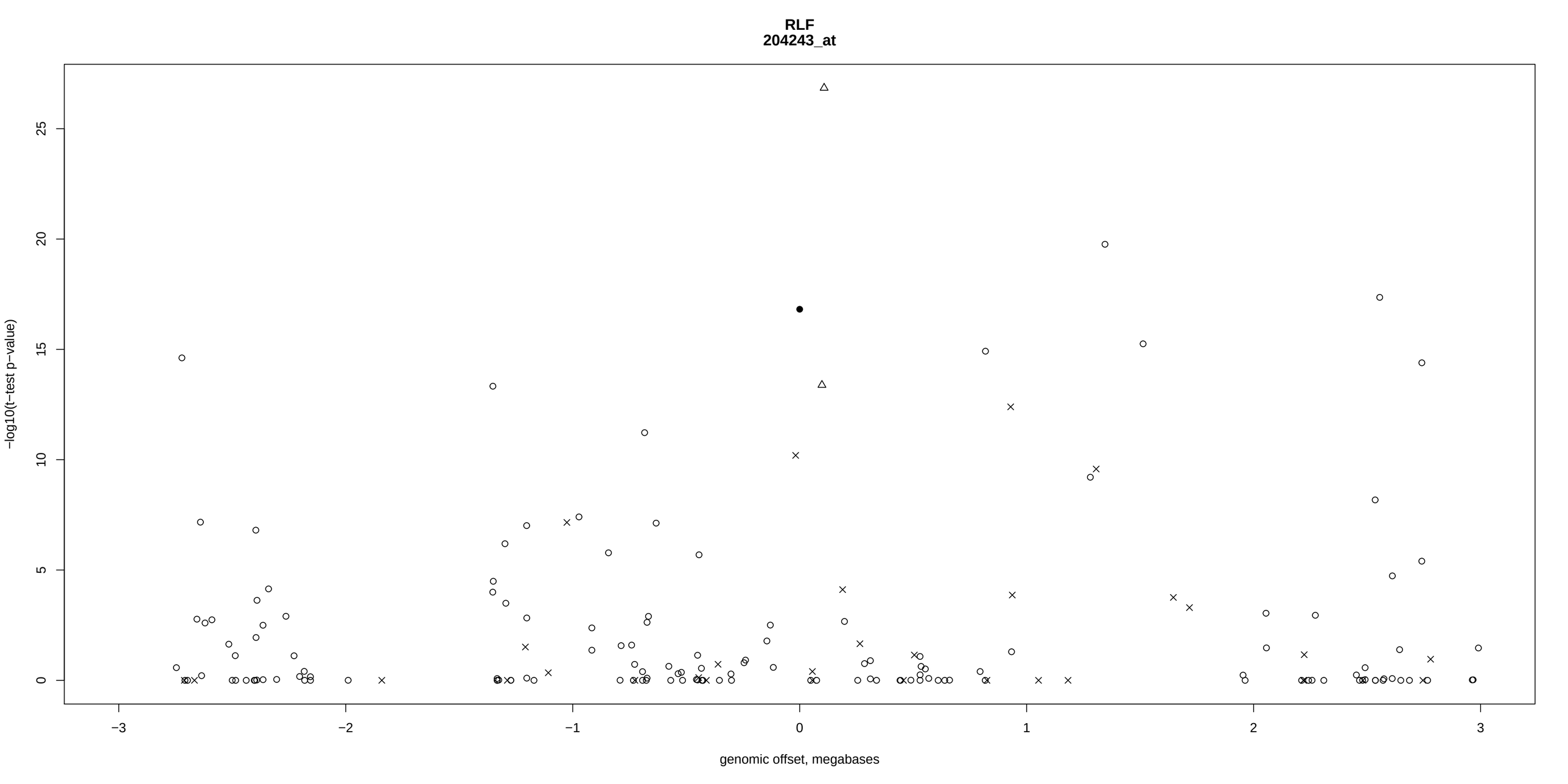

PDPN
204879_at
r
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases

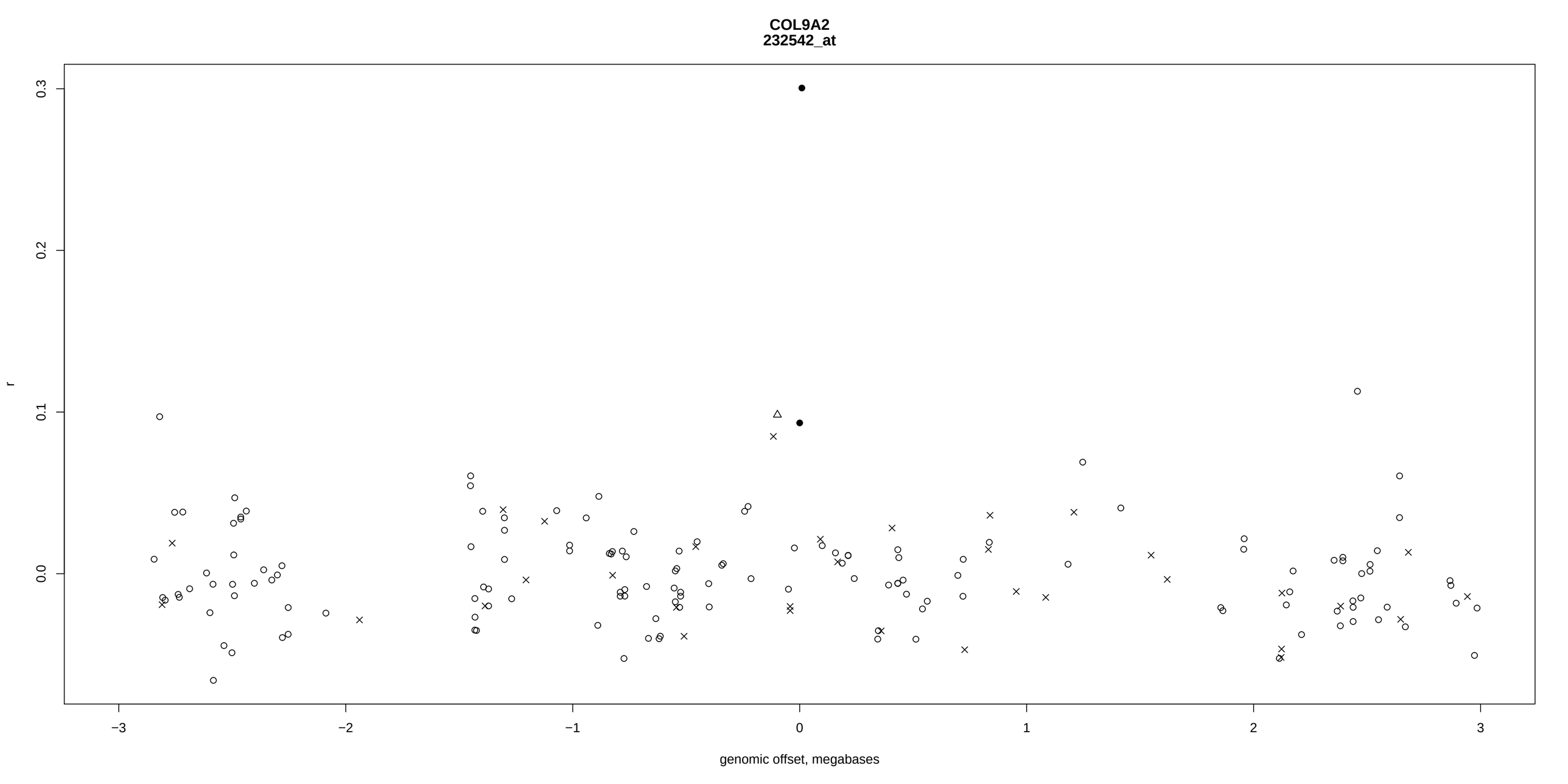

PDPN
204879_at
−log10(t−test p−value)
0
20
40
60
80
−3
−2
−1
0
1
2
3
genomic offset, megabases

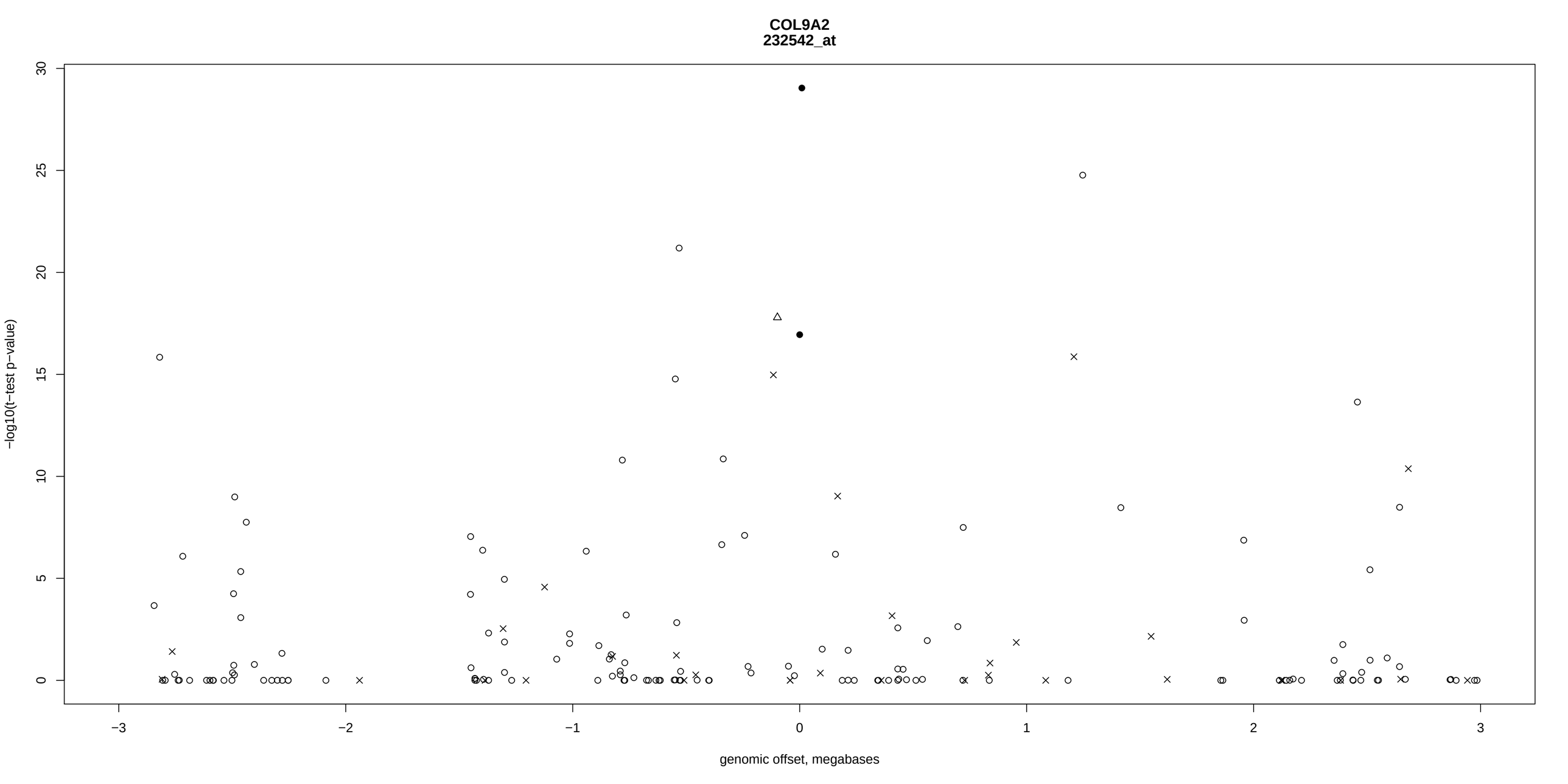

MATN1
206905_s_at
r
0.0
0.1
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases

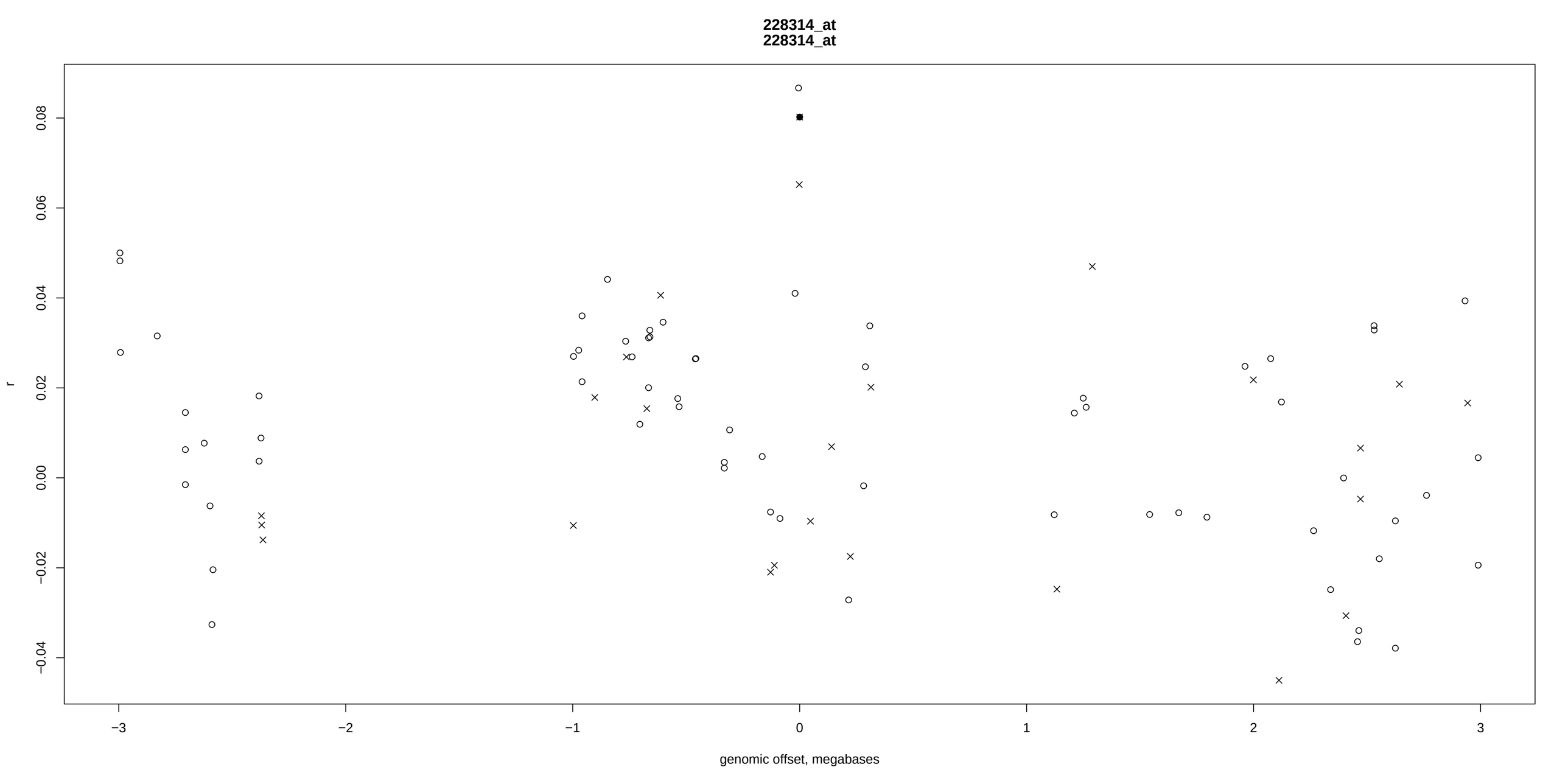

MATN1
206905_s_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

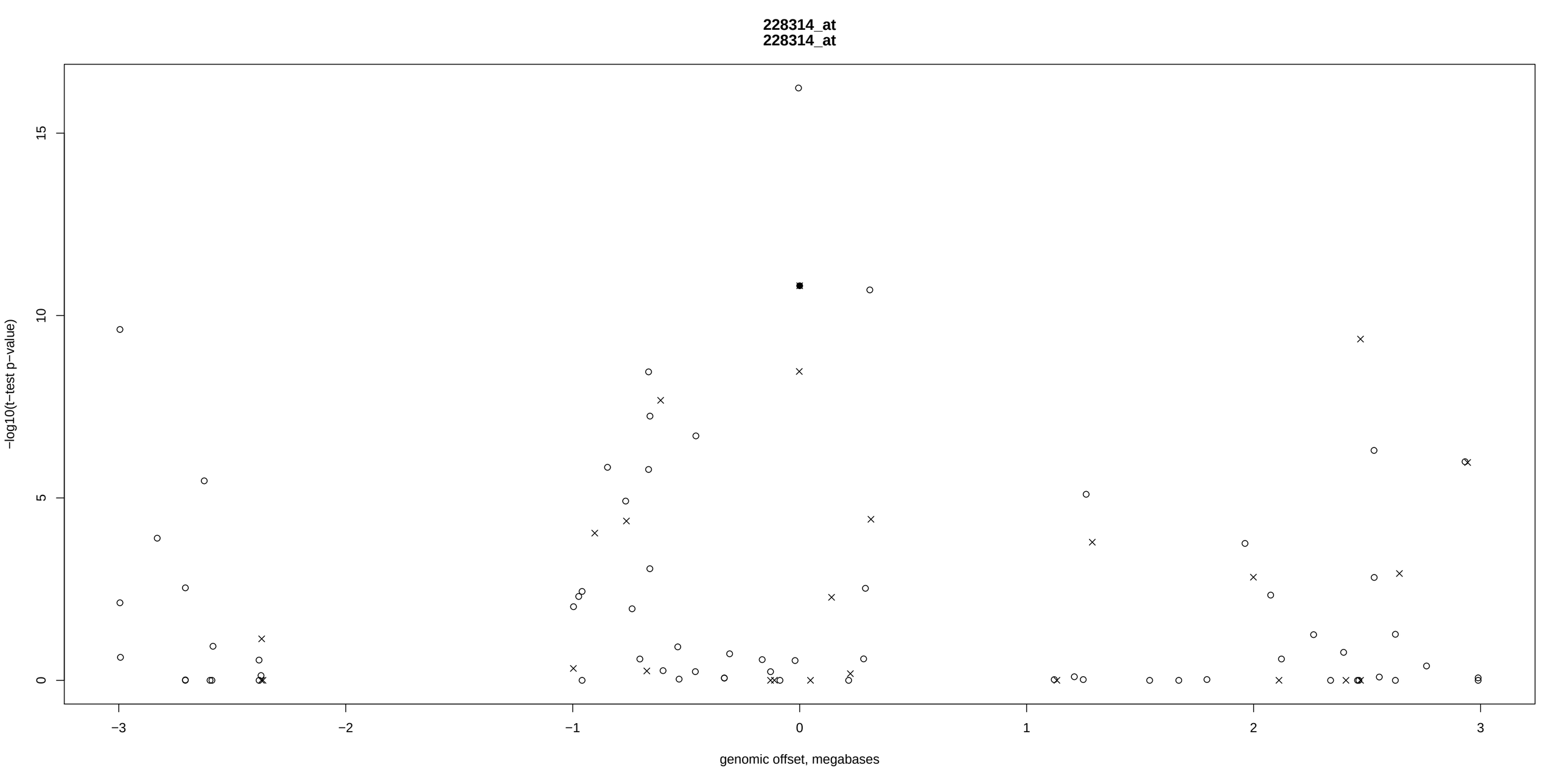

RLF
204243_at
r
0.0
0.1
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases

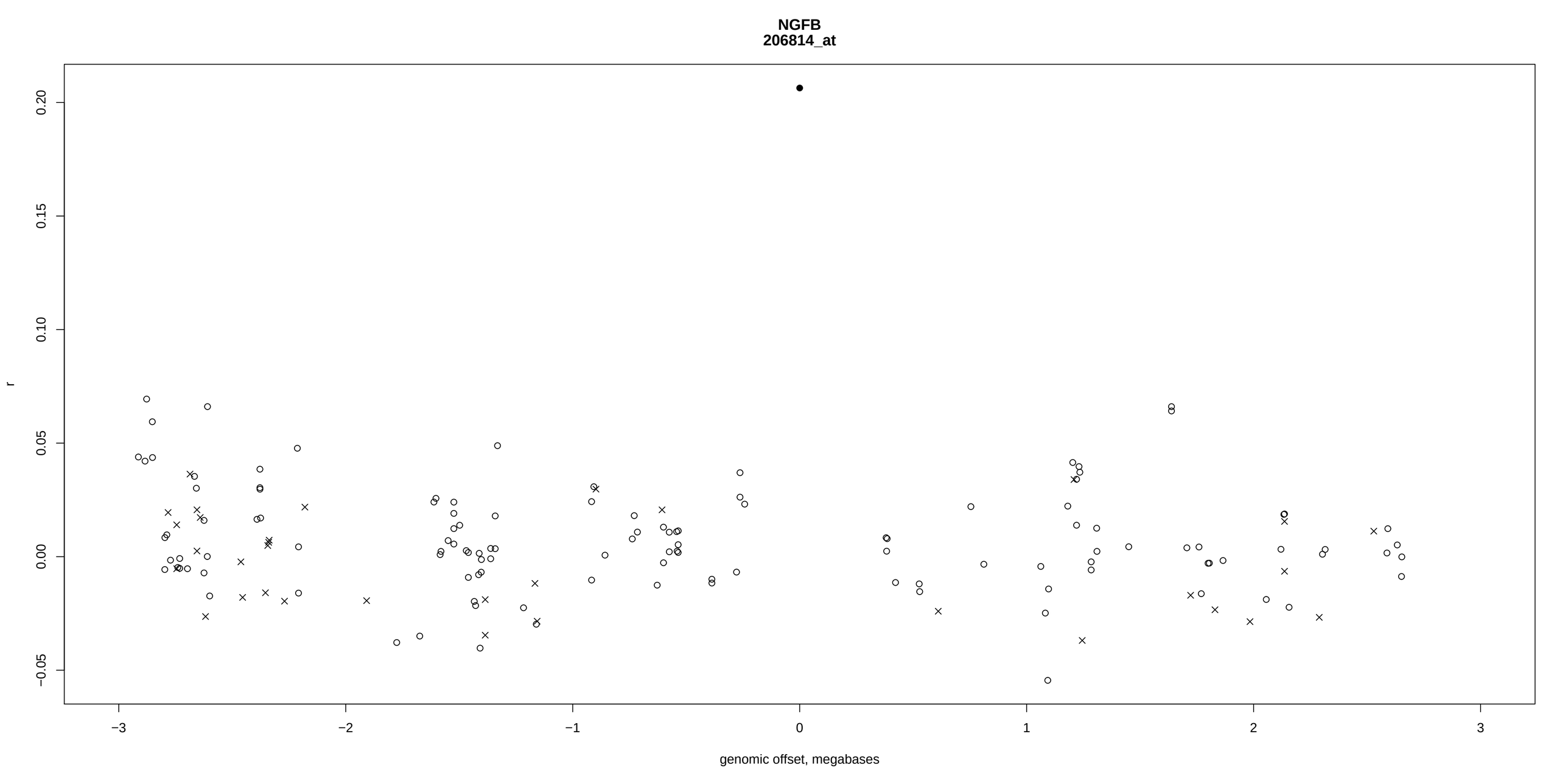

RLF
204243_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

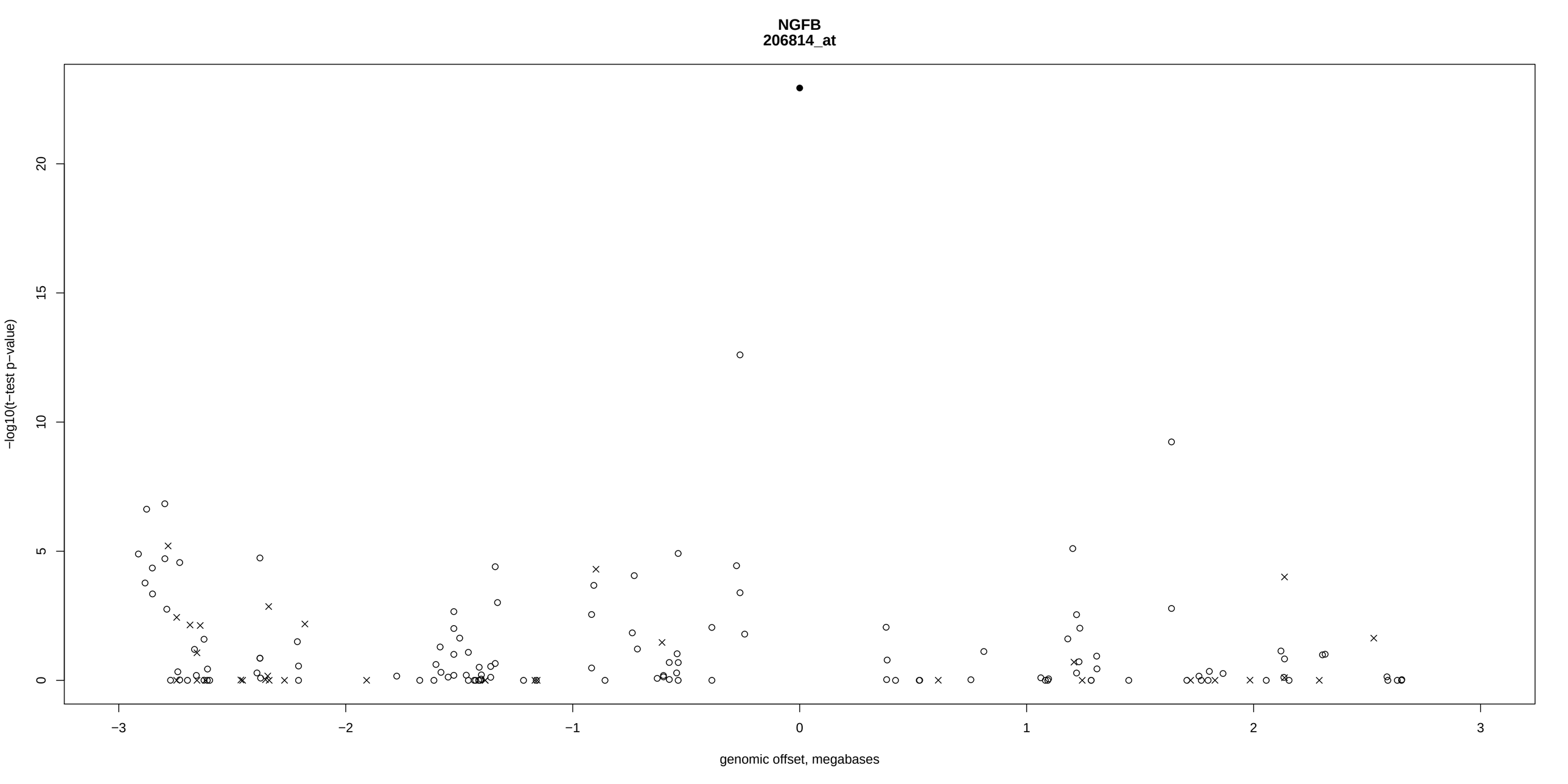

COL9A2
232542_at
r
0.0
0.1
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases

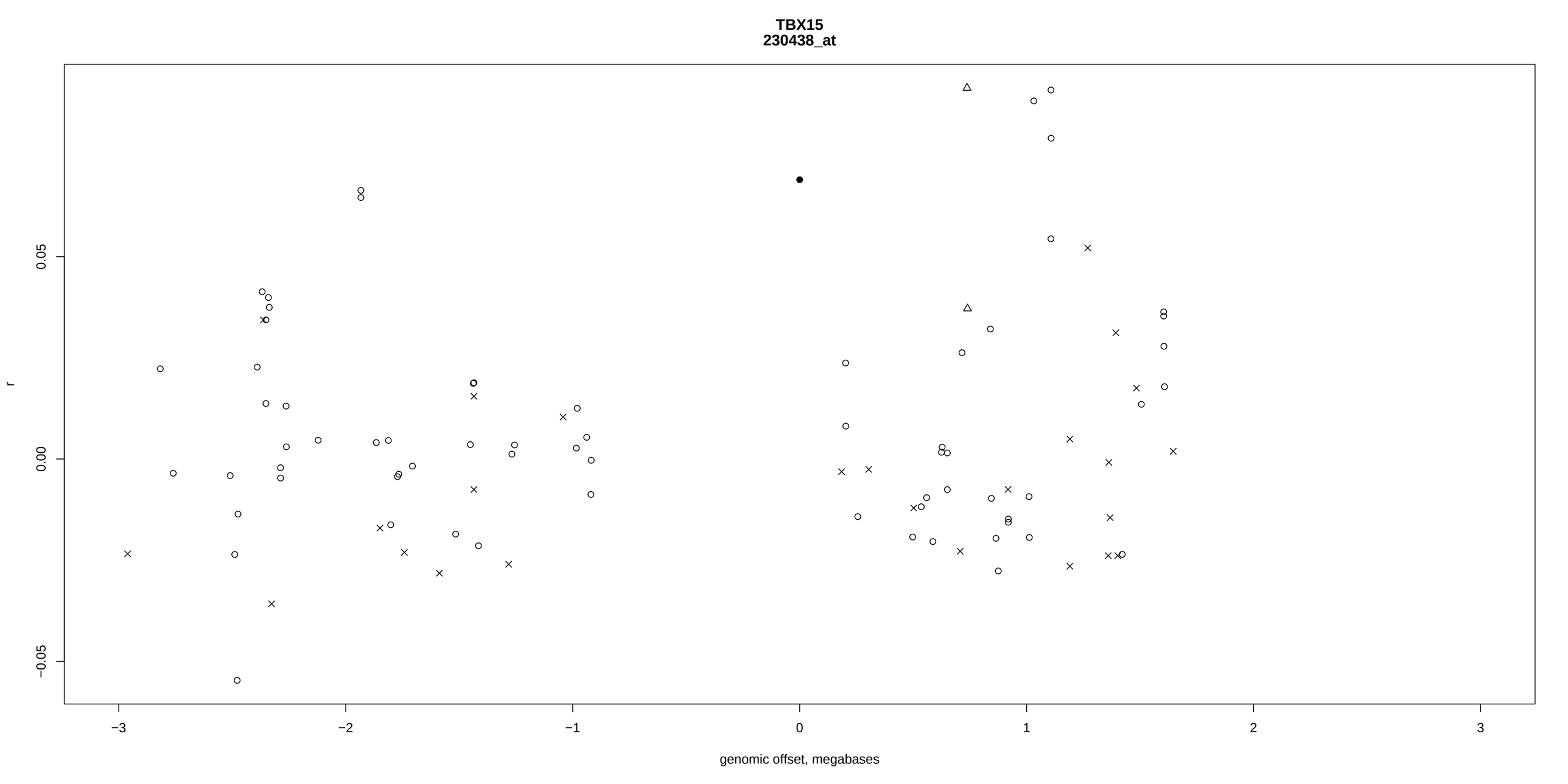

COL9A2
232542_at
−log10(t−test p−value)
genomic offset, megabases
0
5
10
15
20
25
30
−3
−2
−1
0
1
2
3

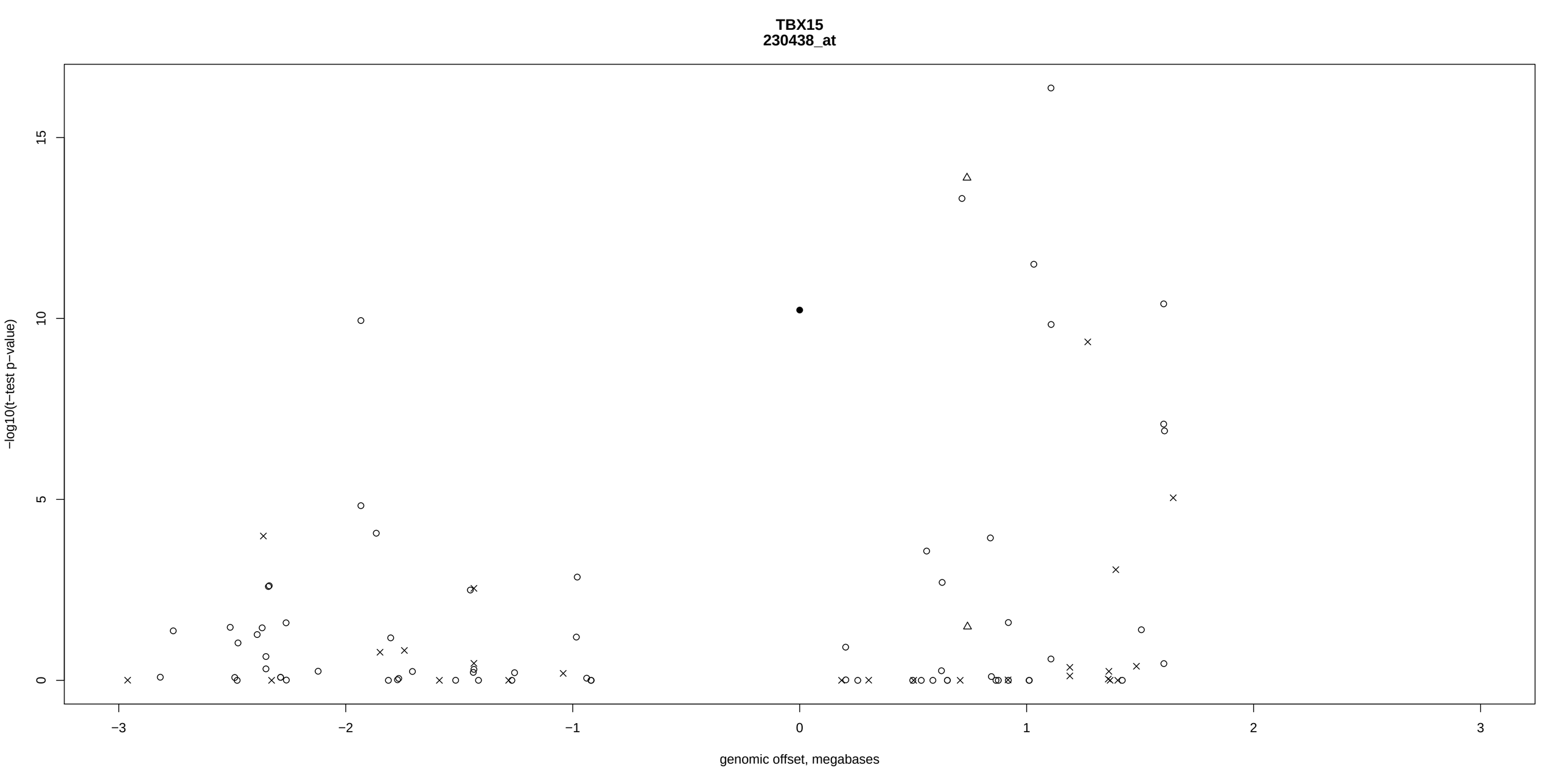

228314_at
228314_at
r
0.08
0.06
0.04
0.02
0.00
−0.02
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases

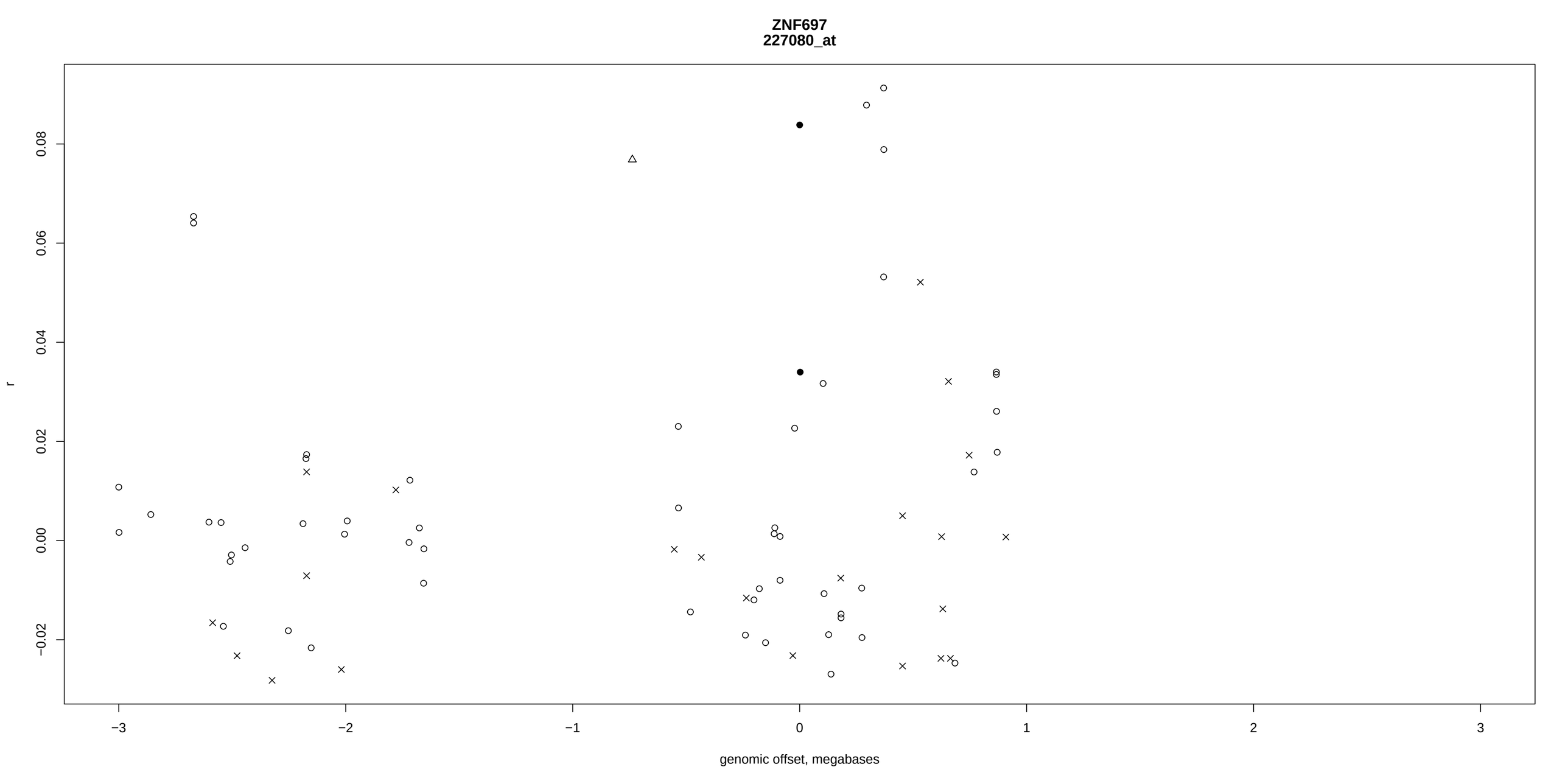

228314_at
228314_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


NGFB
206814_at
r
0.20
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


NGFB
206814_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


TBX15
230438_at
r
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


TBX15
230438_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


ZNF697
227080_at
r
0.08
0.06
0.04
0.02
0.00
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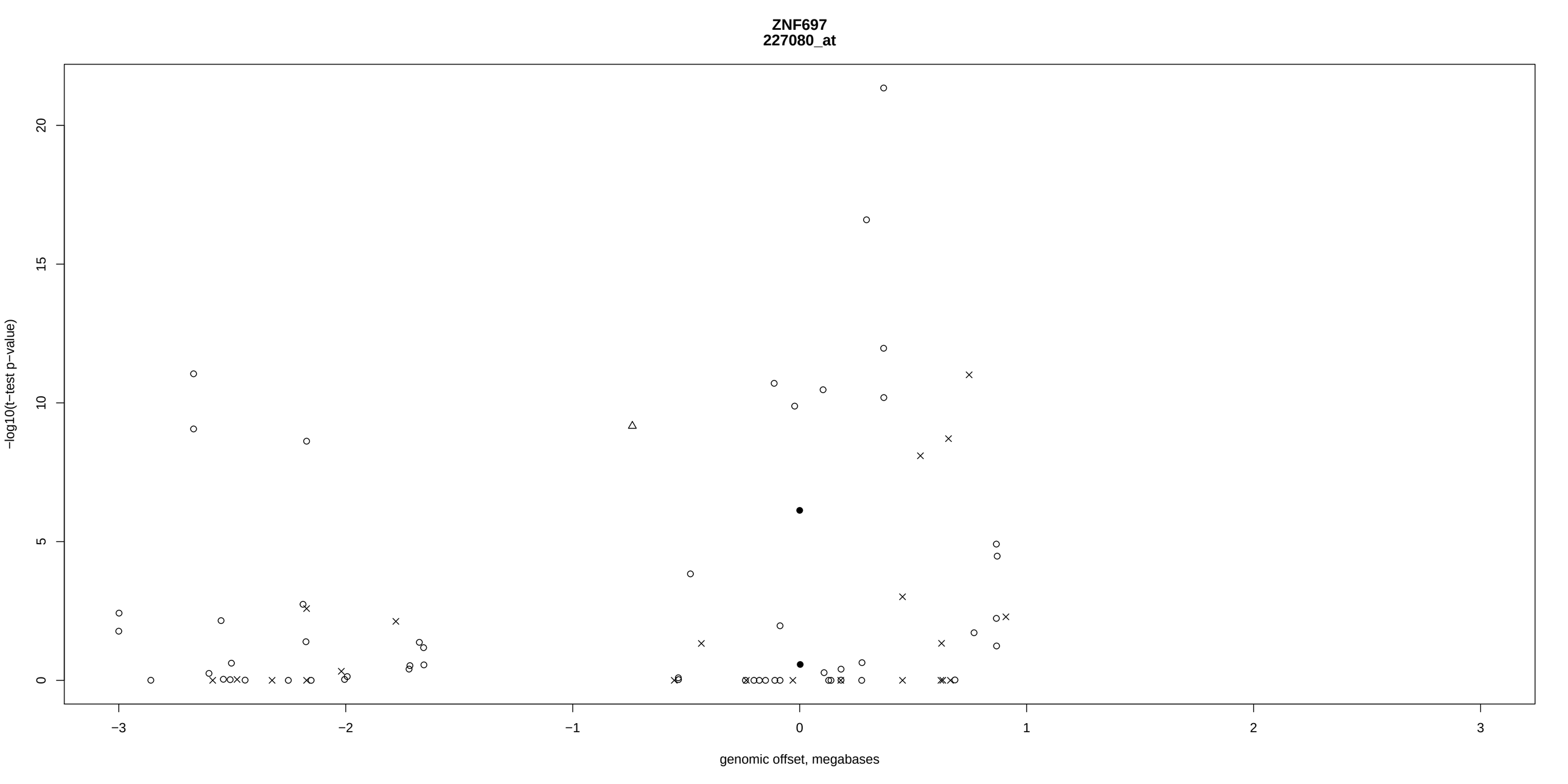
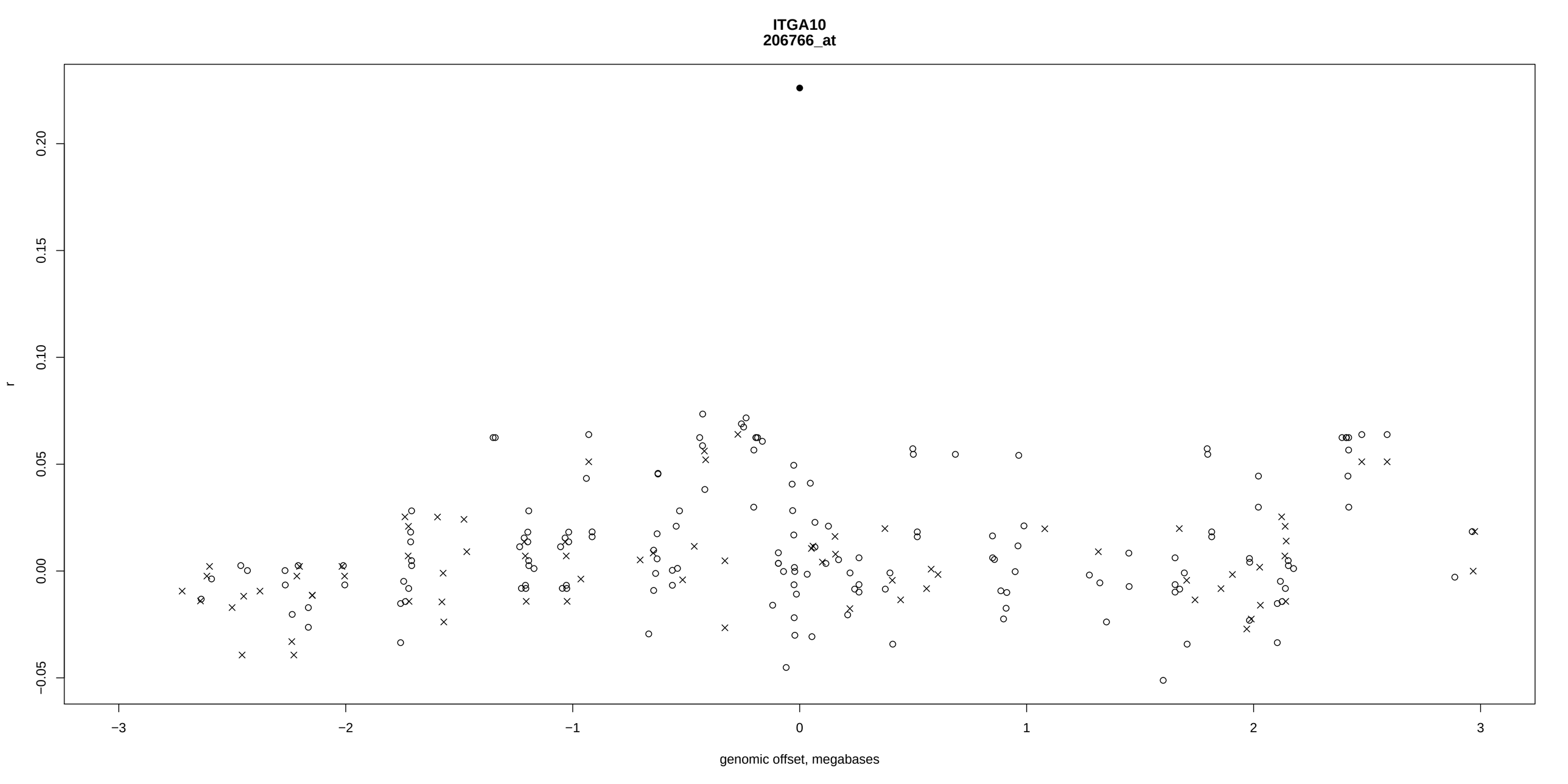
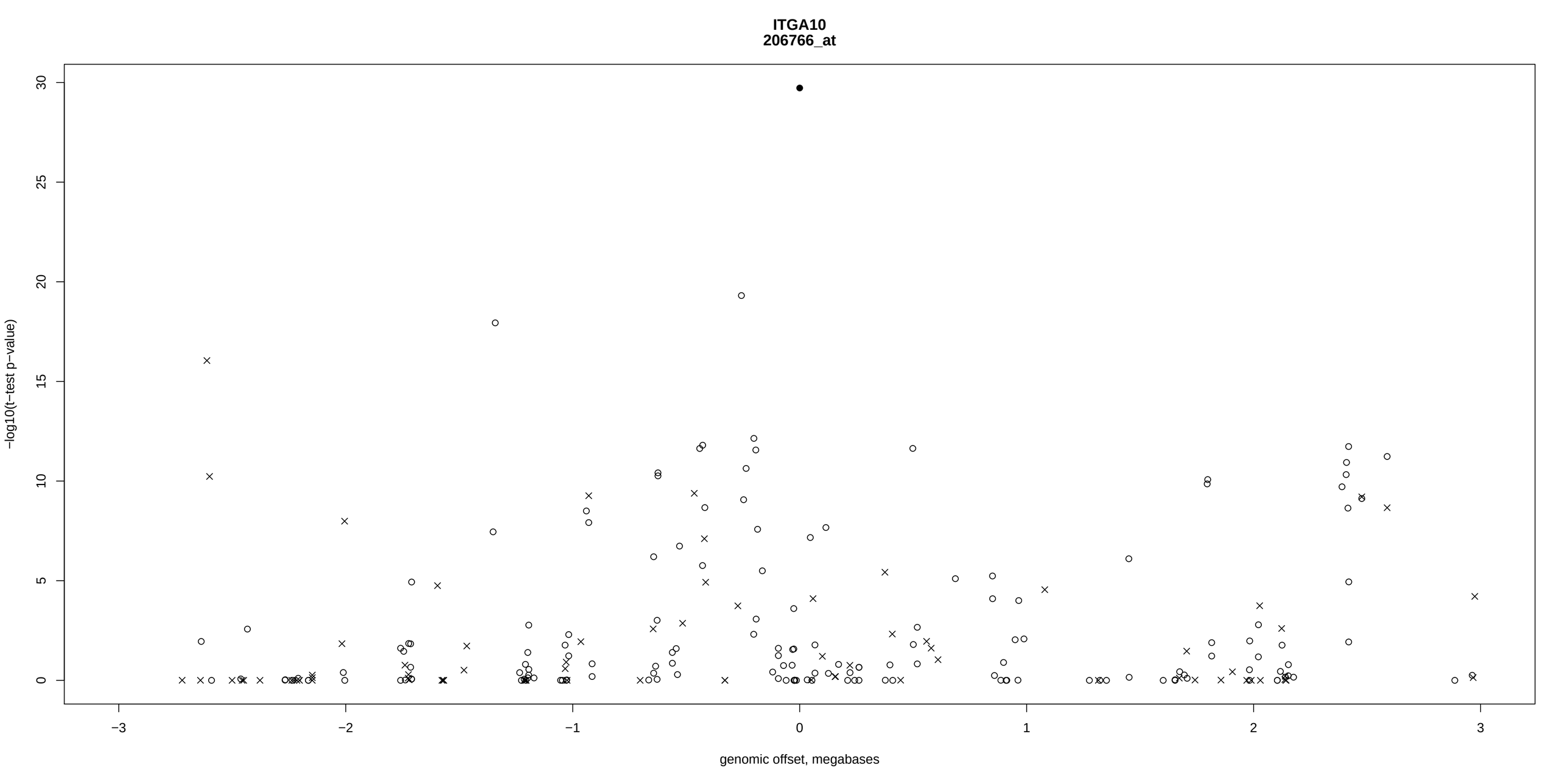
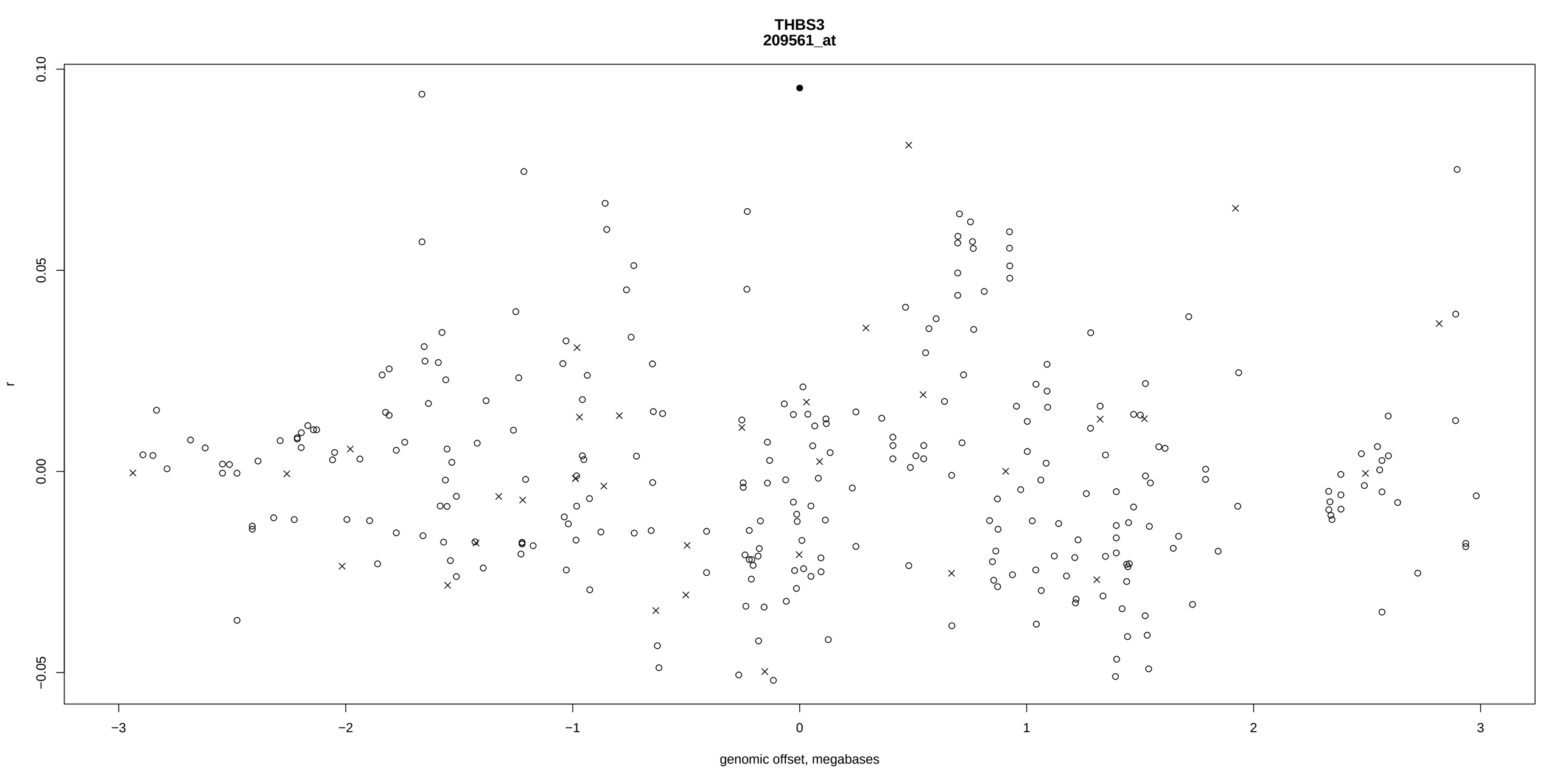
−3
−2
−1
0
1
2
3
genomic offset, megabases


ZNF697
227080_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


ITGA10
206766_at
r
0.20
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


ITGA10
206766_at
−log10(t−test p−value)
0
5
10
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


THBS3
209561_at
0.10
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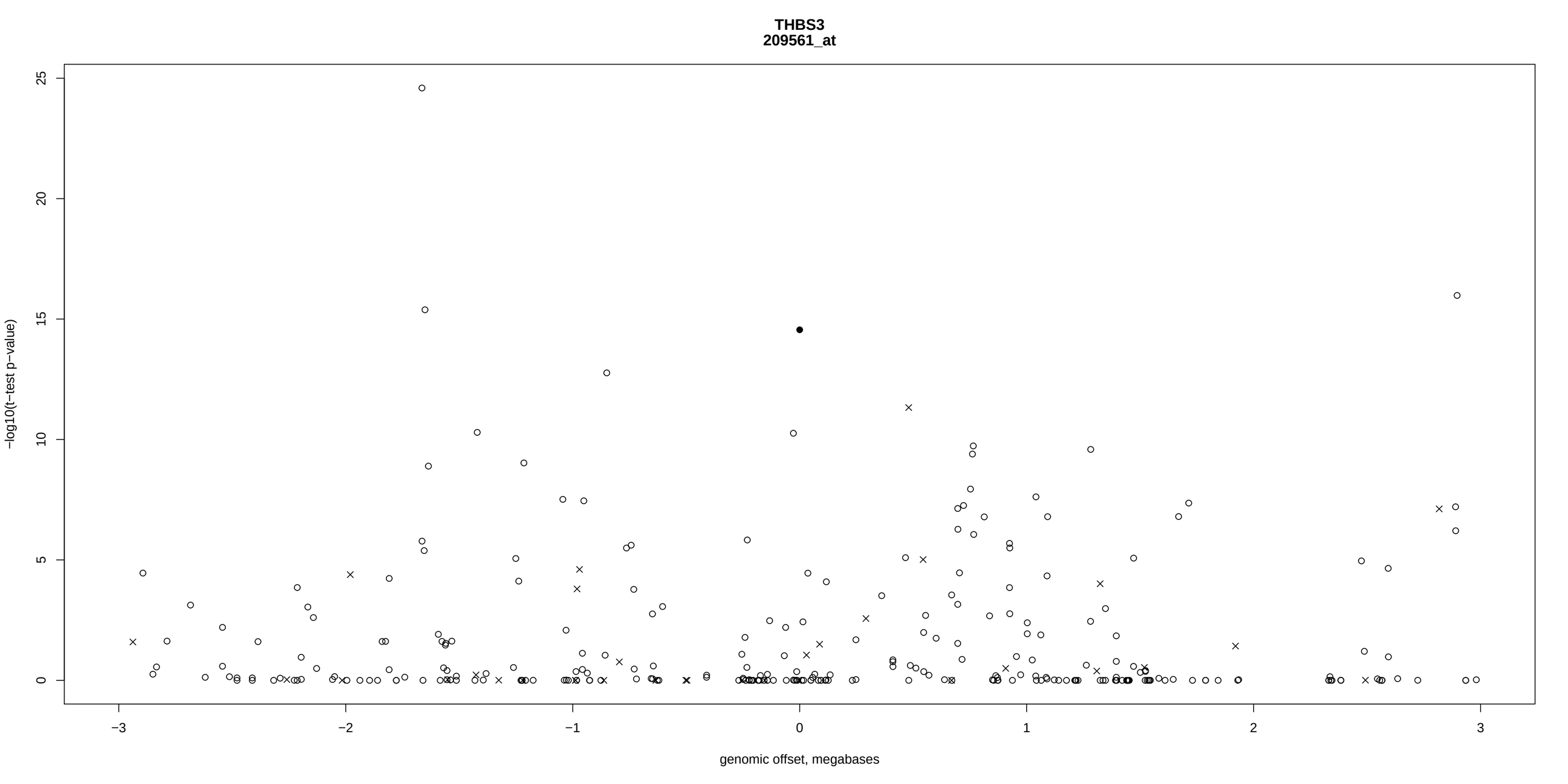
0.00
−0.05
r
−3
−2
−1
0
1
2
3
genomic offset, megabases


THBS3
209561_at
−log10(t−test p−value)
0
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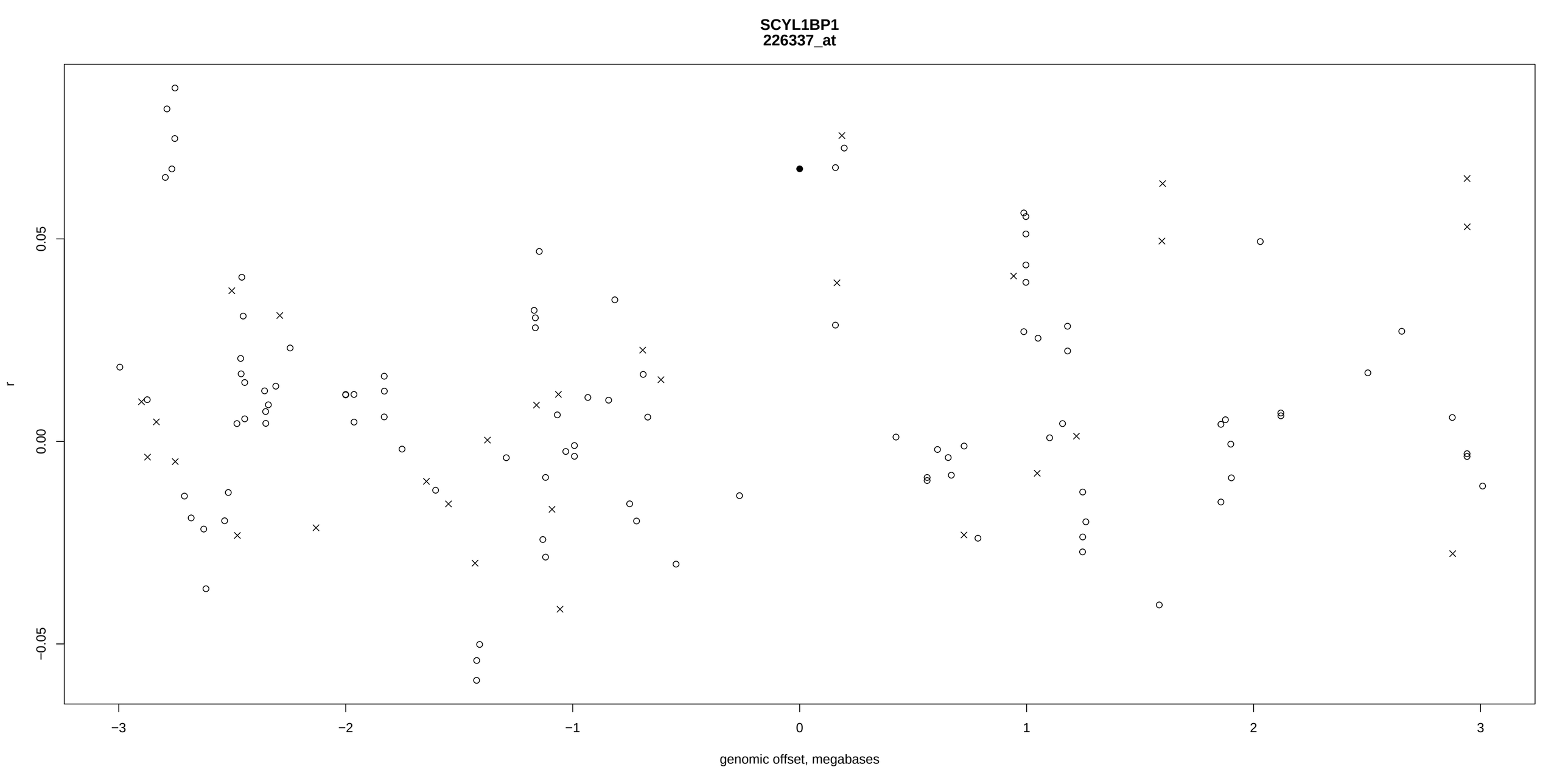
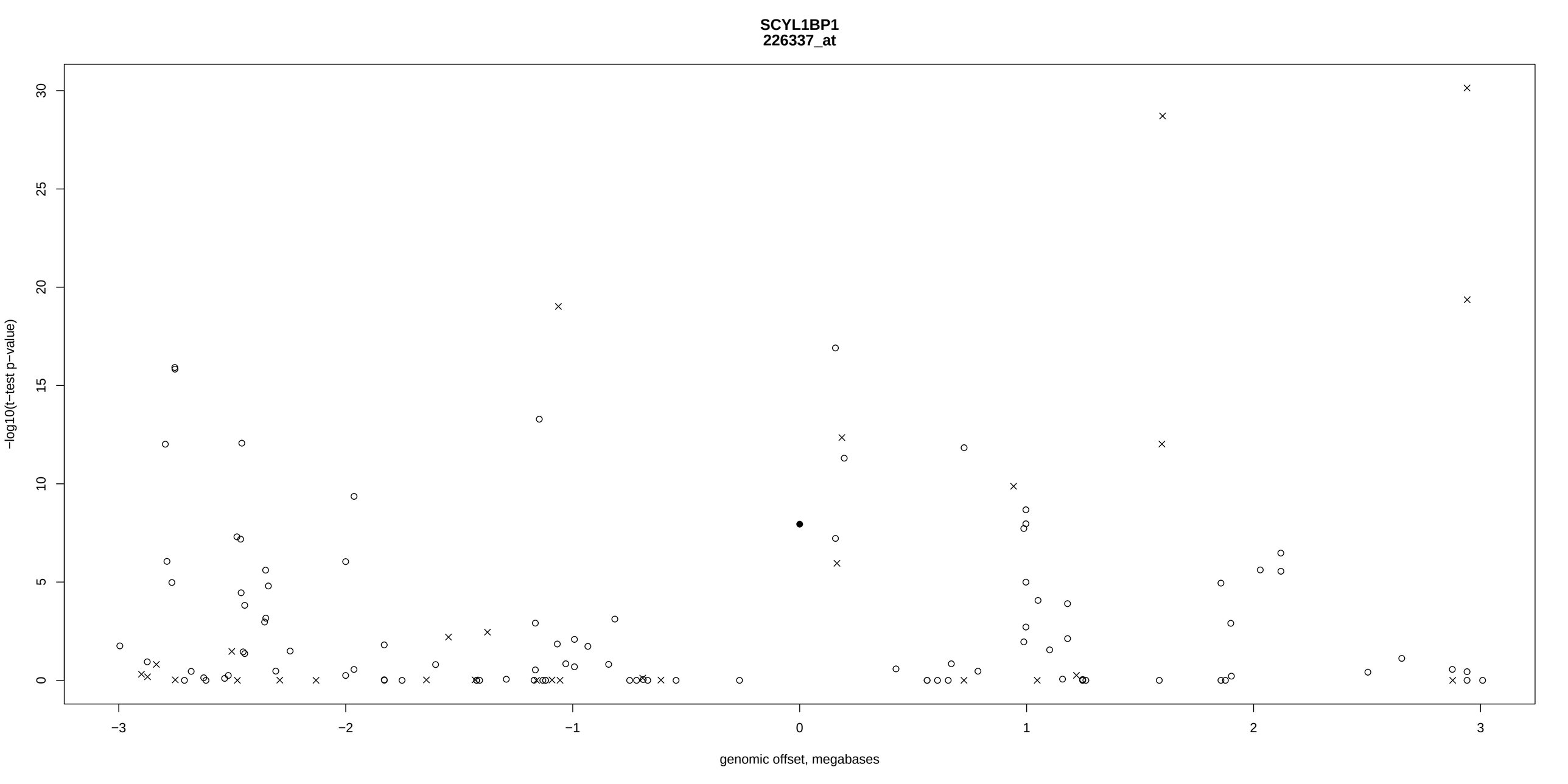
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


SCYL1BP1
226337_at
r
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


SCYL1BP1
226337_at
−log10(t−test p−value)
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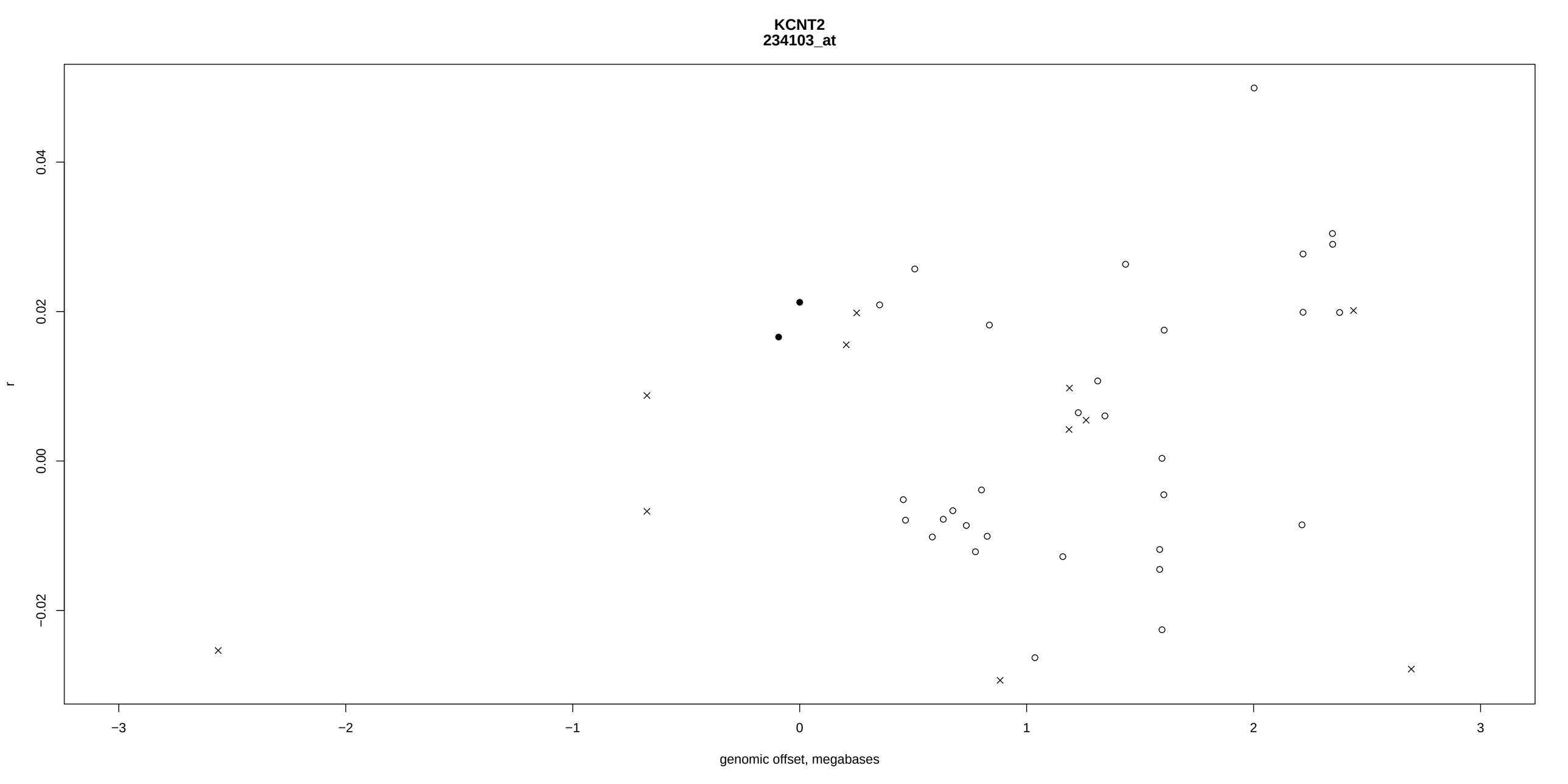
10
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


KCNT2
234103_at
r
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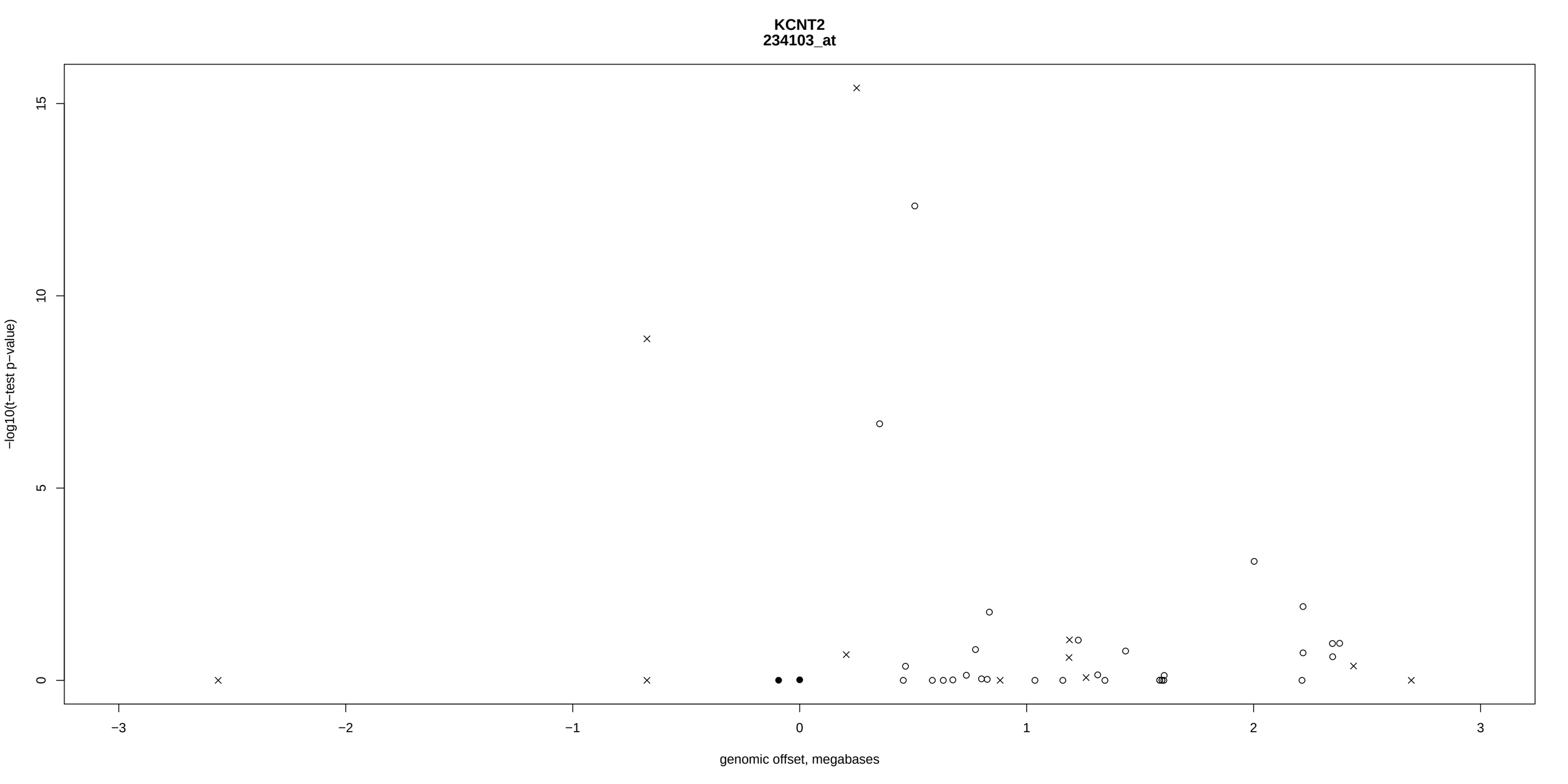
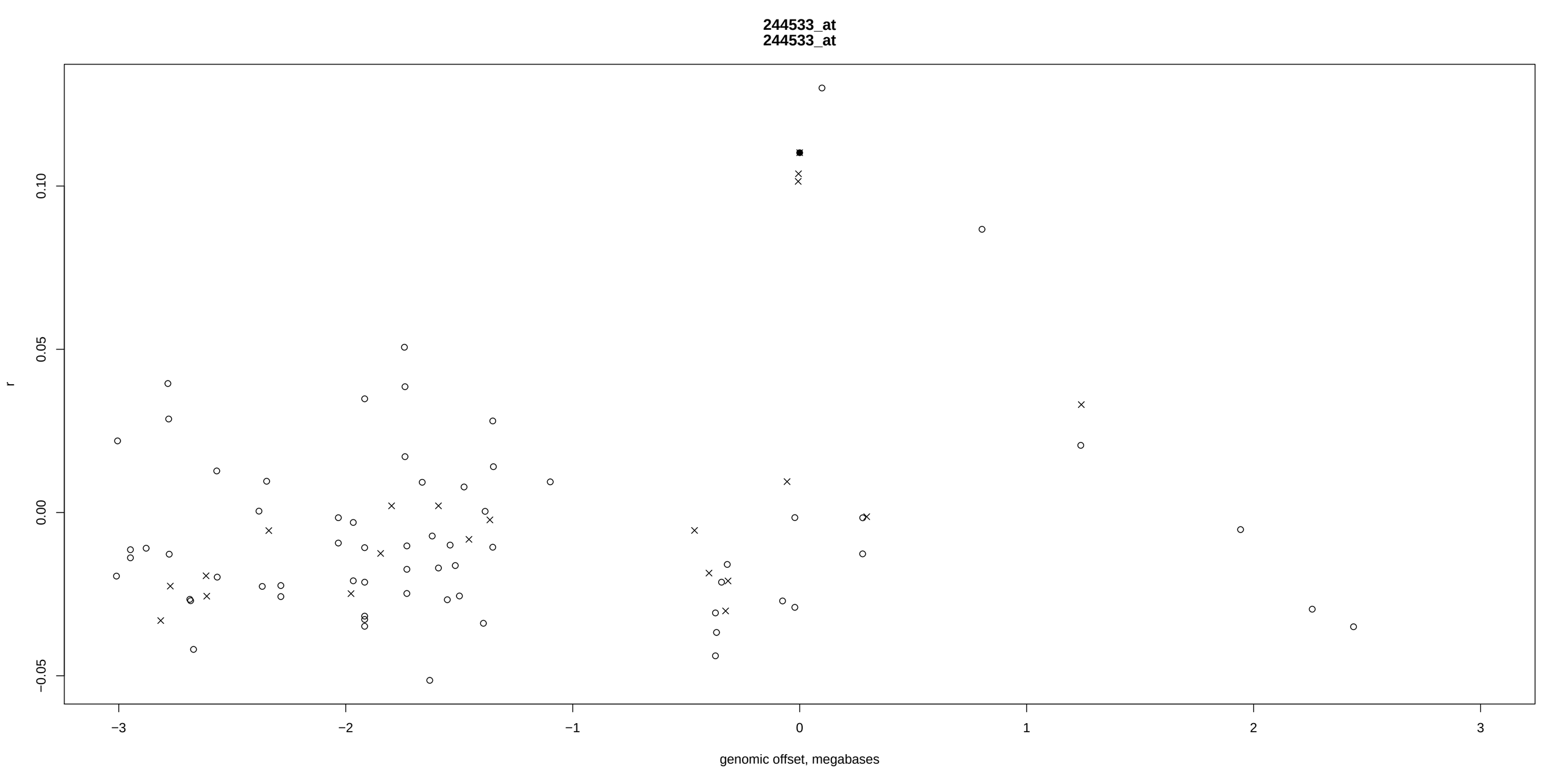
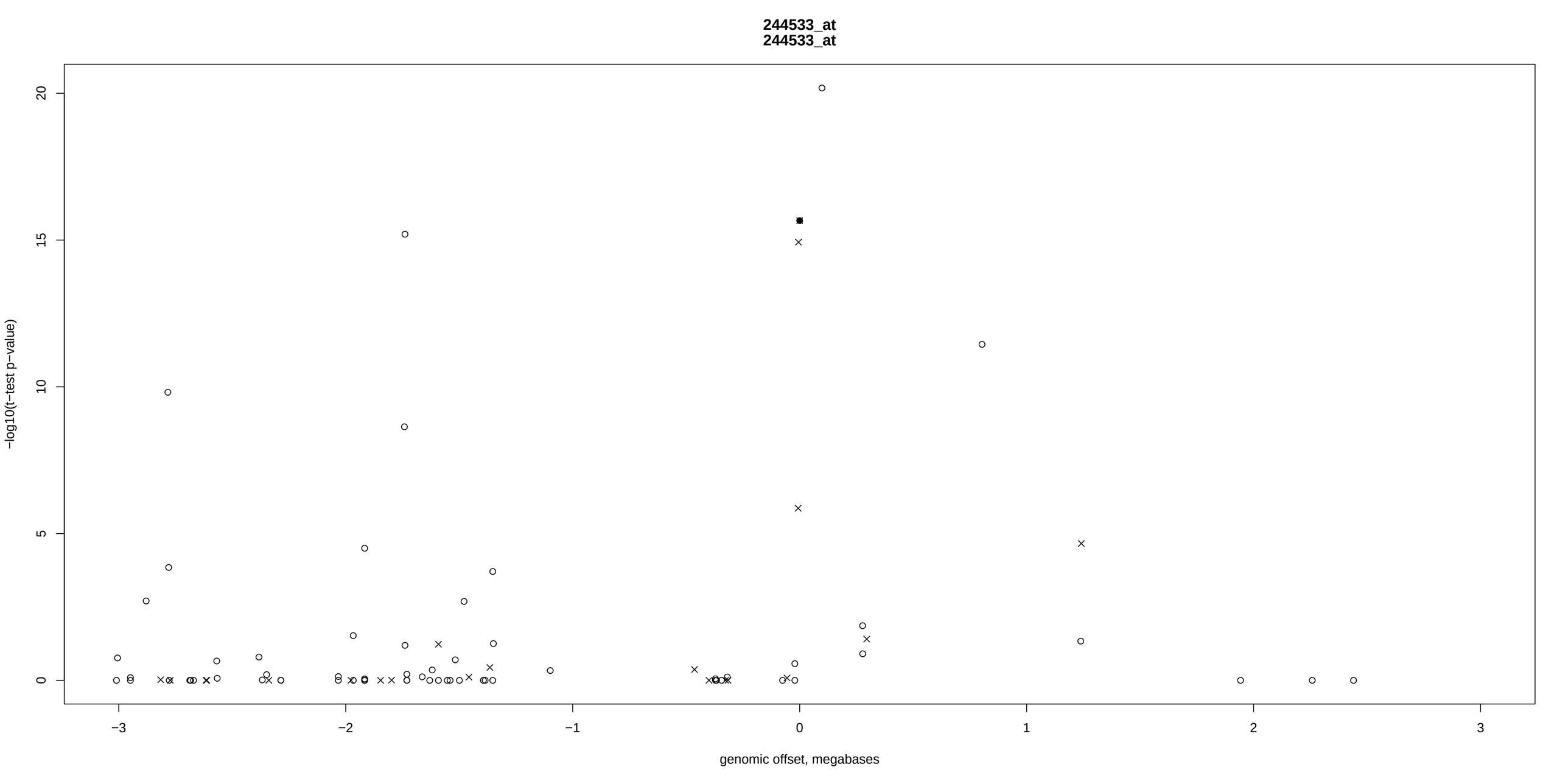
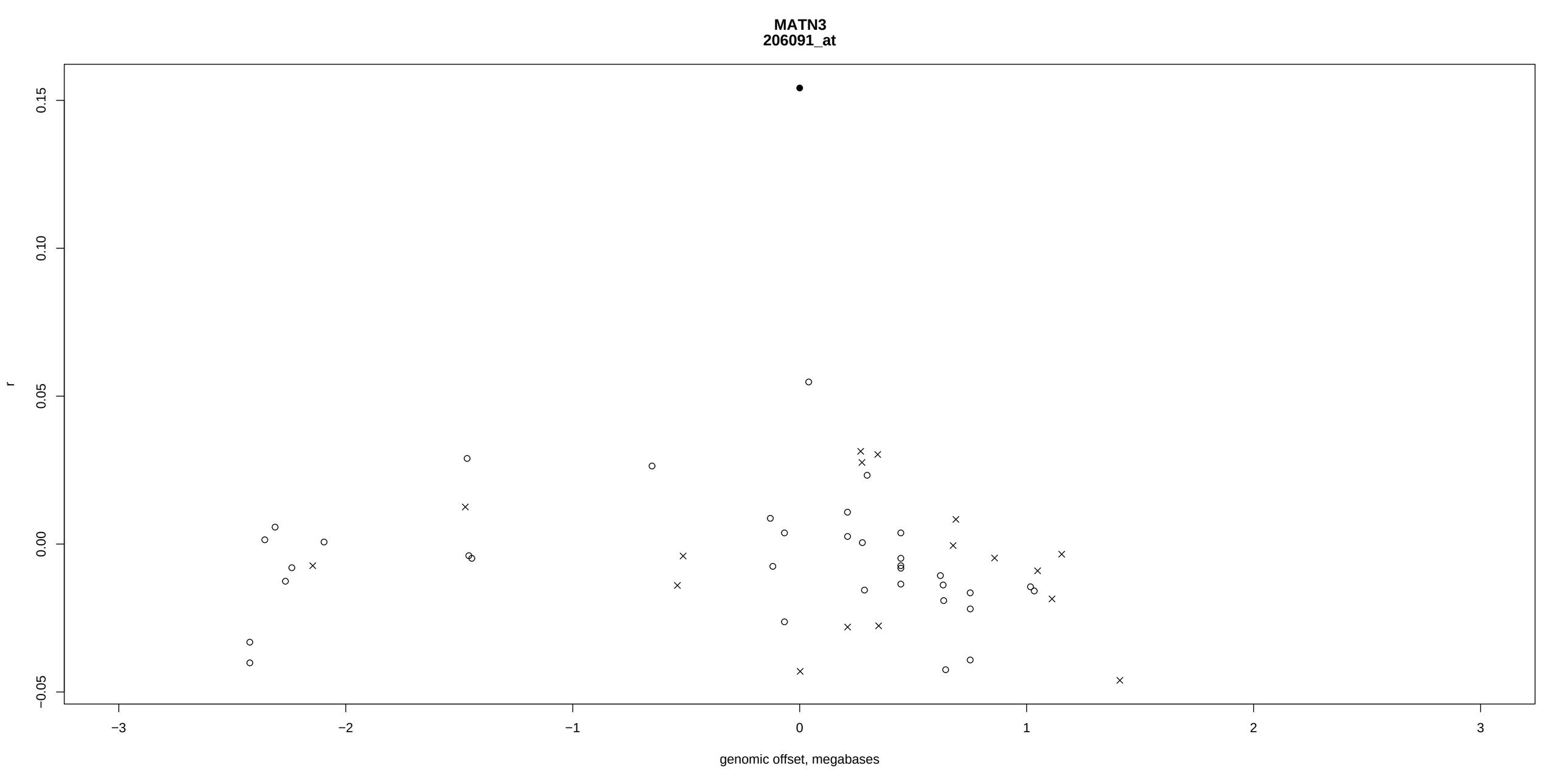
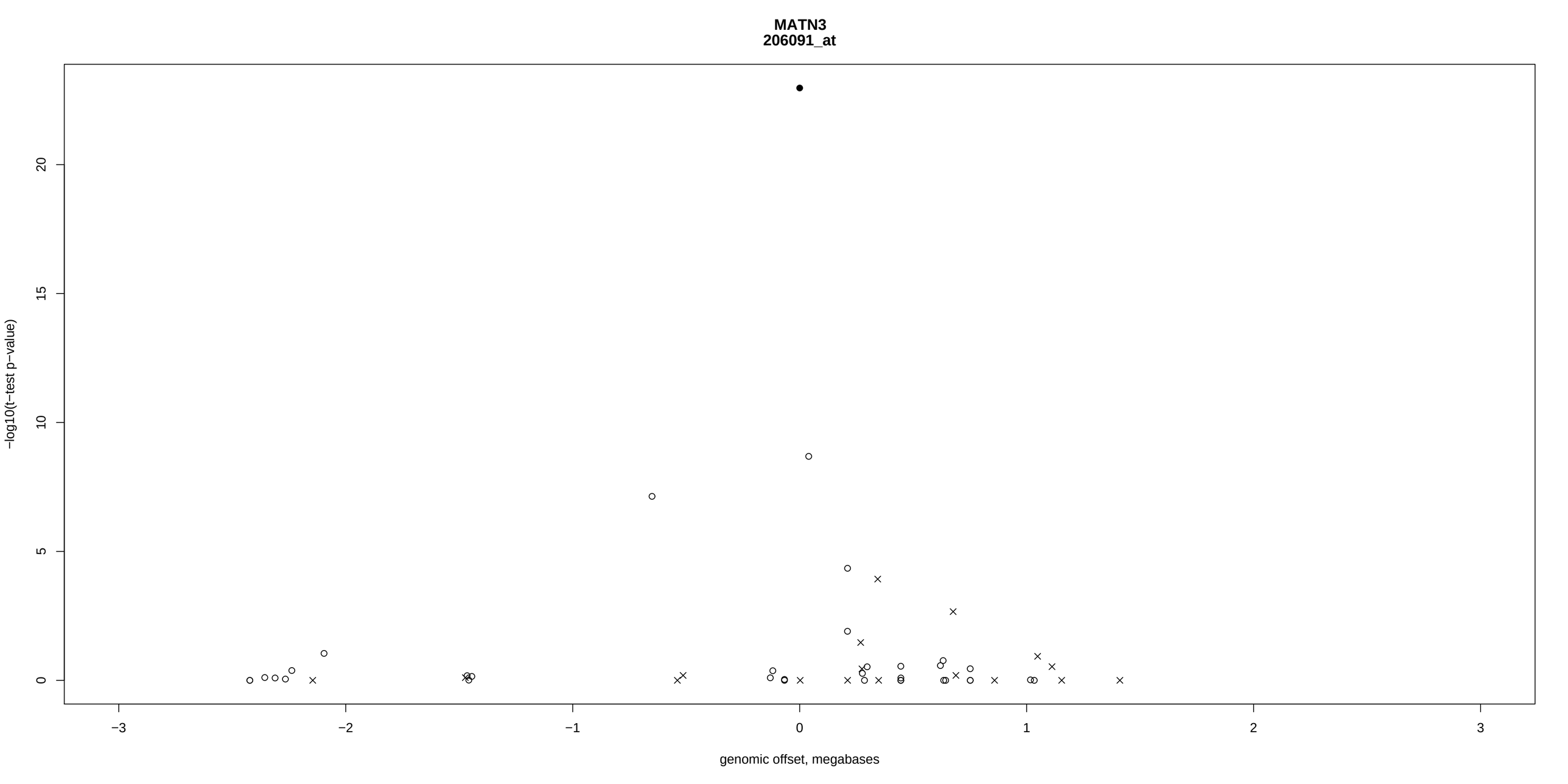
0.02
0.00
−0.02
−3
−2
−1
0
1
2
3
genomic offset, megabases

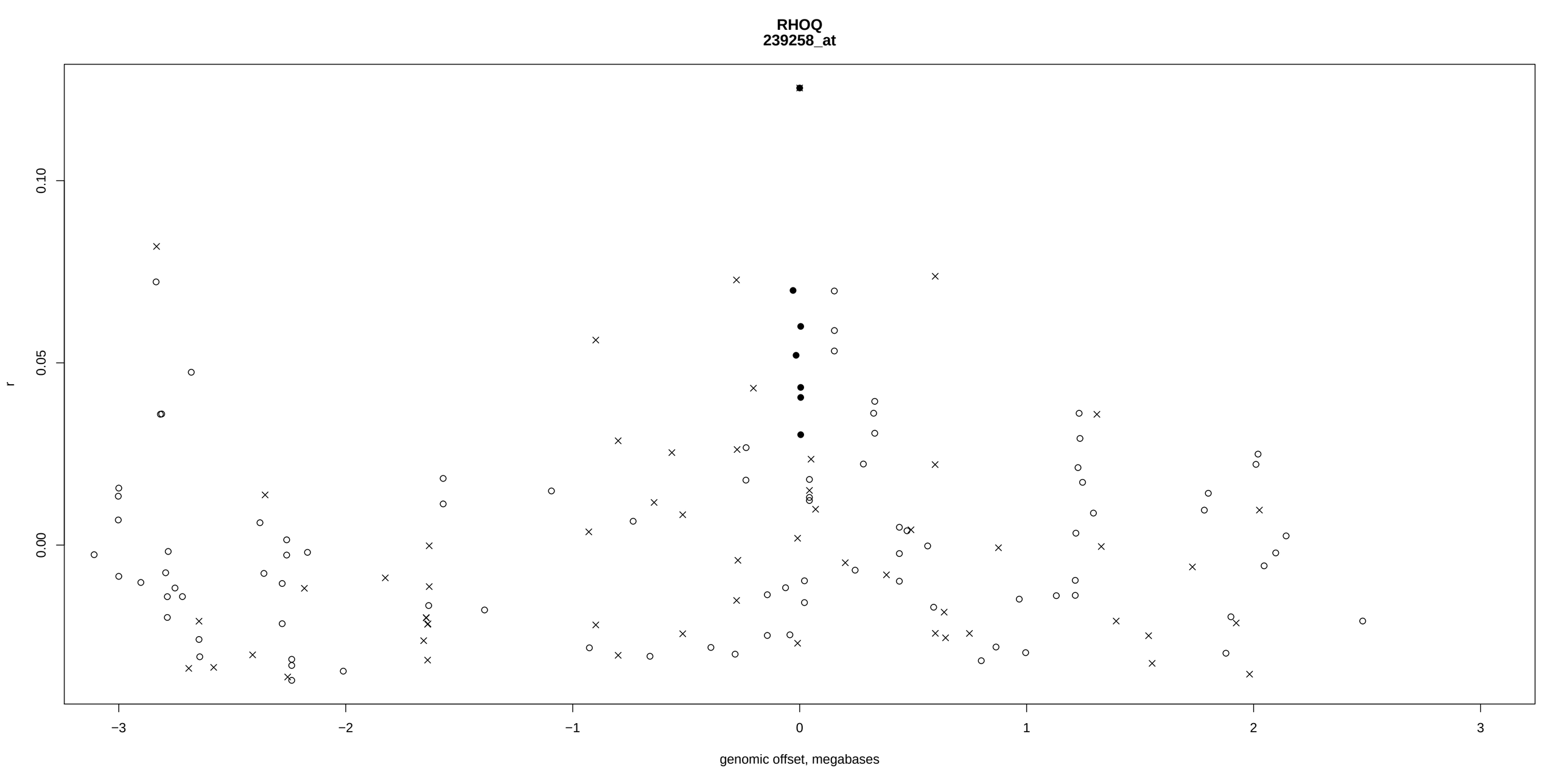

KCNT2
234103_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases

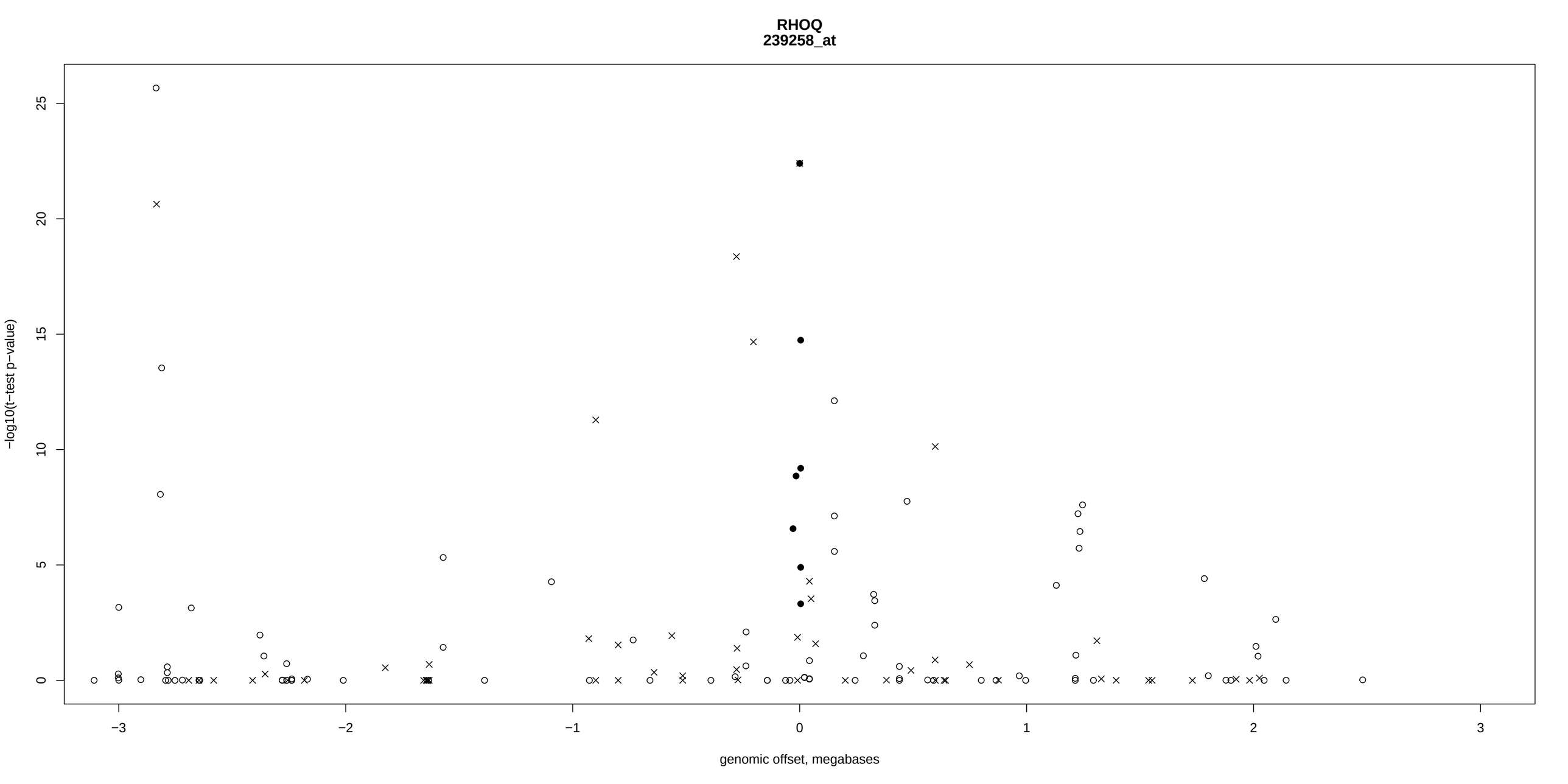

244533_at
244533_at
r
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases

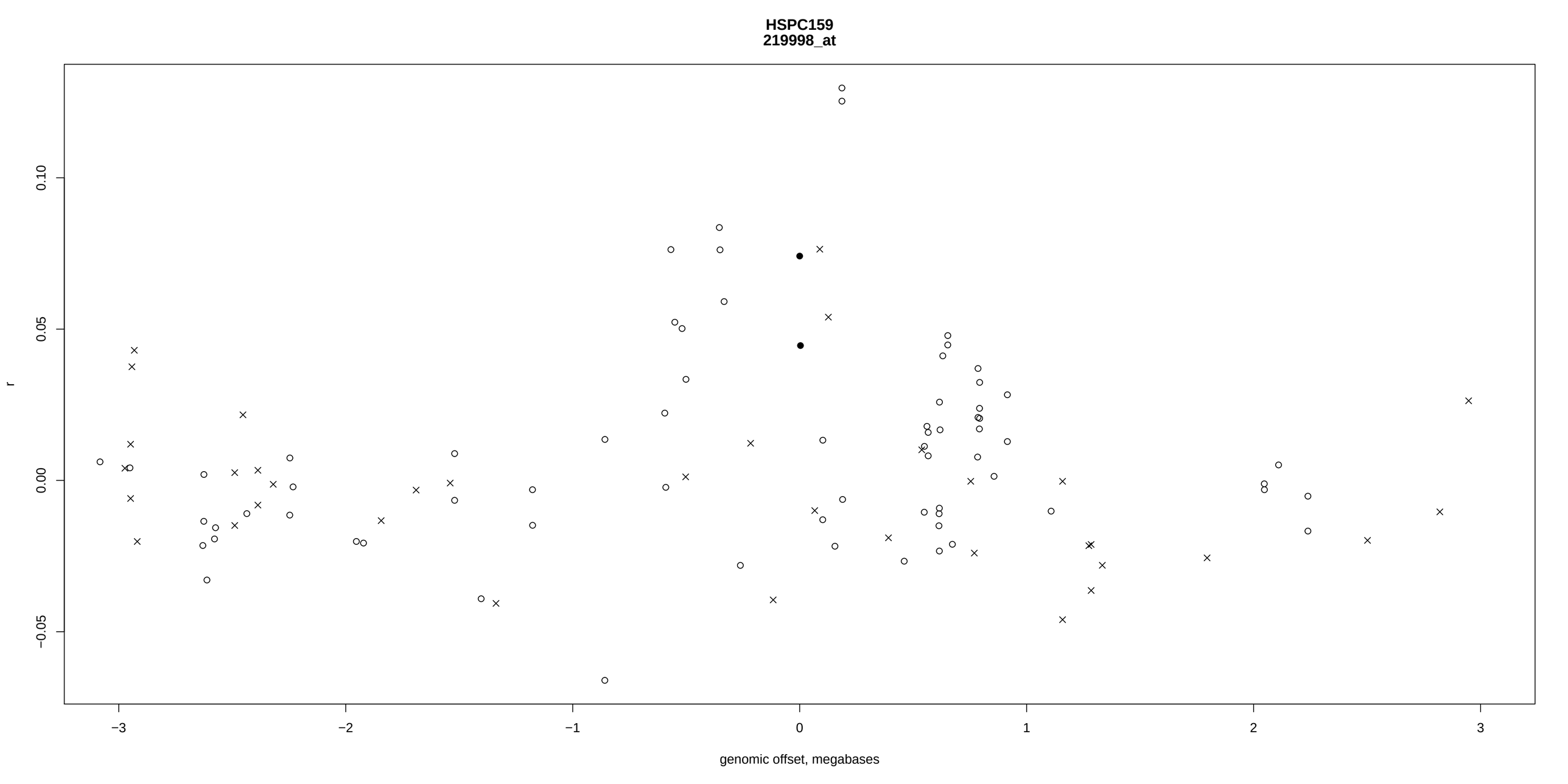

244533_at
244533_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases

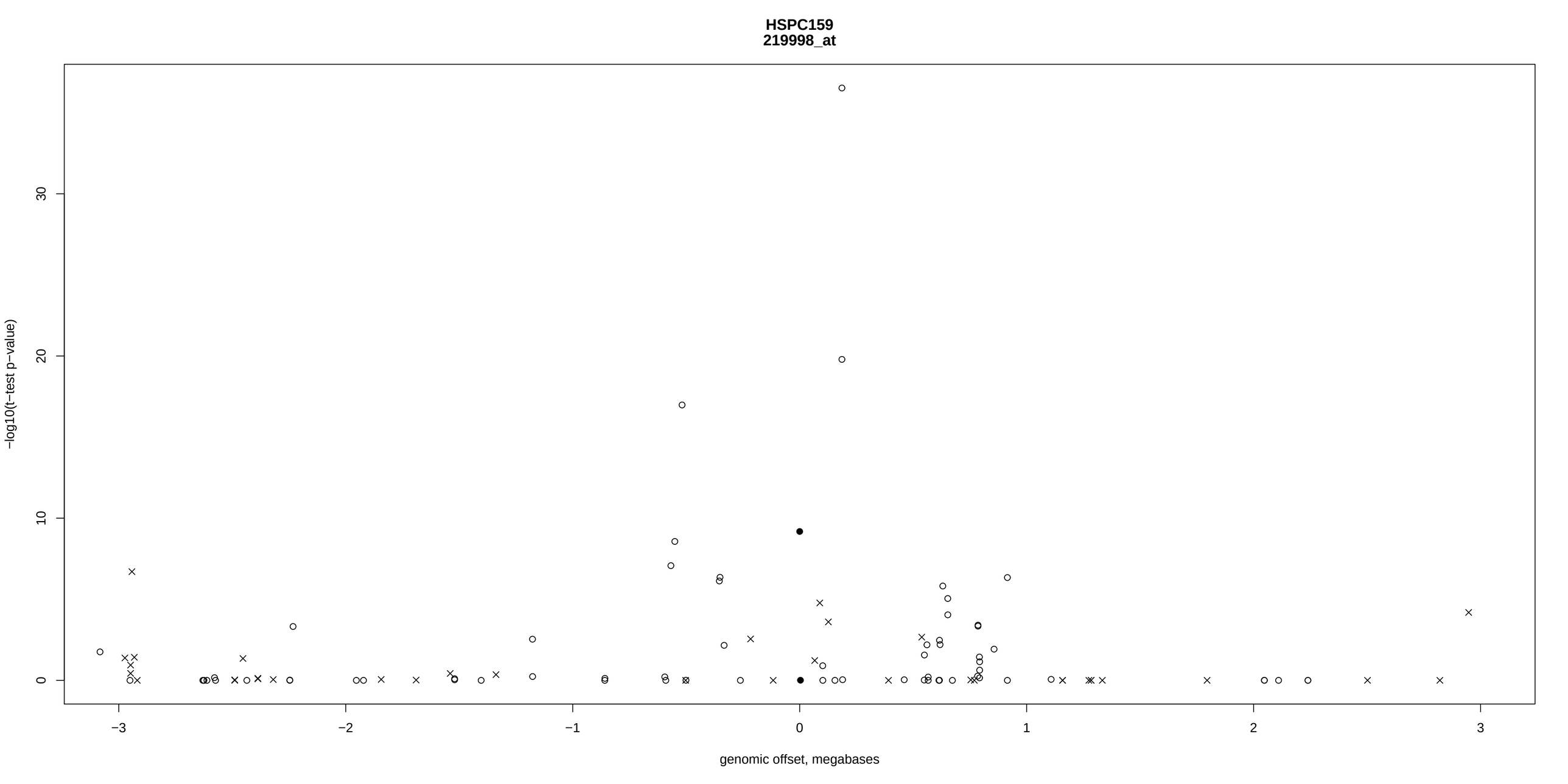

MATN3
206091_at
r
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases

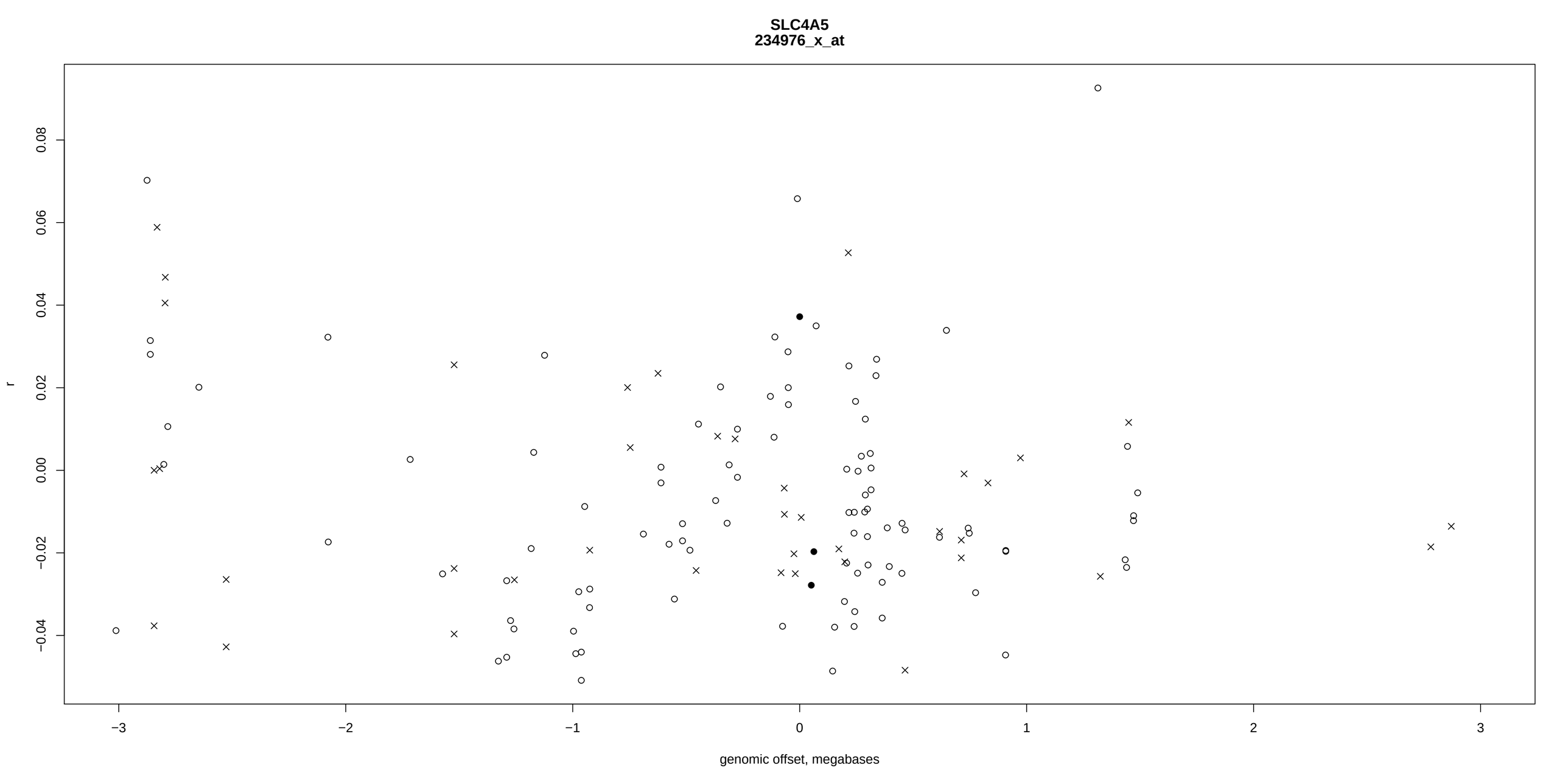

MATN3
206091_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases

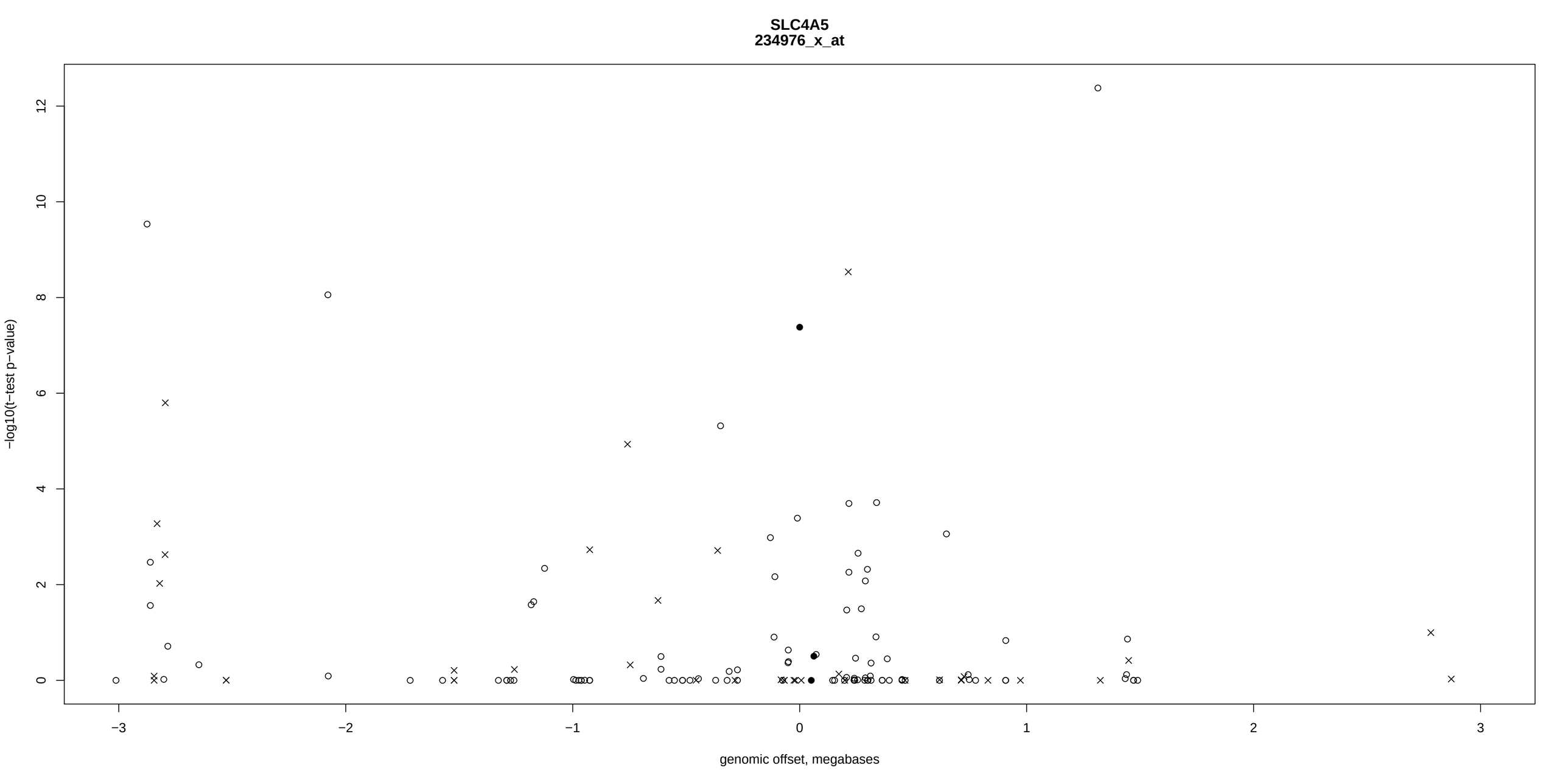

RHOQ
239258_at
r
0.00
0.05
0.10
−3
−2
−1
0
1
2
3
genomic offset, megabases

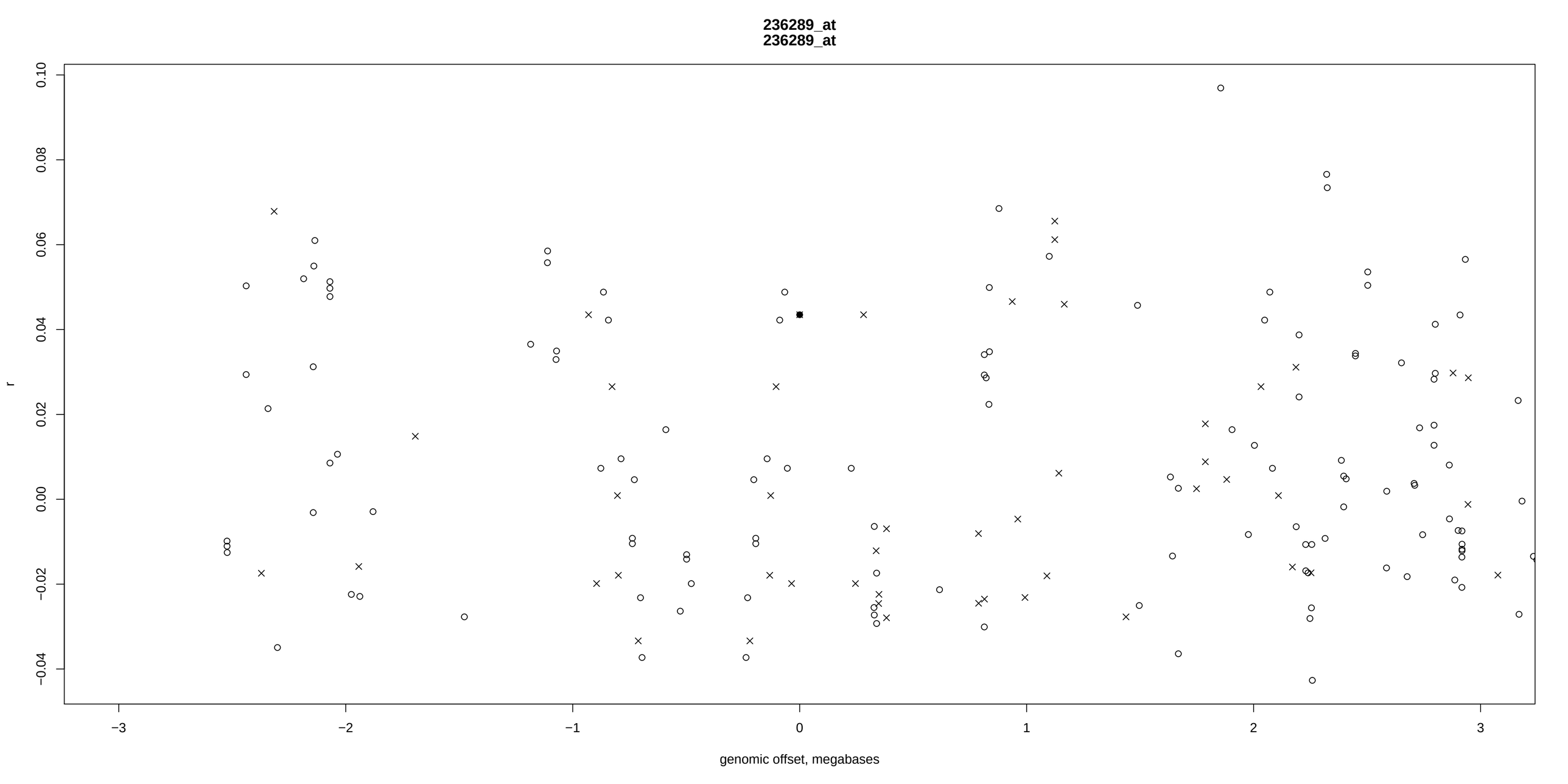

RHOQ
239258_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

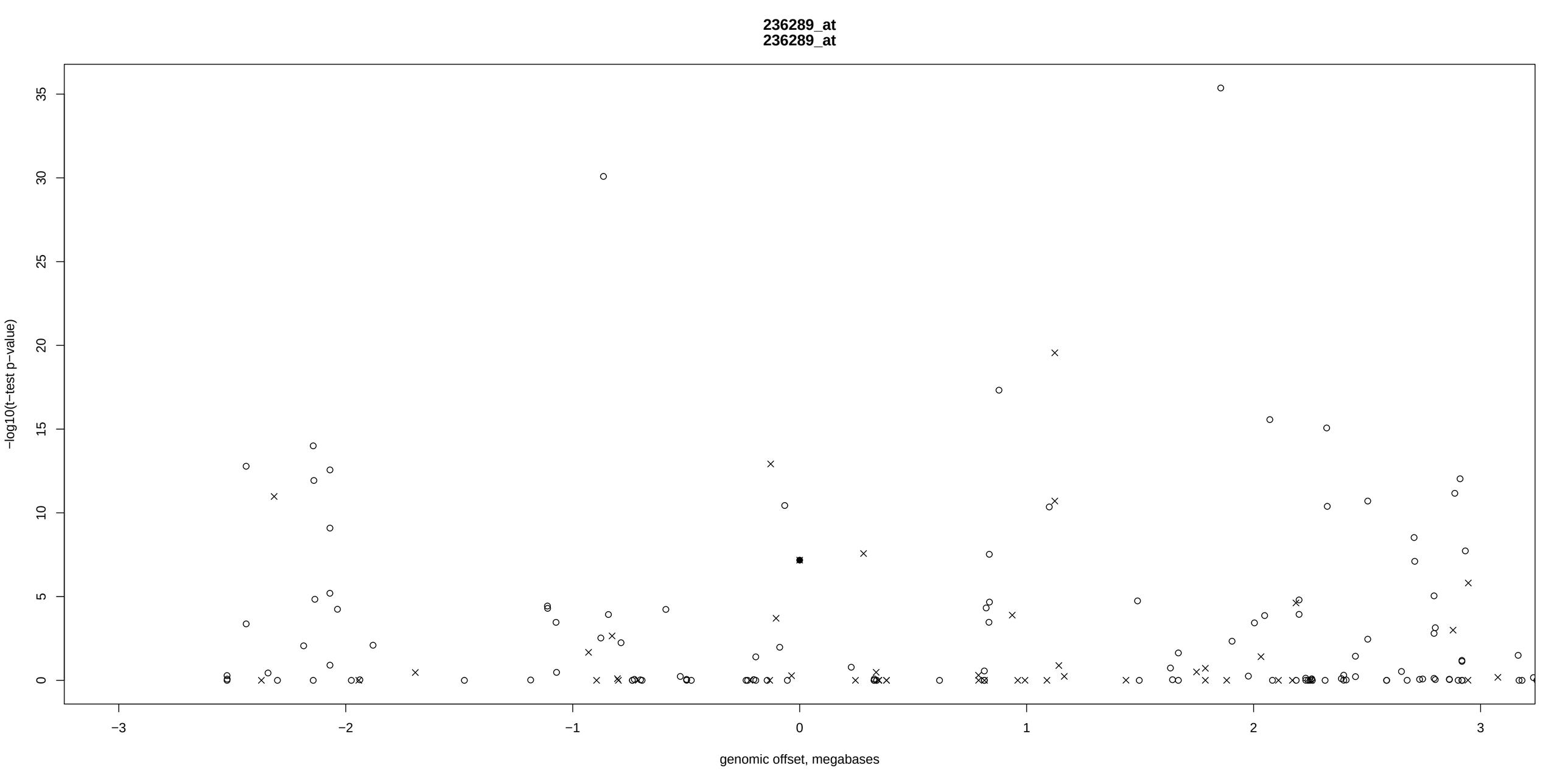

HSPC159
219998_at
r
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases

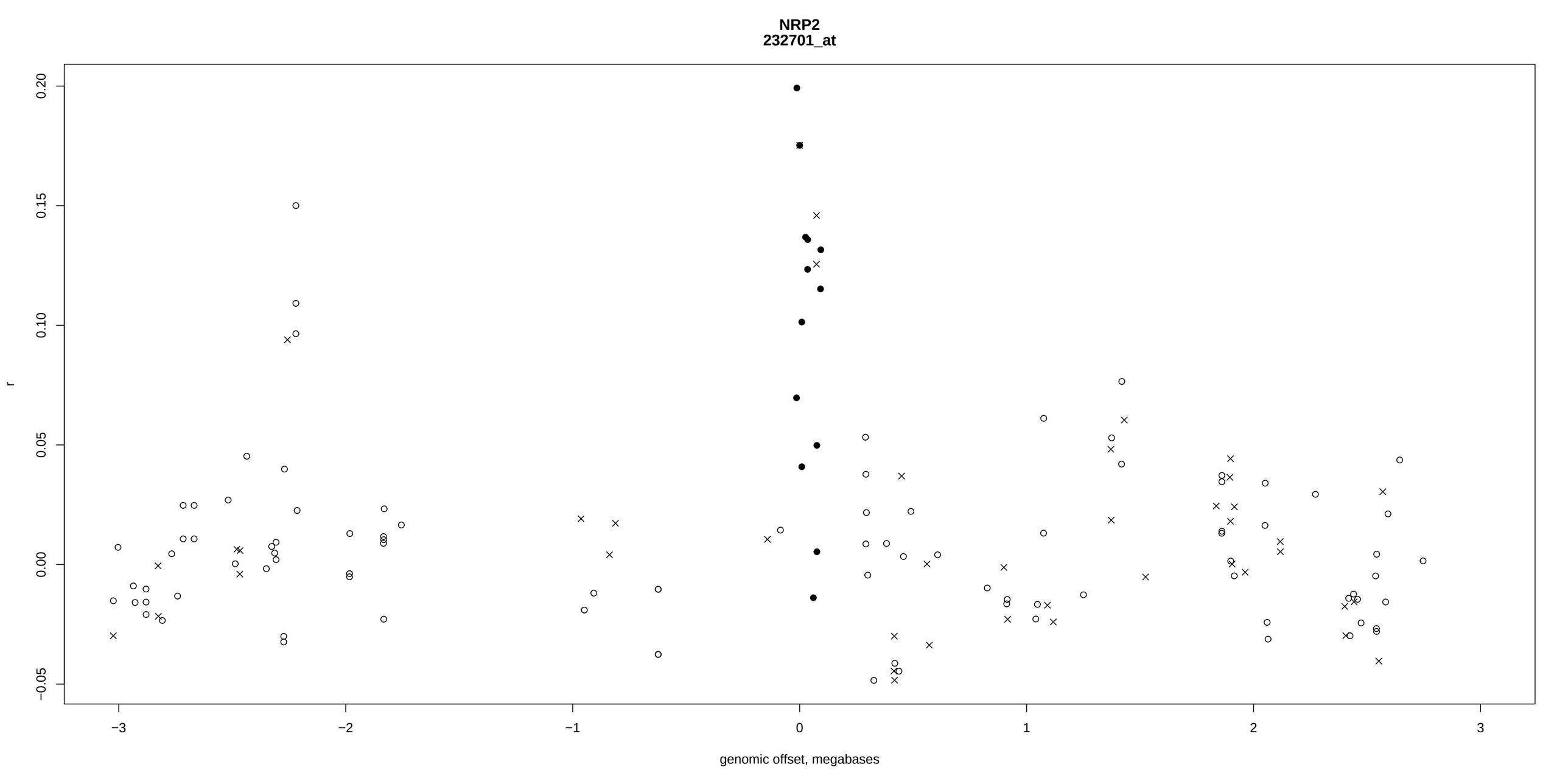

HSPC159
219998_at
−log10(t−test p−value)
0
10
20
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


SLC4A5
234976_x_at
r
0.08
0.06
0.04
0.02
0.00
−0.02
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases


SLC4A5
234976_x_at
−log10(t−test p−value)
0
2
4
6
8
10
12
−3
−2
−1
0
1
2
3
genomic offset, megabases


236289_at
236289_at
r
0.10
0.08
0.06
0.04
0.02
0.00
−0.02
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases


236289_at
236289_at
−log10(t−test p−value)
0
5
10
15
20
25
30
35
−3
−2
−1
0
1
2
3
genomic offset, megabases


NRP2
232701_at
r
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0.05
0.00
−0.05
−3
−2
−1
0
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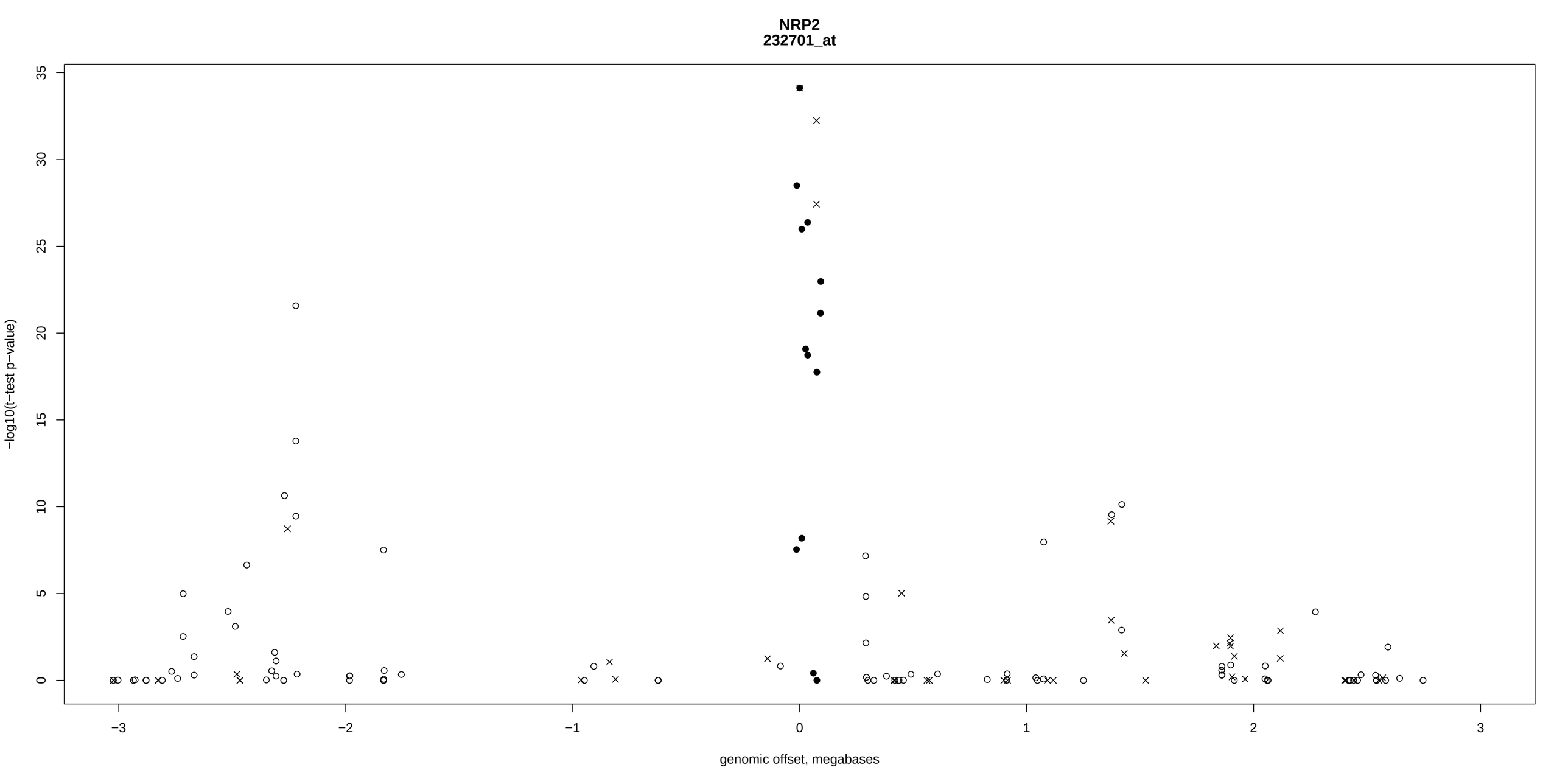
2
3
genomic offset, megabases


NRP2
232701_at
−log10(t−test p−value)
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15
20
25
30
35
−3
−2
−1
0
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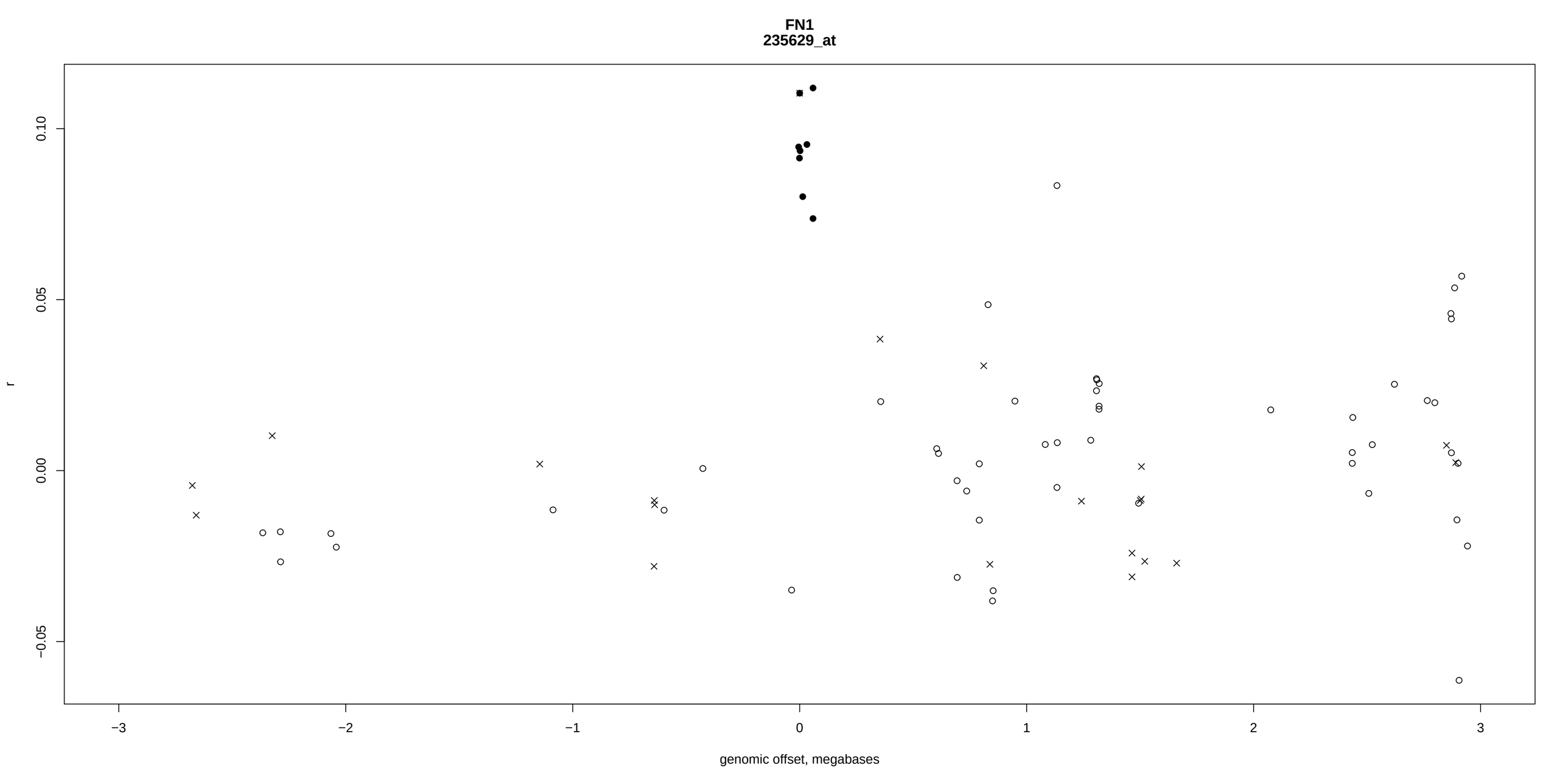
2
3
genomic offset, megabases


FN1
235629_at
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−1
0
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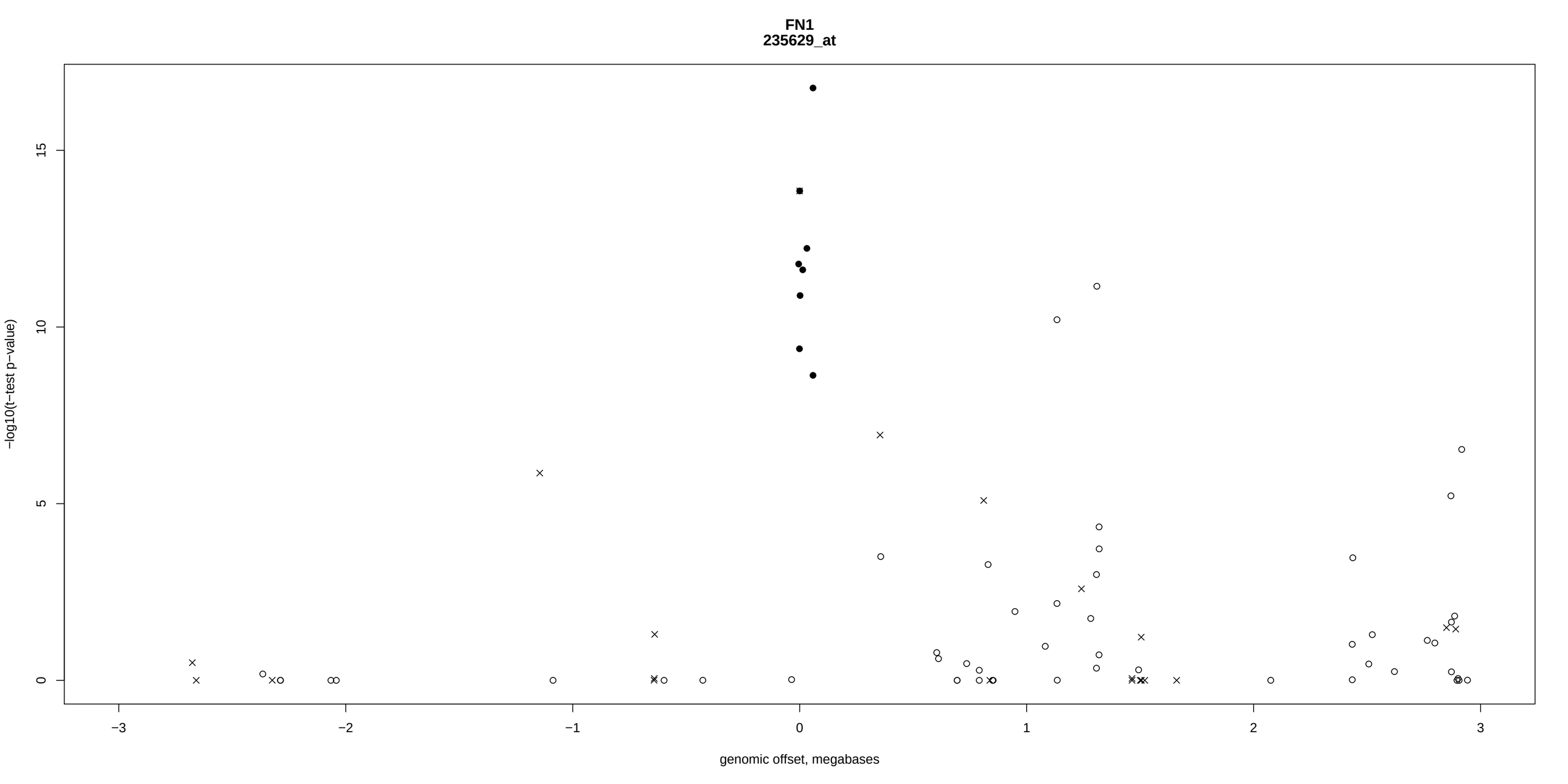
3
genomic offset, megabases


FN1
235629_at
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−3
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−1
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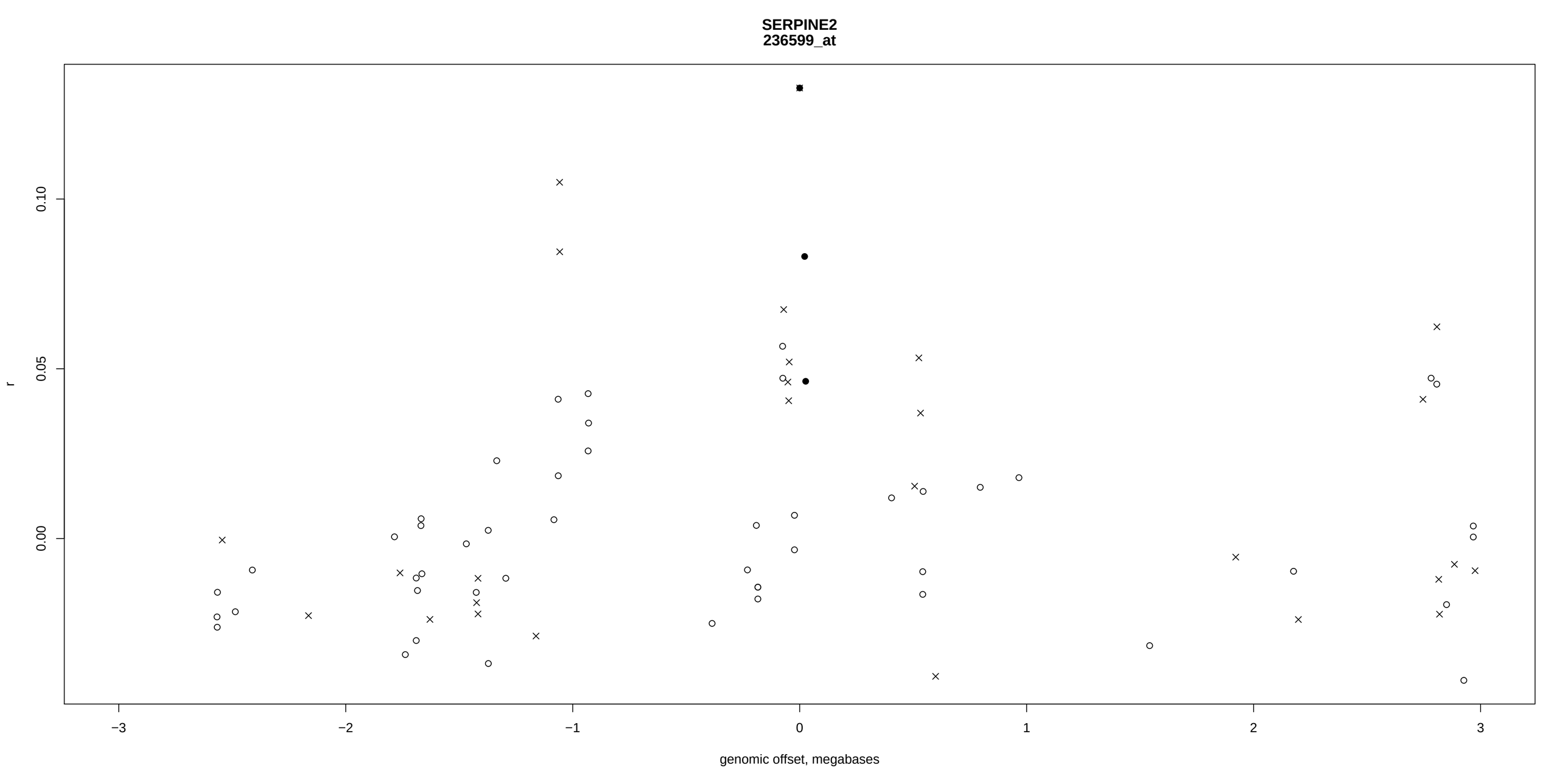
3
genomic offset, megabases


SERPINE2
236599_at
r
0.00
0.05
0.10
−3
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−1
0
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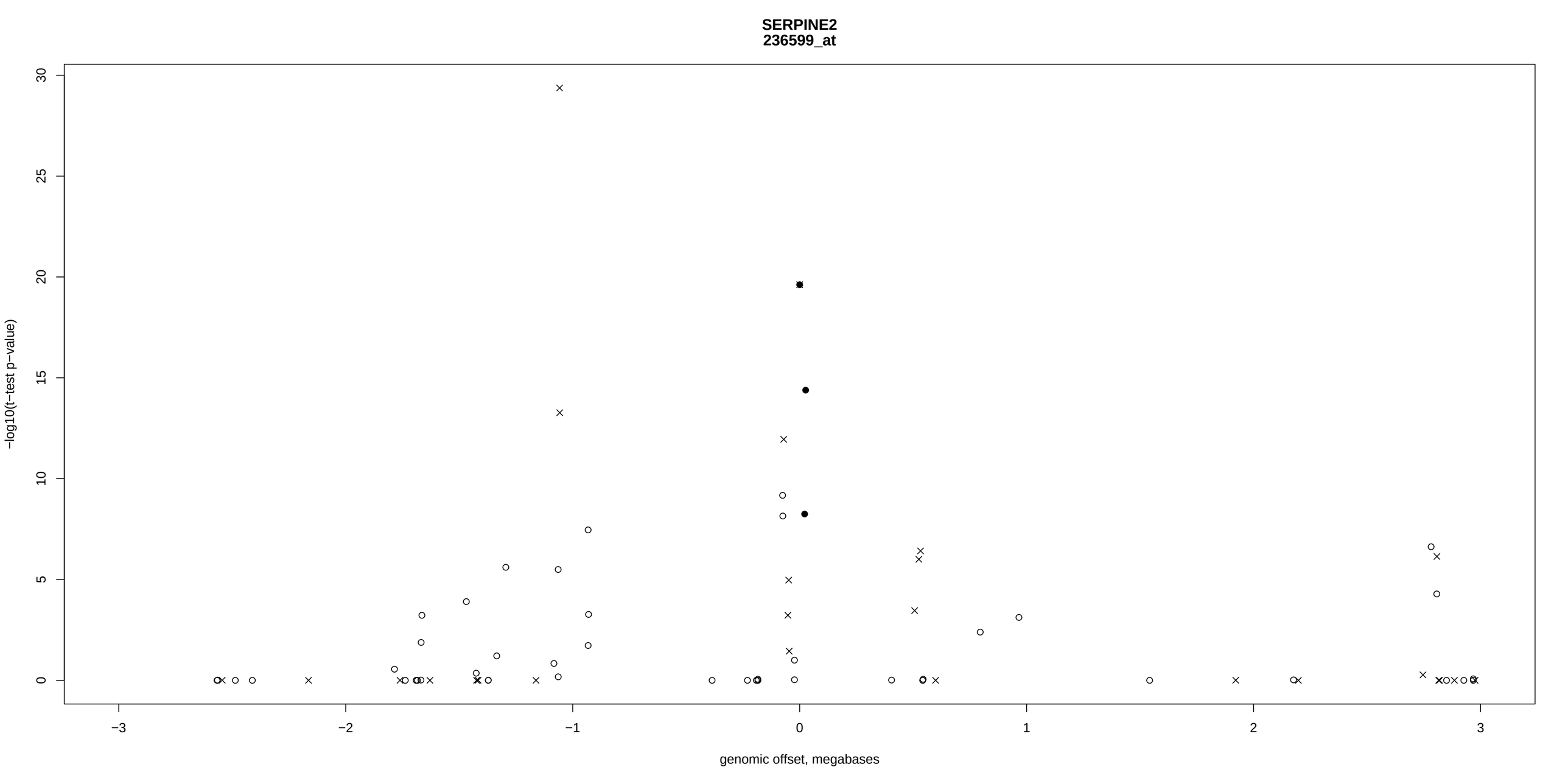
2
3
genomic offset, megabases


SERPINE2
236599_at
−log10(t−test p−value)
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5
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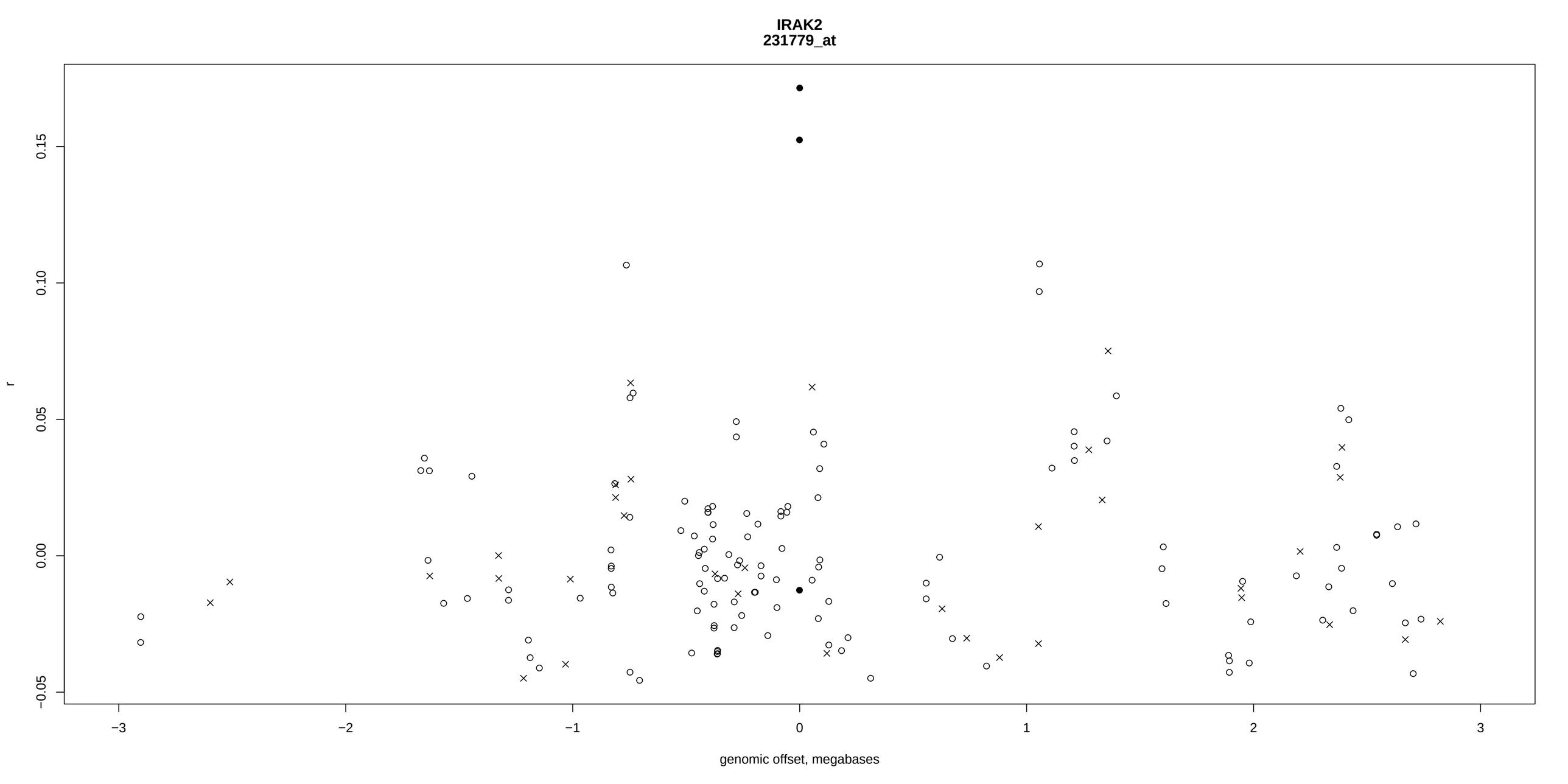
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


IRAK2
231779_at
r
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0.05
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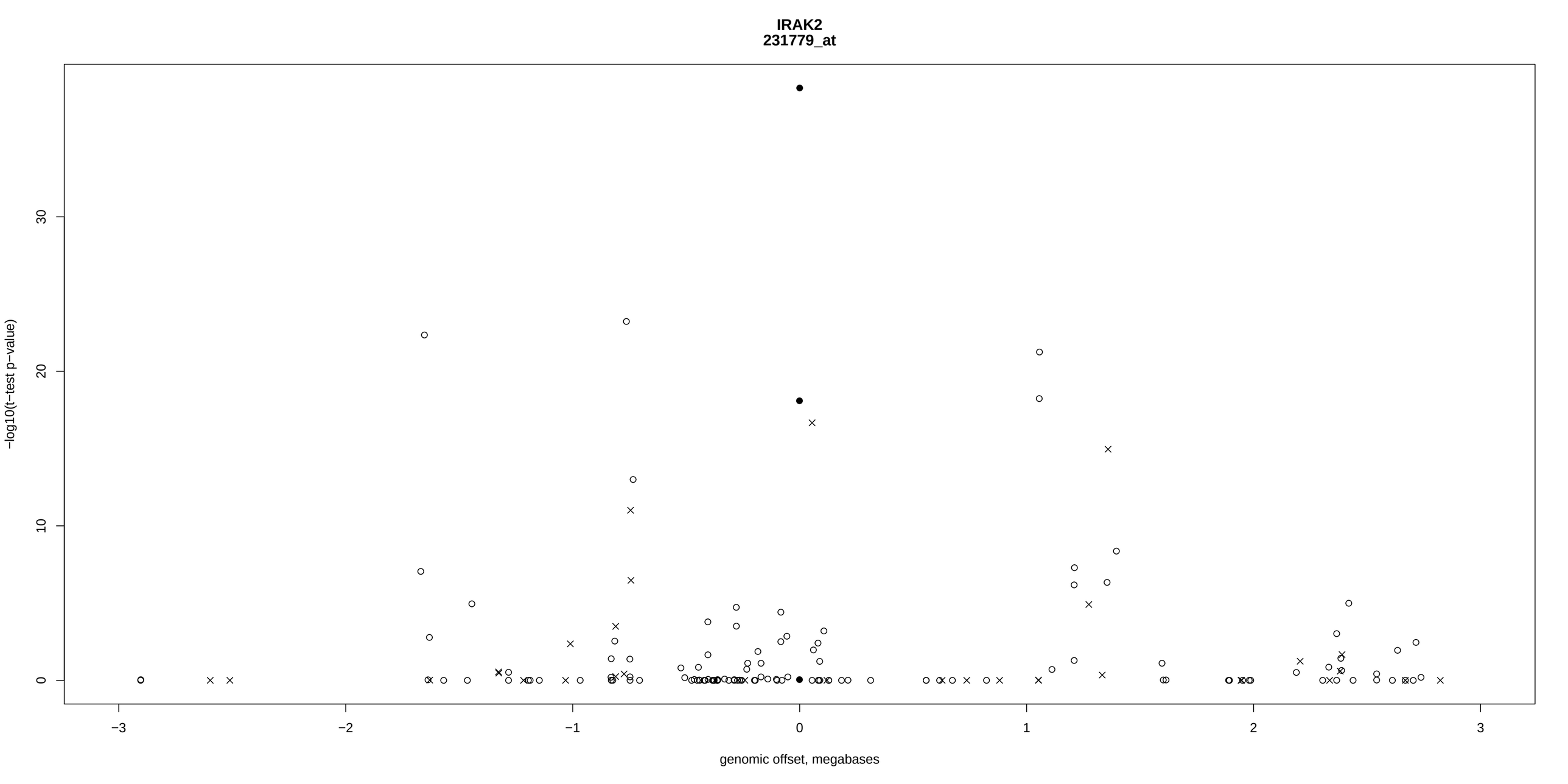
3
genomic offset, megabases


IRAK2
231779_at
−log10(t−test p−value)
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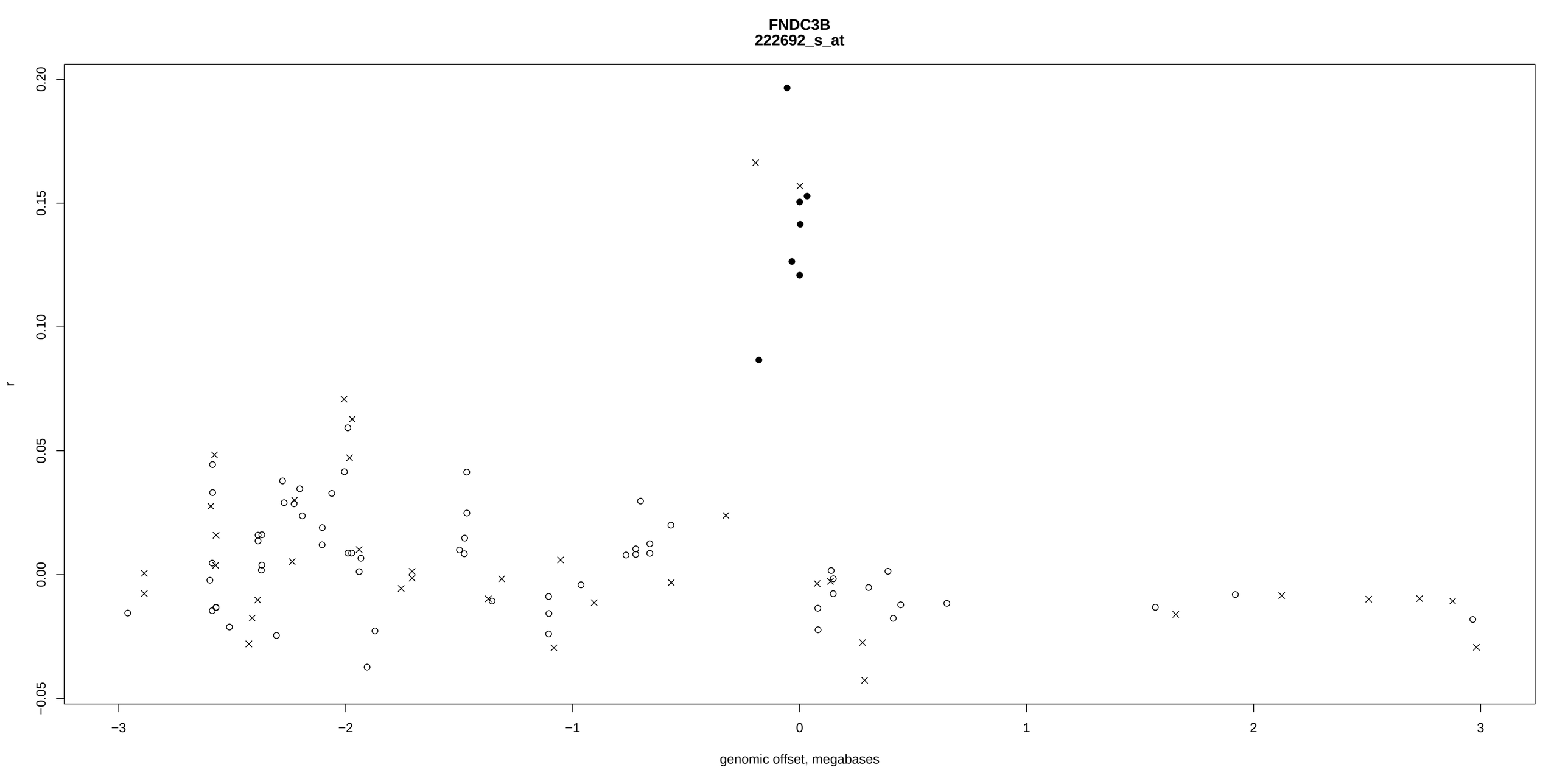
10
20
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


FNDC3B
222692_s_at
r
0.20
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0.05
0.00
−0.05
−3
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−1
0
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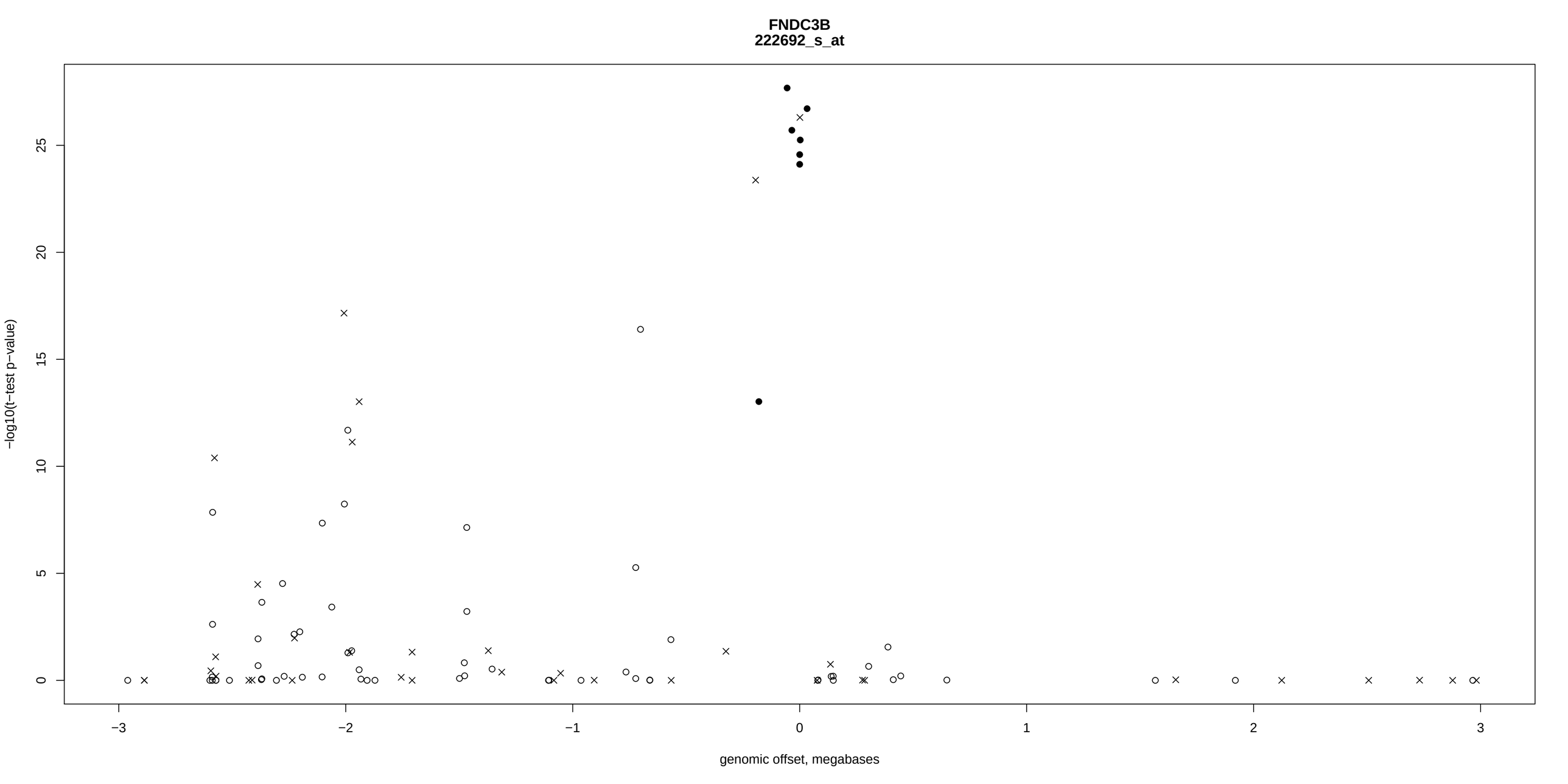
3
genomic offset, megabases


FNDC3B
222692_s_at
−log10(t−test p−value)
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10
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−3
−2
−1
0
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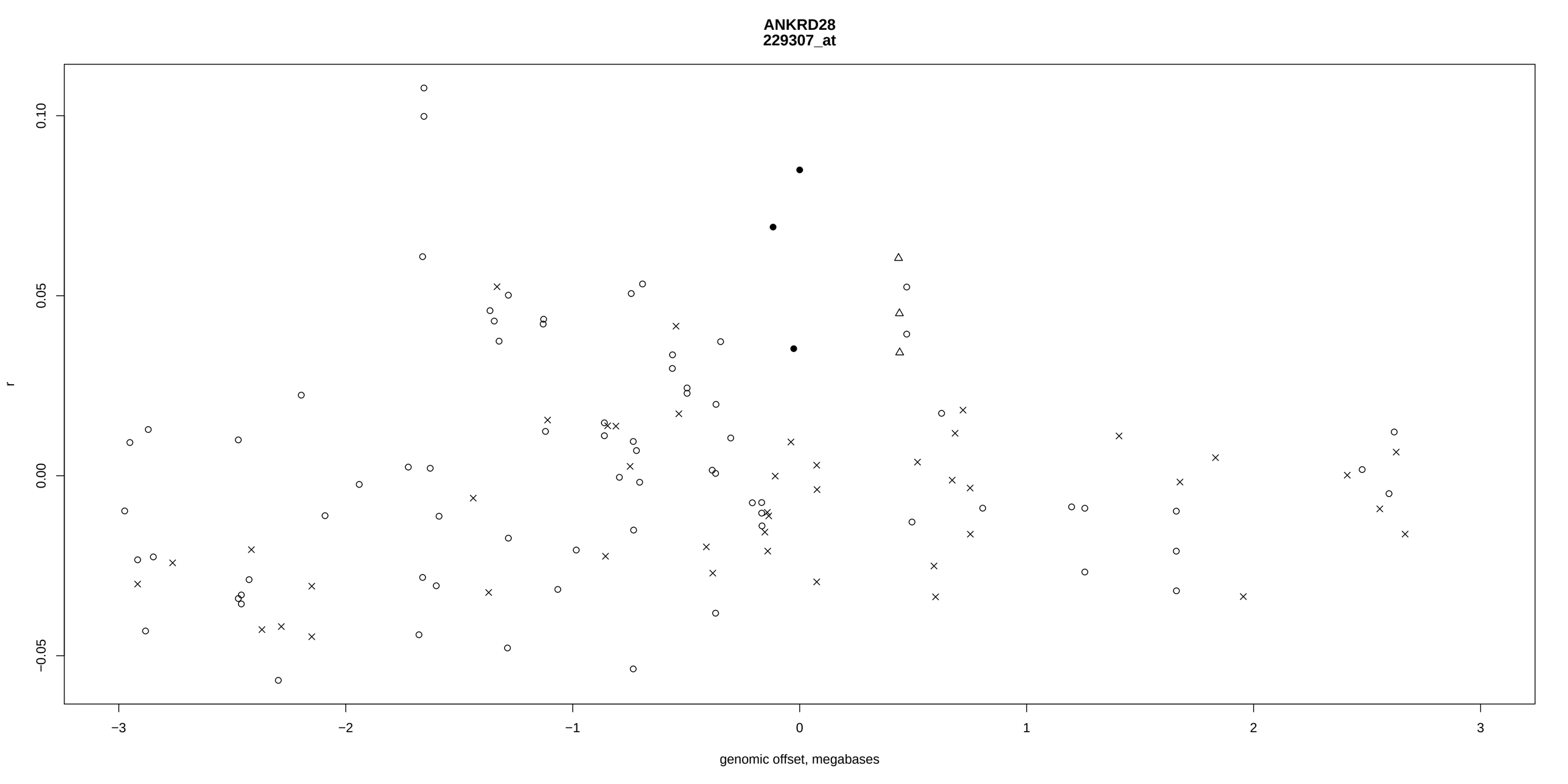
2
3
genomic offset, megabases


ANKRD28
229307_at
0.10
0.05
0.00
−0.05
r
−3
−2
−1
0
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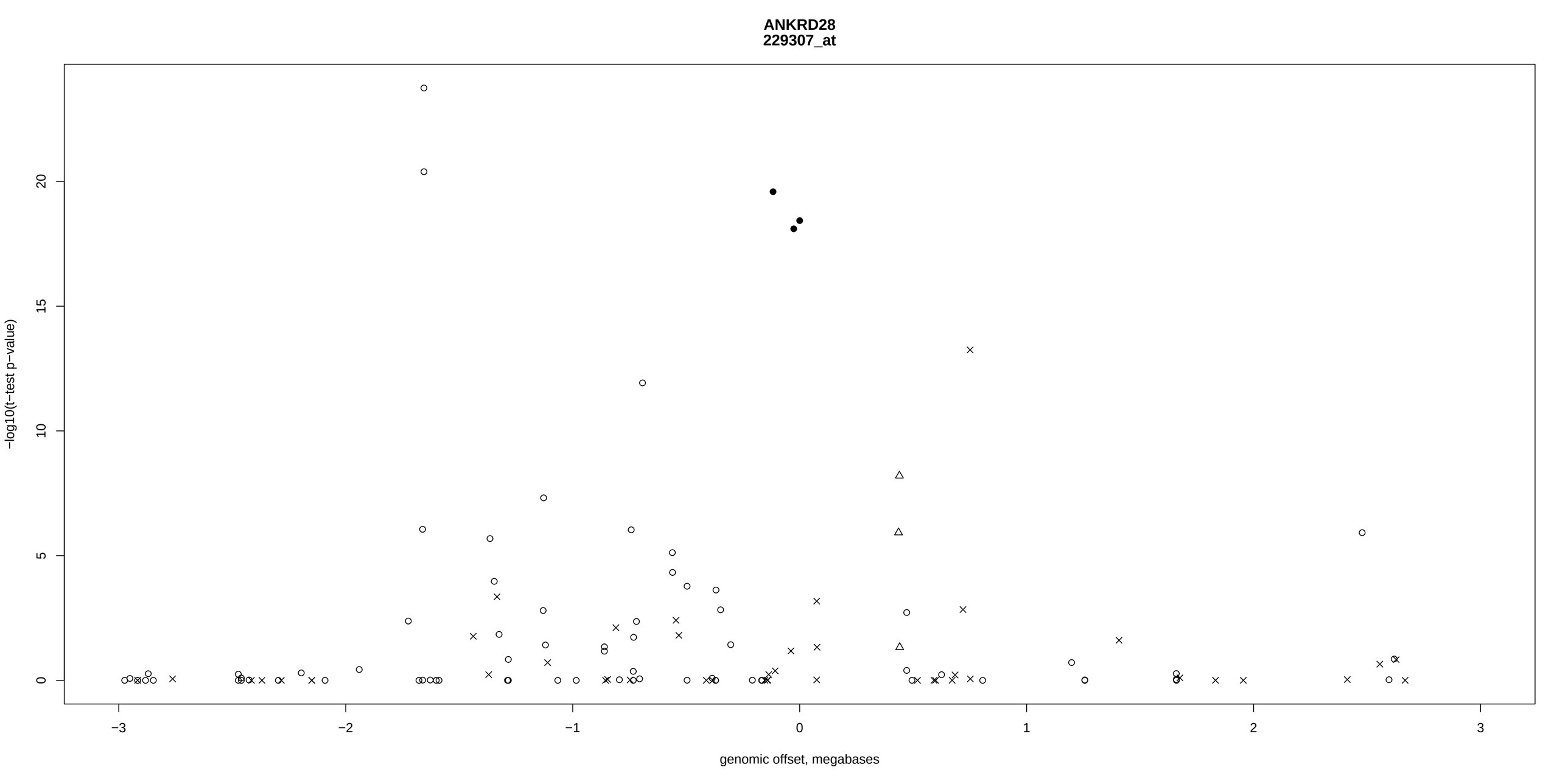
2
3
genomic offset, megabases


ANKRD28
229307_at
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5
10
15
20
−3
−2
−1
0
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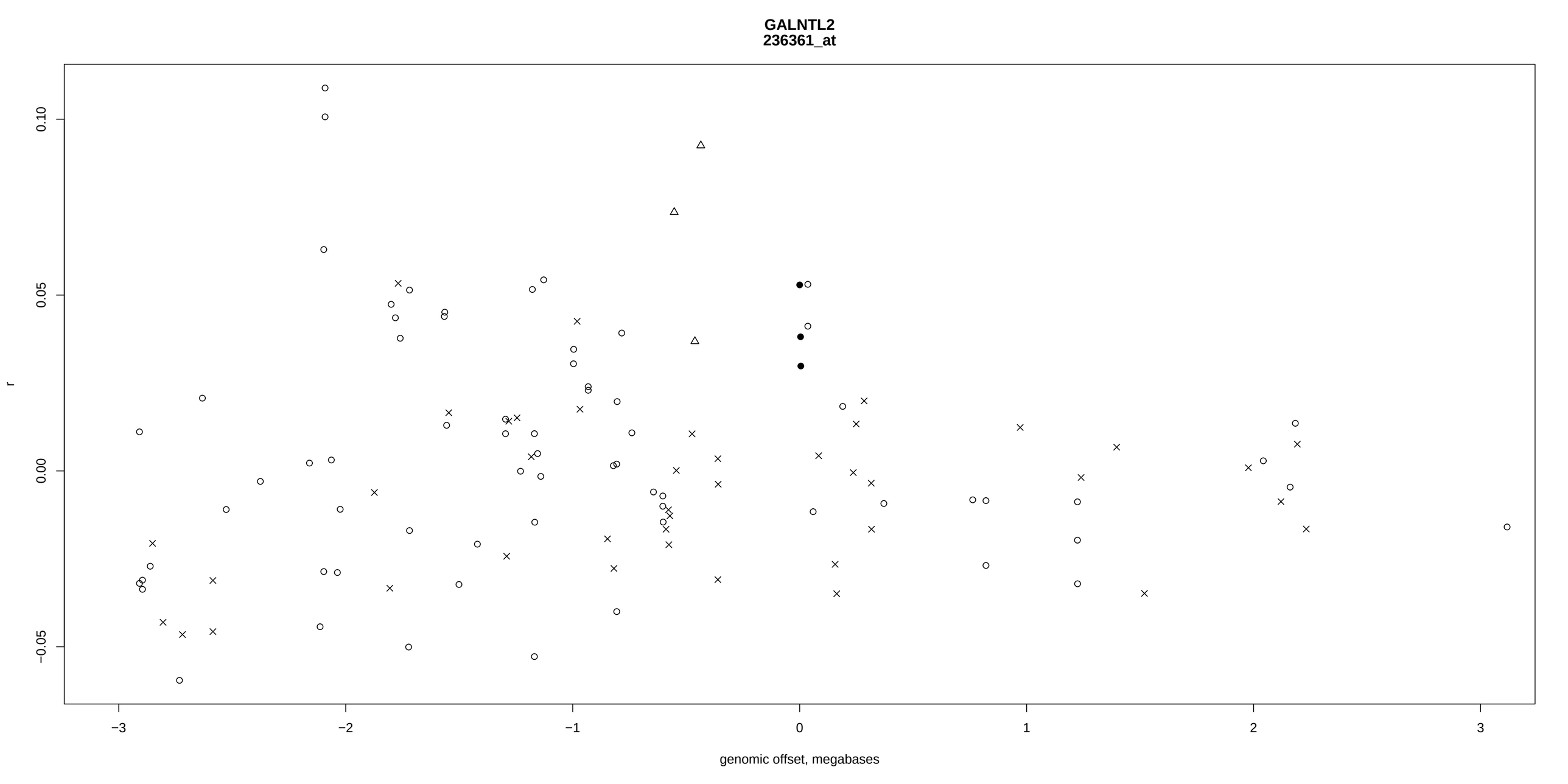
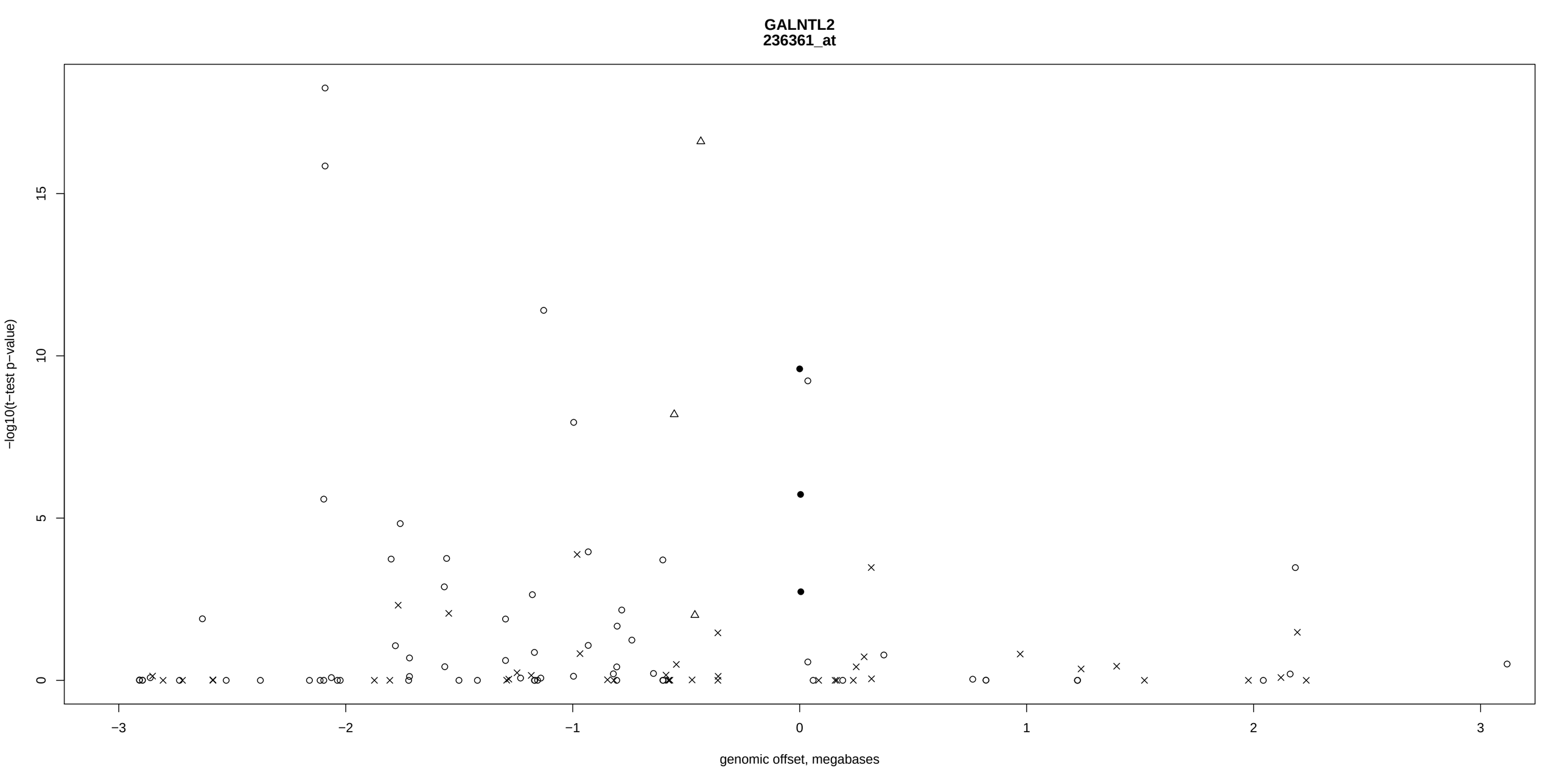
2
3
genomic offset, megabases


GALNTL2
236361_at
0.10
0.05
0.00
−0.05
r
−3
−2
−1
0
1
2
3
genomic offset, megabases


GALNTL2
236361_at
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−3
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−1
0
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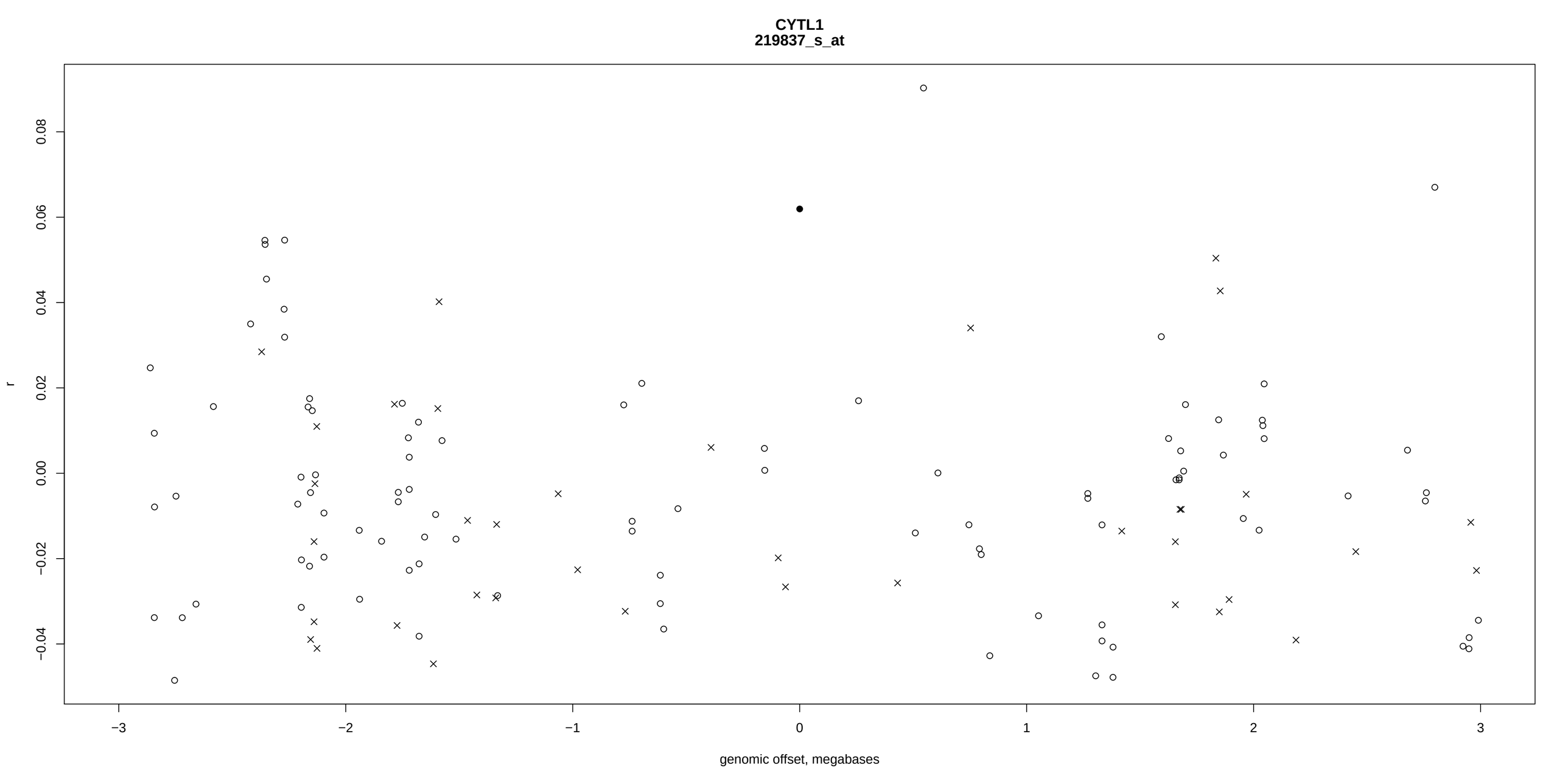
2
3
genomic offset, megabases


CYTL1
219837_s_at
r
0.08
0.06
0.04
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0.00
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−0.04
−3
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−1
0
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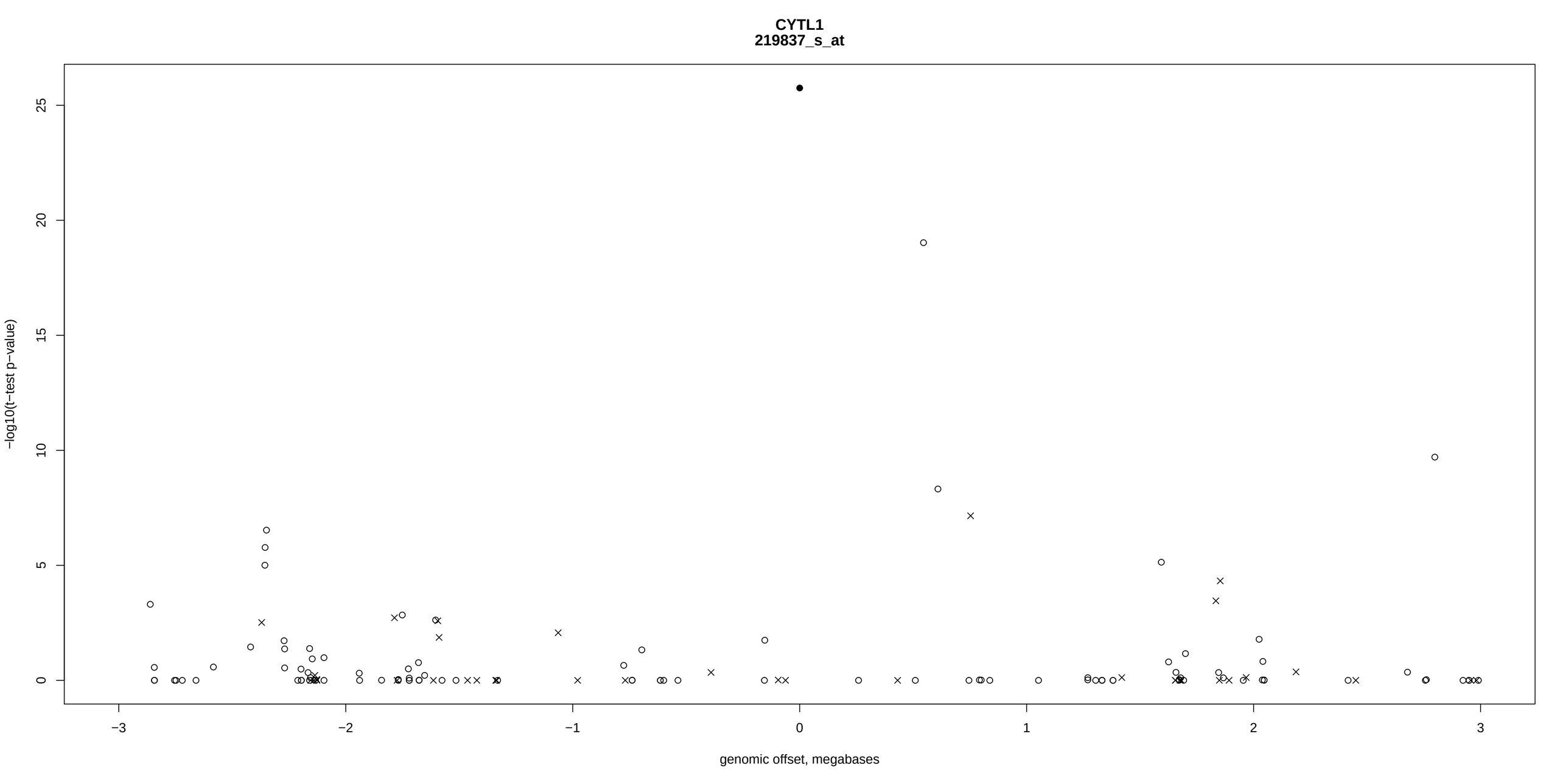
2
3
genomic offset, megabases


CYTL1
219837_s_at
−log10(t−test p−value)
0
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10
15
20
25
−3
−2
−1
0
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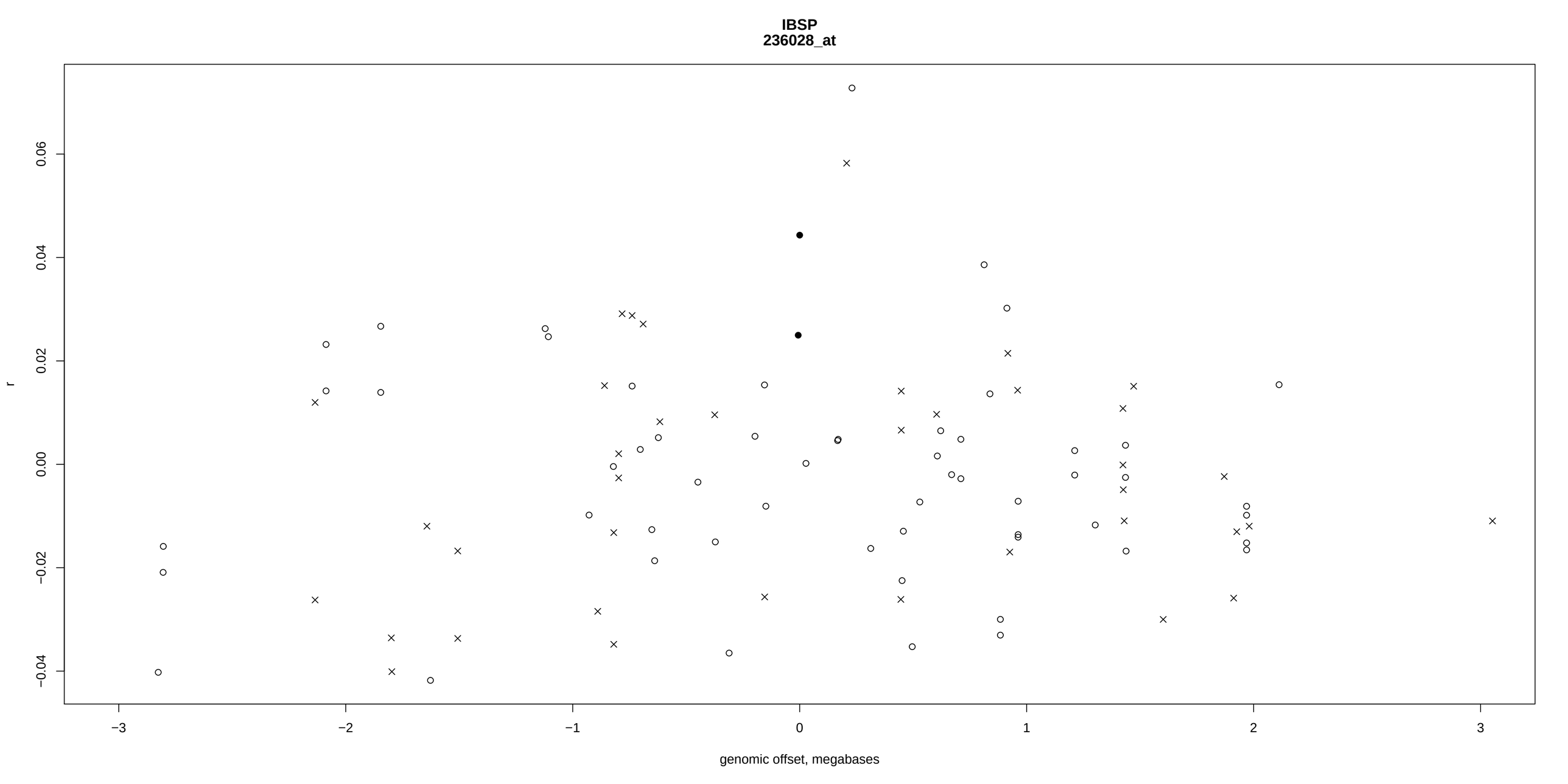
2
3
genomic offset, megabases


IBSP
236028_at
r
0.06
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−0.04
−3
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−1
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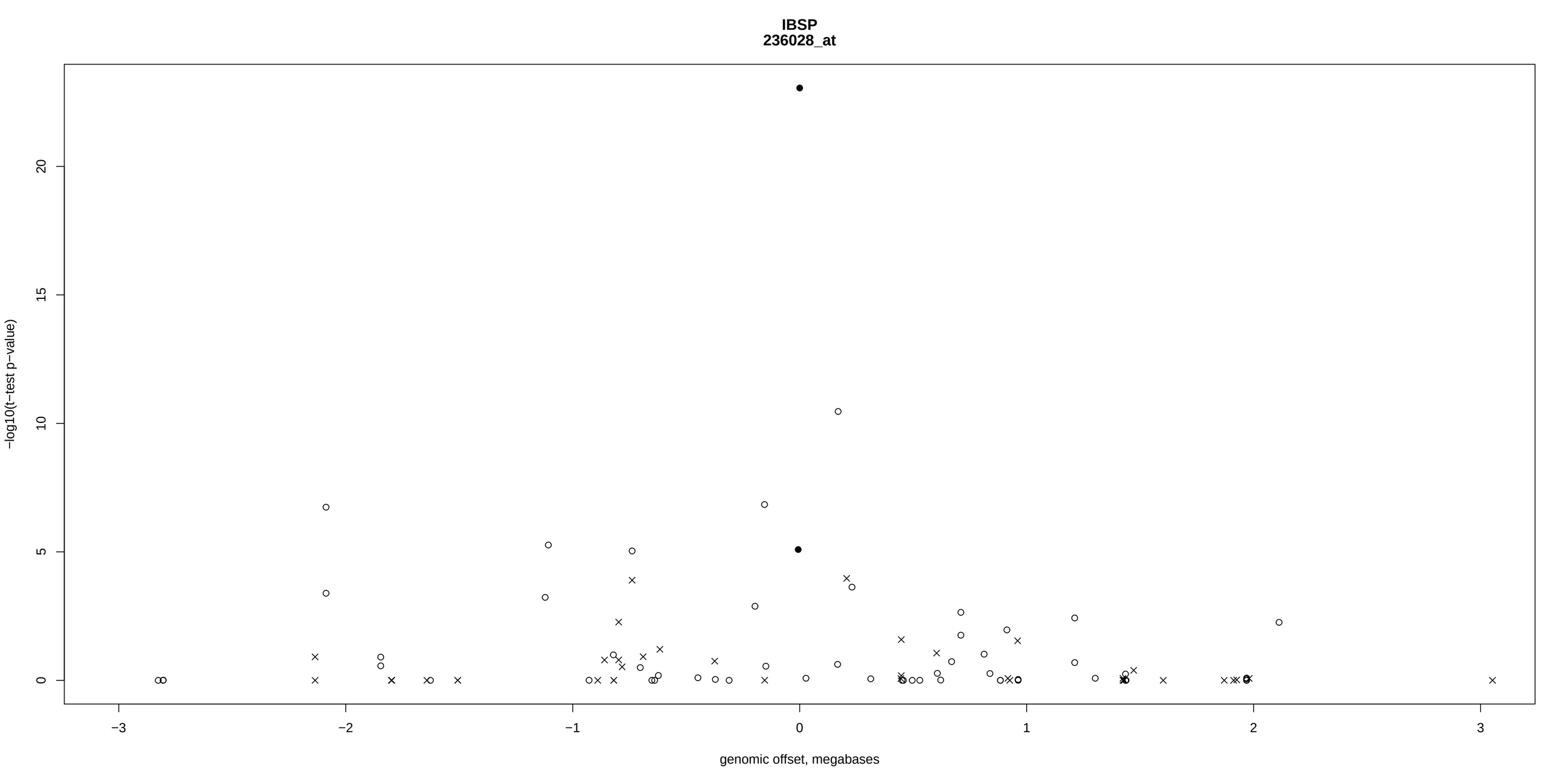
2
3
genomic offset, megabases


IBSP
236028_at
−log10(t−test p−value)
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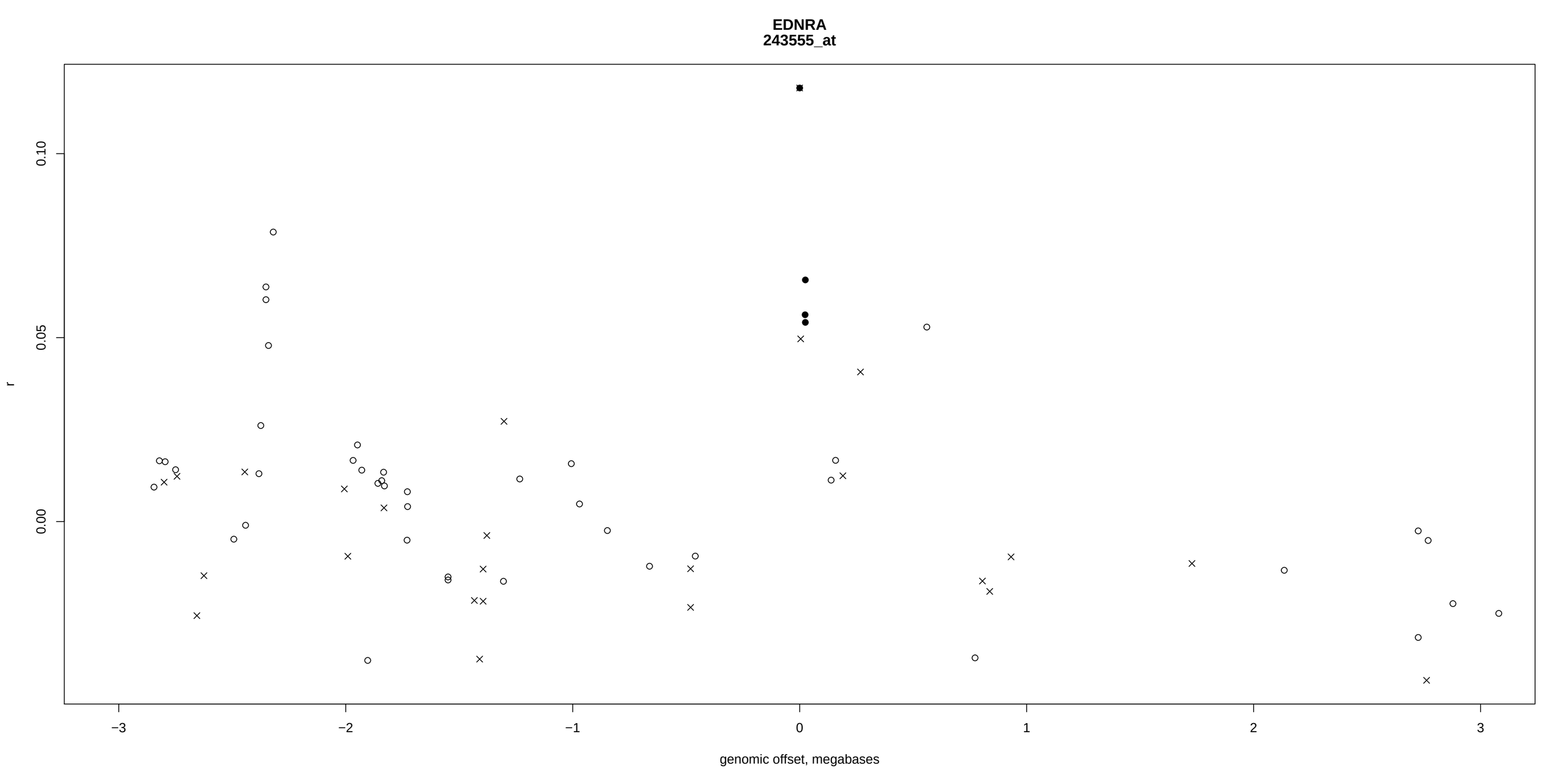
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


EDNRA
243555_at
r
0.10
0.05
0.00
−3
−2
−1
0
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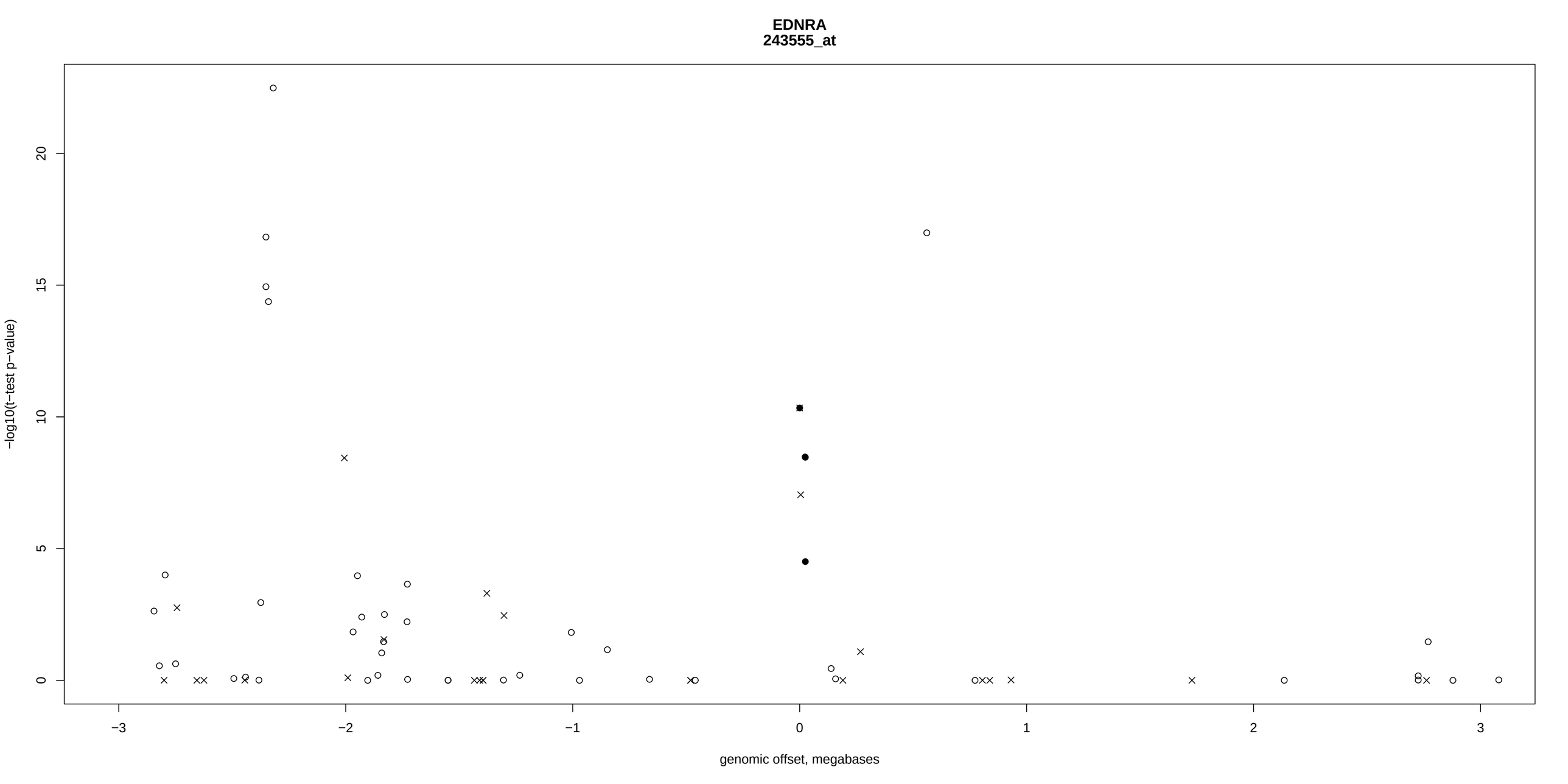
2
3
genomic offset, megabases


EDNRA
243555_at
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0
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20
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−1
0
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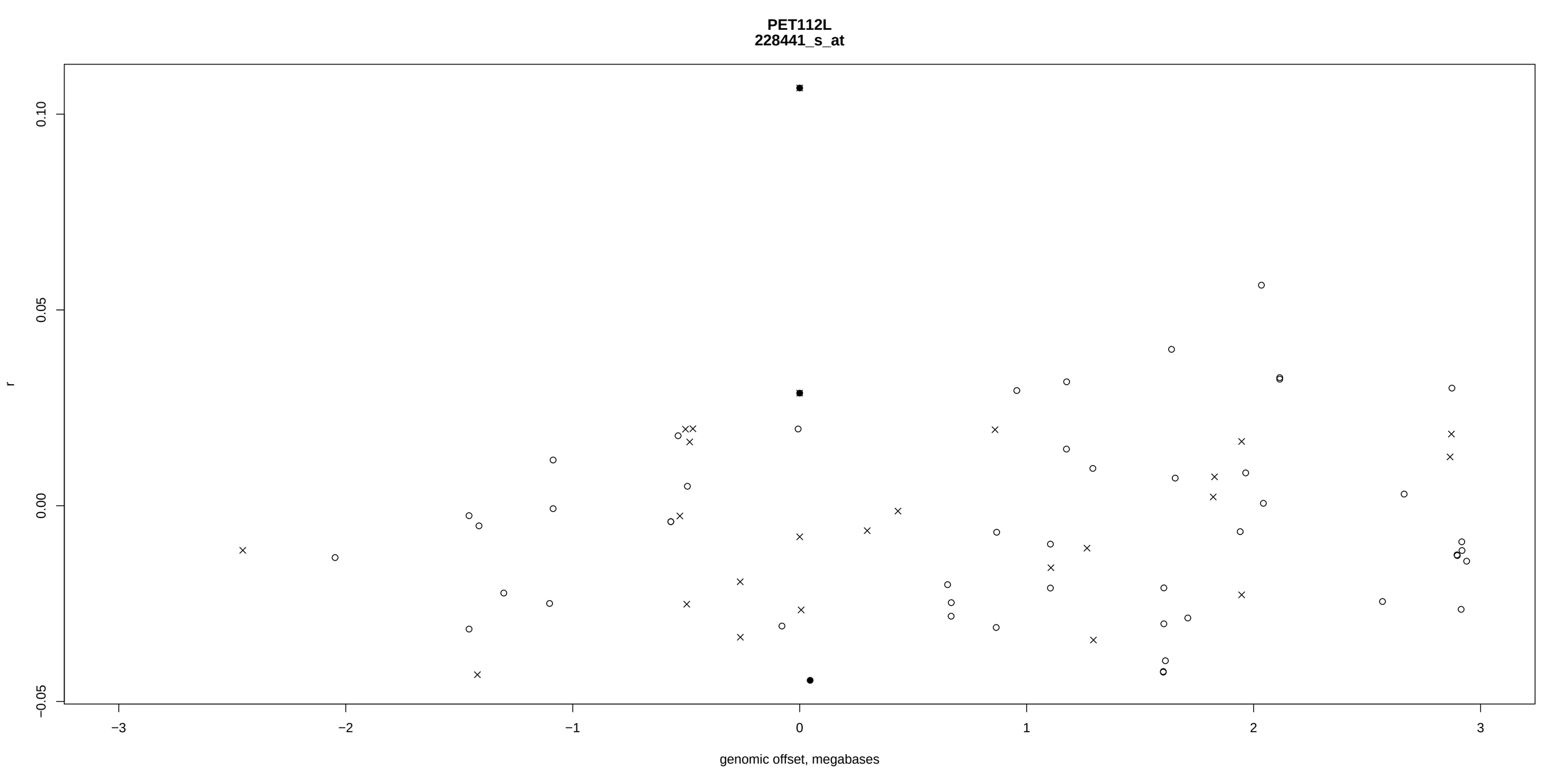
2
3
genomic offset, megabases


PET112L
228441_s_at
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0.00
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−1
0
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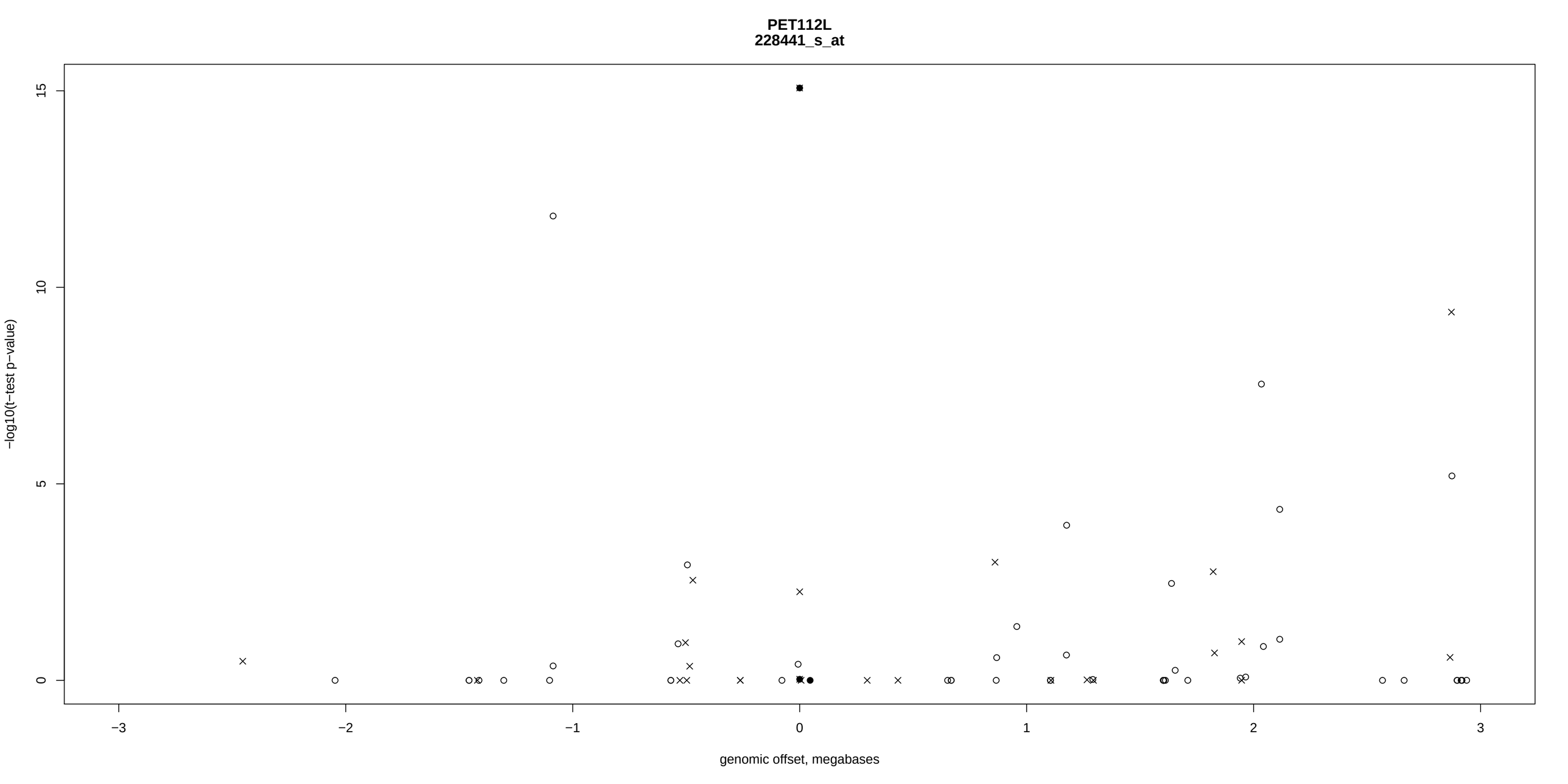
2
3
genomic offset, megabases


PET112L
228441_s_at
−log10(t−test p−value)
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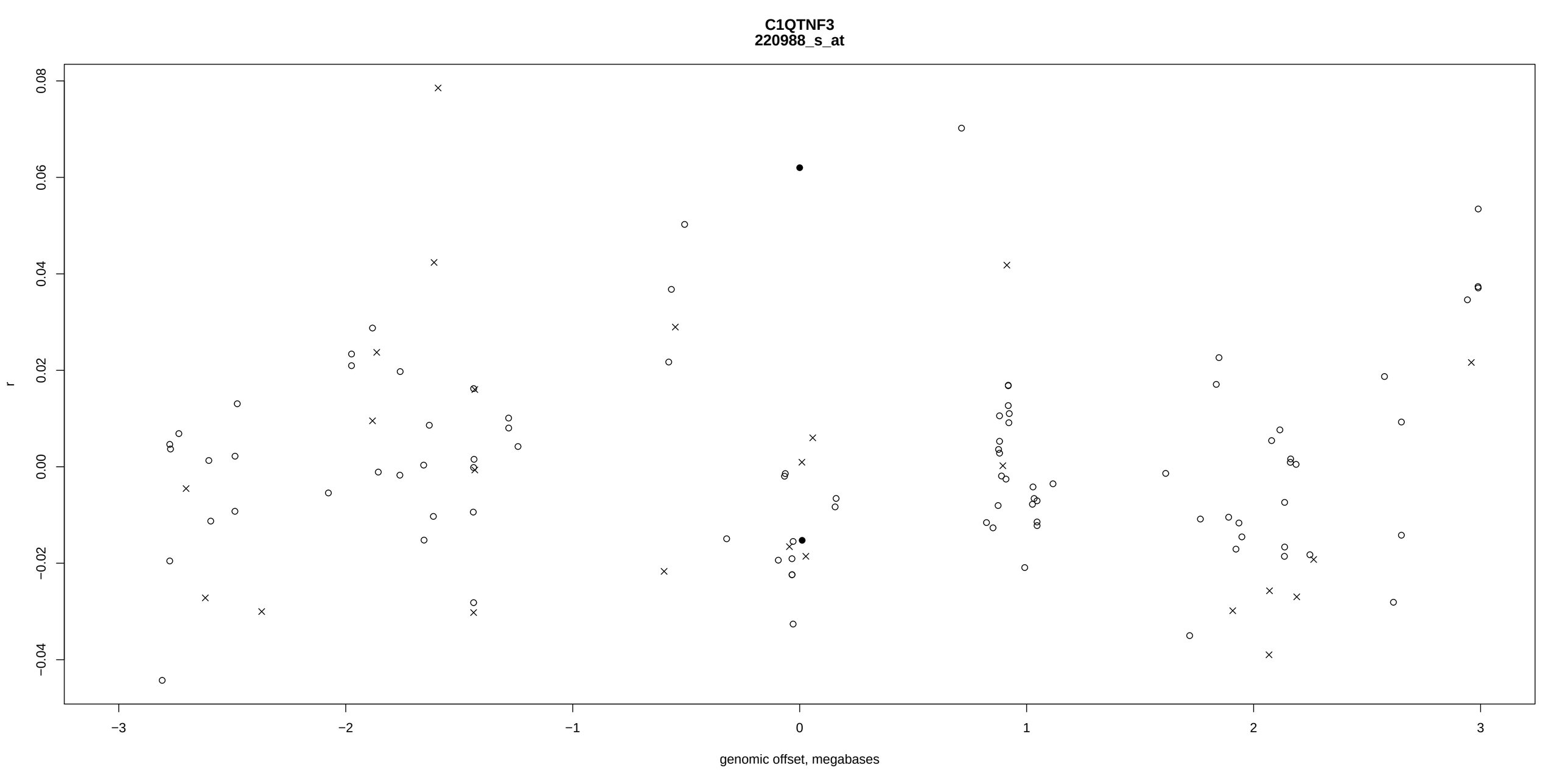
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


C1QTNF3
220988_s_at
r
0.08
0.06
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−0.04
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−1
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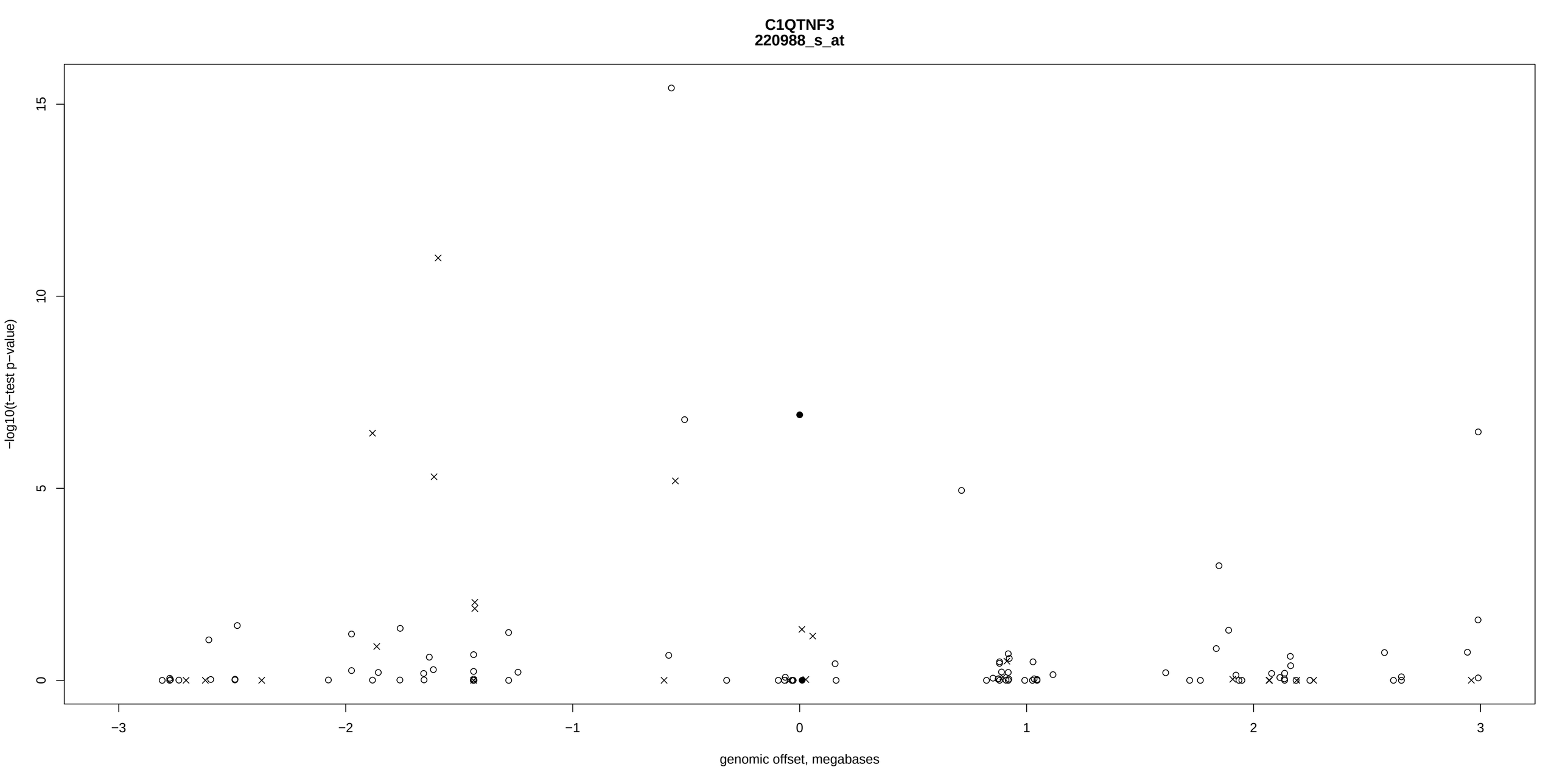
2
3
genomic offset, megabases


C1QTNF3
220988_s_at
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−1
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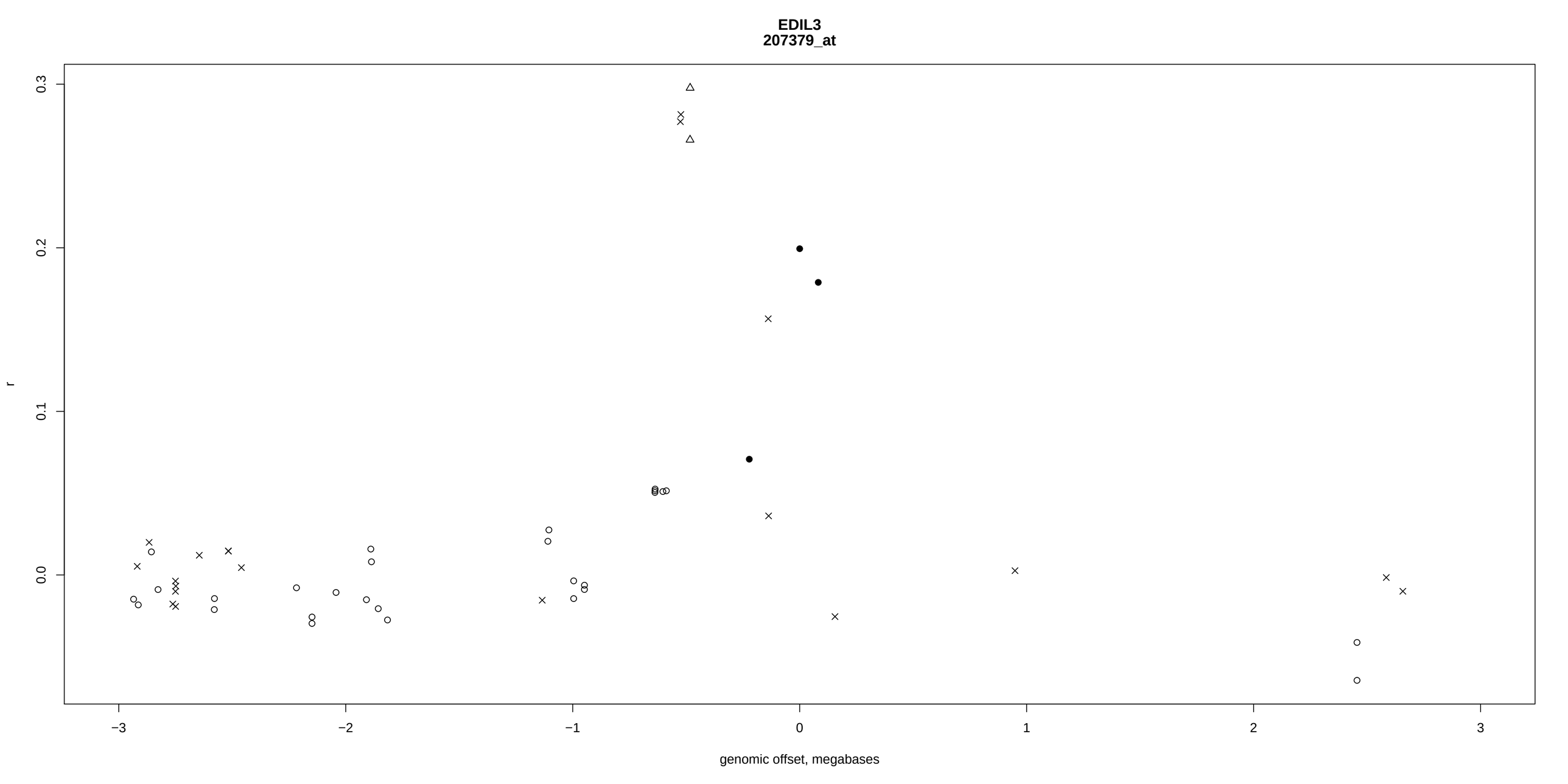
3
genomic offset, megabases


EDIL3
207379_at
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−1
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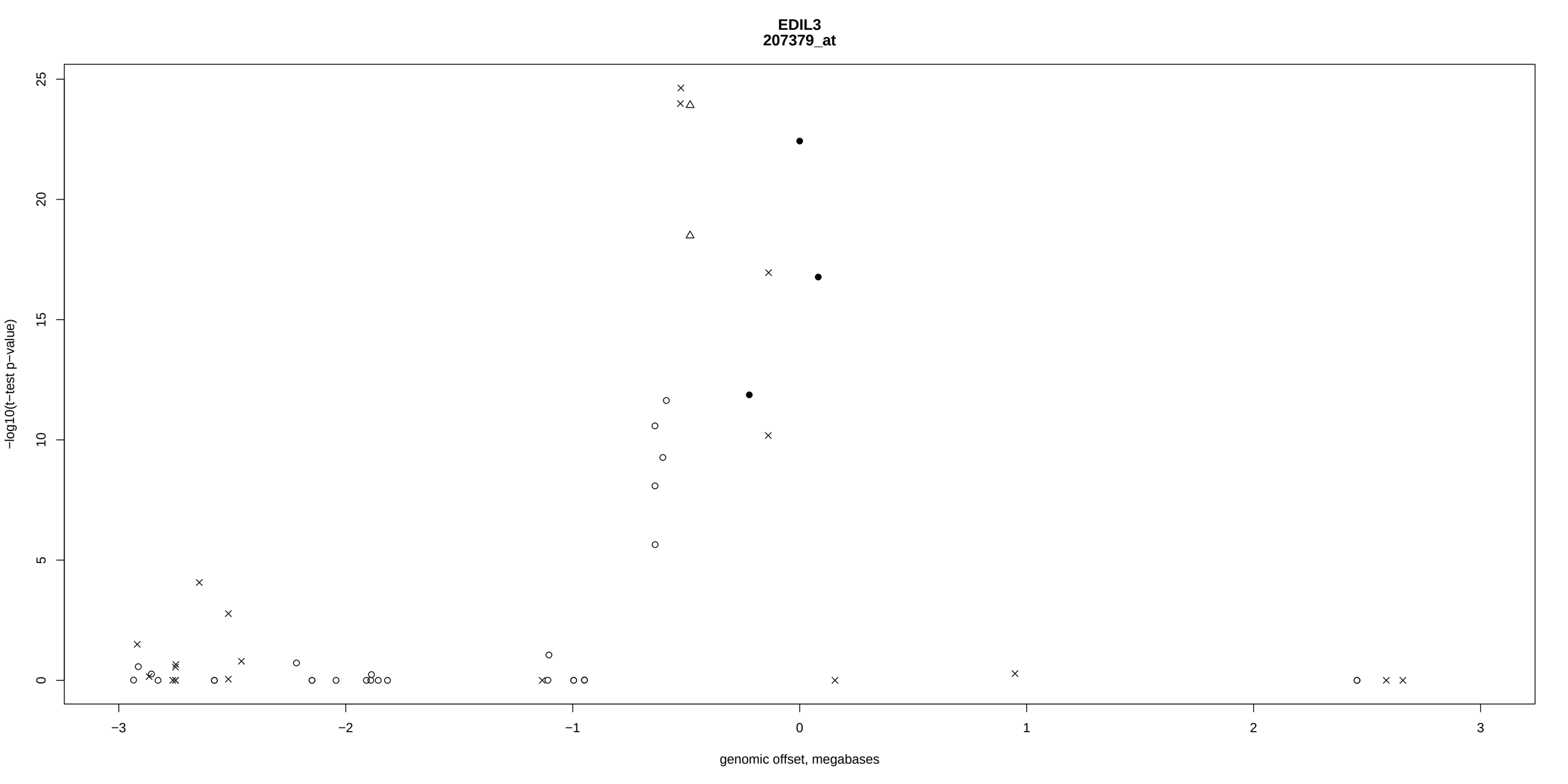
2
3
genomic offset, megabases


EDIL3
207379_at
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10
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20
25
−3
−2
−1
0
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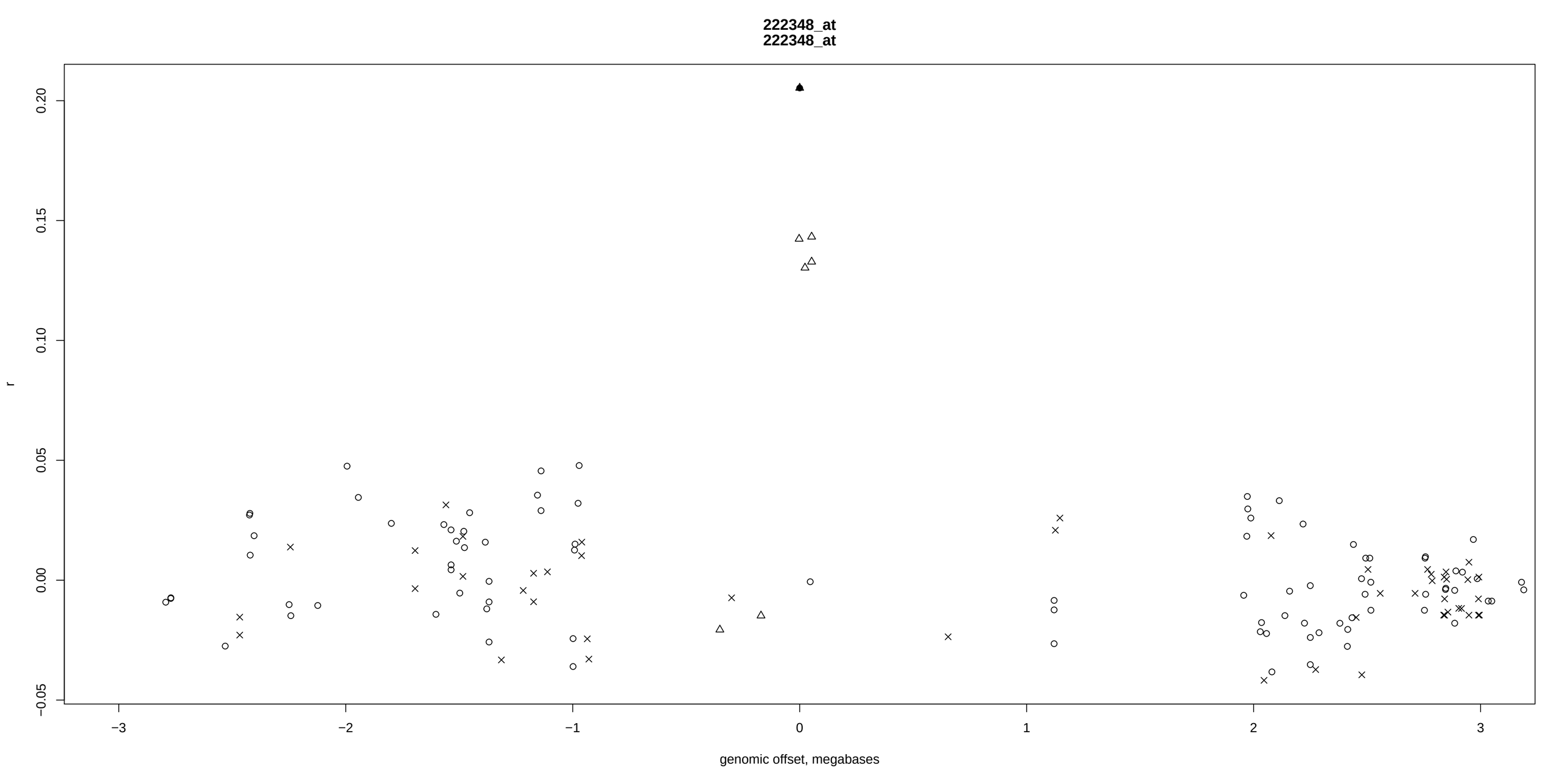
2
3
genomic offset, megabases


222348_at
222348_at
r
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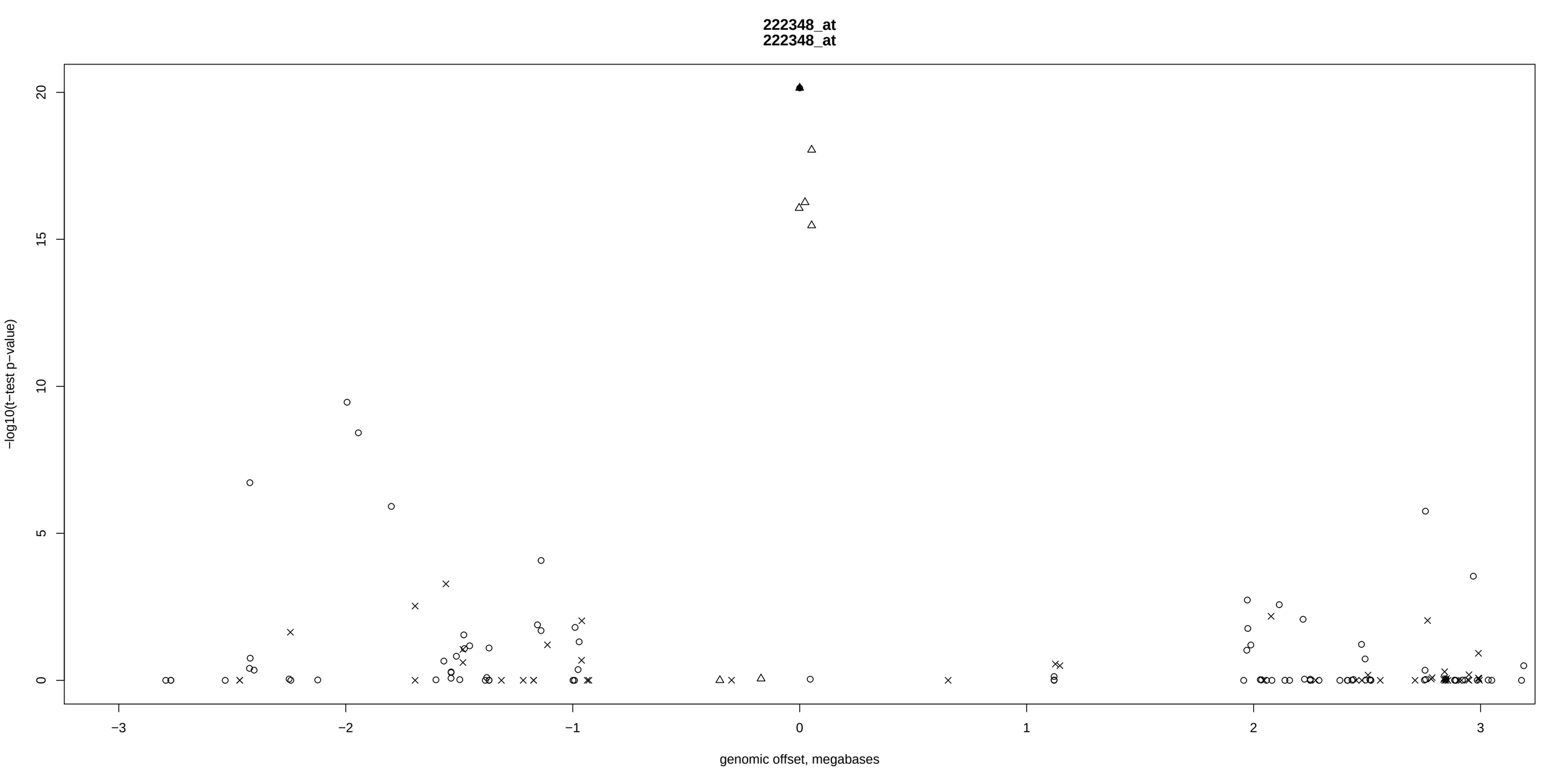
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


222348_at
222348_at
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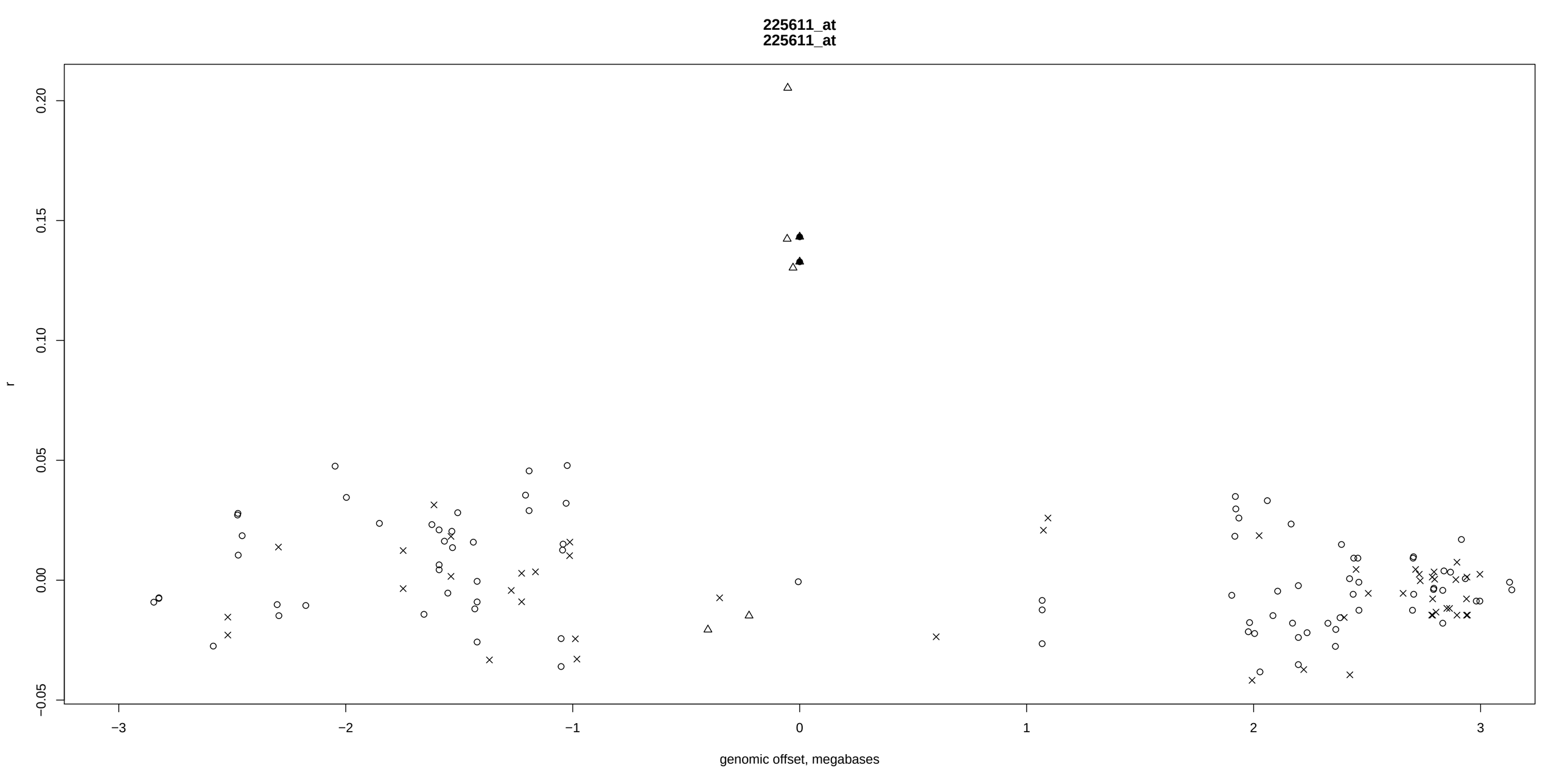
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


225611_at
225611_at
r
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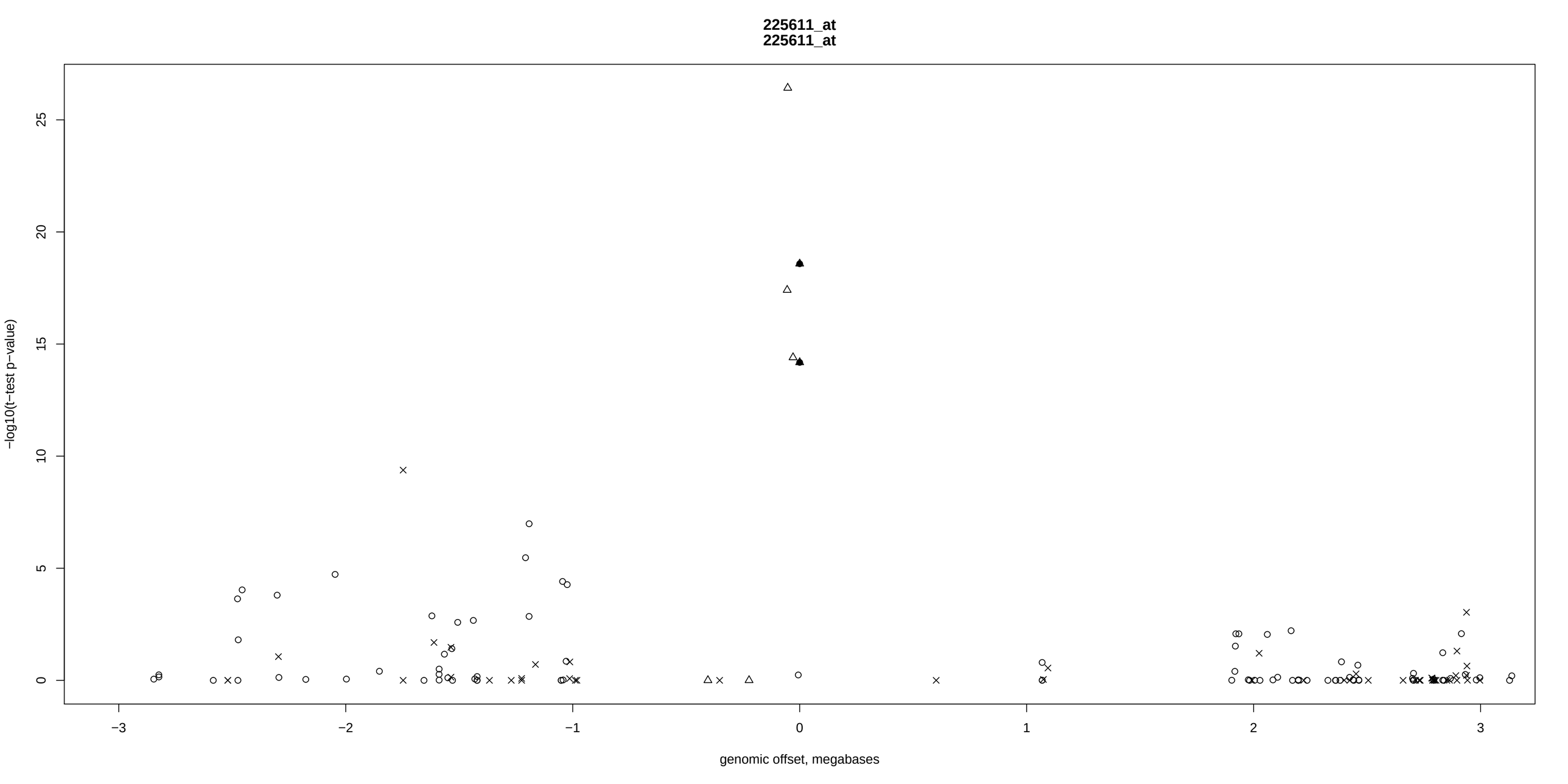
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


225611_at
225611_at
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0
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10
15
20
25
−3
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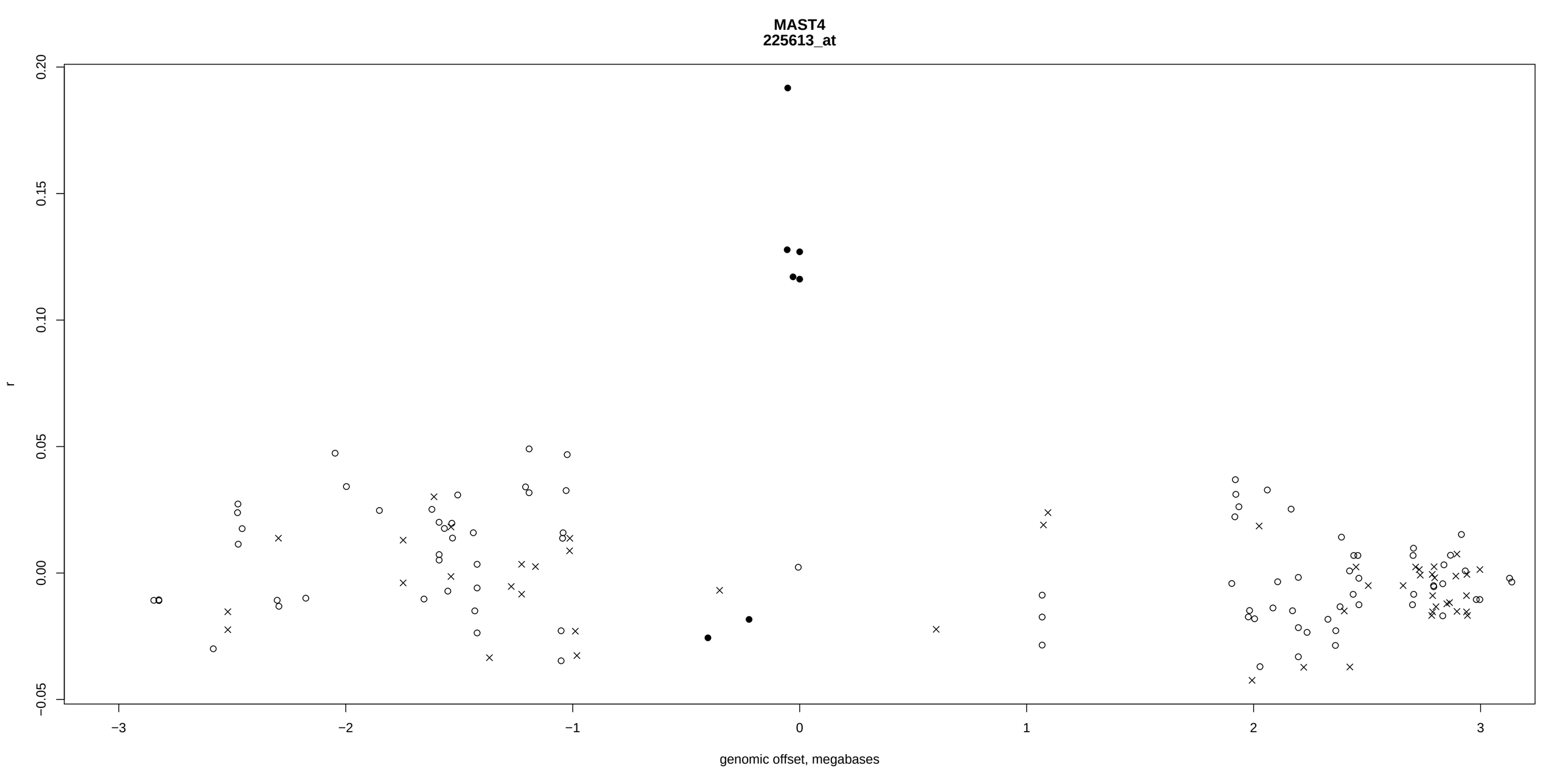
−1
0
1
2
3
genomic offset, megabases


MAST4
225613_at
r
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−3
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−1
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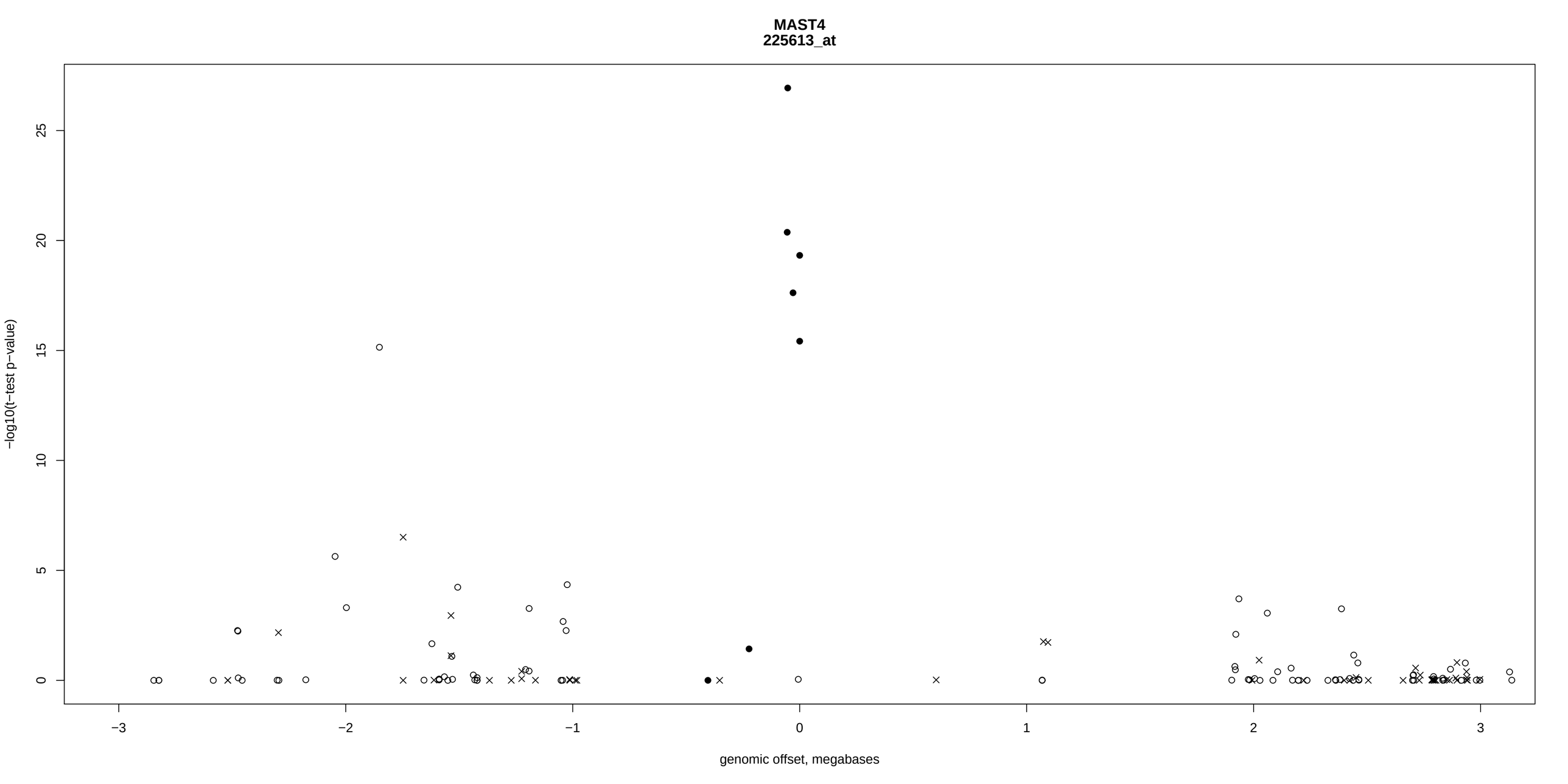
3
genomic offset, megabases


MAST4
225613_at
−log10(t−test p−value)
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10
15
20
25
−3
−2
−1
0
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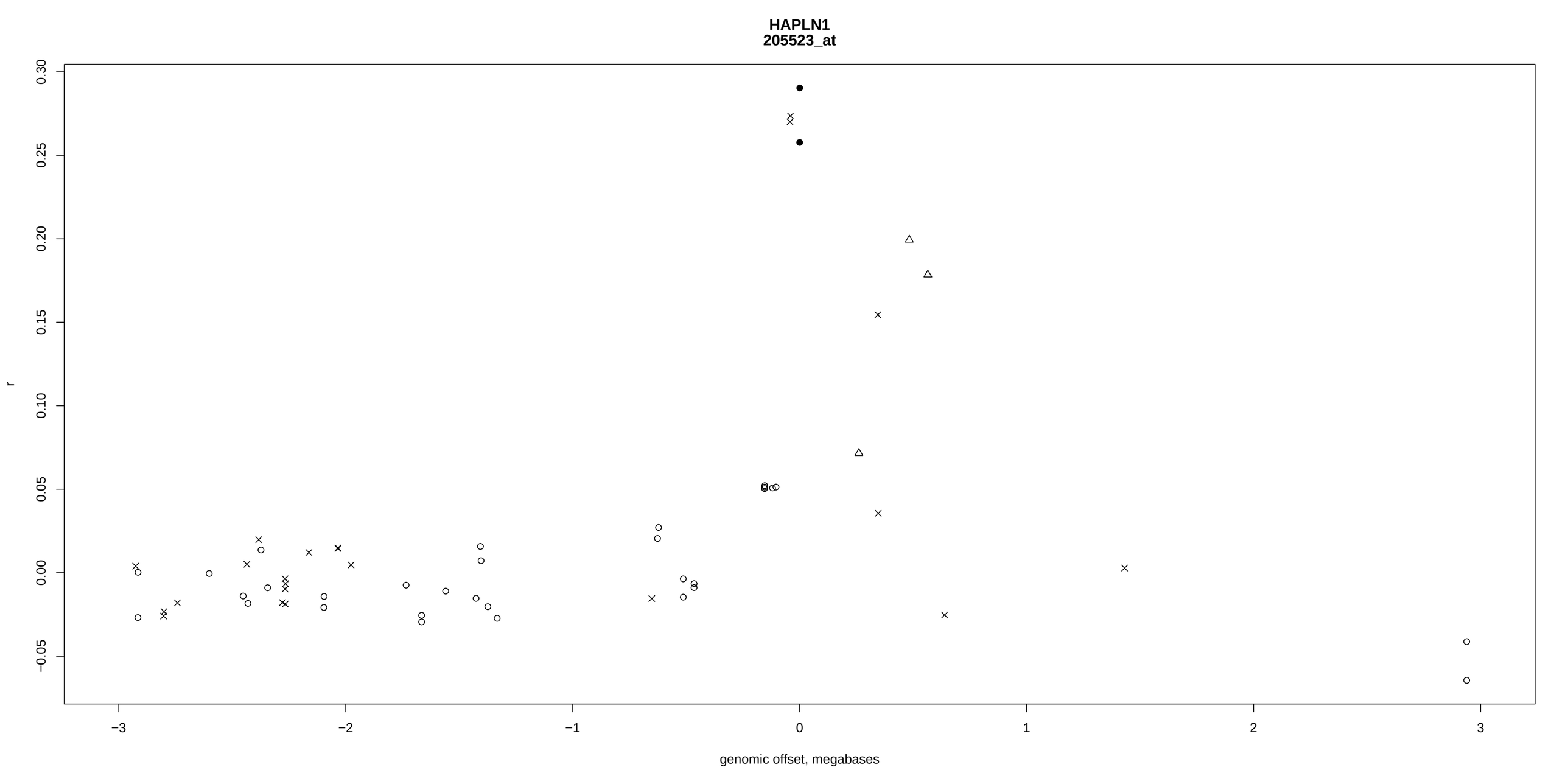
2
3
genomic offset, megabases


HAPLN1
205523_at
r
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0.25
0.20
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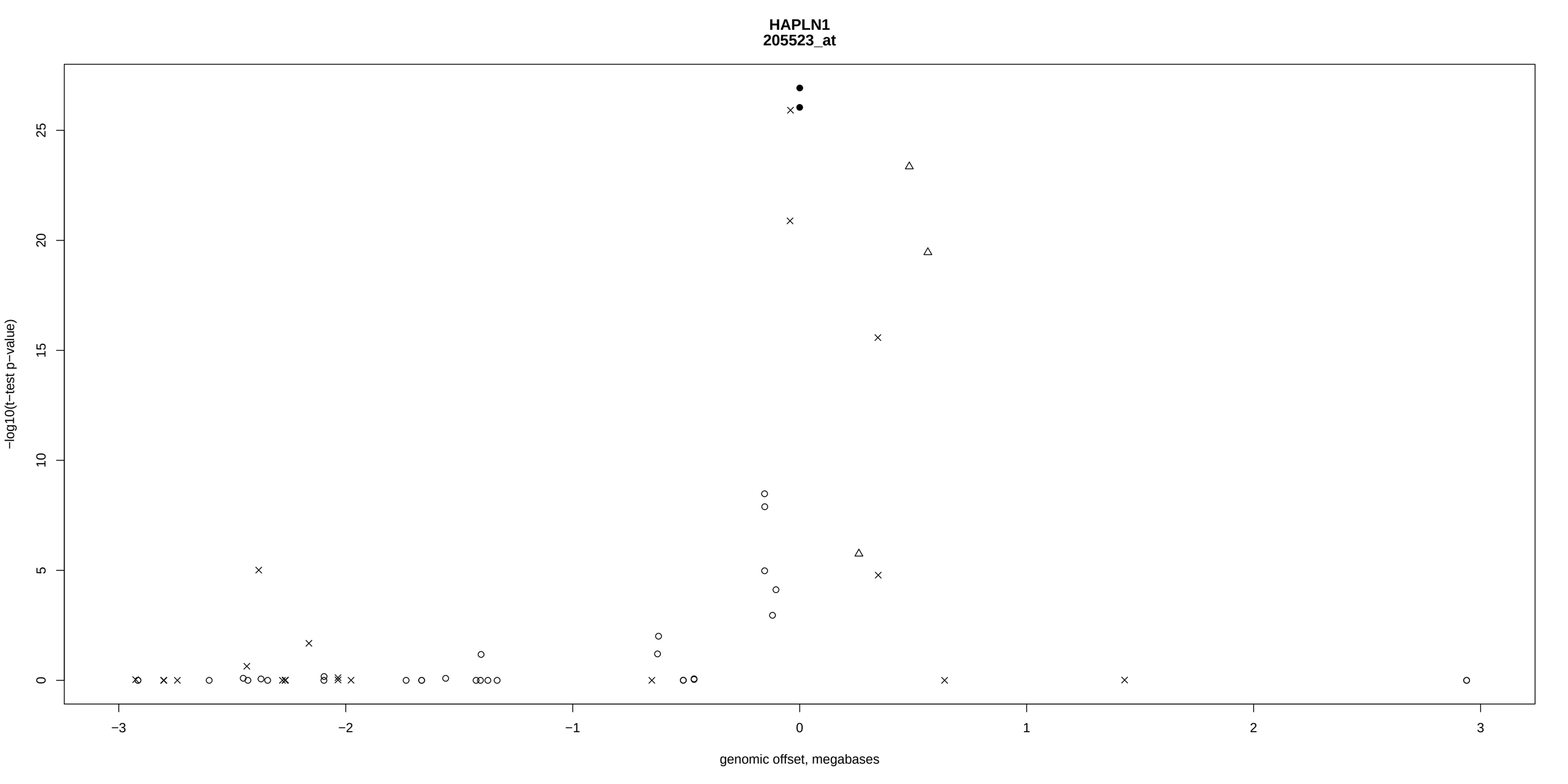
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


HAPLN1
205523_at
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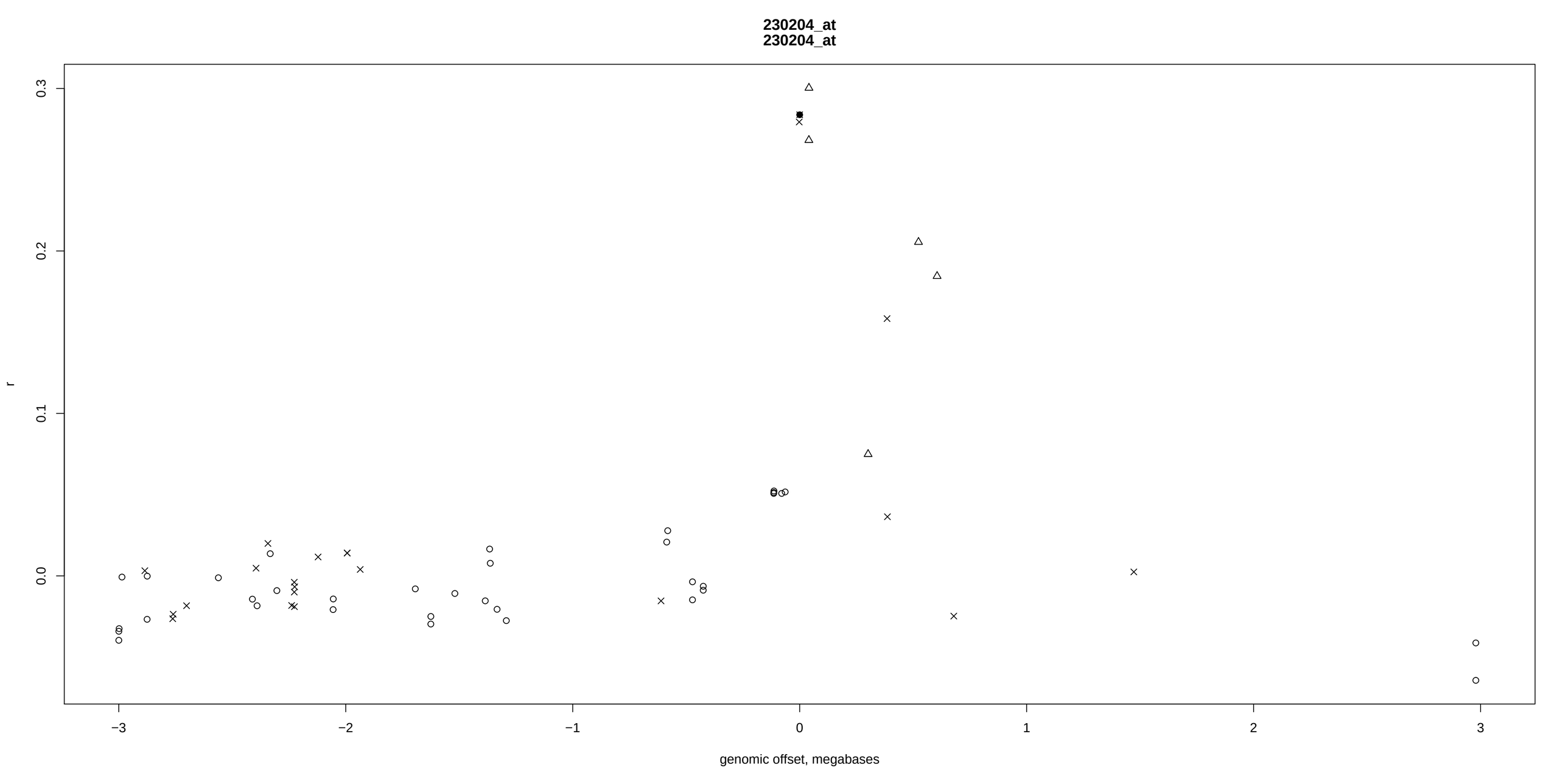
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


230204_at
230204_at
r
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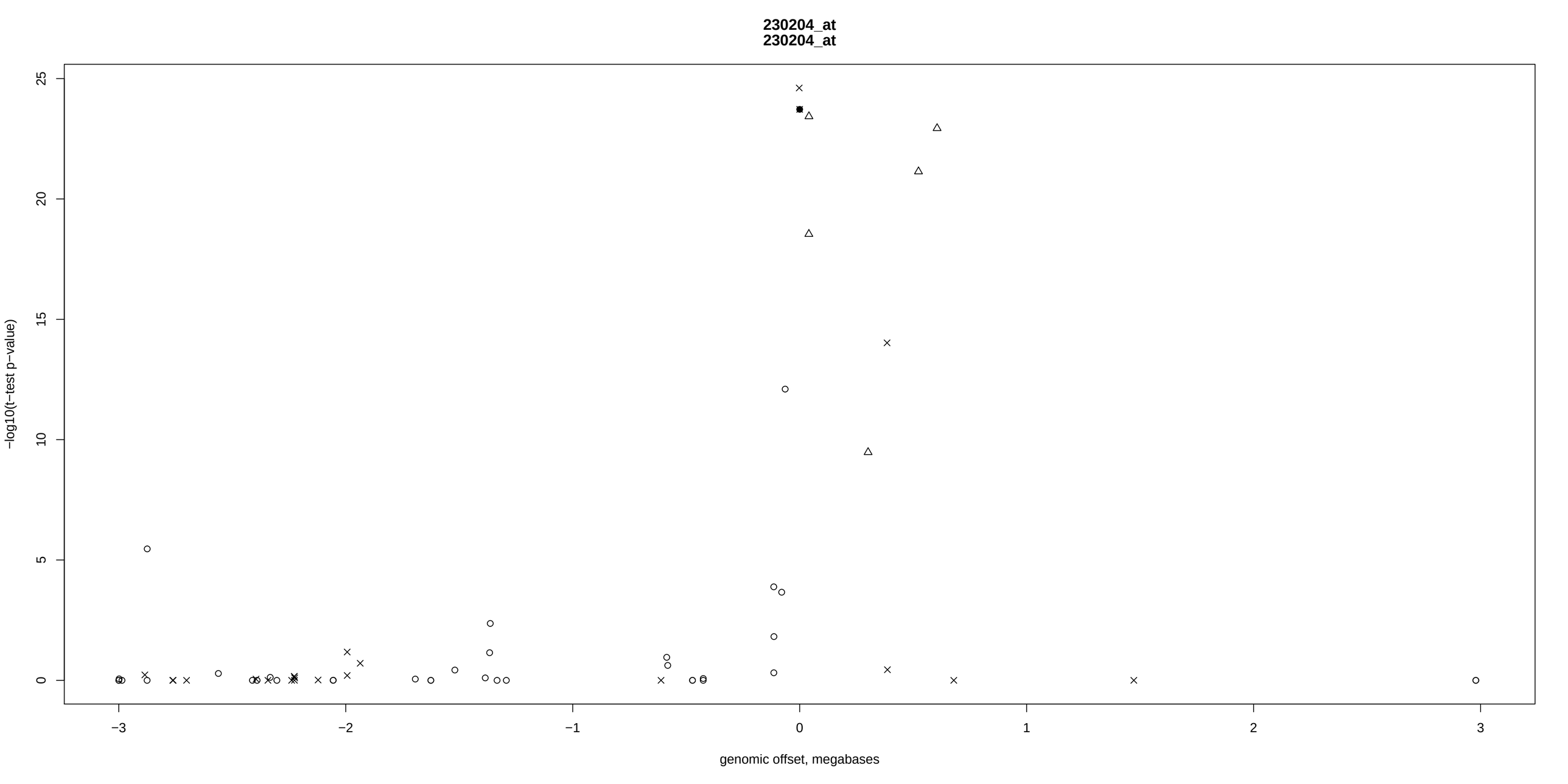
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases


230204_at
230204_at
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0
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10
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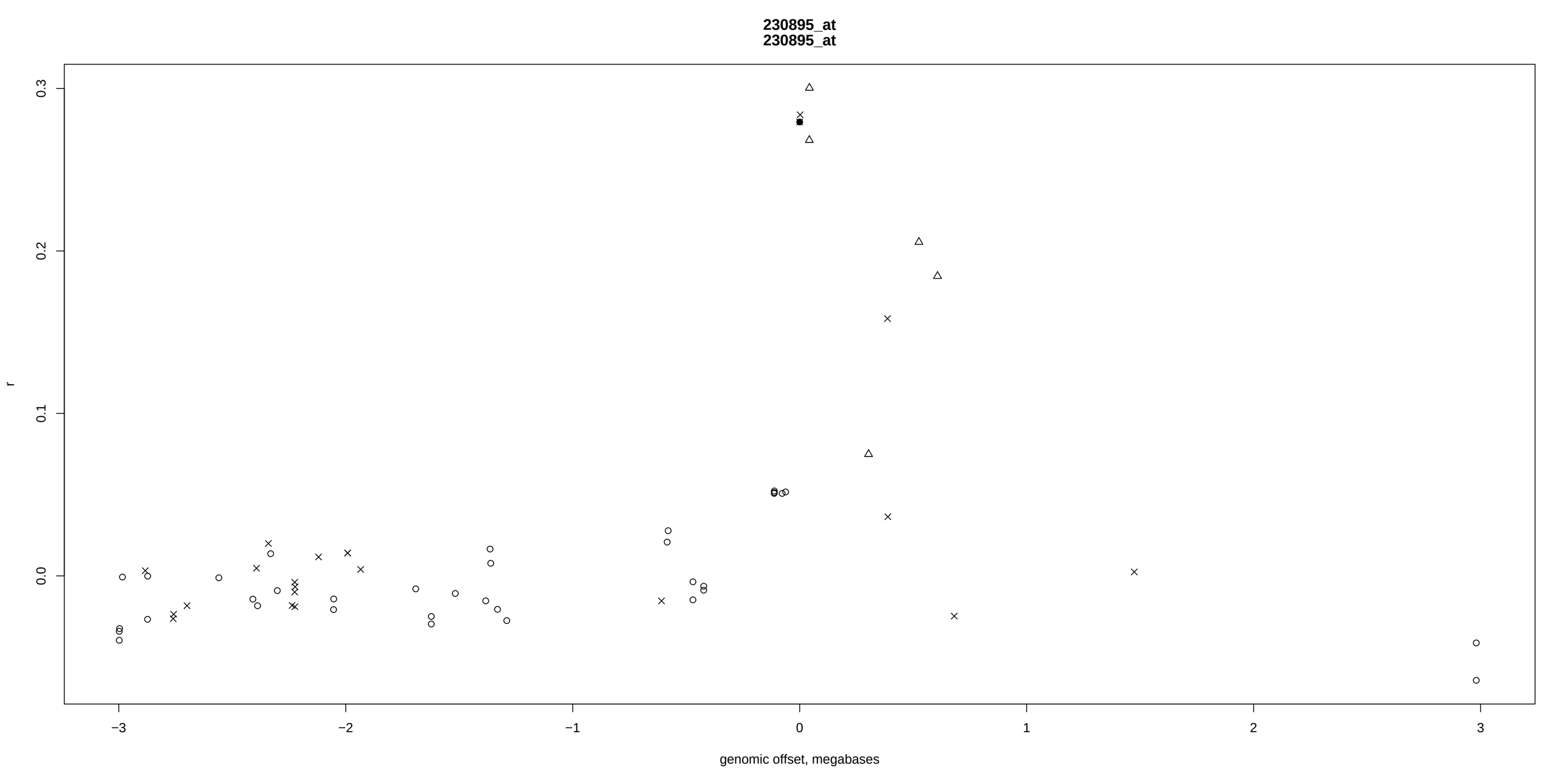
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


230895_at
230895_at
r
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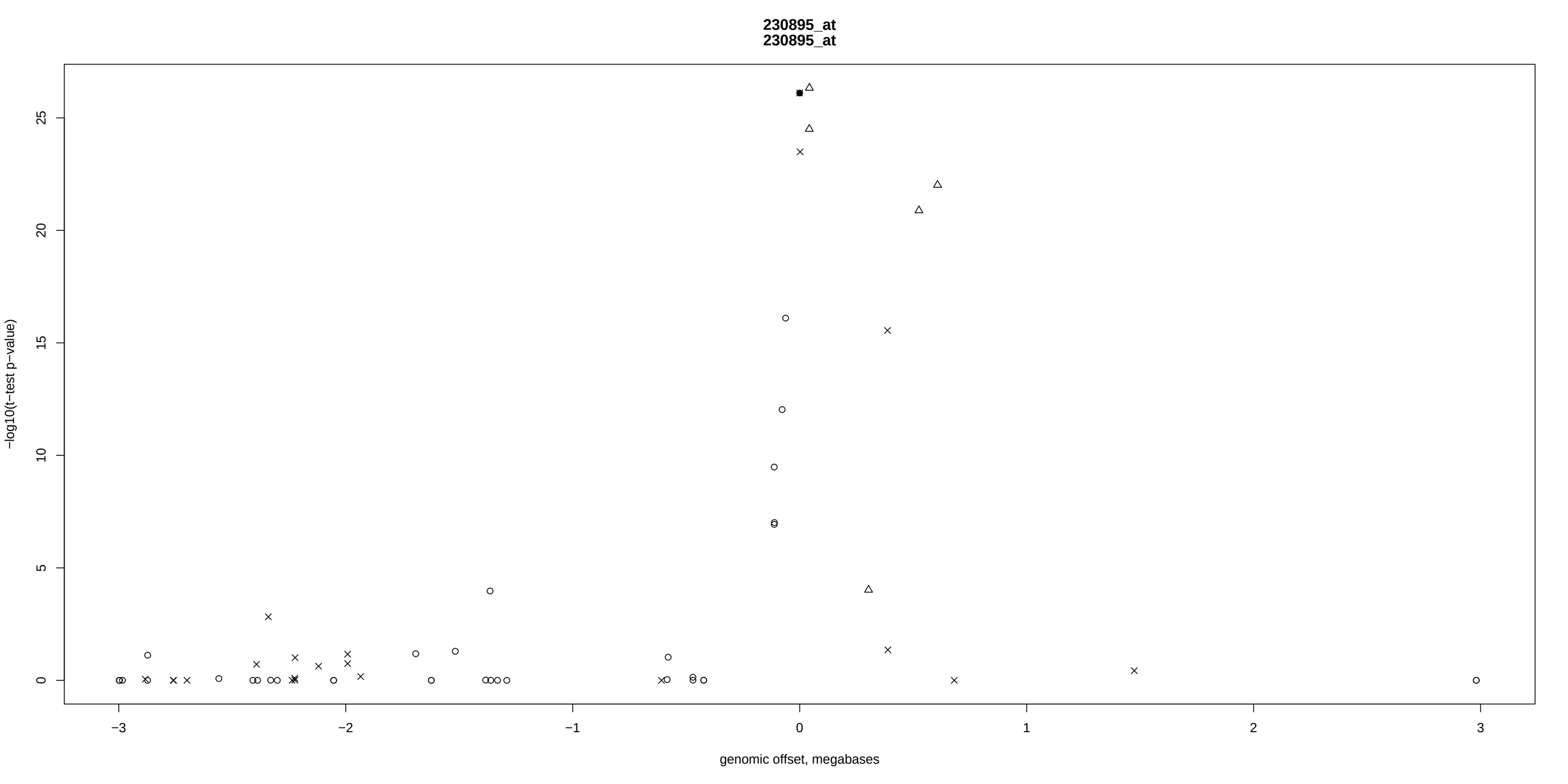
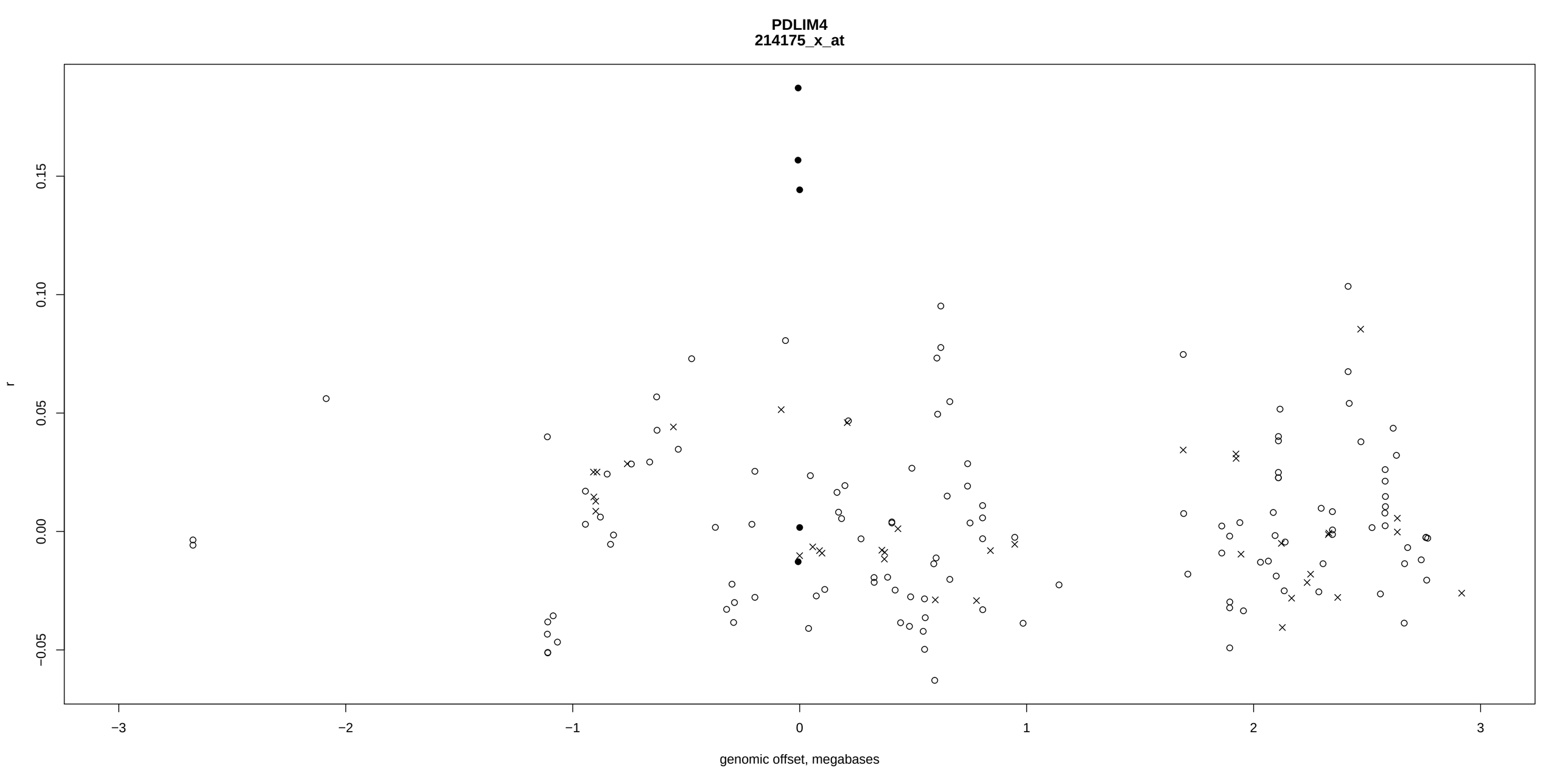
0.1
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases


230895_at
230895_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


PDLIM4
214175_x_at
r
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0.10
0.05
0.00
−0.05
−3
−2
−1
0
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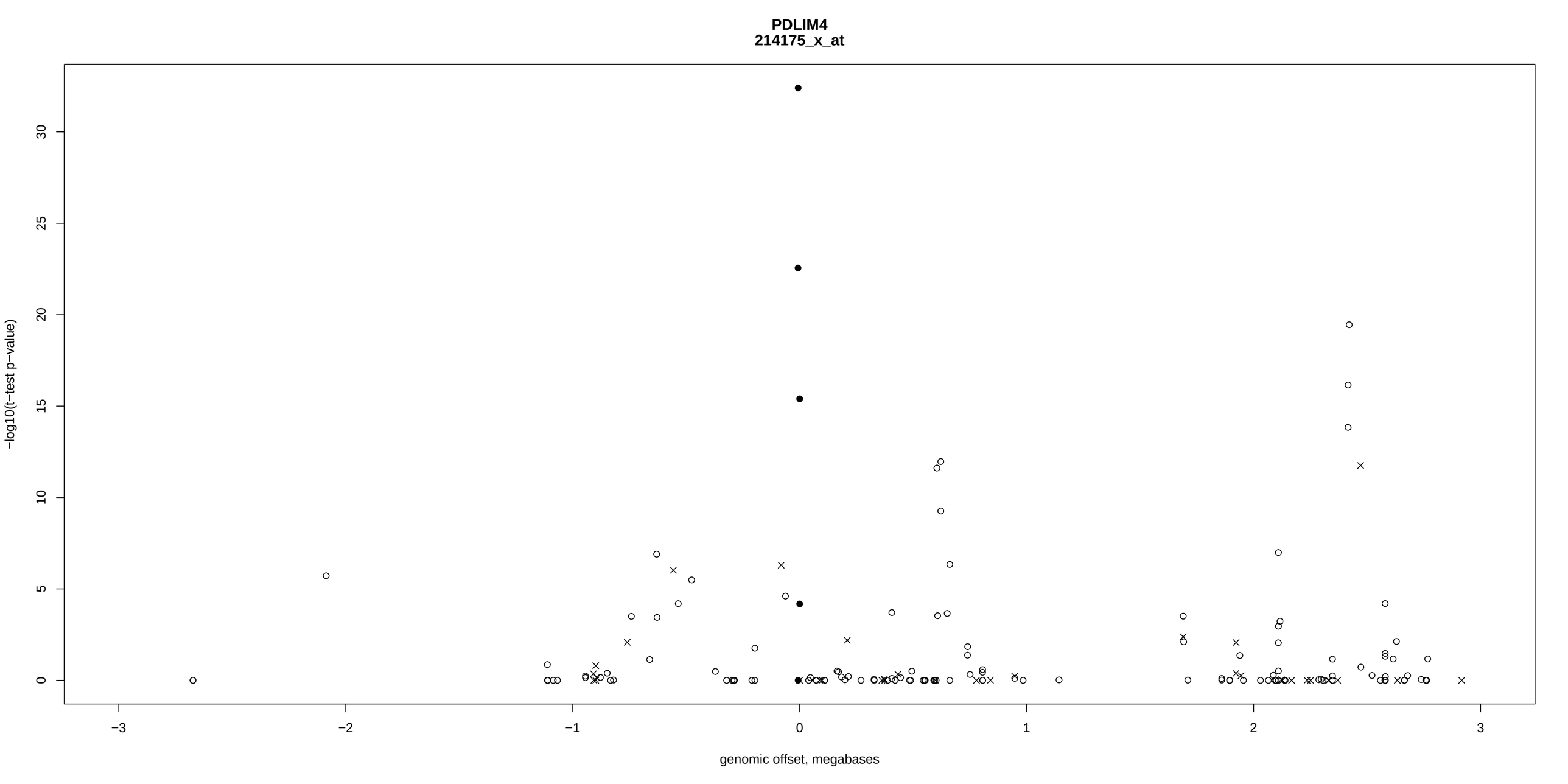
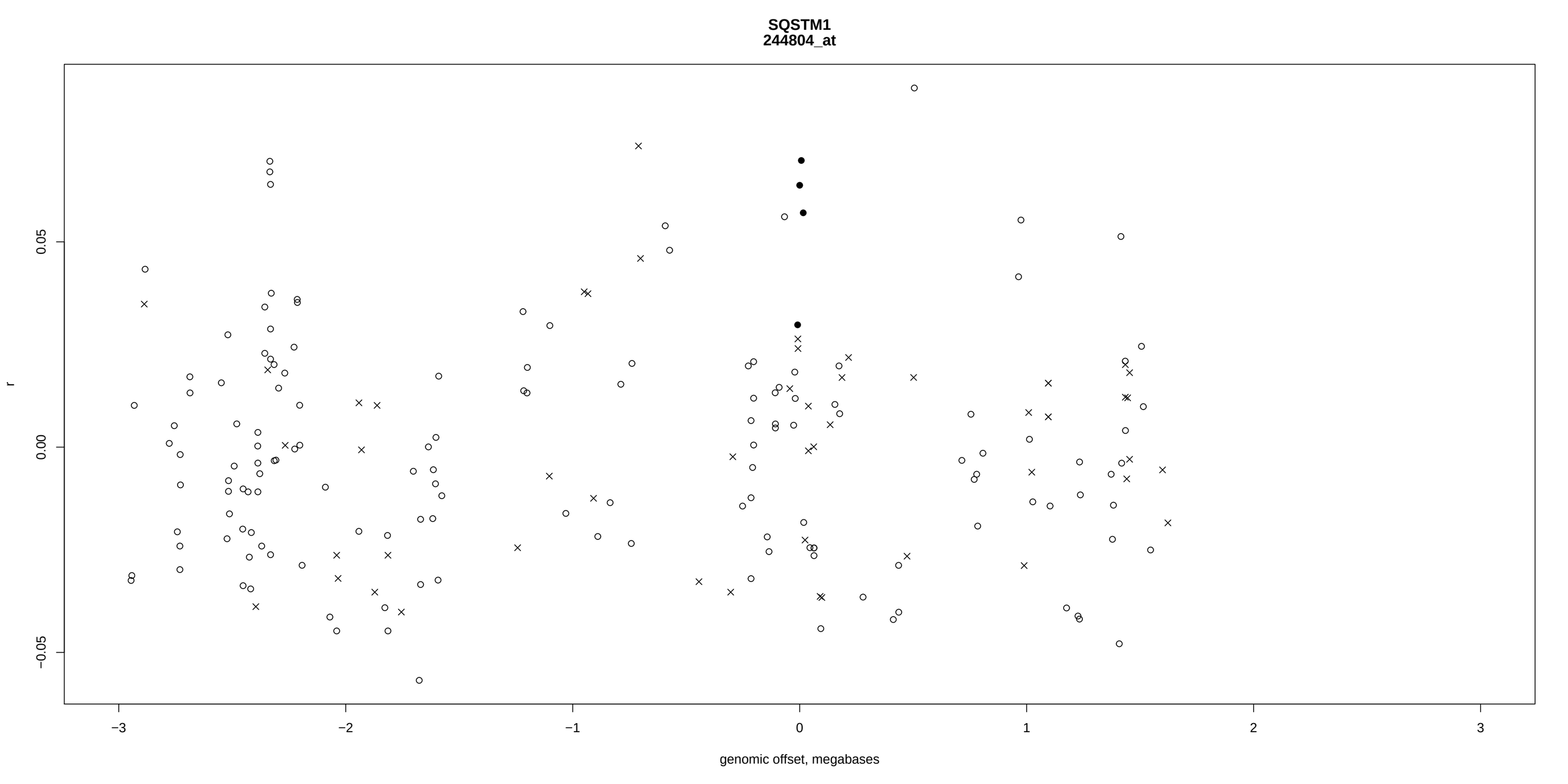
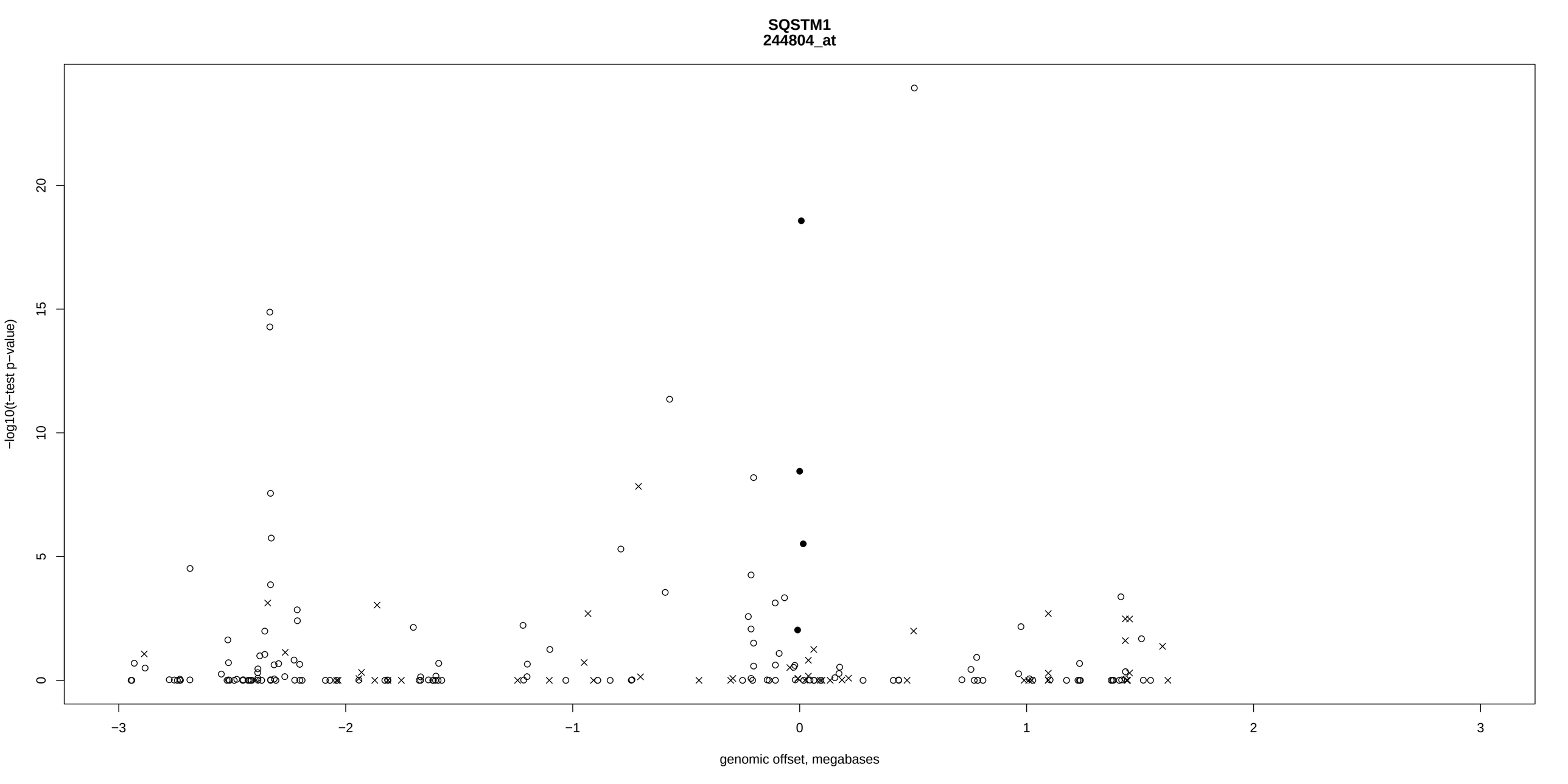
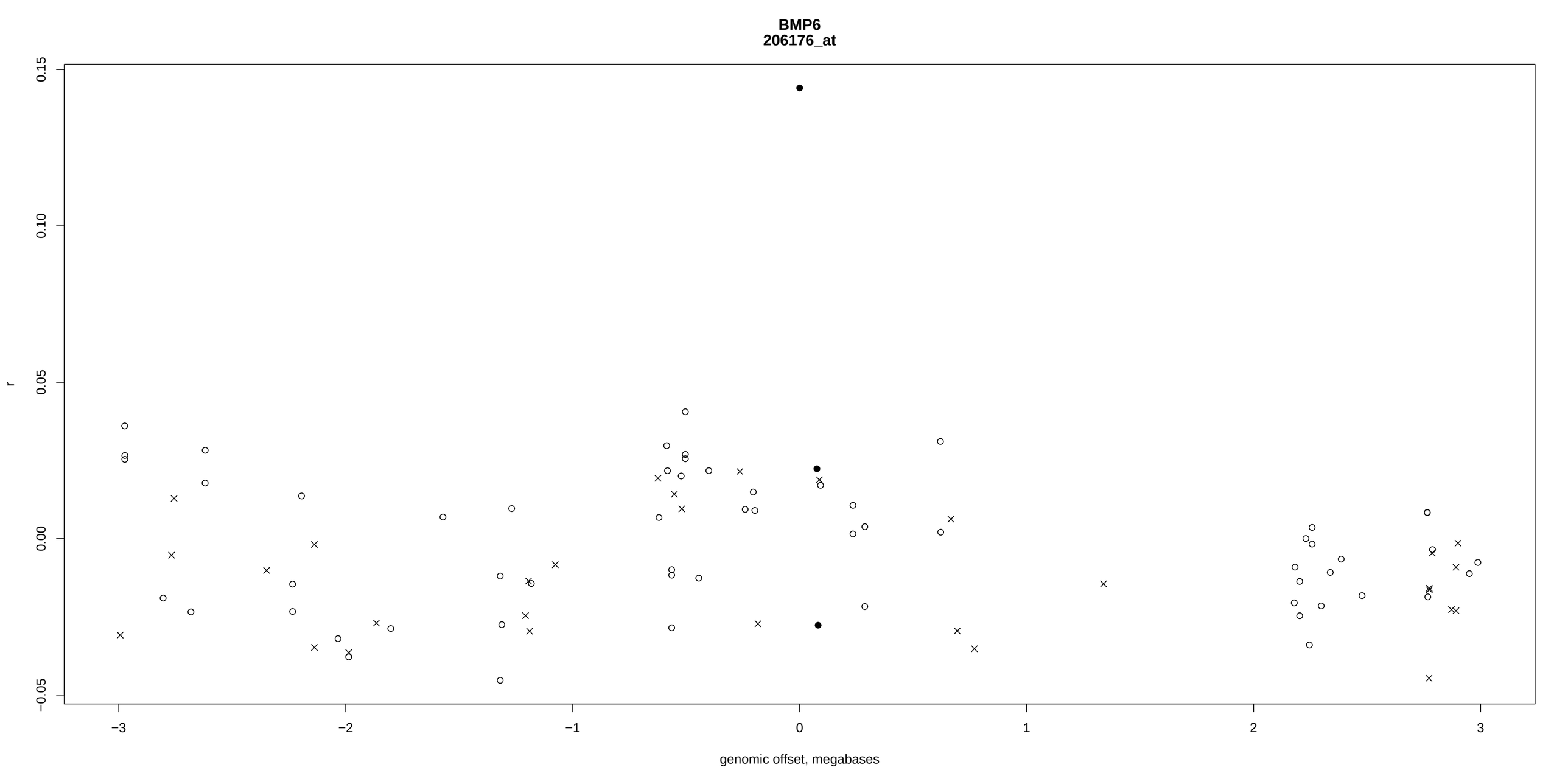
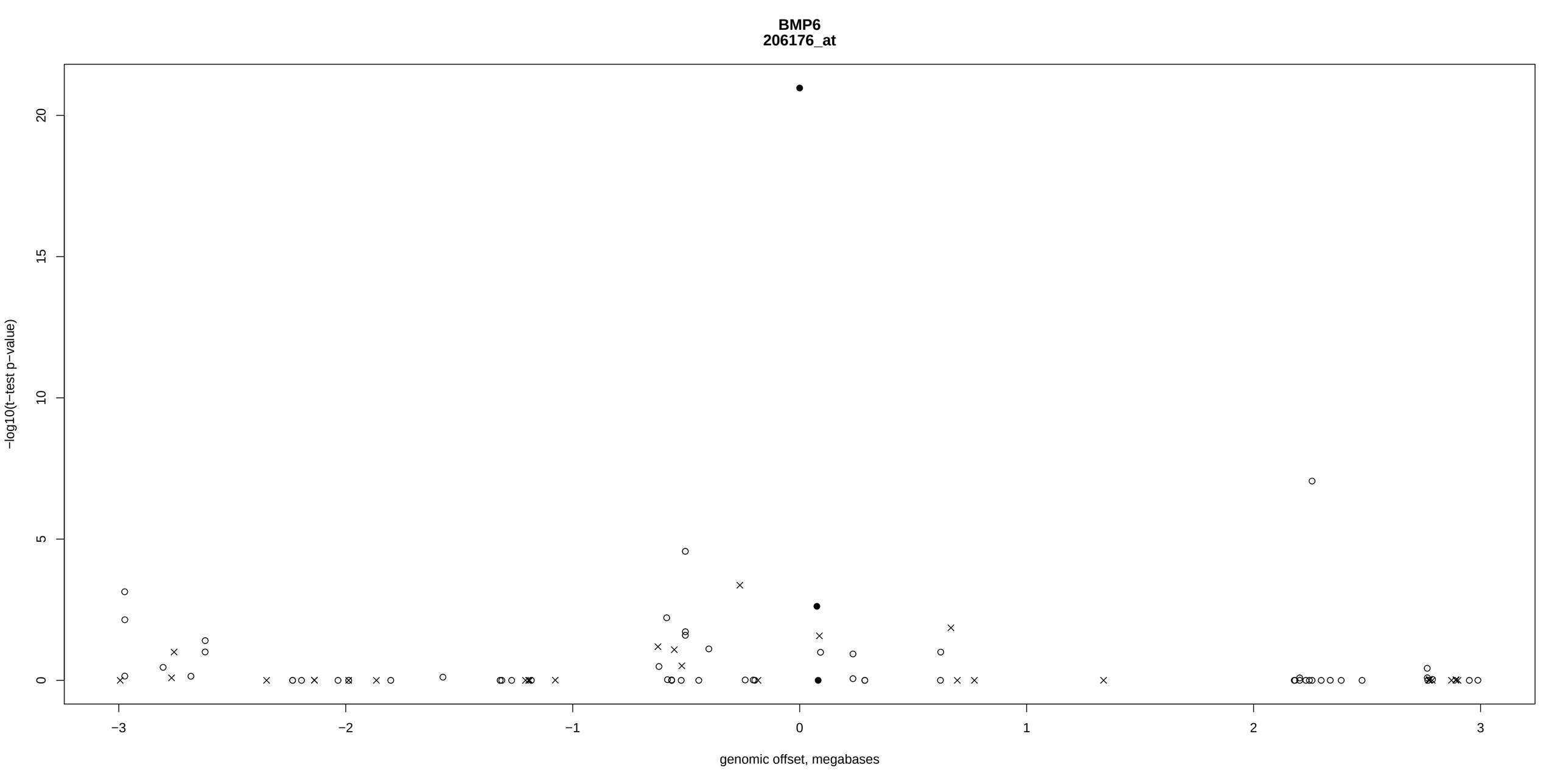
2
3
genomic offset, megabases

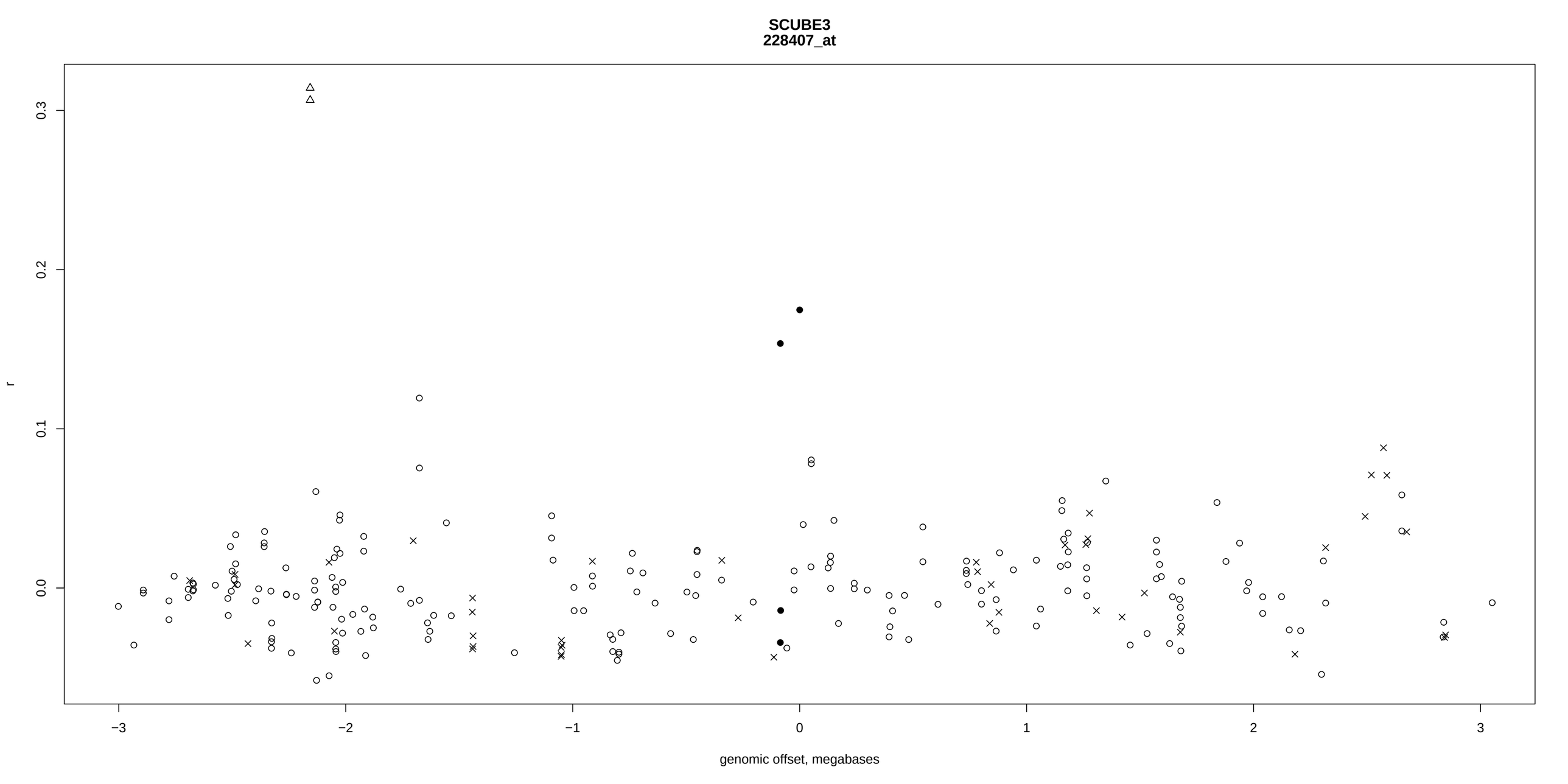

PDLIM4
214175_x_at
−log10(t−test p−value)
0
5
10
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases

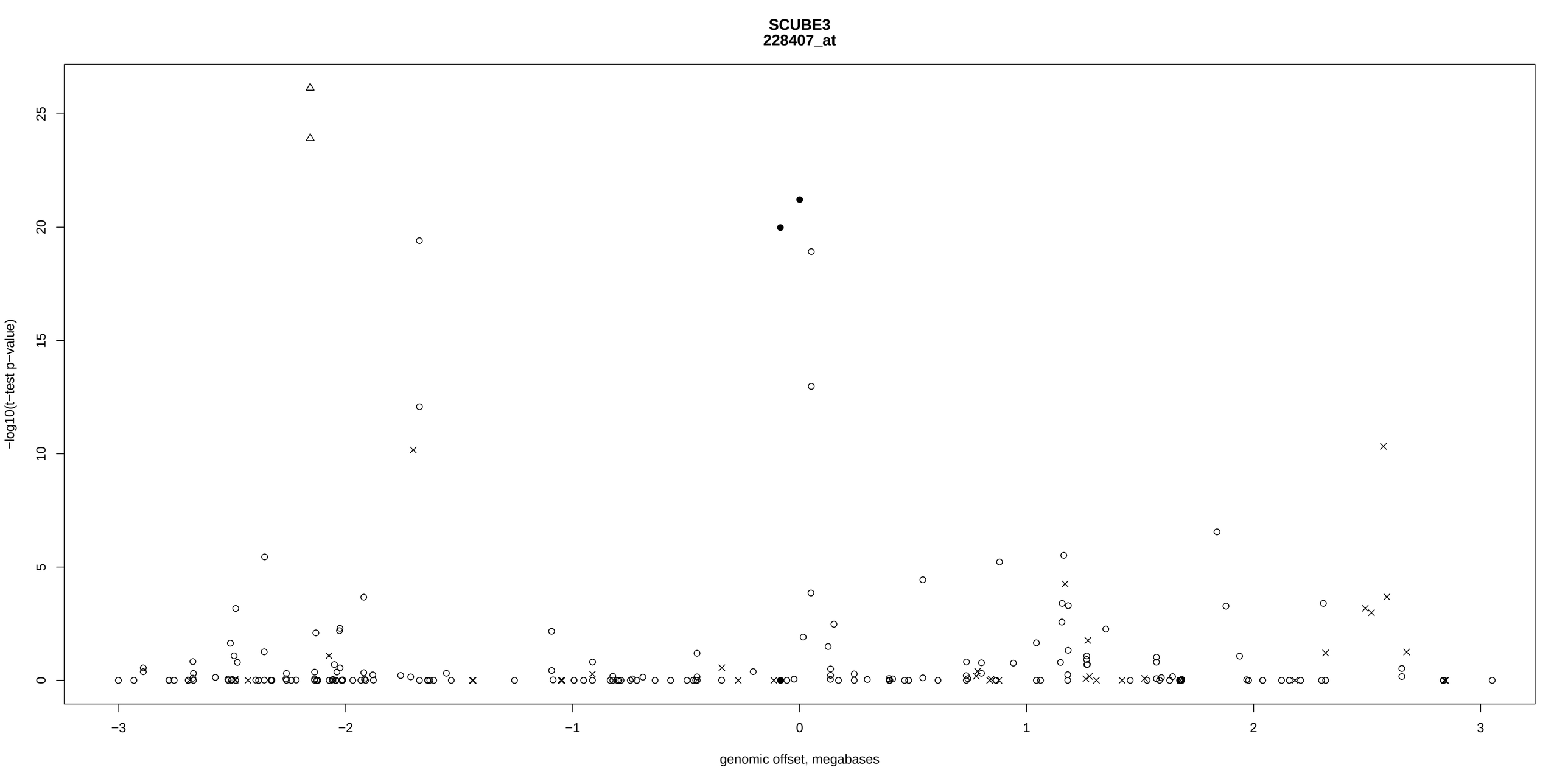

SQSTM1
244804_at
r
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


SQSTM1
244804_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


BMP6
206176_at
r
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


BMP6
206176_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


SCUBE3
228407_at
r
0.0
0.1
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases


SCUBE3
228407_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
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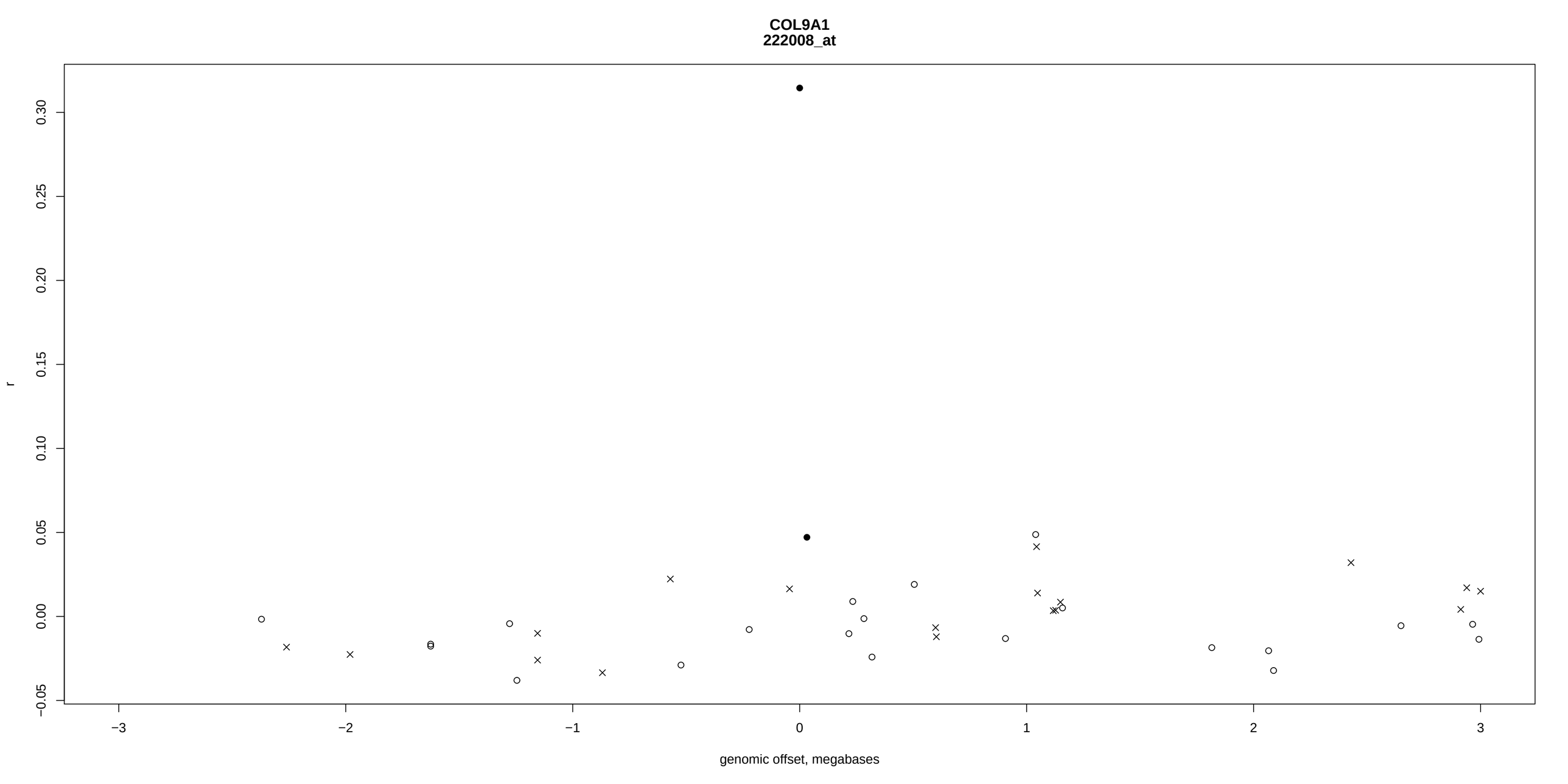
2
3
genomic offset, megabases


COL9A1
222008_at
r
0.30
0.25
0.20
0.15
0.10
0.05
0.00
−0.05
−3
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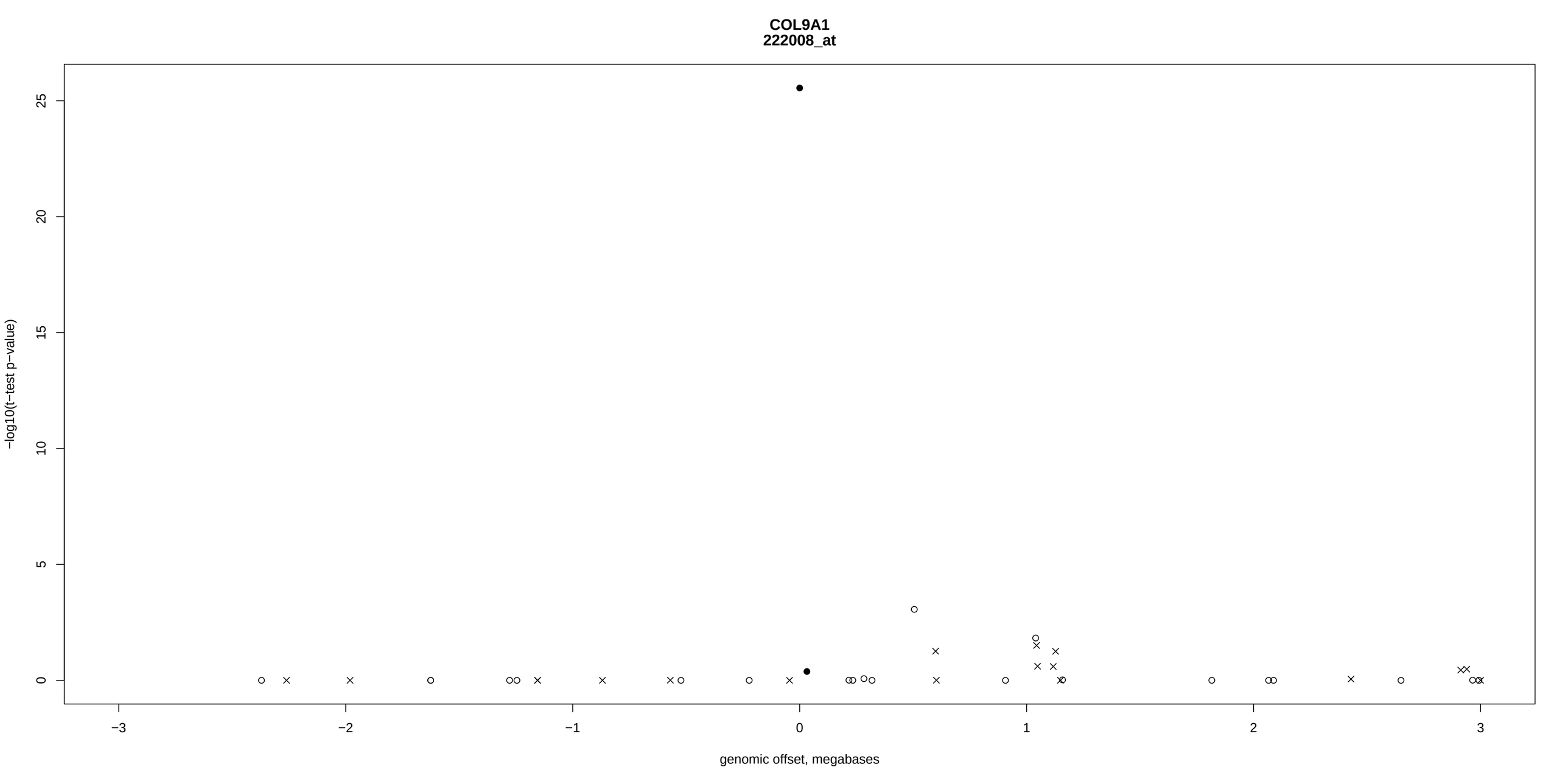
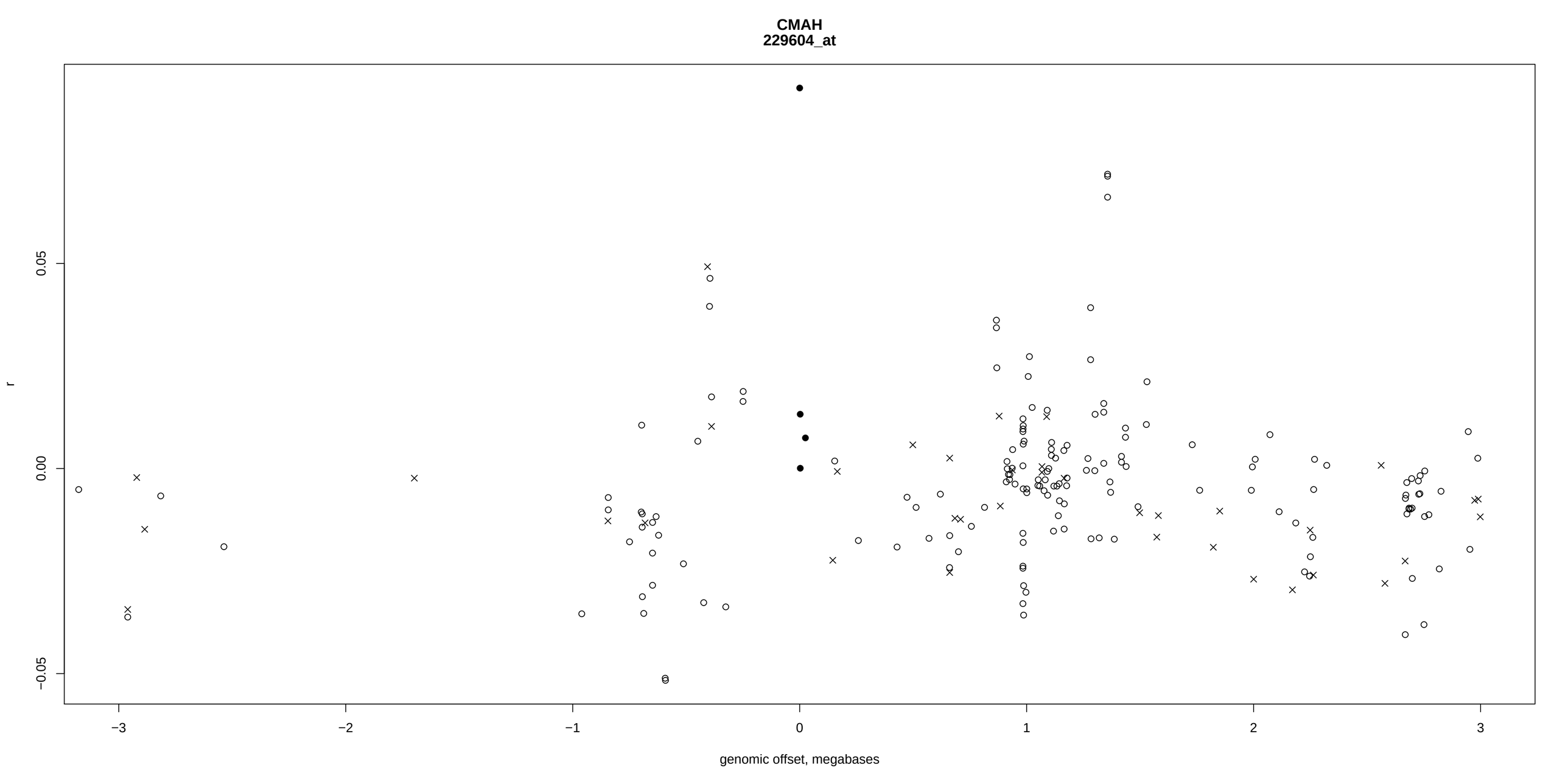
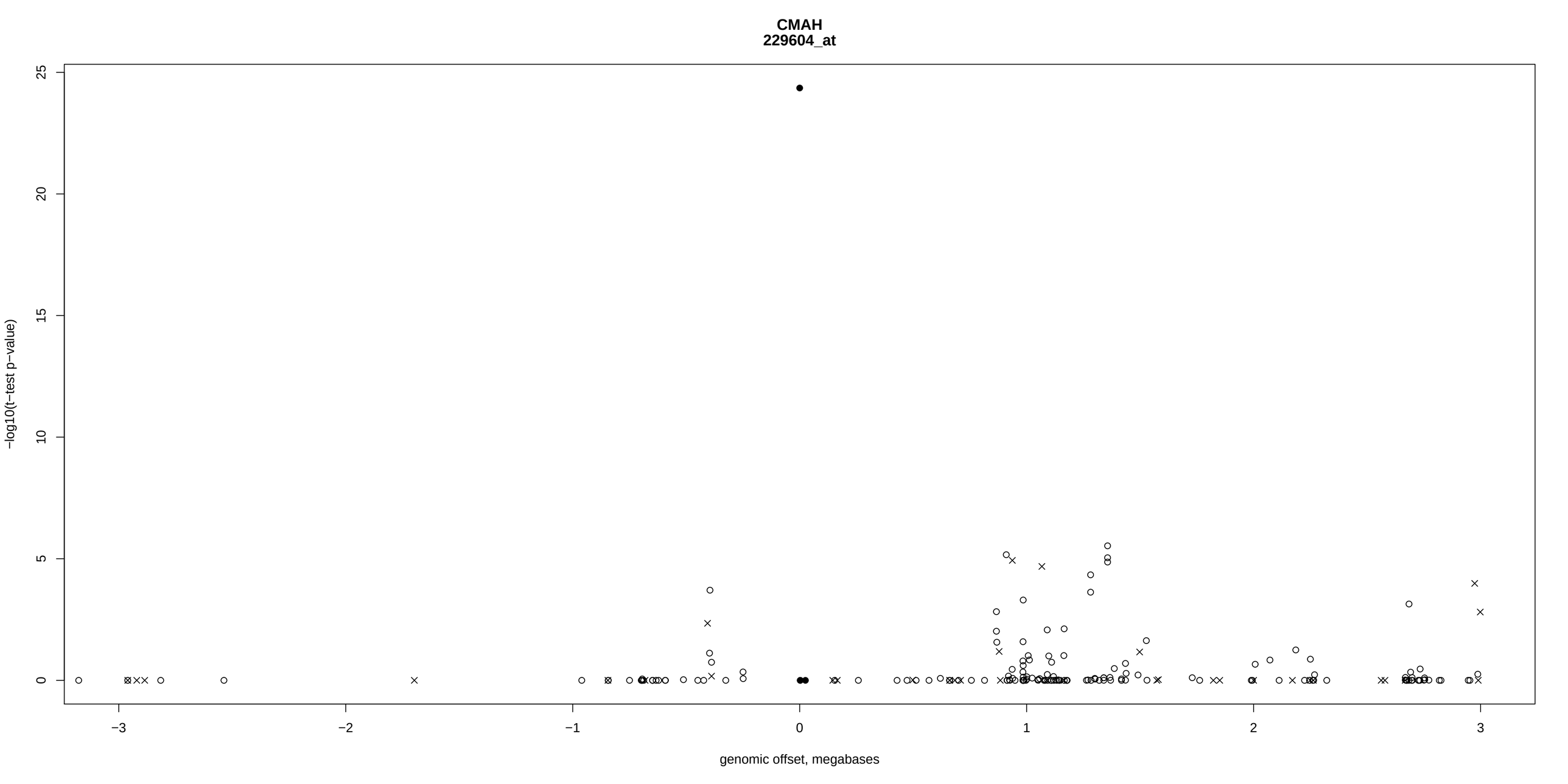
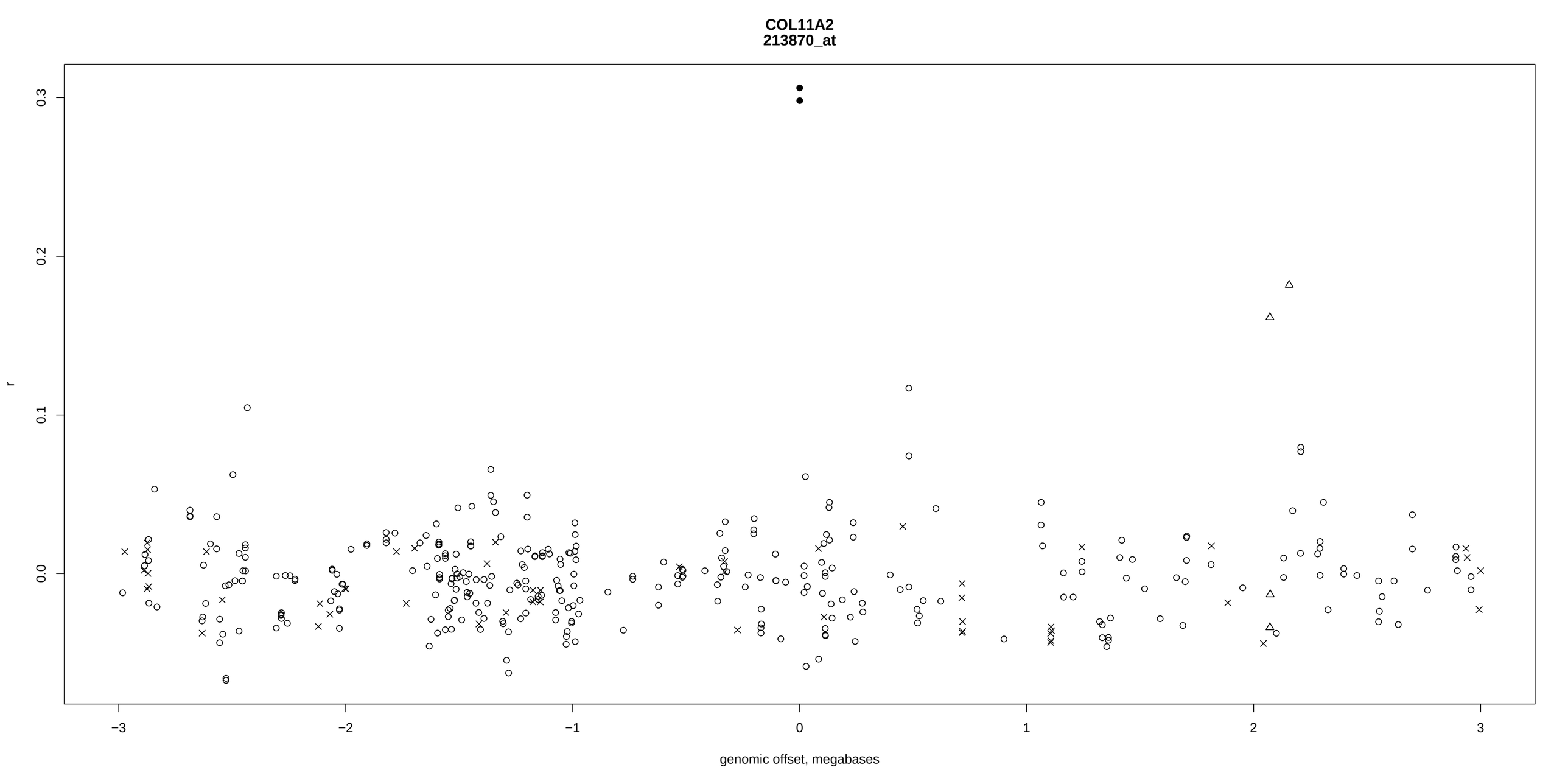
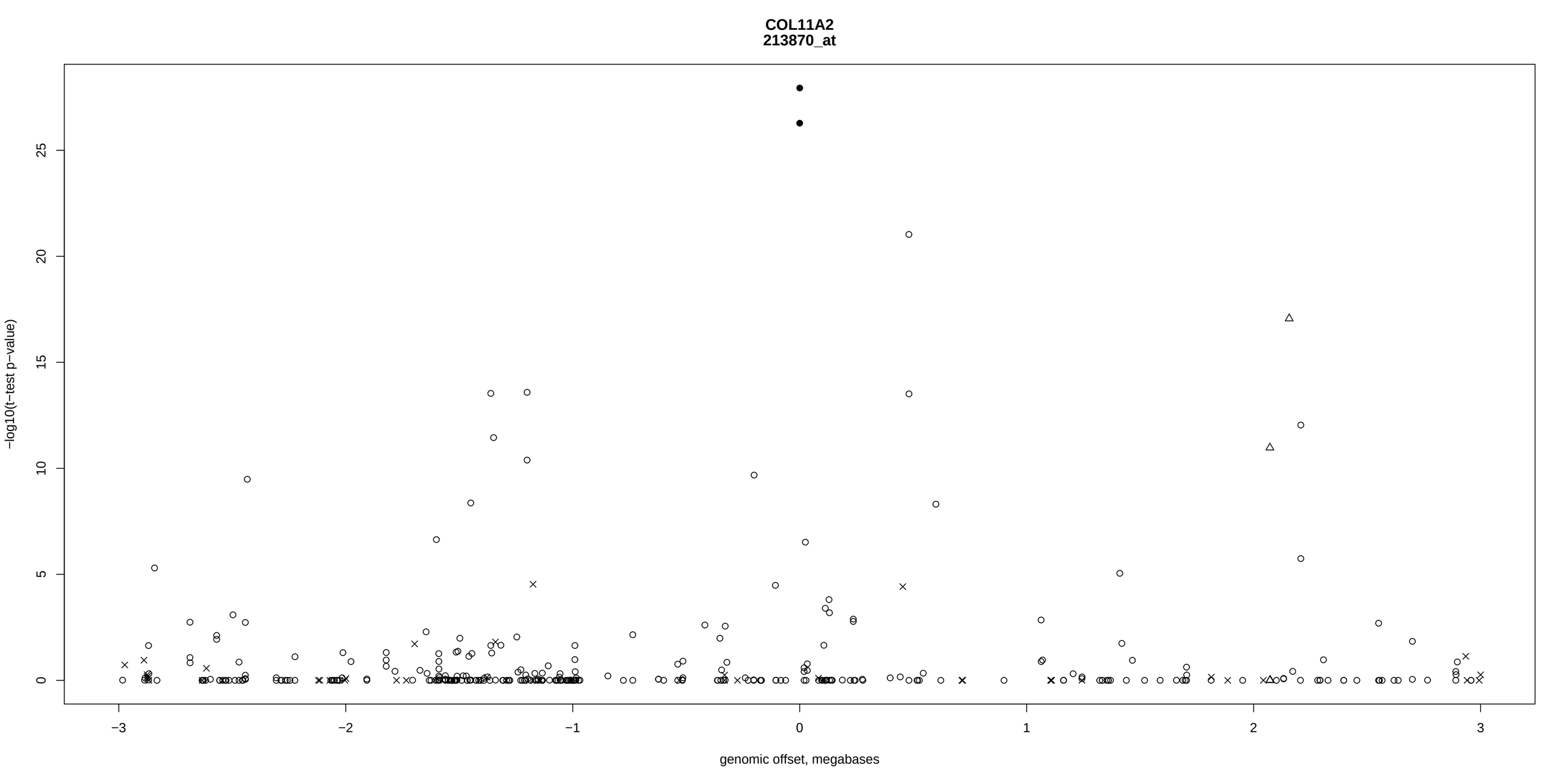
−1
0
1
2
3
genomic offset, megabases

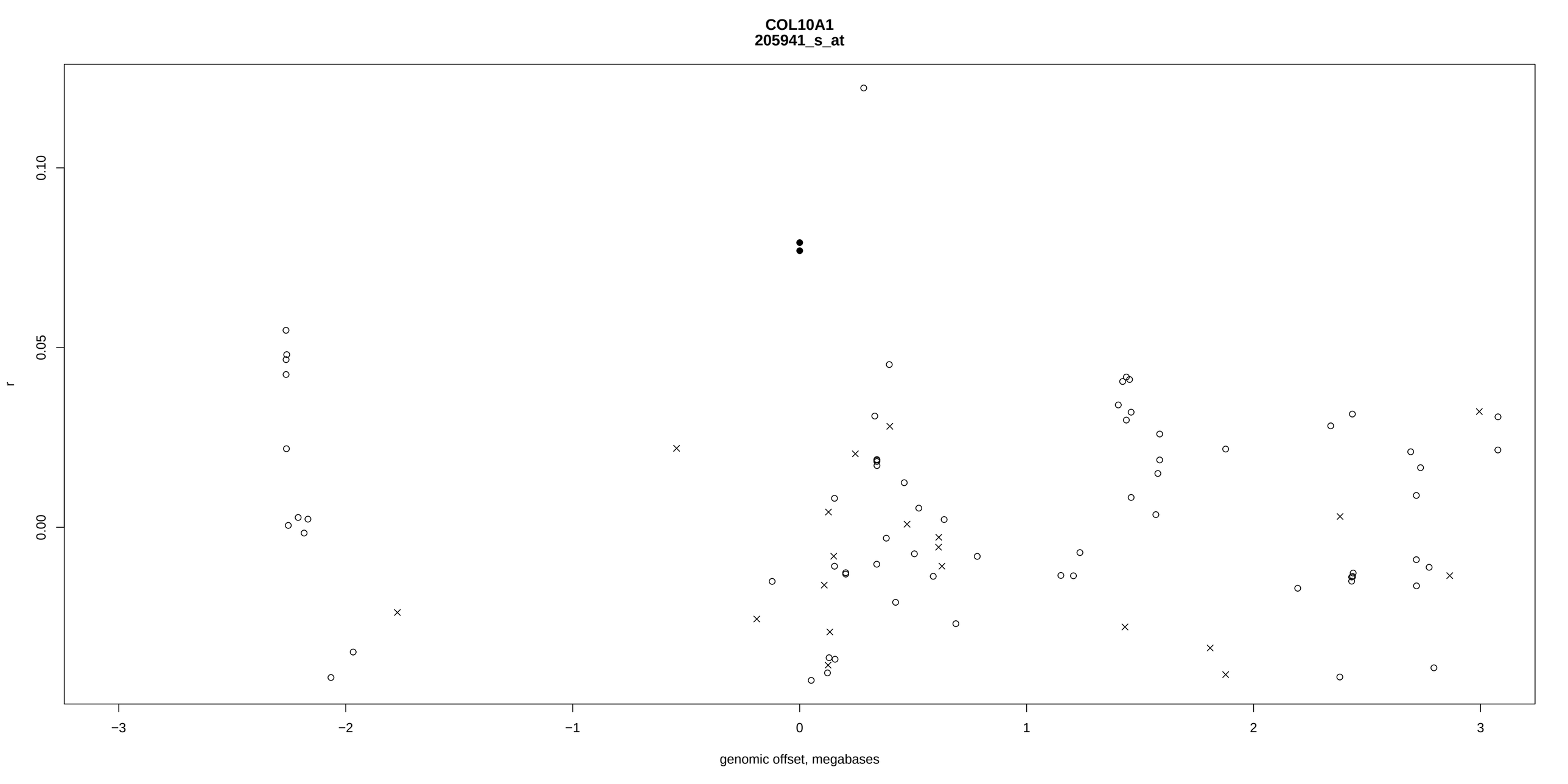

COL9A1
222008_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

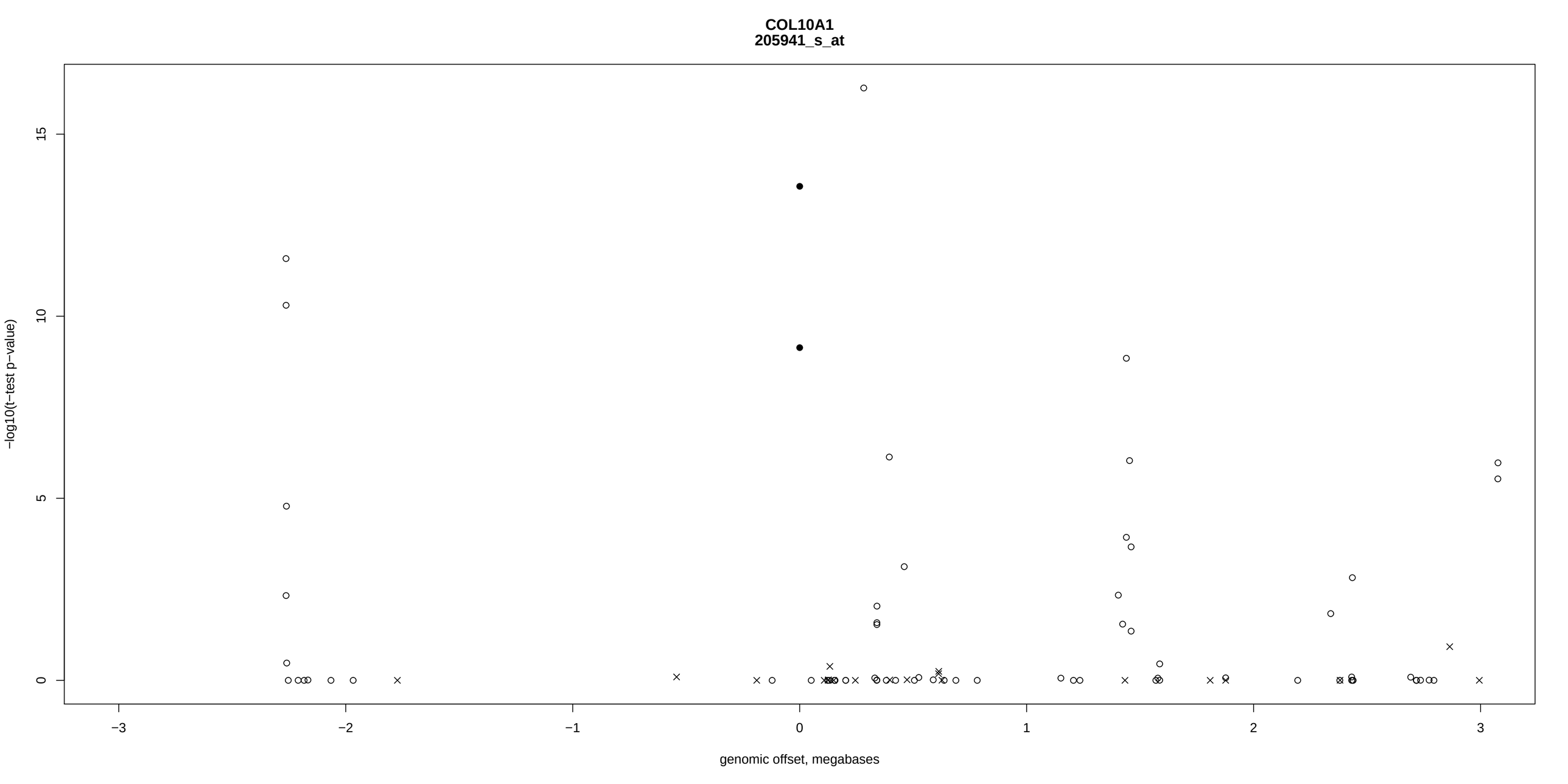

CMAH
229604_at
r
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases

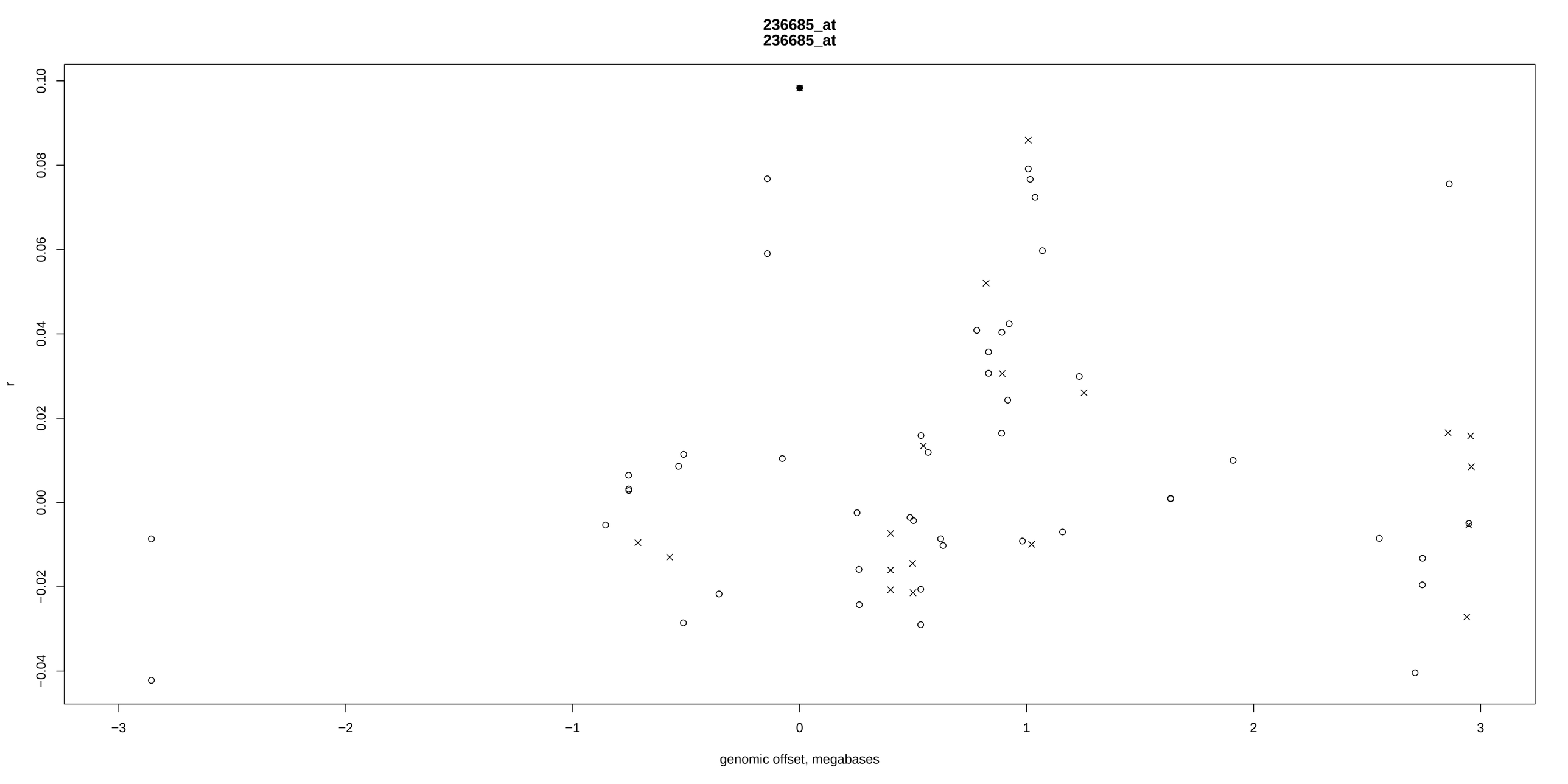

CMAH
229604_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

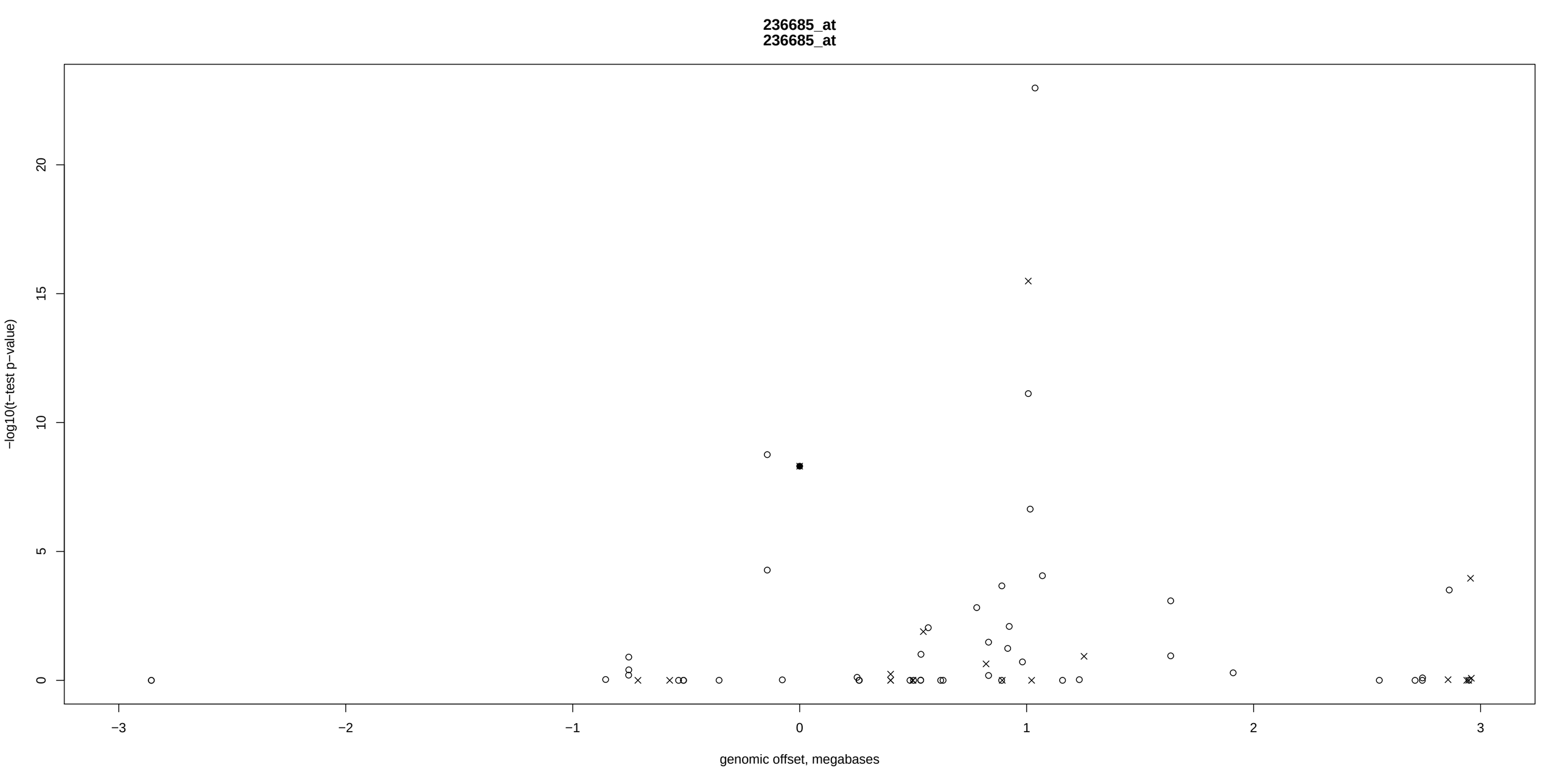

COL11A2
213870_at
r
0.0
0.1
0.2
0.3
−3
−2
−1
0
1
2
3
genomic offset, megabases

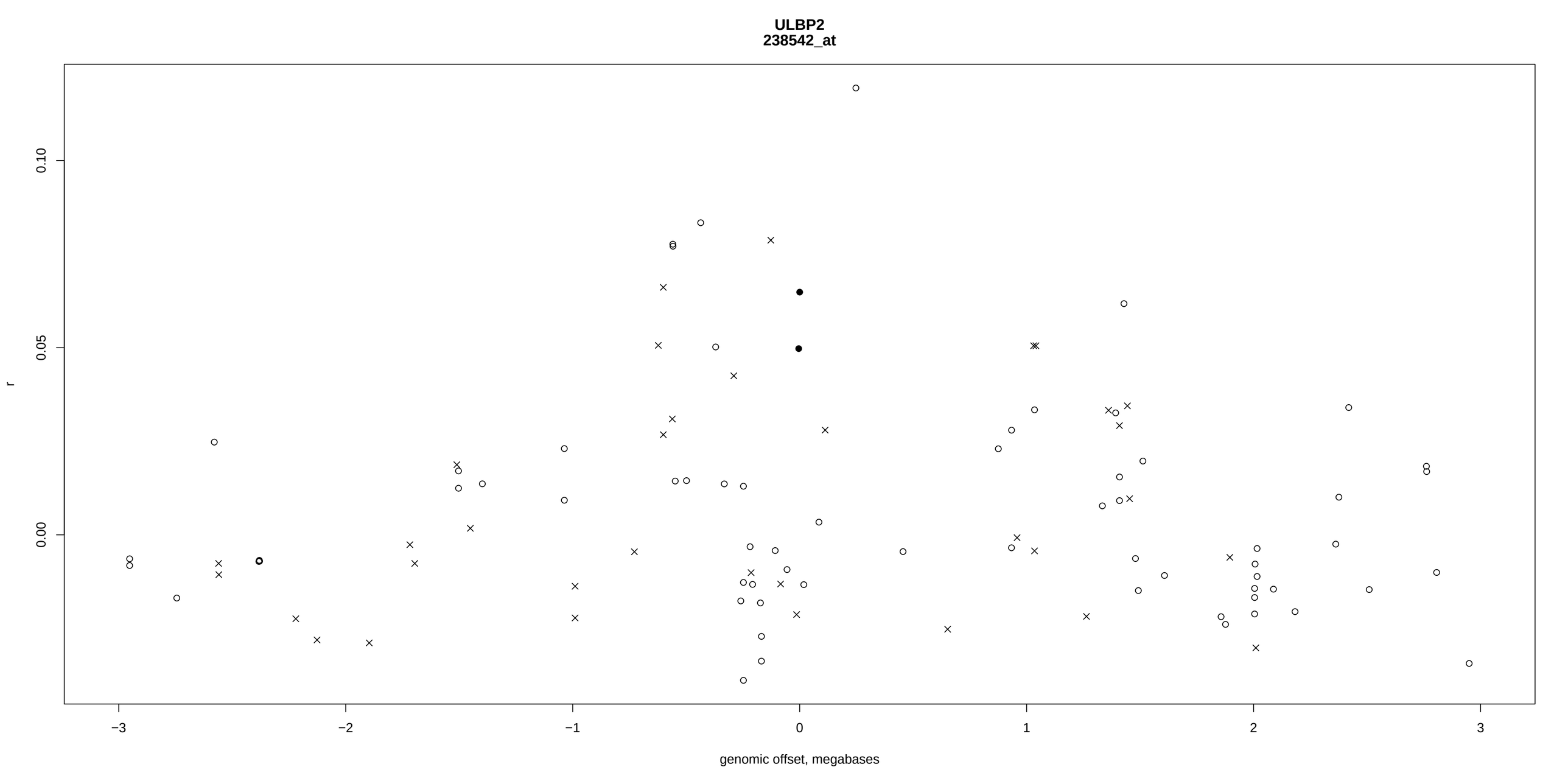

COL11A2
213870_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases

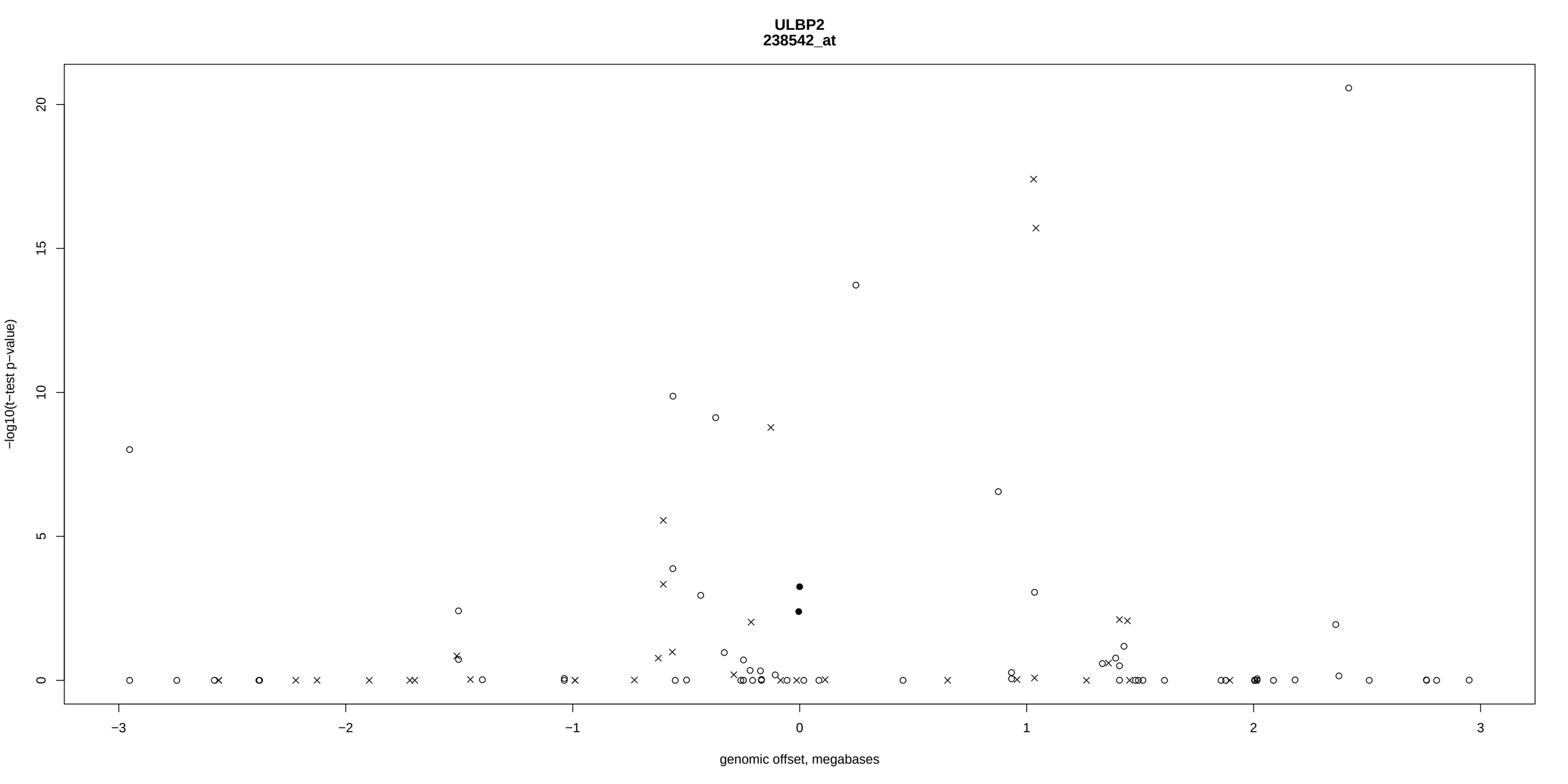

COL10A1
205941_s_at
r
0.10
0.05
0.00
−3
−2
−1
0
1
2
3
genomic offset, megabases

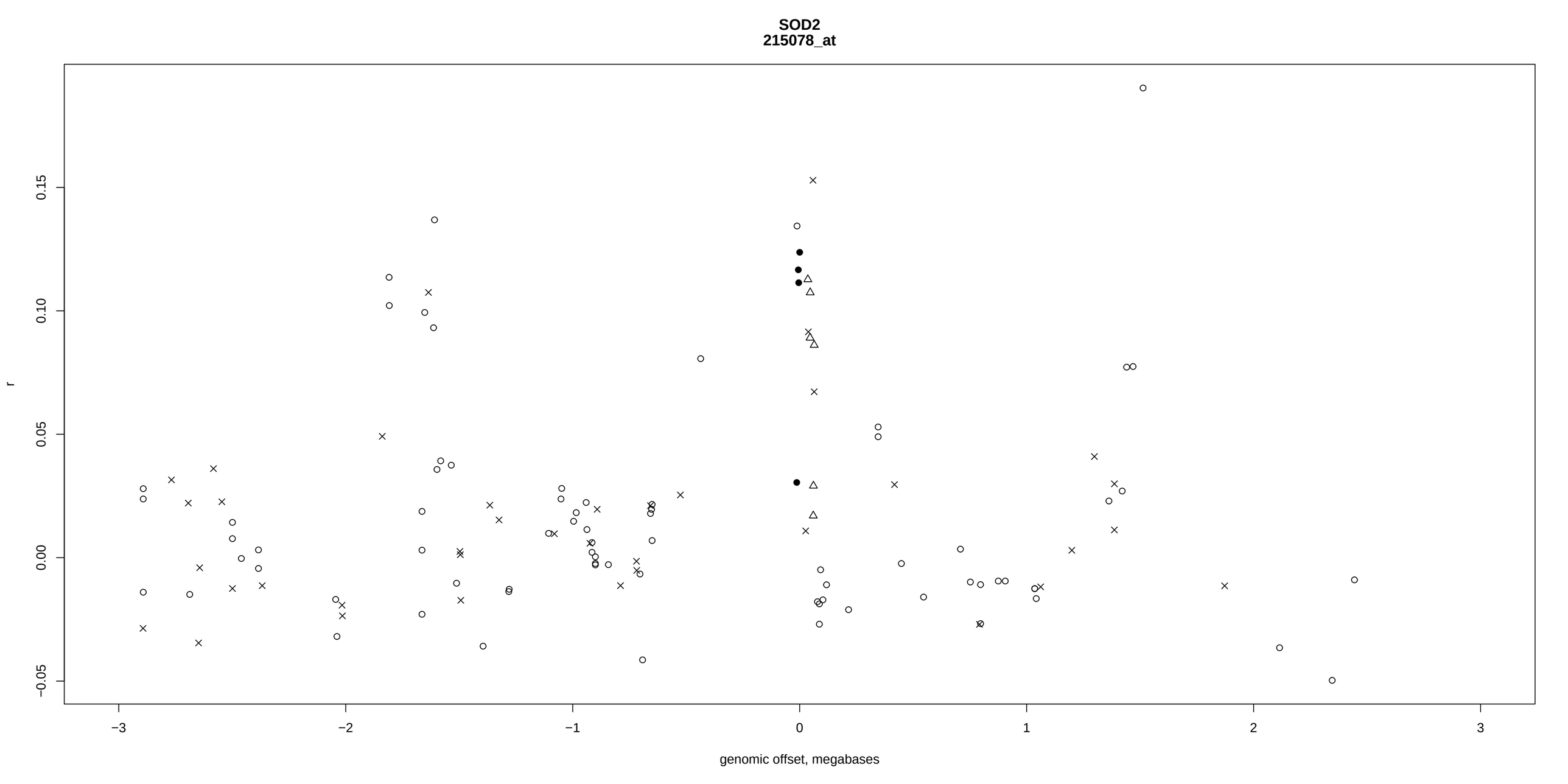

COL10A1
205941_s_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases

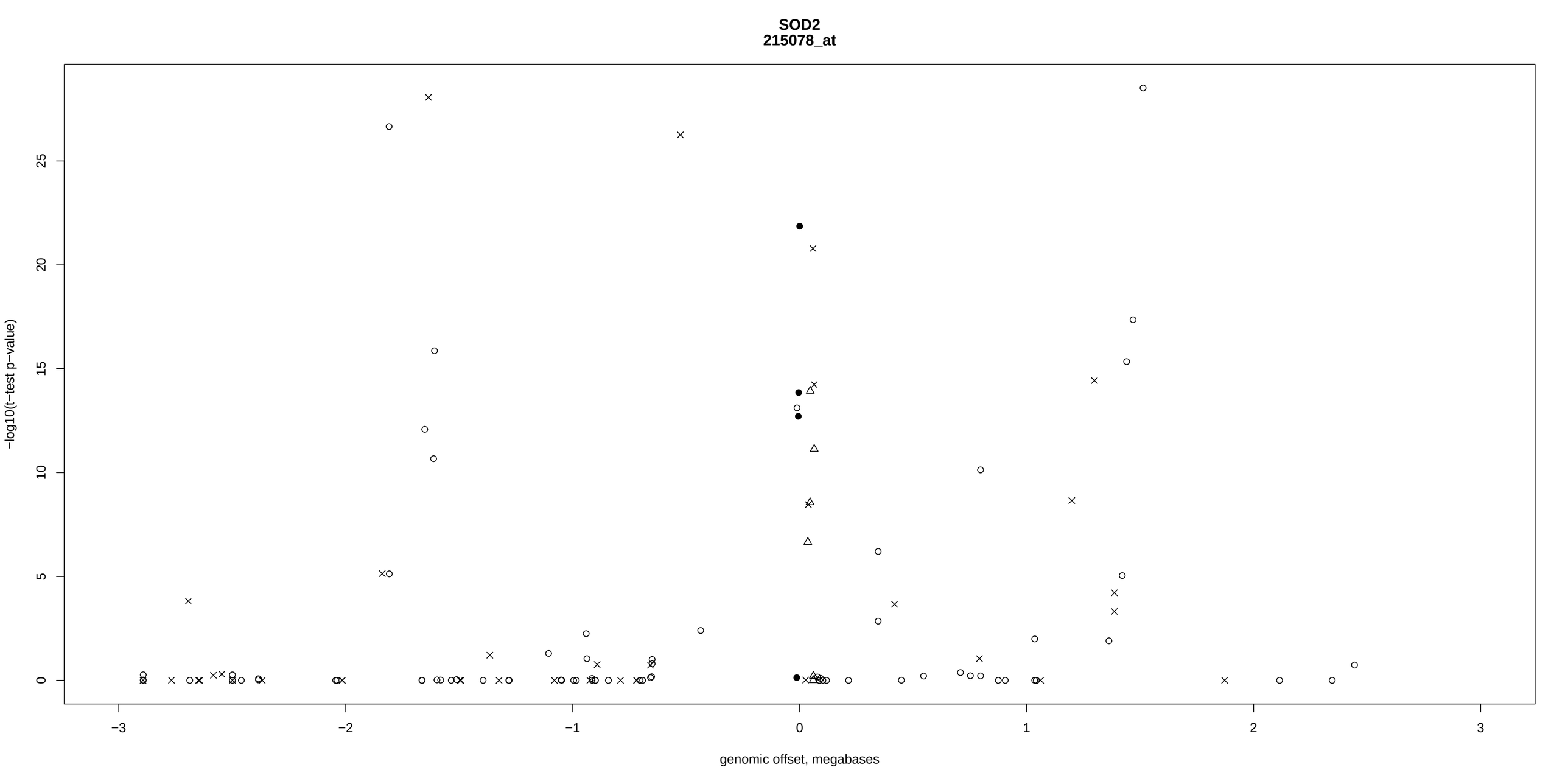

236685_at
236685_at
r
0.10
0.08
0.06
0.04
0.02
0.00
−0.02
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases

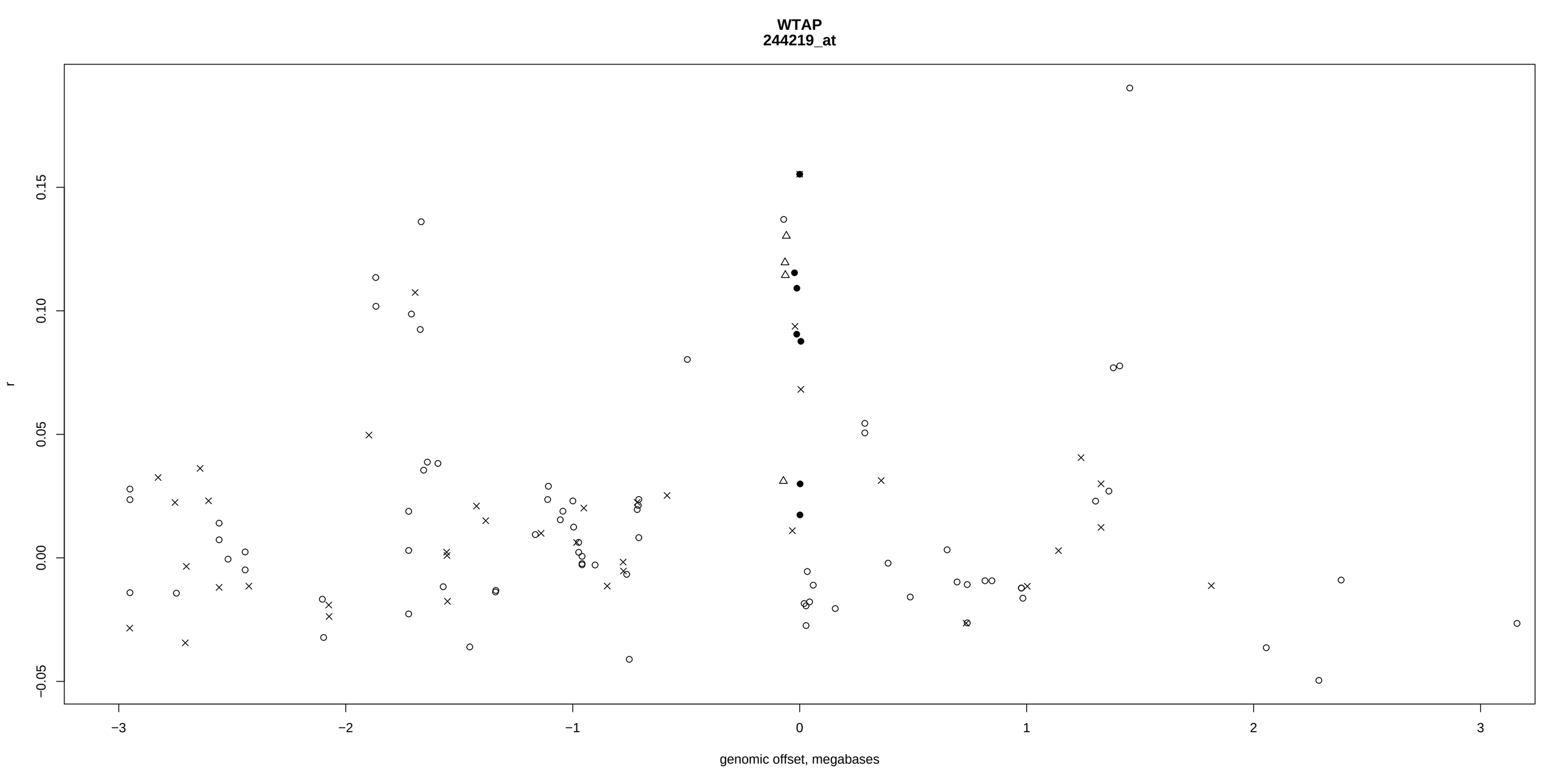

236685_at
236685_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


ULBP2
238542_at
r
0.10
0.05
0.00
−3
−2
−1
0
1
2
3
genomic offset, megabases


ULBP2
238542_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


SOD2
215078_at
r
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


SOD2
215078_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


WTAP
244219_at
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0.05
0.00
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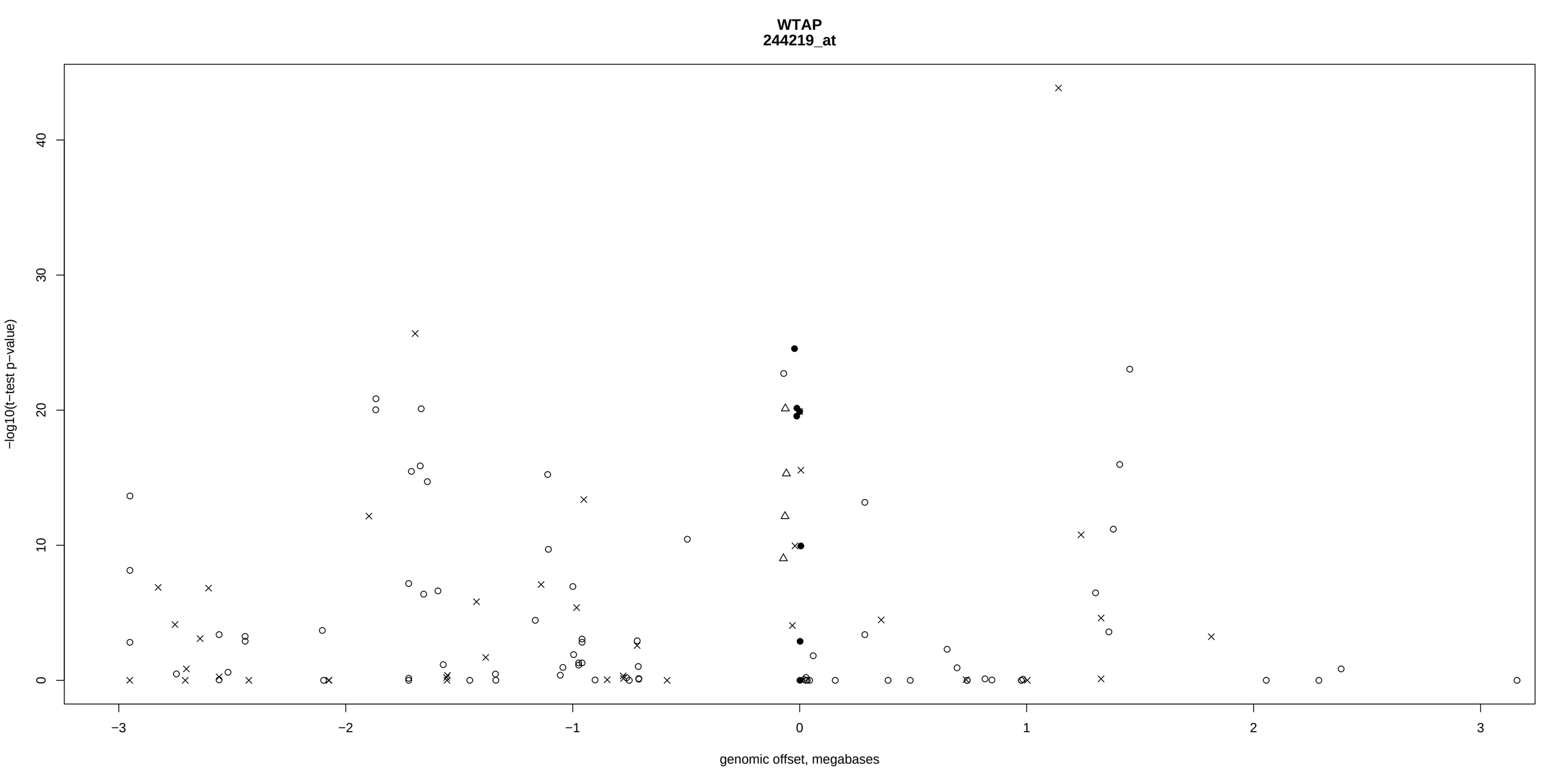
−3
−2
−1
0
1
2
3
genomic offset, megabases


WTAP
244219_at
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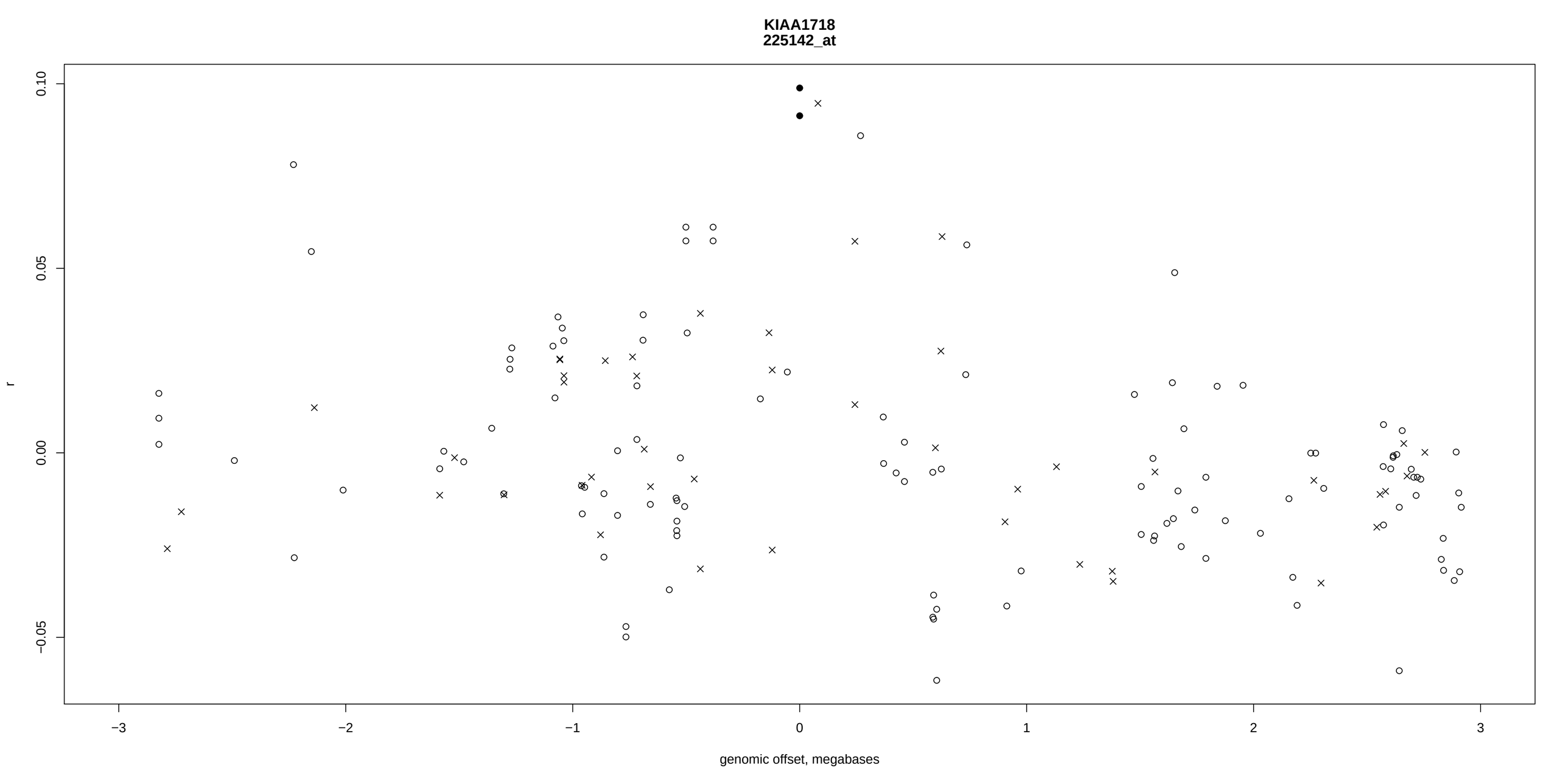
0
10
20
30
40
−3
−2
−1
0
1
2
3
genomic offset, megabases


KIAA1718
225142_at
0.10
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0.00
−0.05
r
−3
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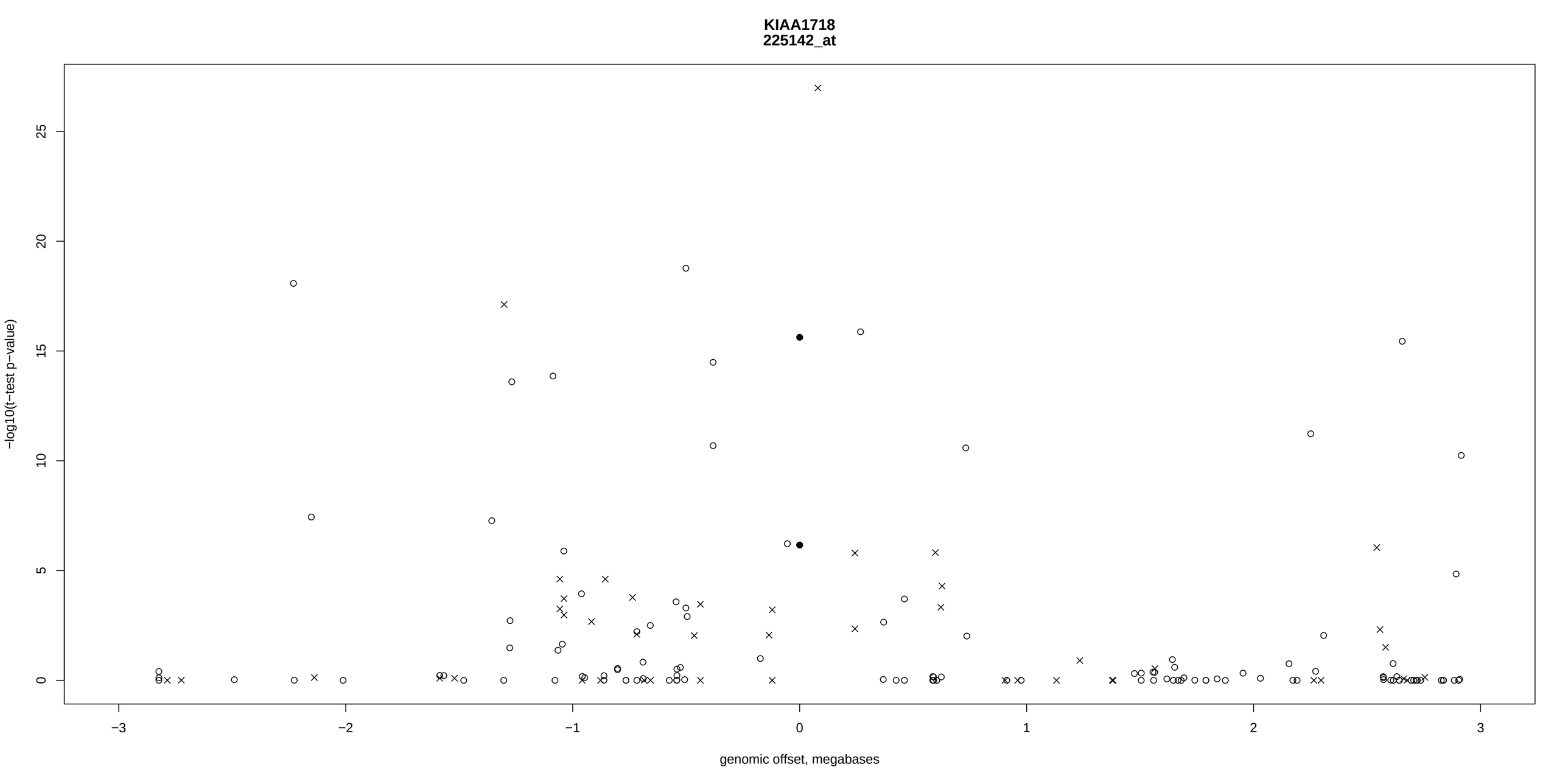
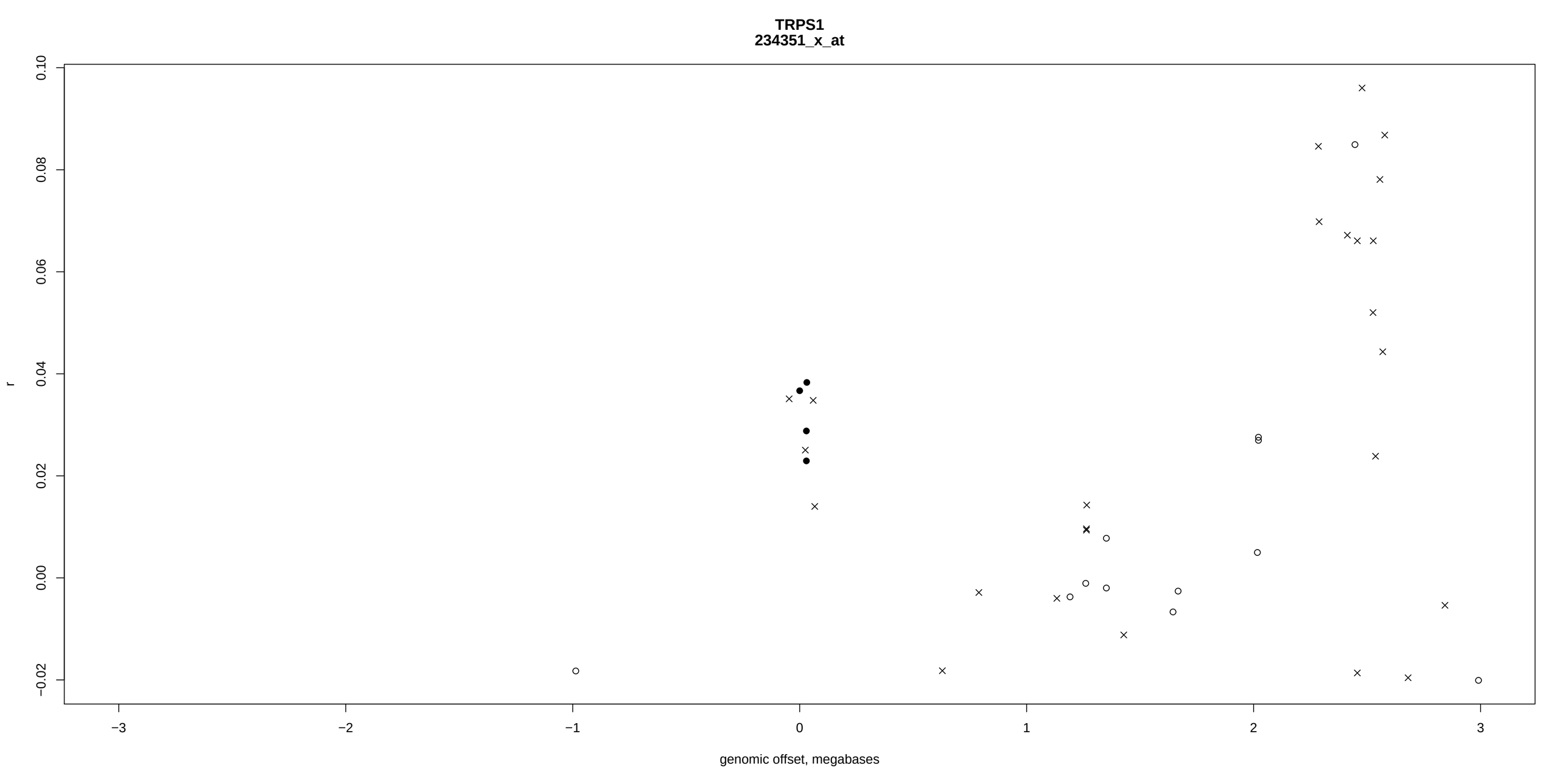
−1
0
1
2
3
genomic offset, megabases


KIAA1718
225142_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


TRPS1
234351_x_at
r
0.10
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0.06
0.04
0.02
0.00
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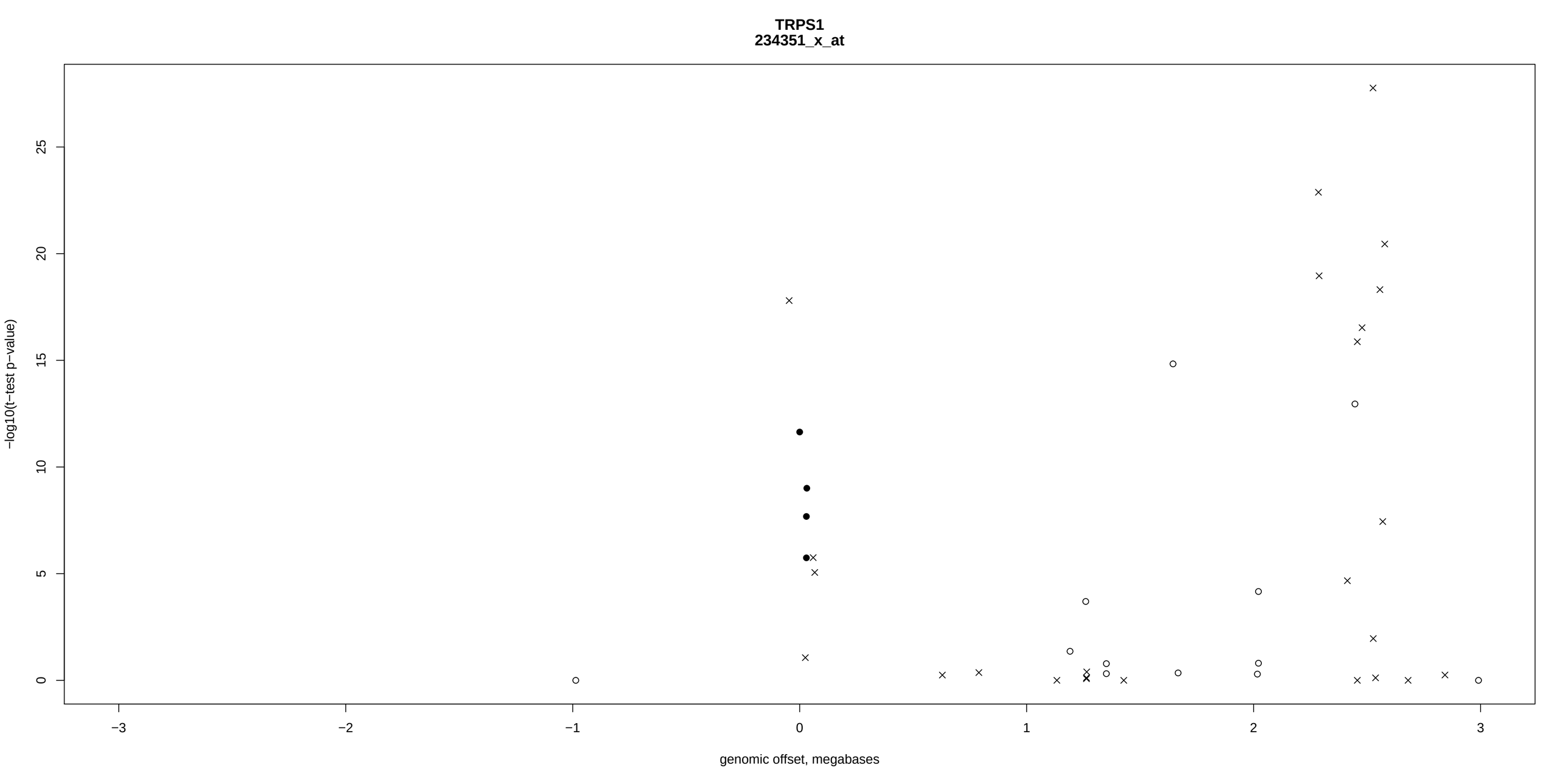
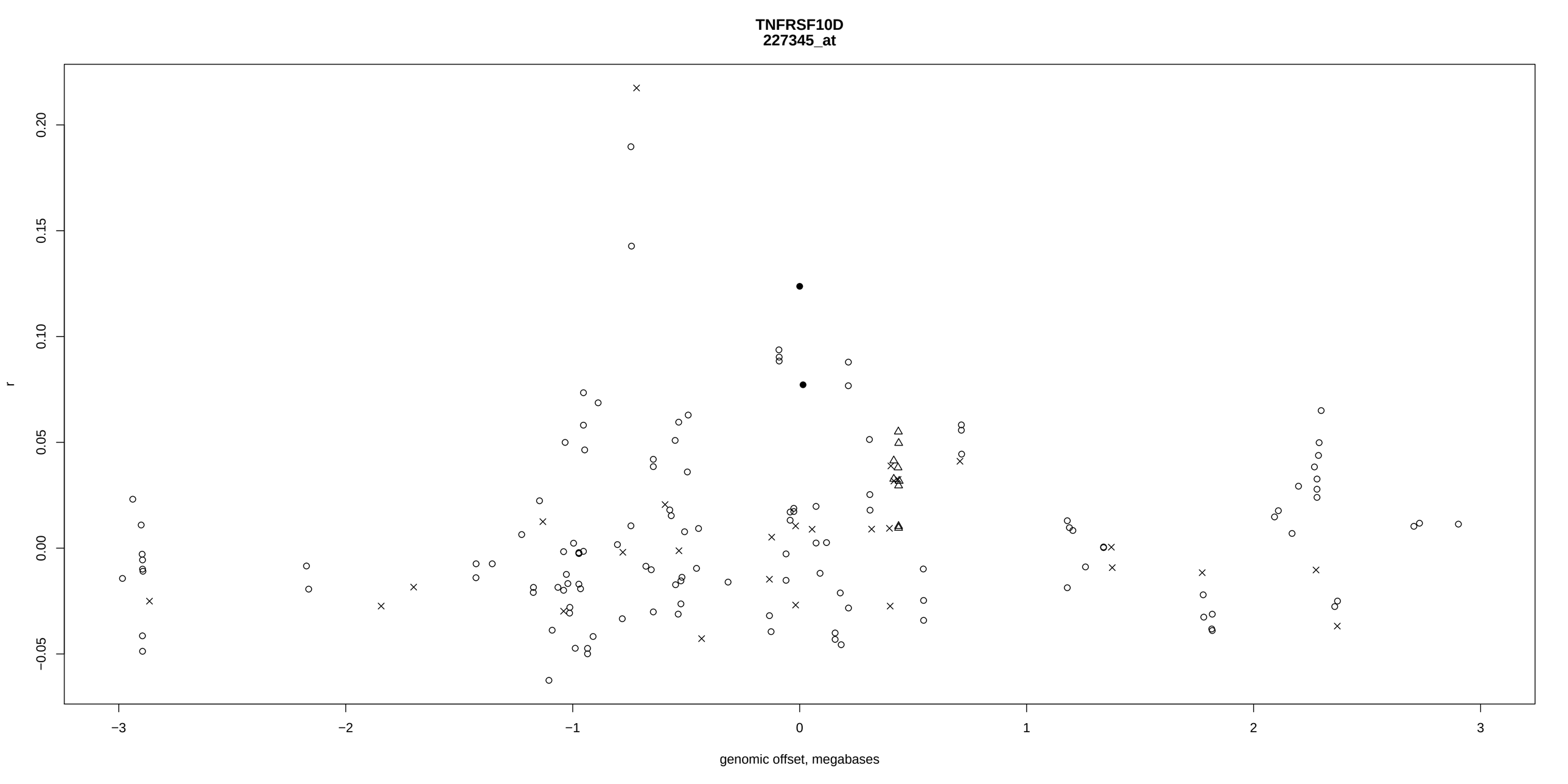
−3
−2
−1
0
1
2
3
genomic offset, megabases


TRPS1
234351_x_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


TNFRSF10D
227345_at
r
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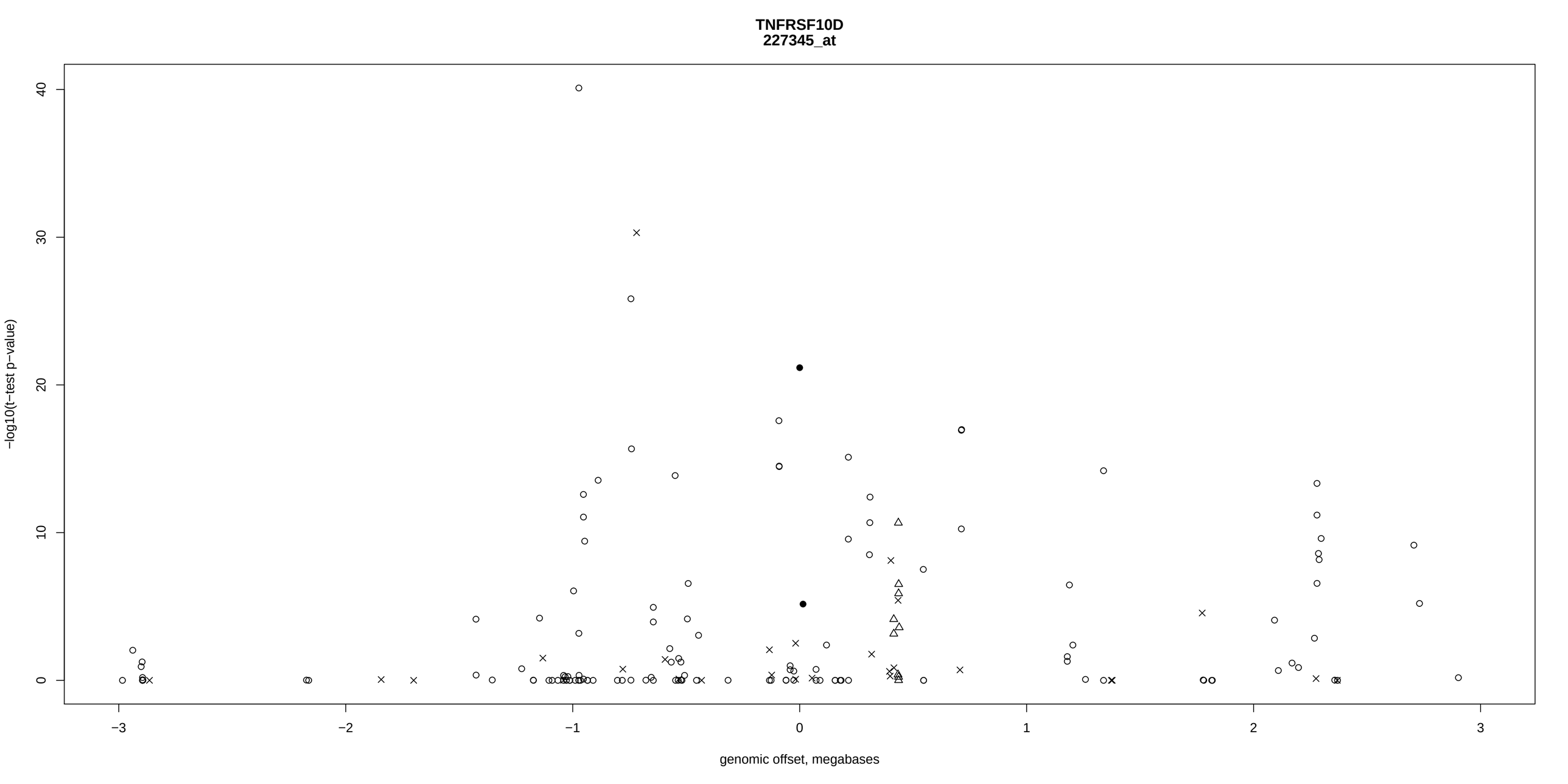
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


TNFRSF10D
227345_at
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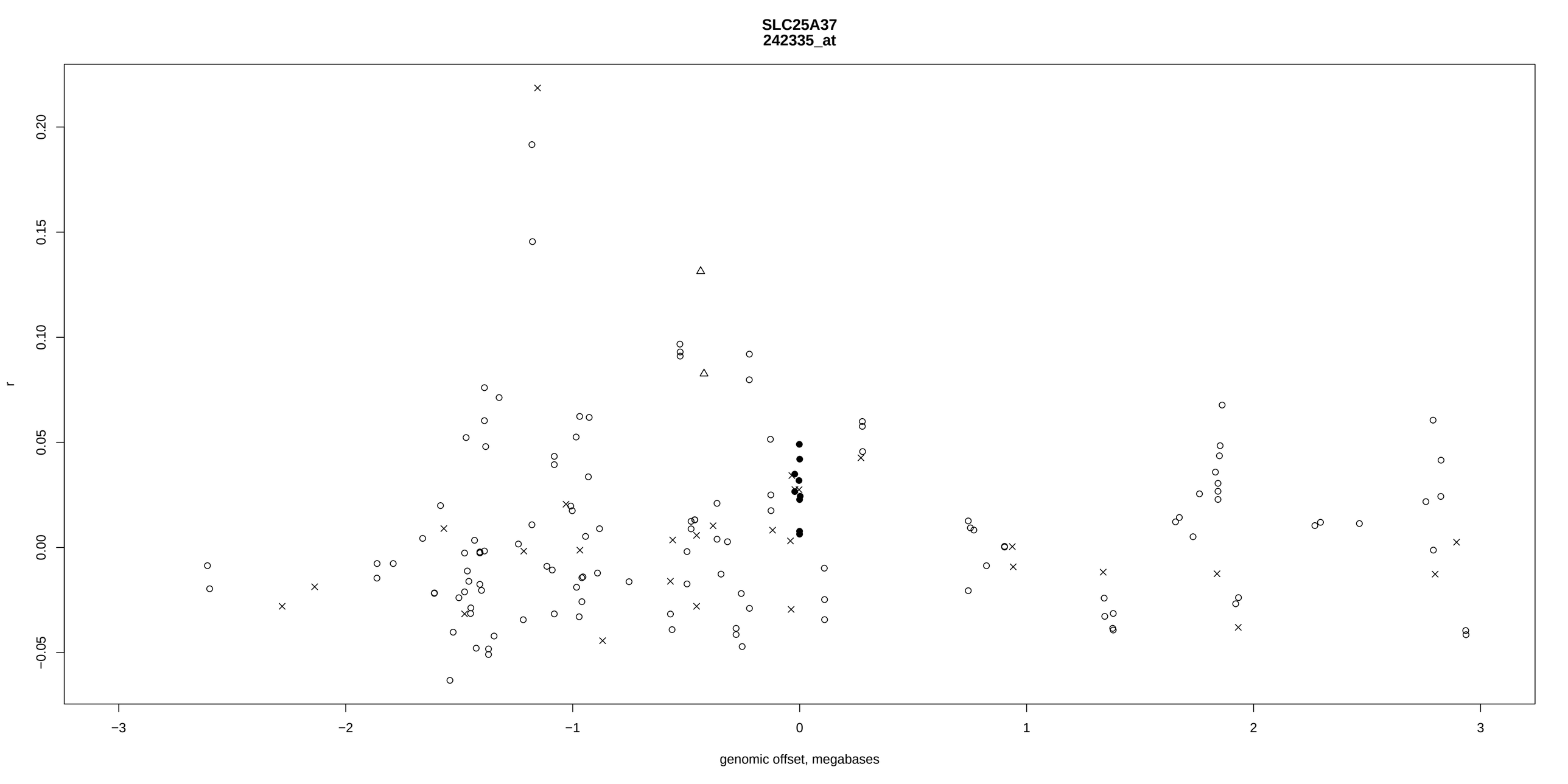
0
10
20
30
40
−3
−2
−1
0
1
2
3
genomic offset, megabases


SLC25A37
242335_at
r
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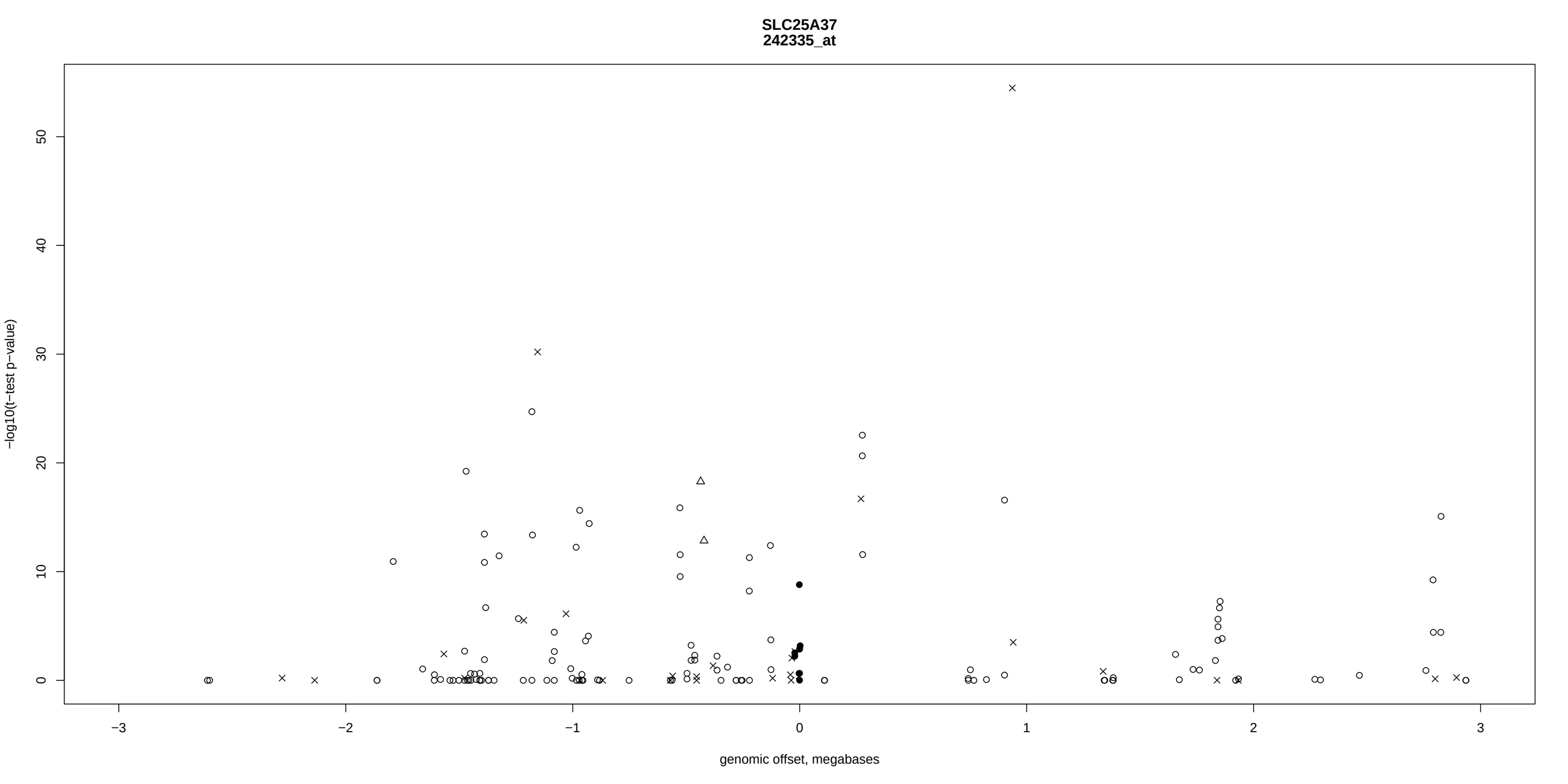
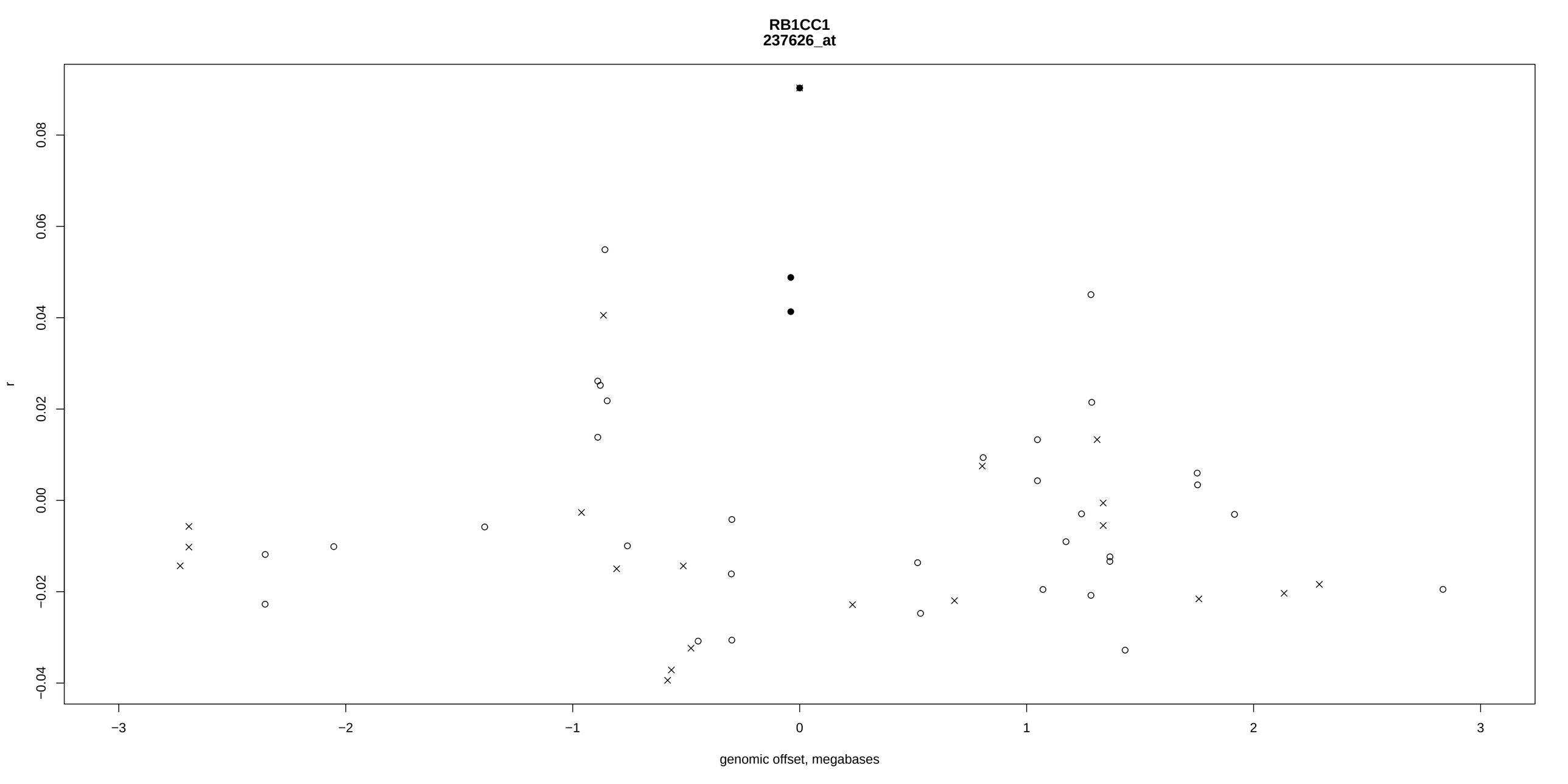
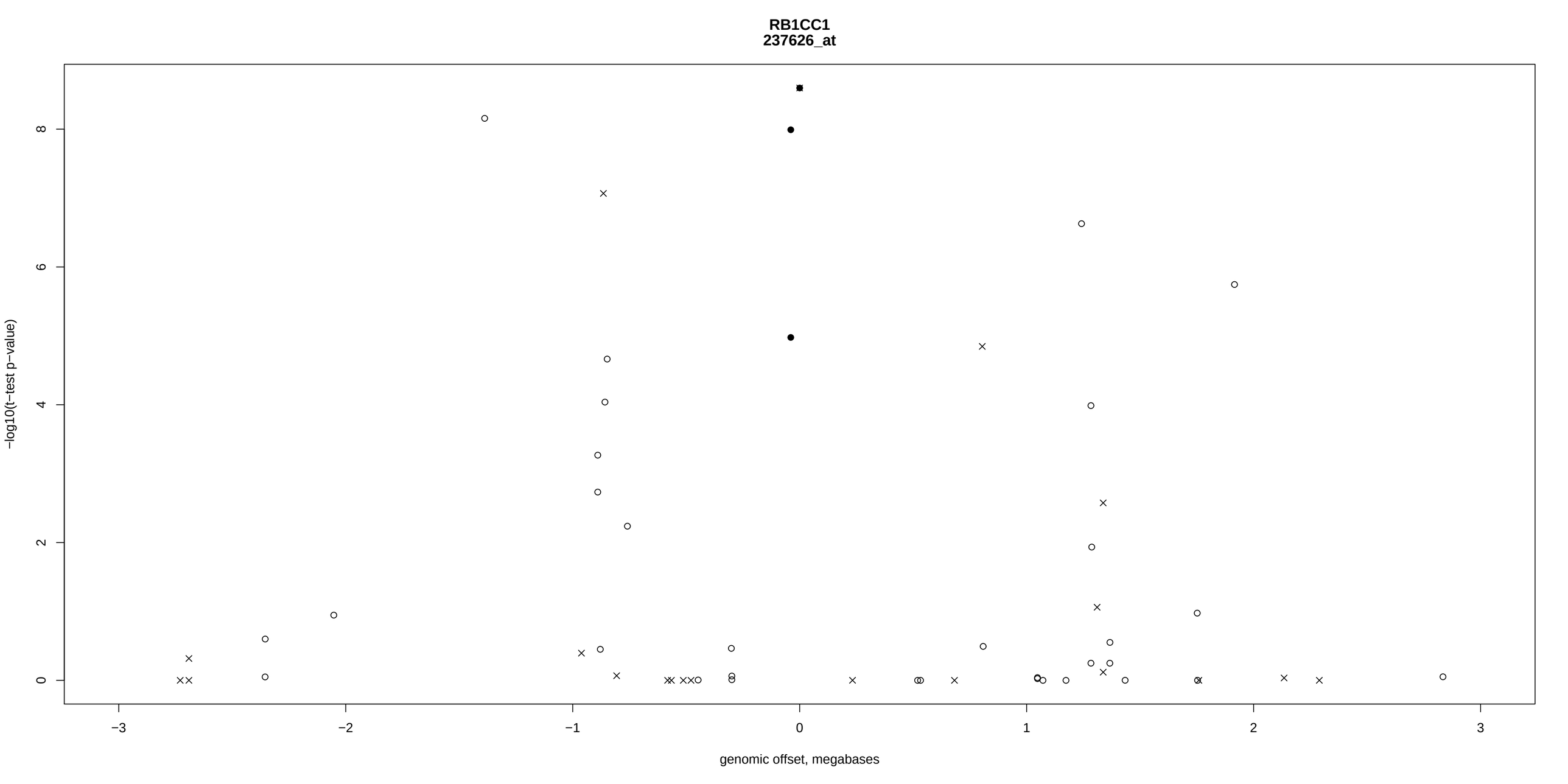
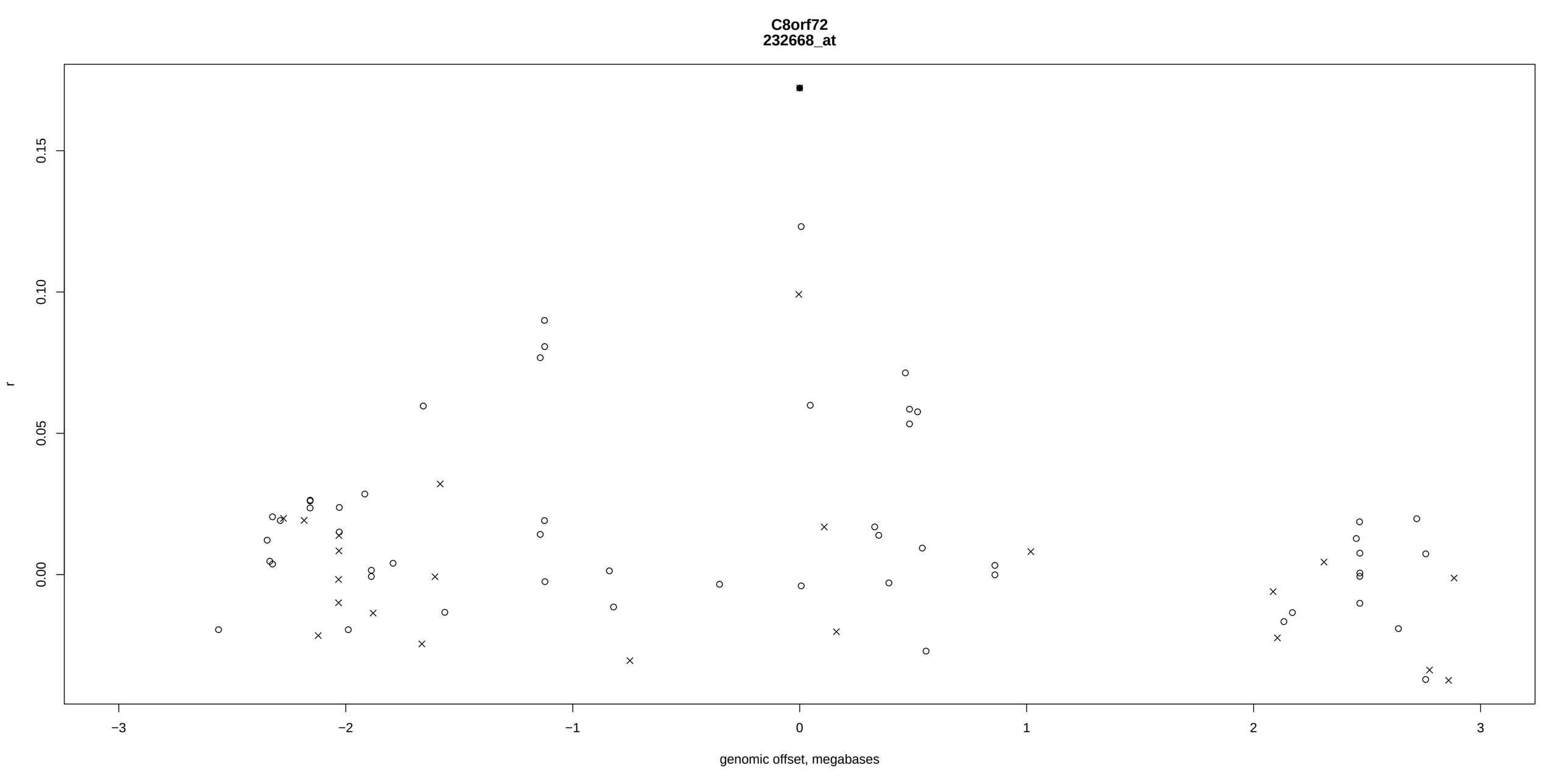
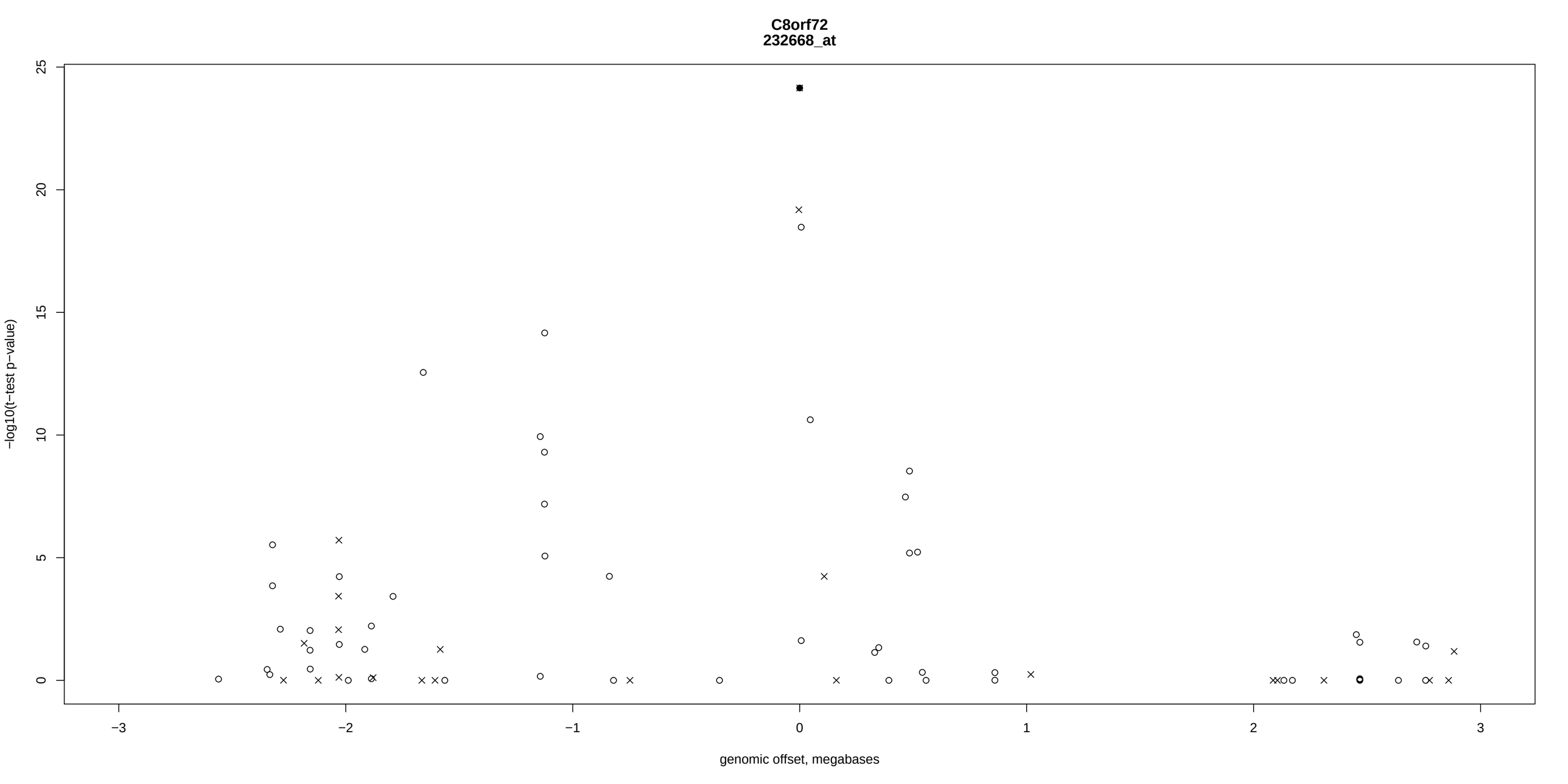
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases

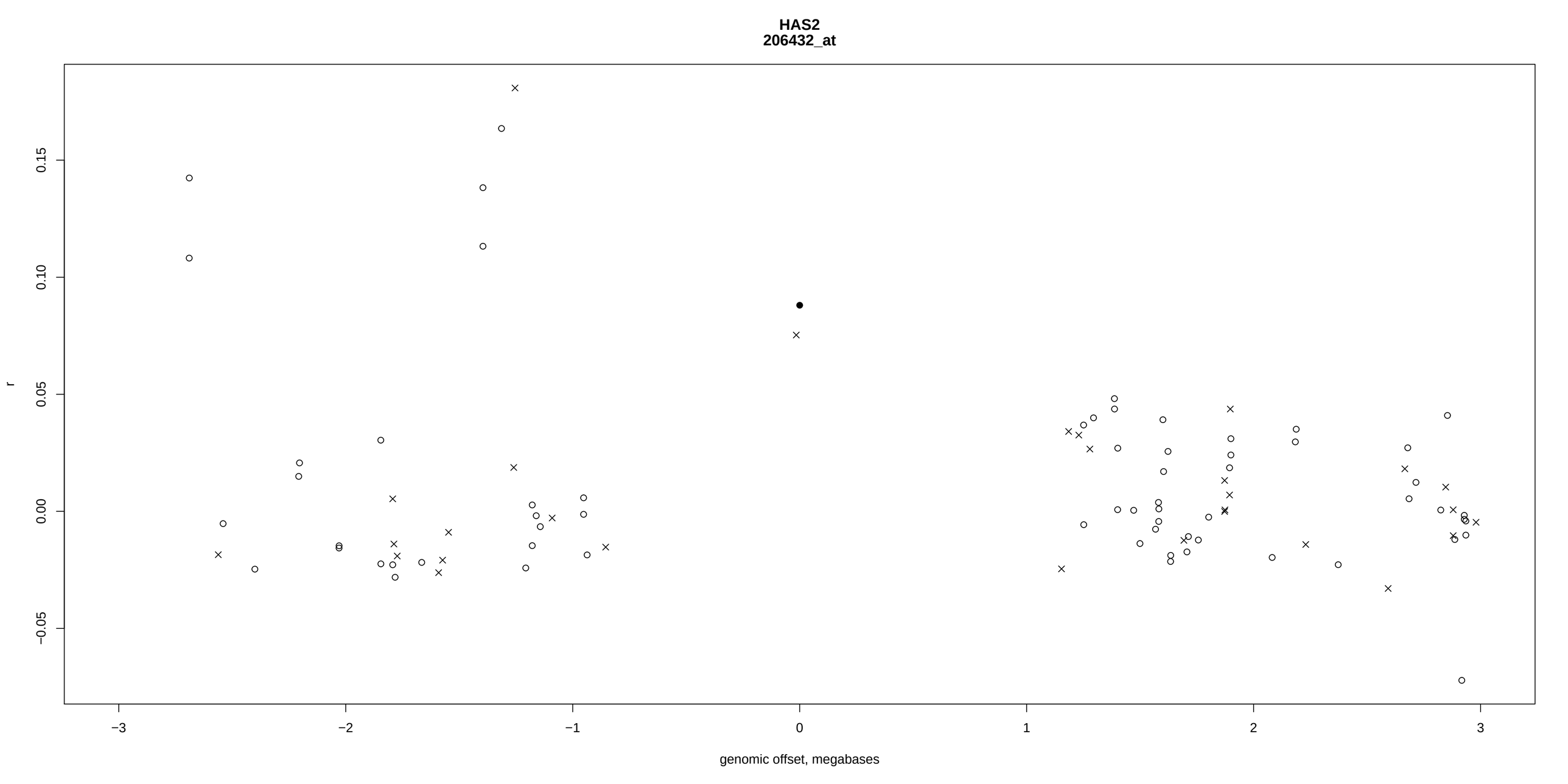

SLC25A37
242335_at
−log10(t−test p−value)
0
10
20
30
40
50
−3
−2
−1
0
1
2
3
genomic offset, megabases

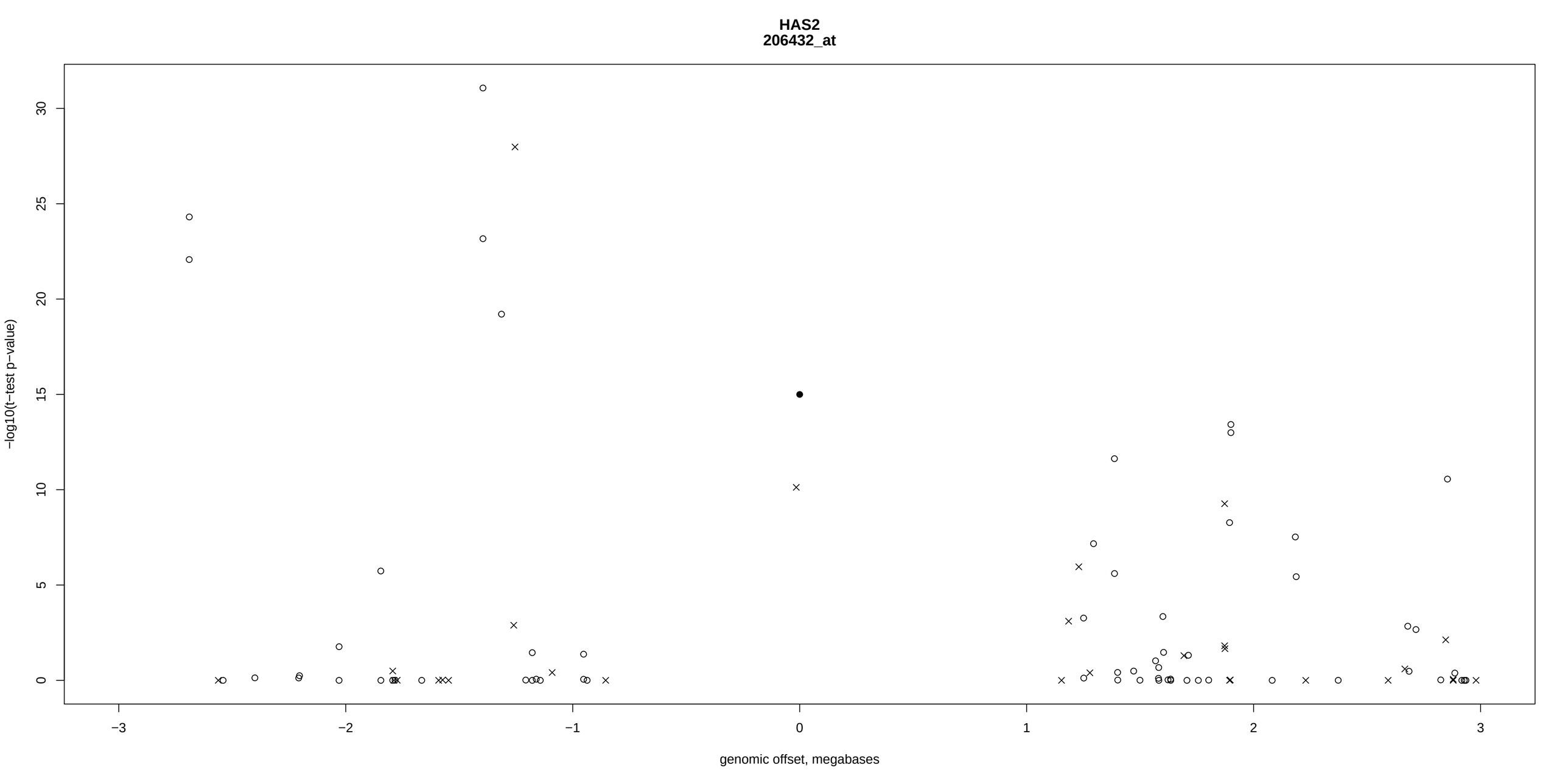

RB1CC1
237626_at
r
0.08
0.06
0.04
0.02
0.00
−0.02
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases

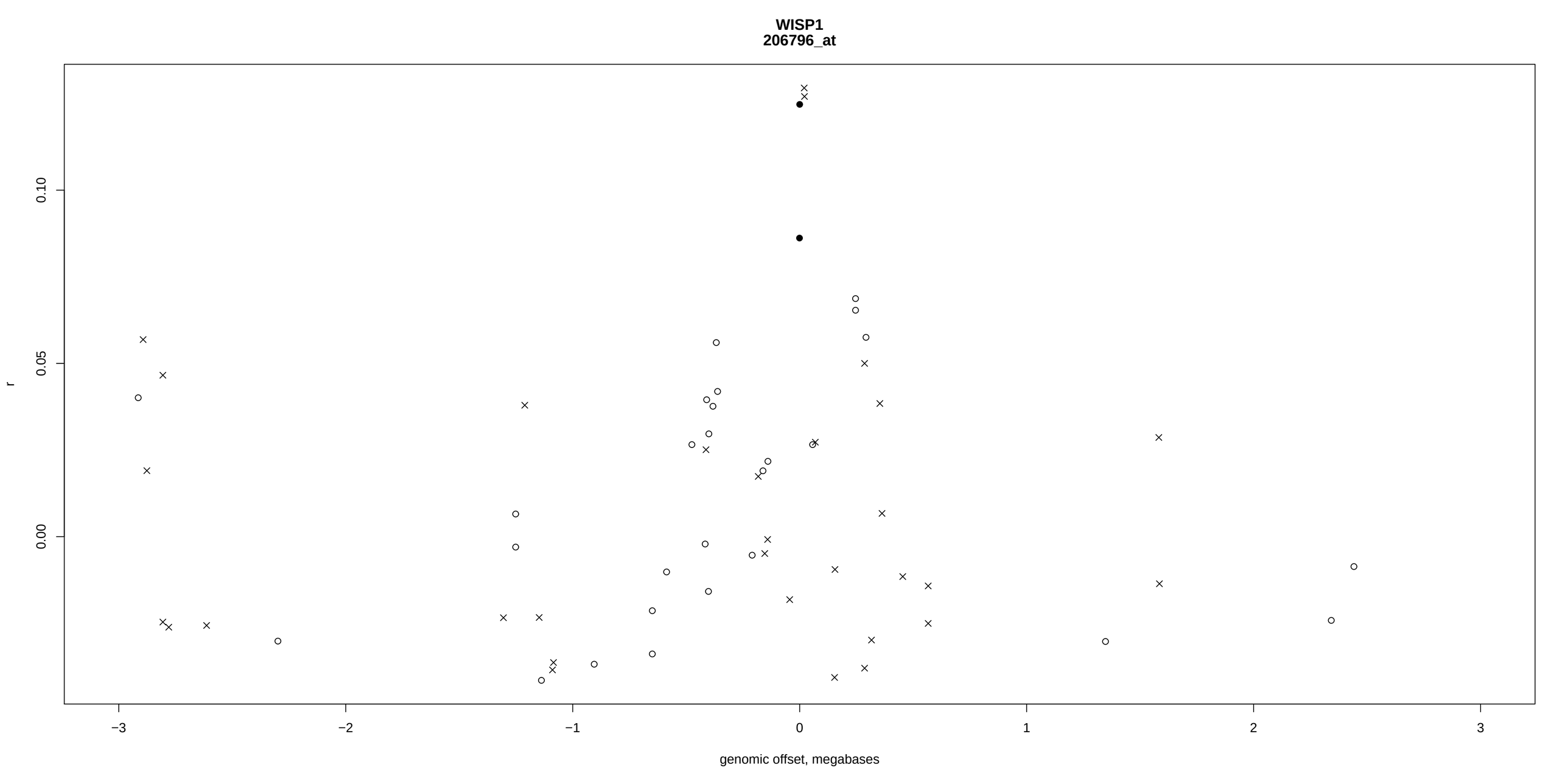

RB1CC1
237626_at
−log10(t−test p−value)
0
2
4
6
8
−3
−2
−1
0
1
2
3
genomic offset, megabases


C8orf72
232668_at
r
0.00
0.05
0.10
0.15
−3
−2
−1
0
1
2
3
genomic offset, megabases


C8orf72
232668_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


HAS2
206432_at
r
0.15
0.10
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


HAS2
206432_at
−log10(t−test p−value)
0
5
10
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


WISP1
206796_at
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0.10
0.05
0.00
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−1
0
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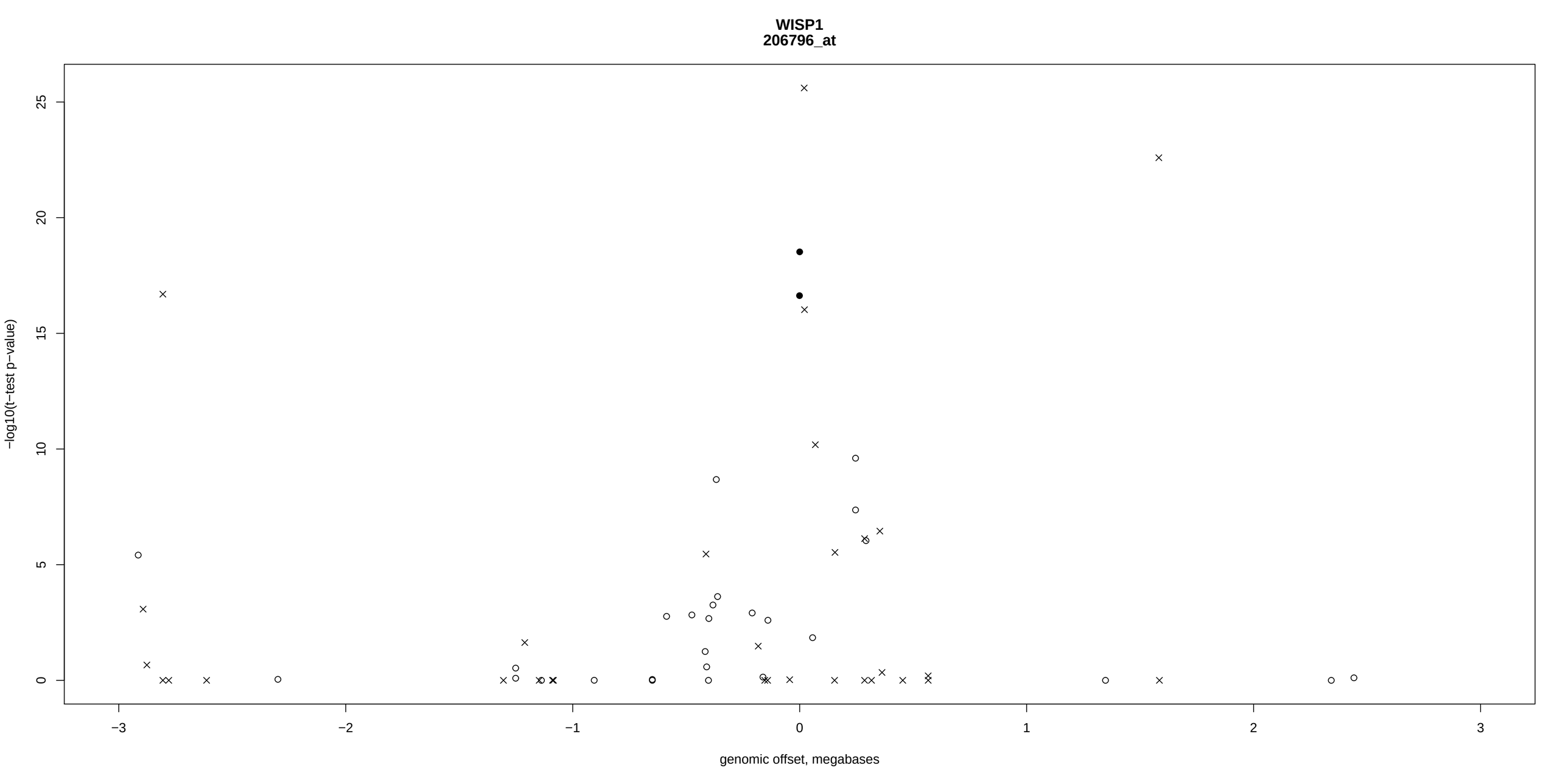
2
3
genomic offset, megabases


WISP1
206796_at
−log10(t−test p−value)
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5
10
15
20
25
−3
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−1
0
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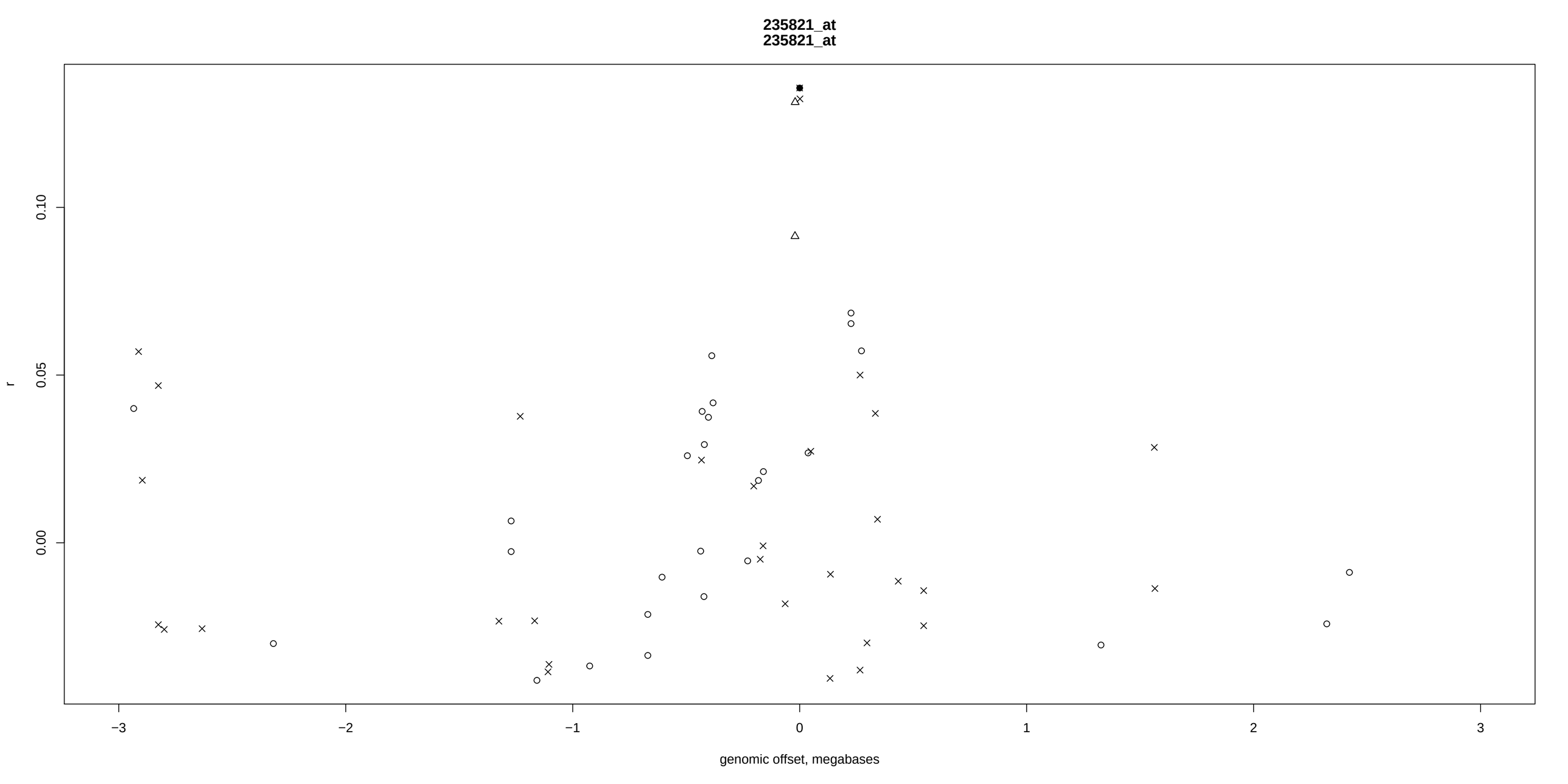
2
3
genomic offset, megabases


235821_at
235821_at
r
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0.05
0.10
−3
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−1
0
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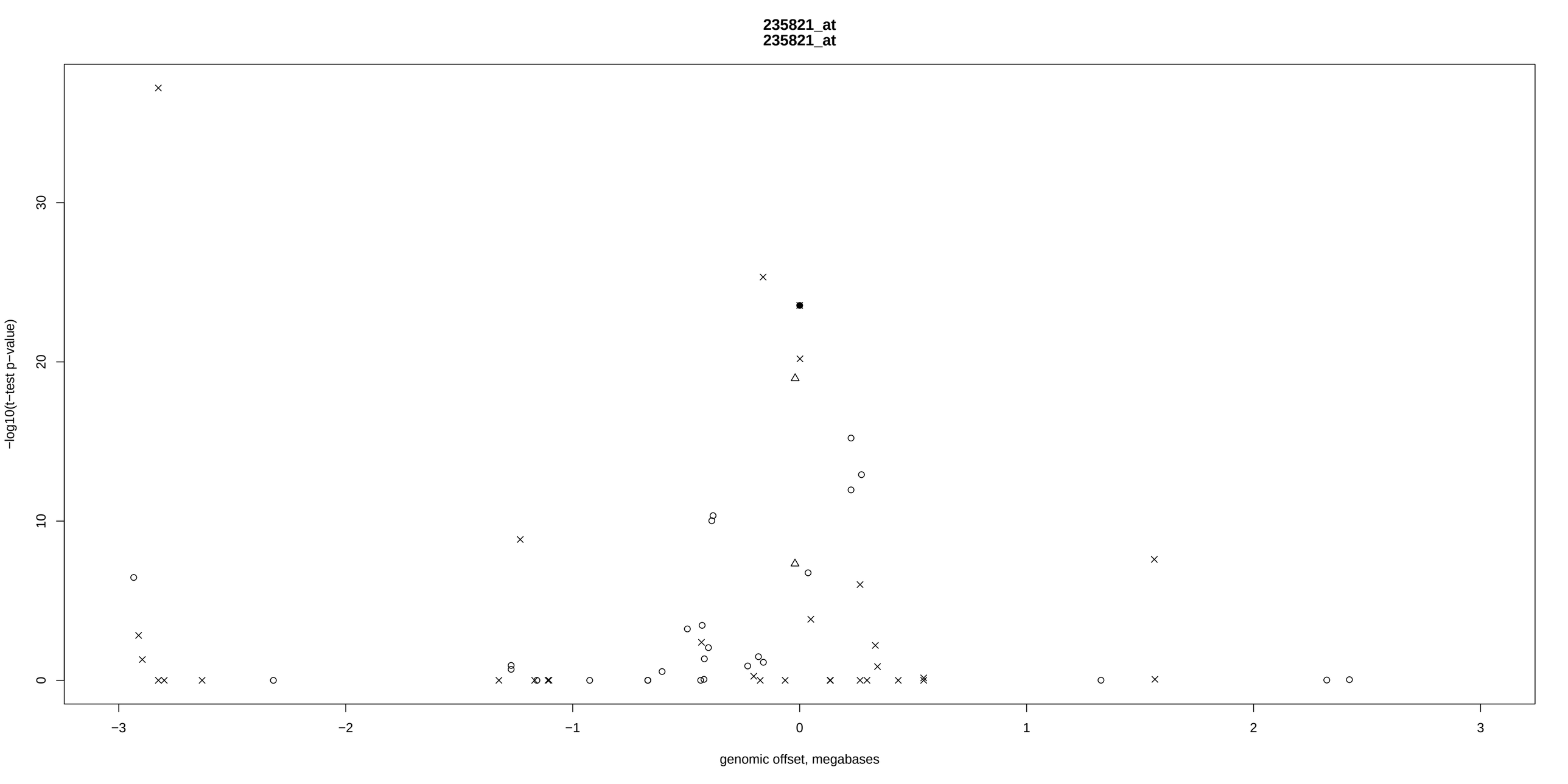
2
3
genomic offset, megabases


235821_at
235821_at
−log10(t−test p−value)
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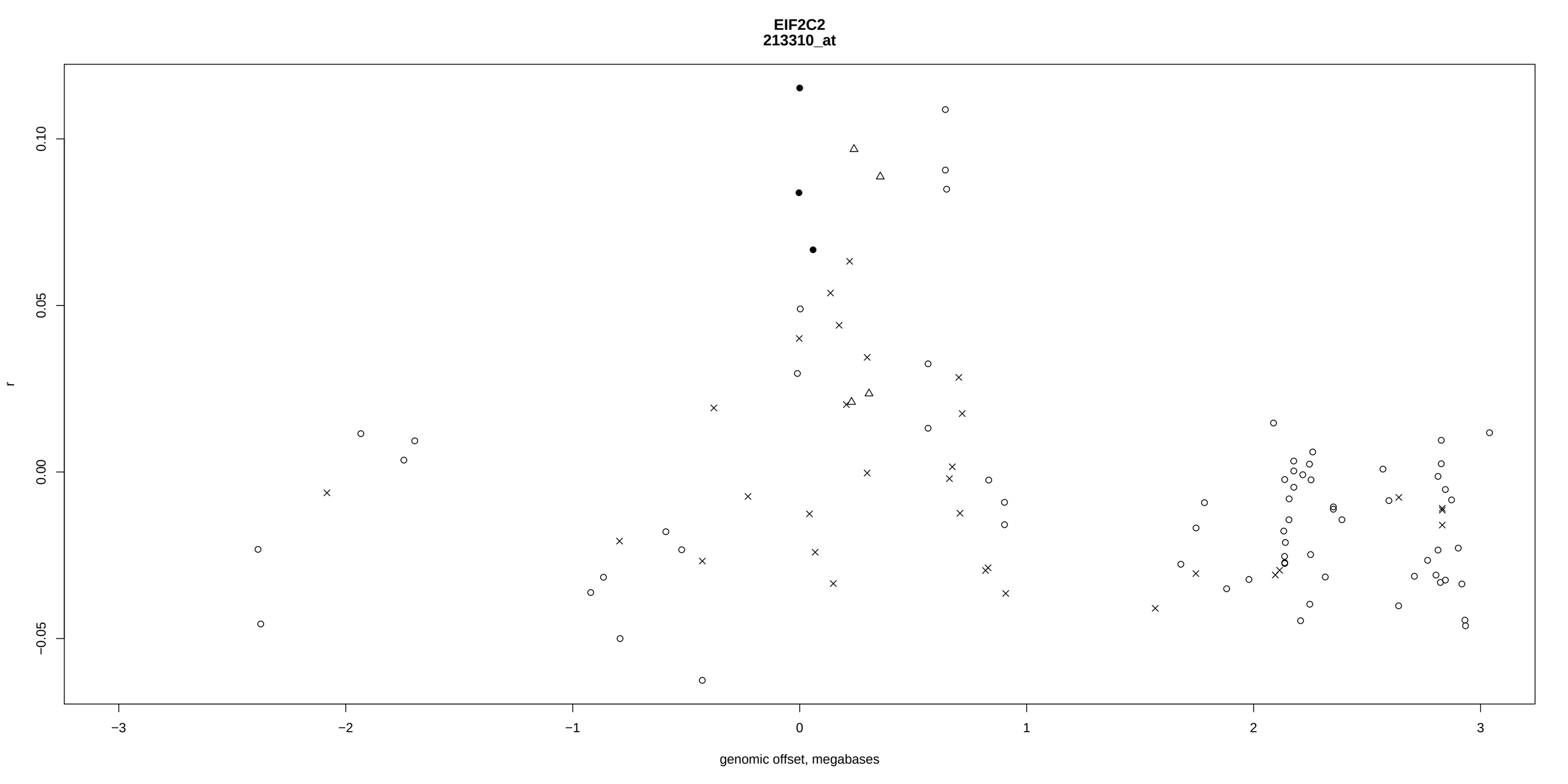
10
20
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


EIF2C2
213310_at
r
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0.05
0.00
−0.05
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−1
0
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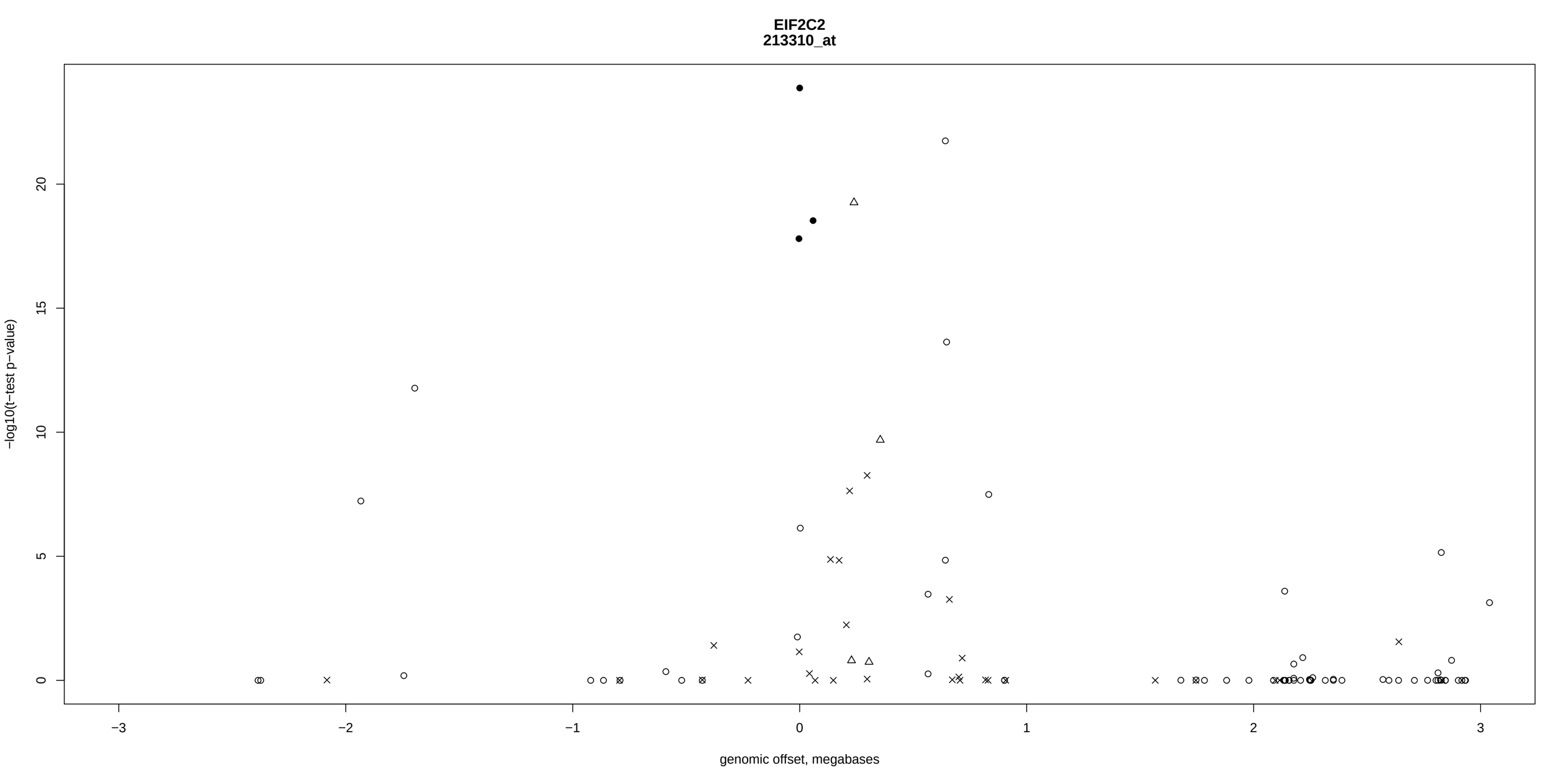
3
genomic offset, megabases


EIF2C2
213310_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
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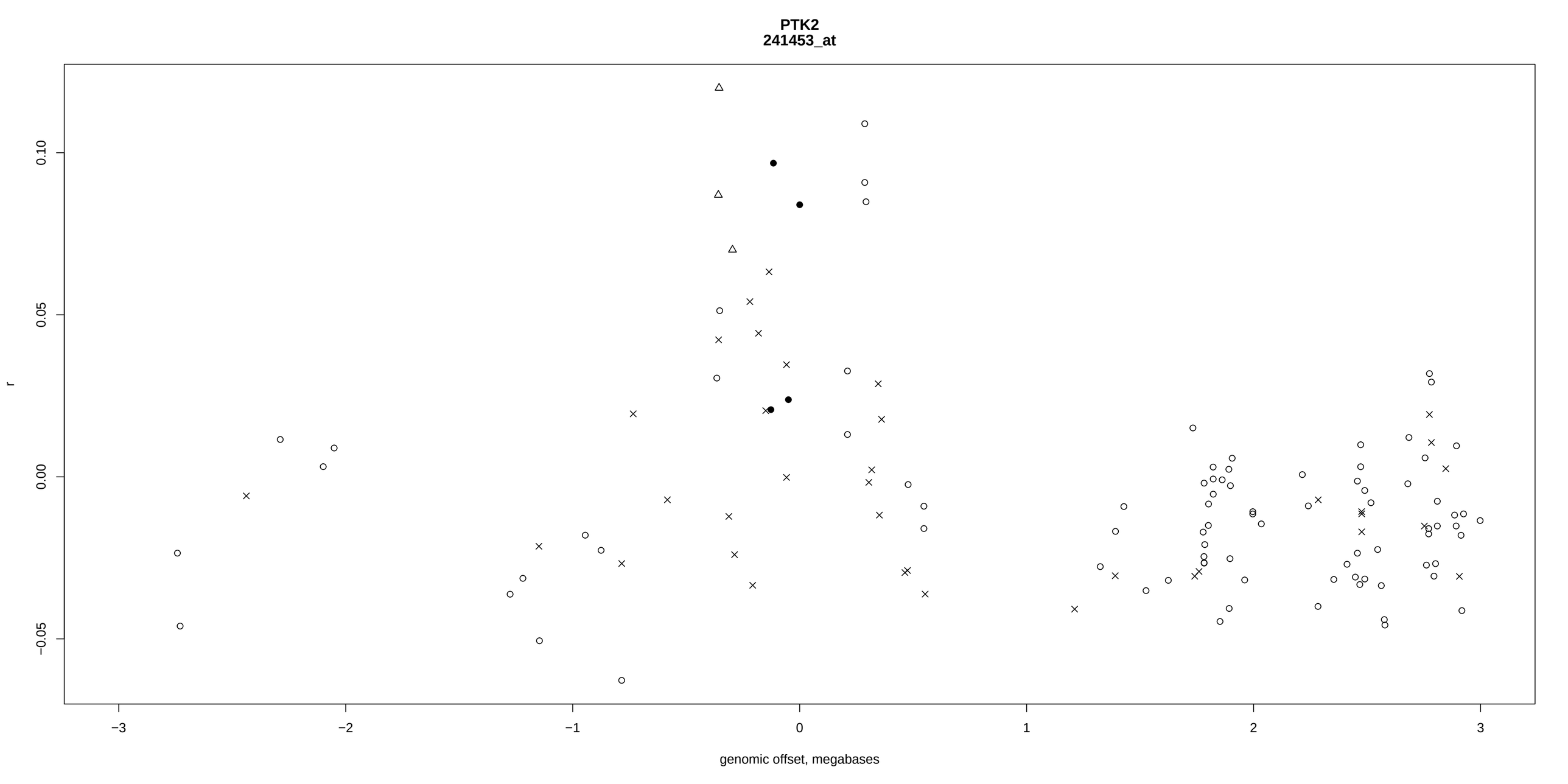
2
3
genomic offset, megabases


PTK2
241453_at
r
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0.05
0.00
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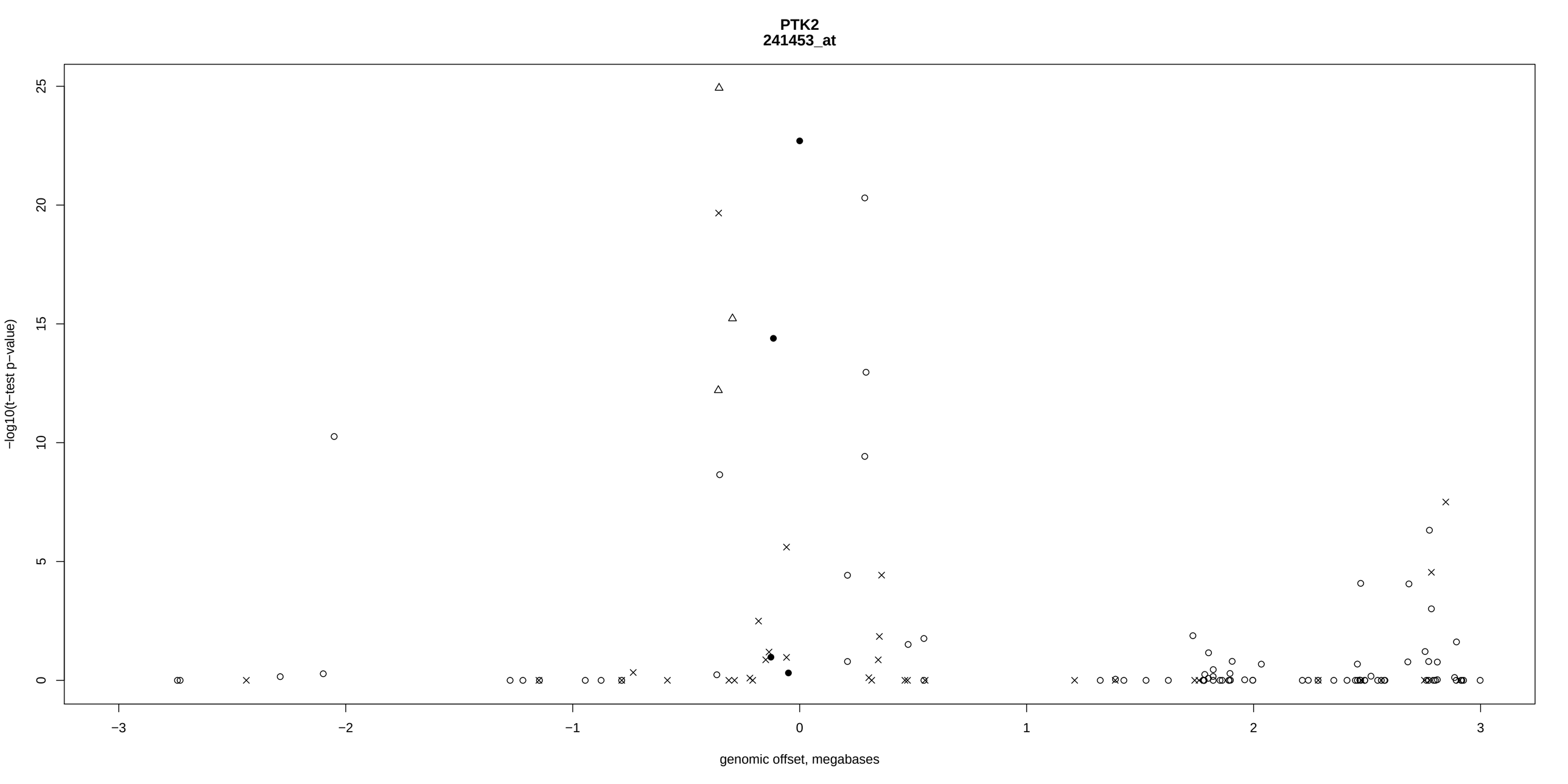
−3
−2
−1
0
1
2
3
genomic offset, megabases


PTK2
241453_at
−log10(t−test p−value)
0
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10
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20
25
−3
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−1
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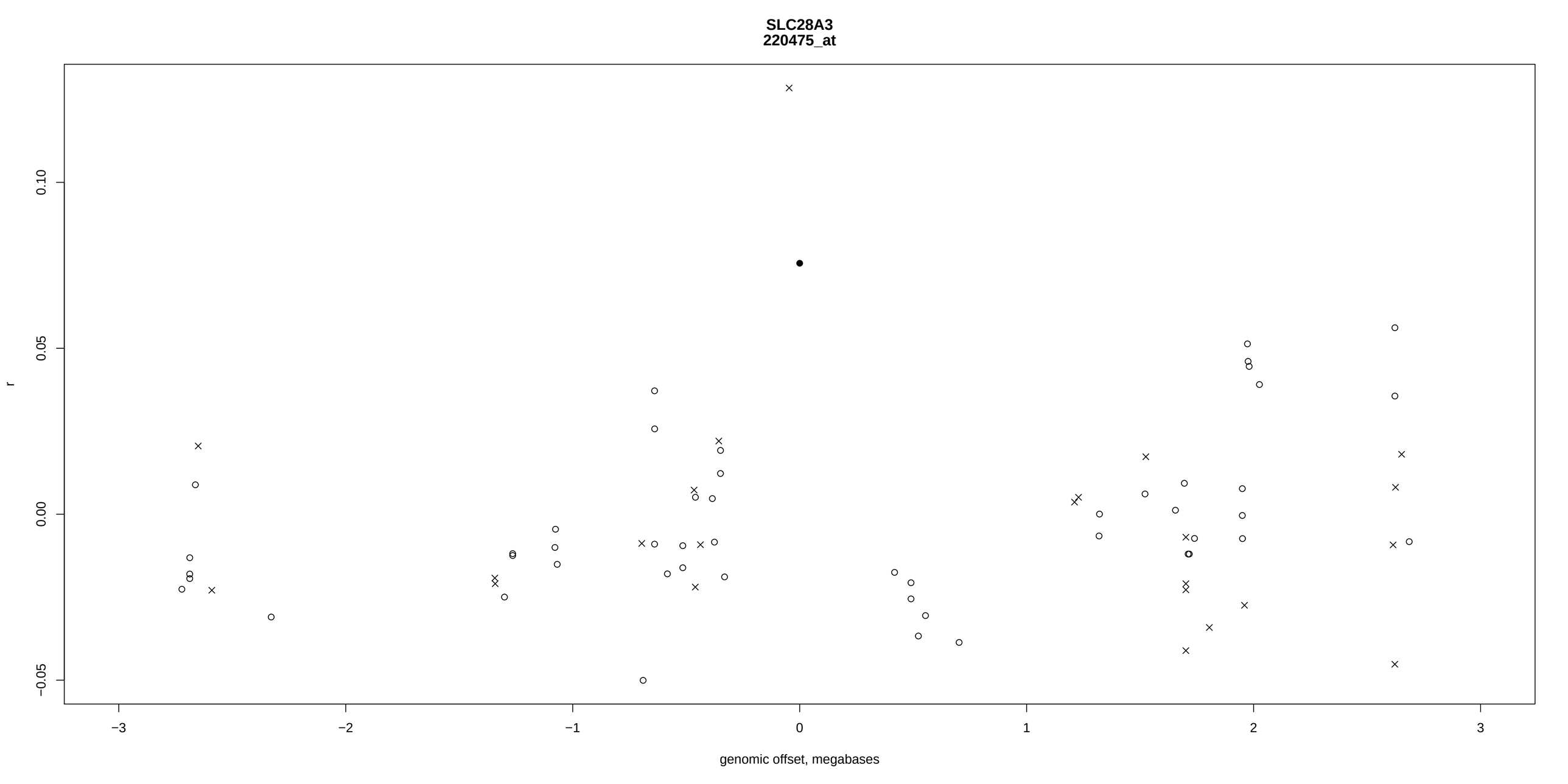
1
2
3
genomic offset, megabases


SLC28A3
220475_at
r
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0.05
0.00
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−3
−2
−1
0
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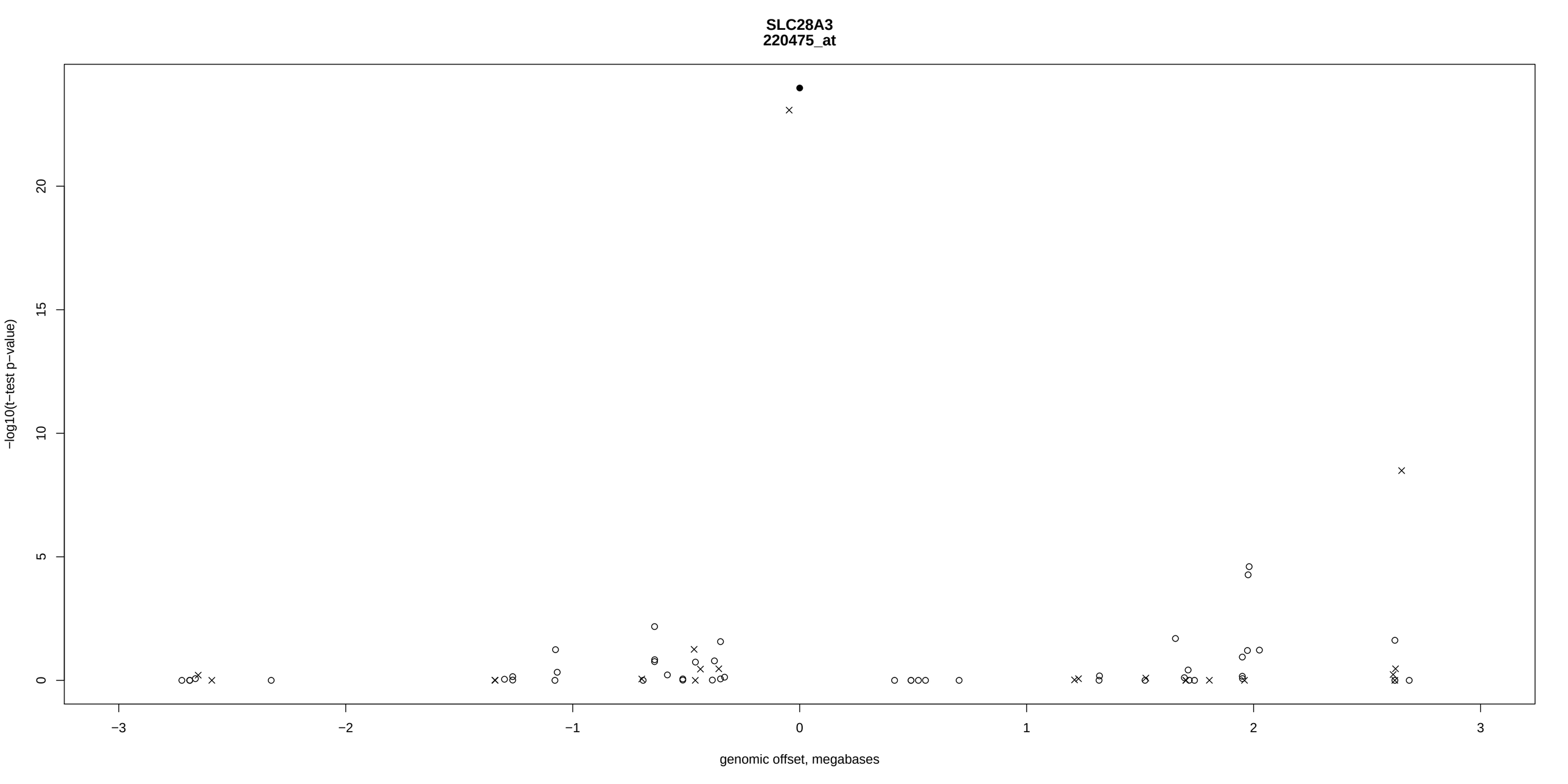
2
3
genomic offset, megabases


SLC28A3
220475_at
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0
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10
15
20
−3
−2
−1
0
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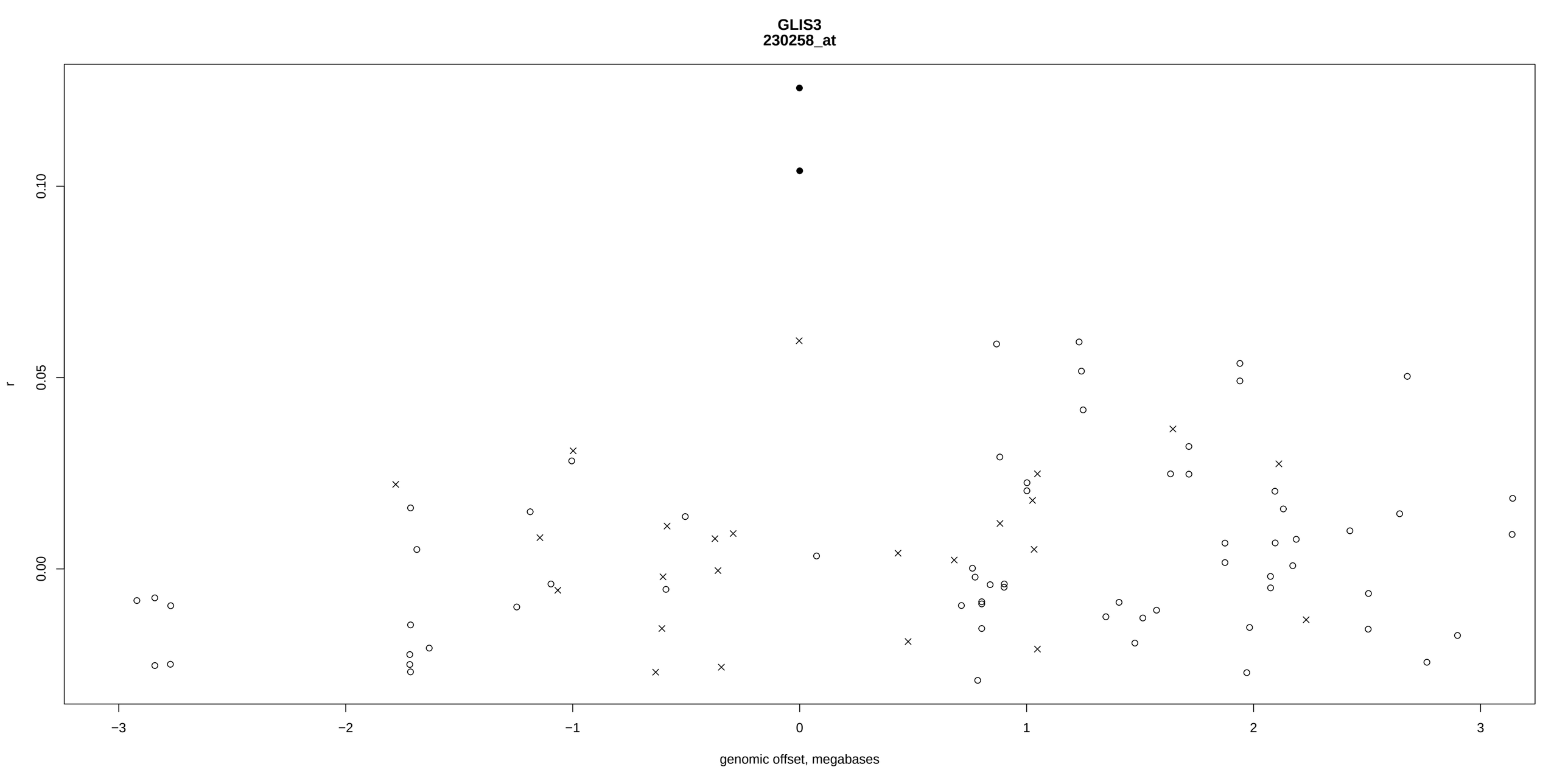
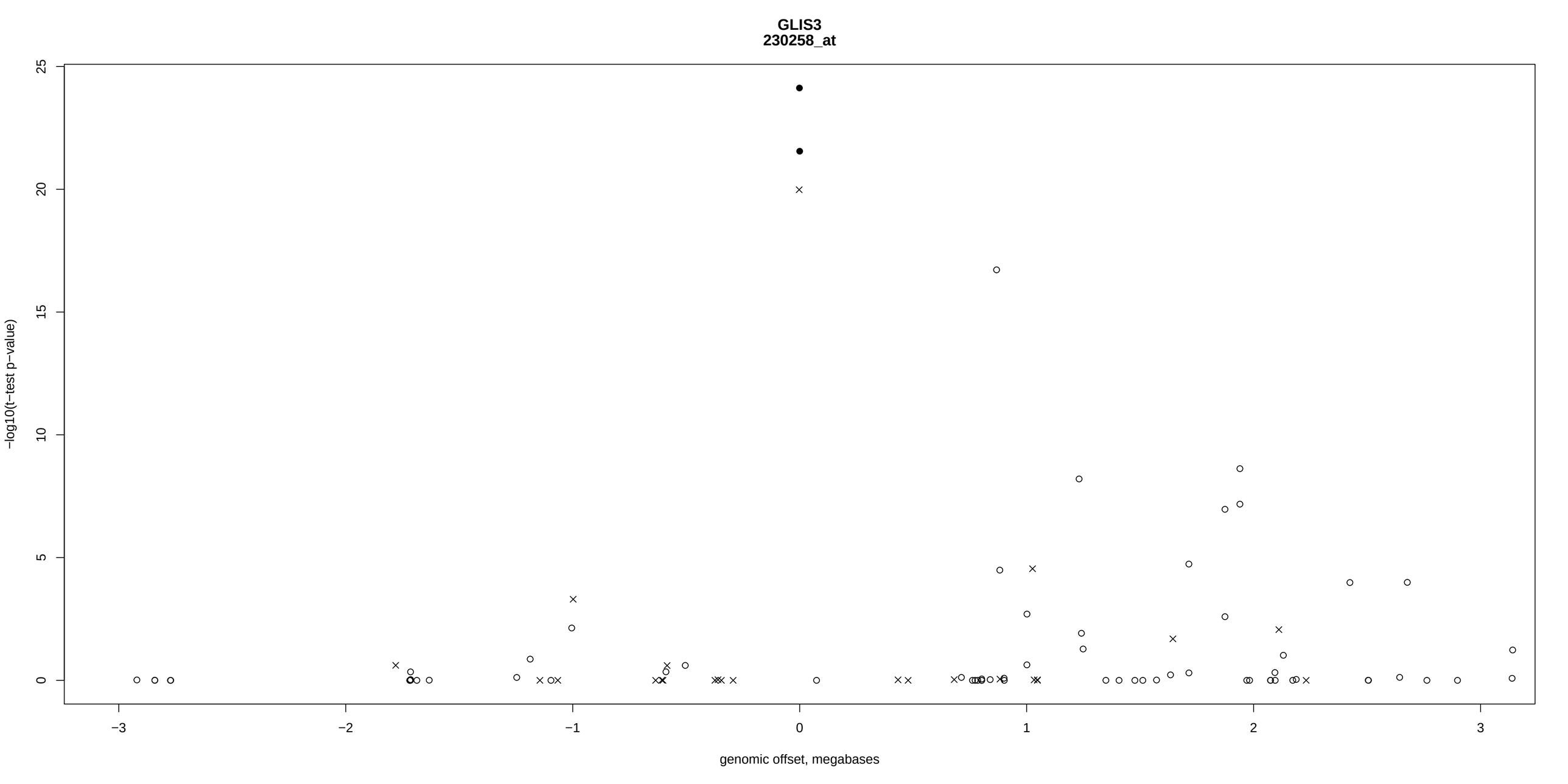
2
3
genomic offset, megabases


GLIS3
230258_at
r
0.00
0.05
0.10
−3
−2
−1
0
1
2
3
genomic offset, megabases


GLIS3
230258_at
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0
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10
15
20
25
−3
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−1
0
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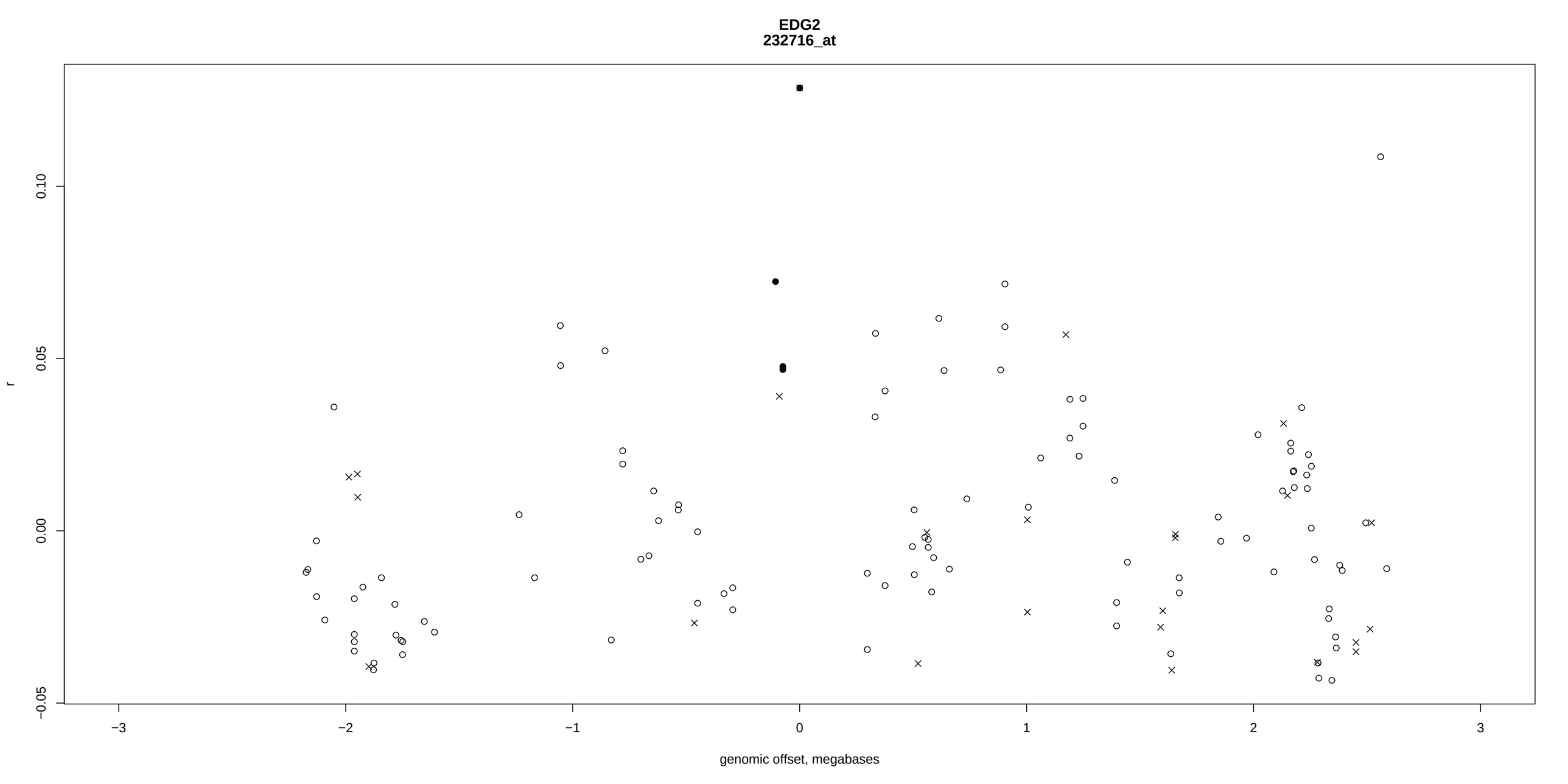
2
3
genomic offset, megabases


EDG2
232716_at
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0.05
0.00
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−3
−2
−1
0
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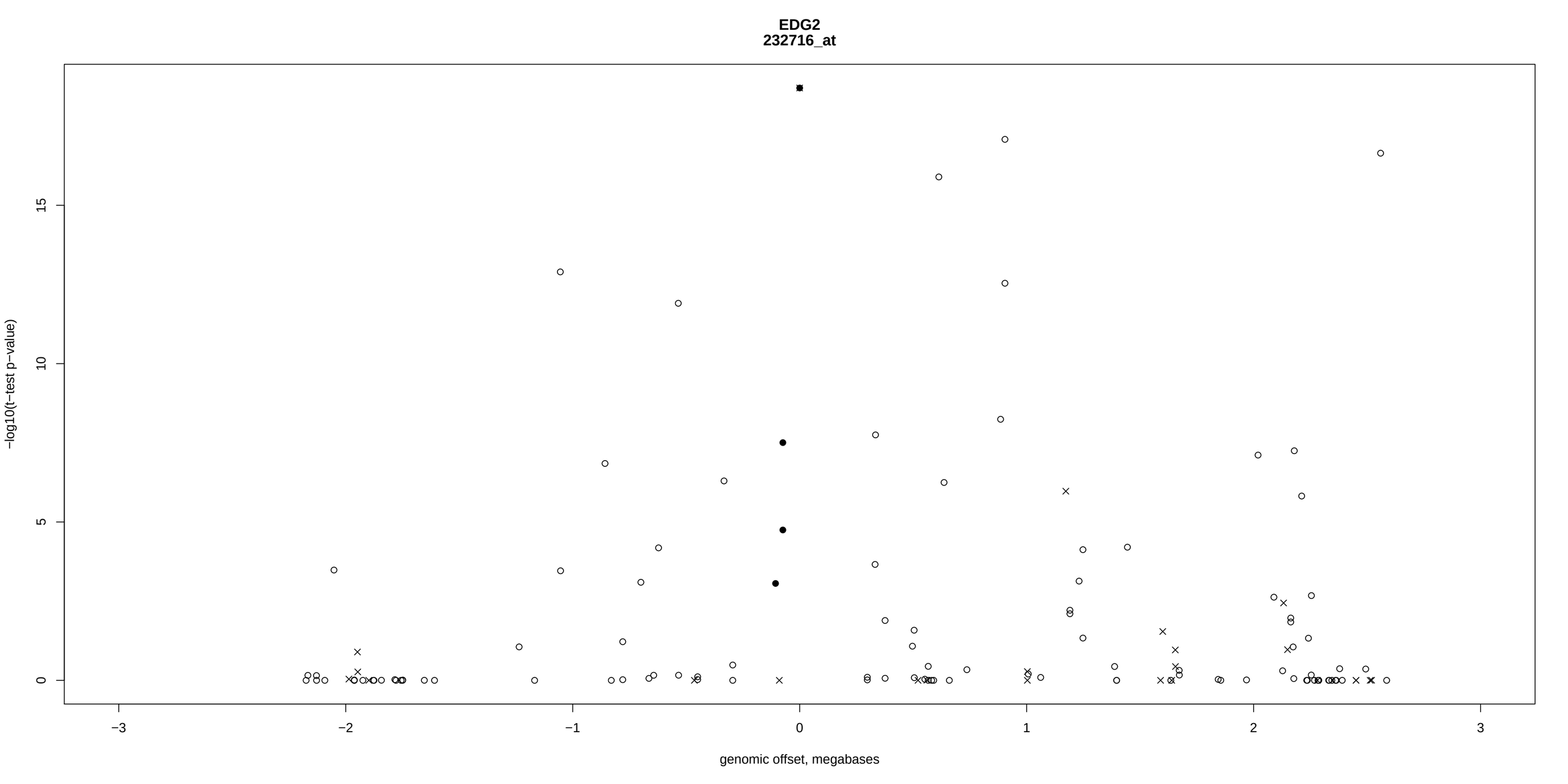
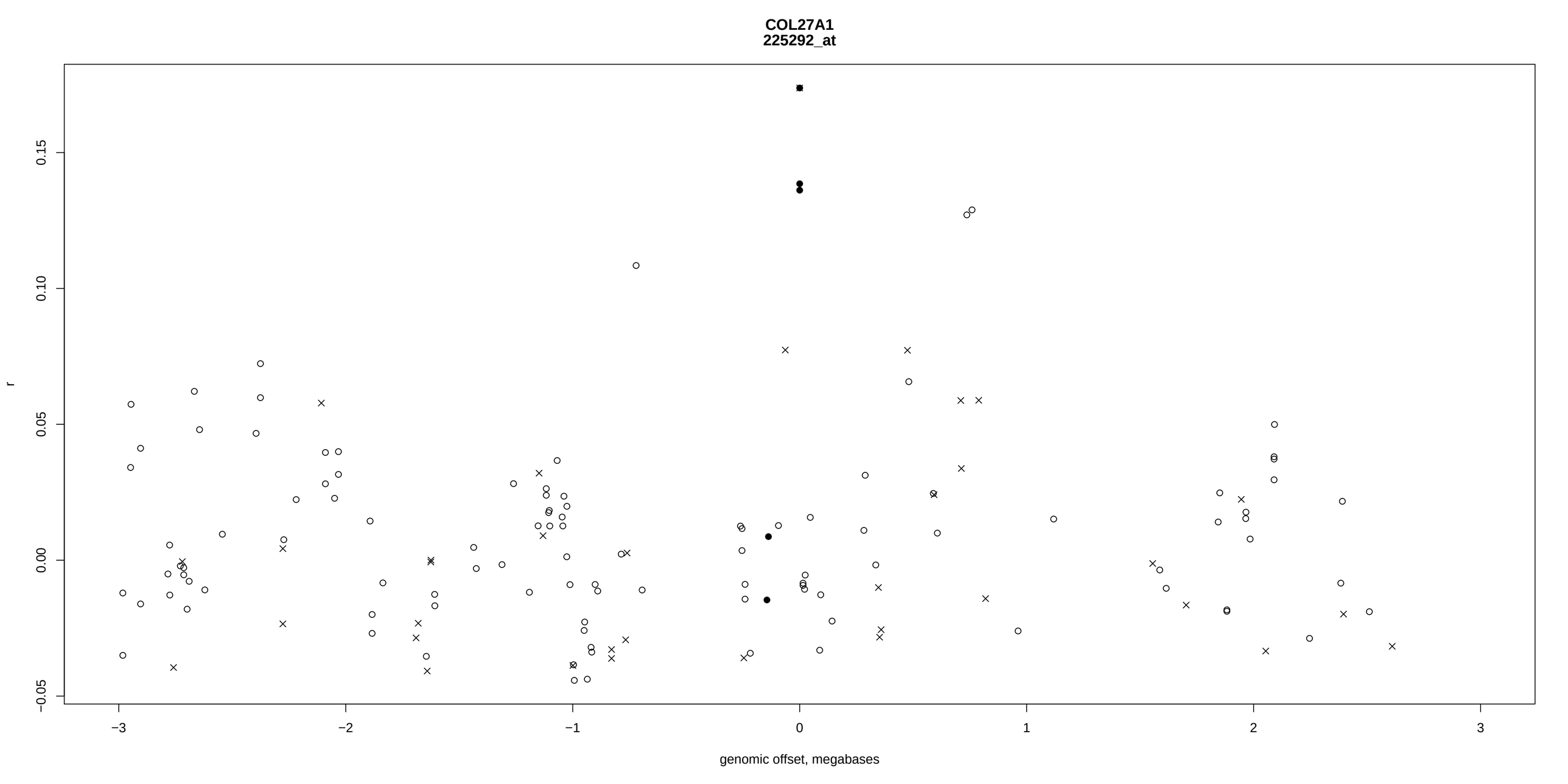
2
3
genomic offset, megabases


EDG2
232716_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


COL27A1
225292_at
r
0.15
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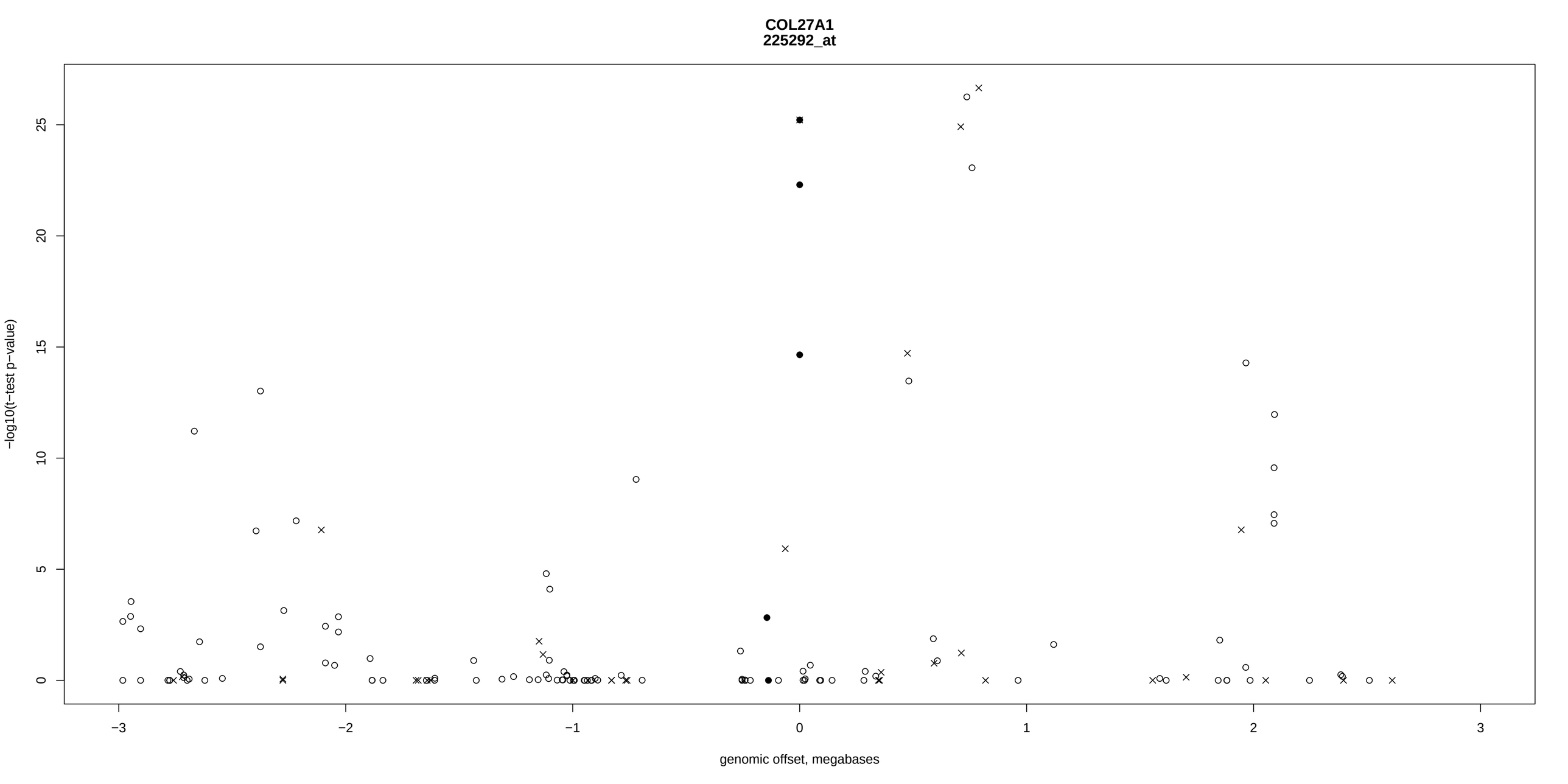
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


COL27A1
225292_at
−log10(t−test p−value)
0
5
10
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20
25
−3
−2
−1
0
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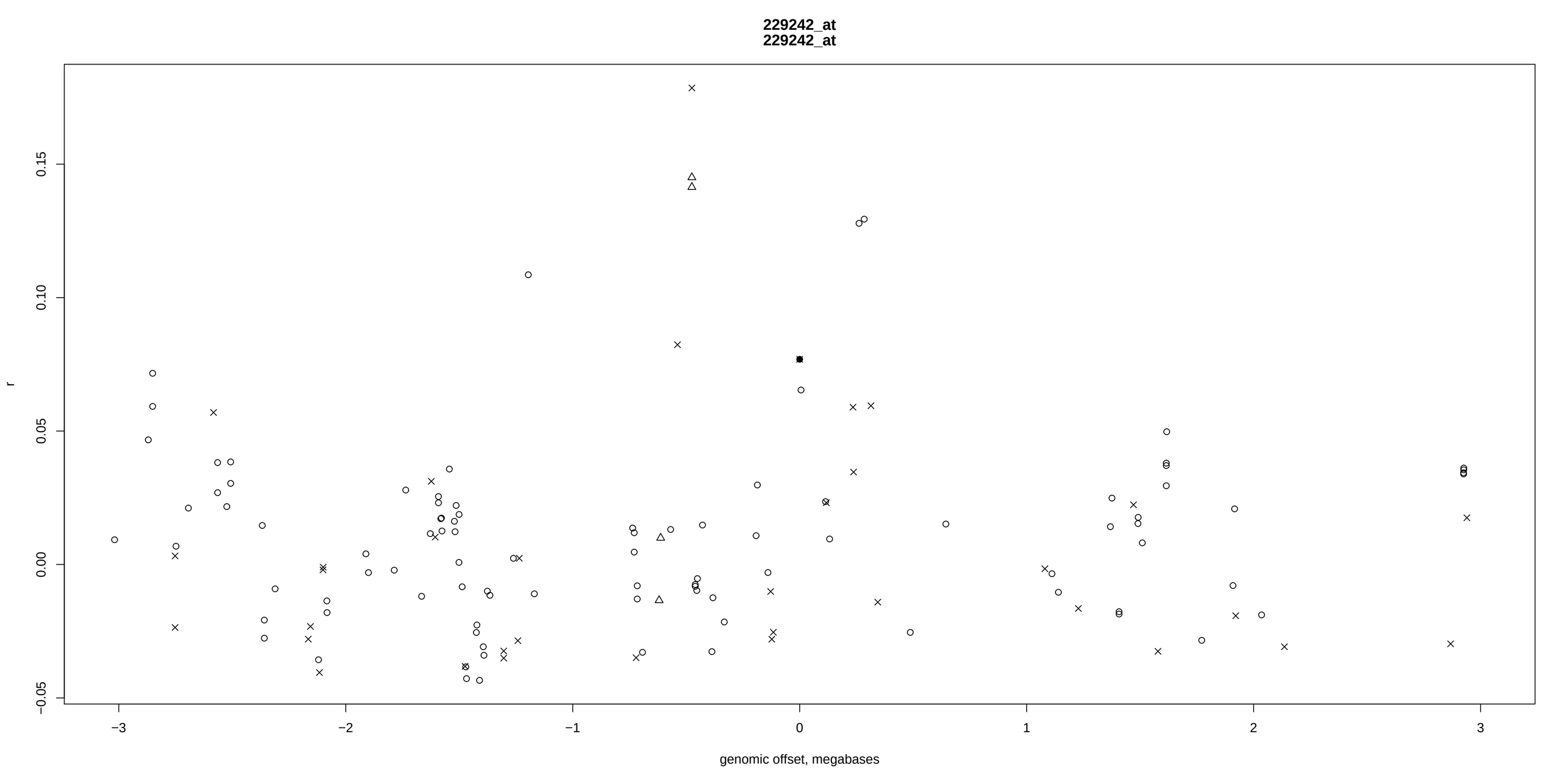
2
3
genomic offset, megabases


229242_at
229242_at
r
0.15
0.10
0.05
0.00
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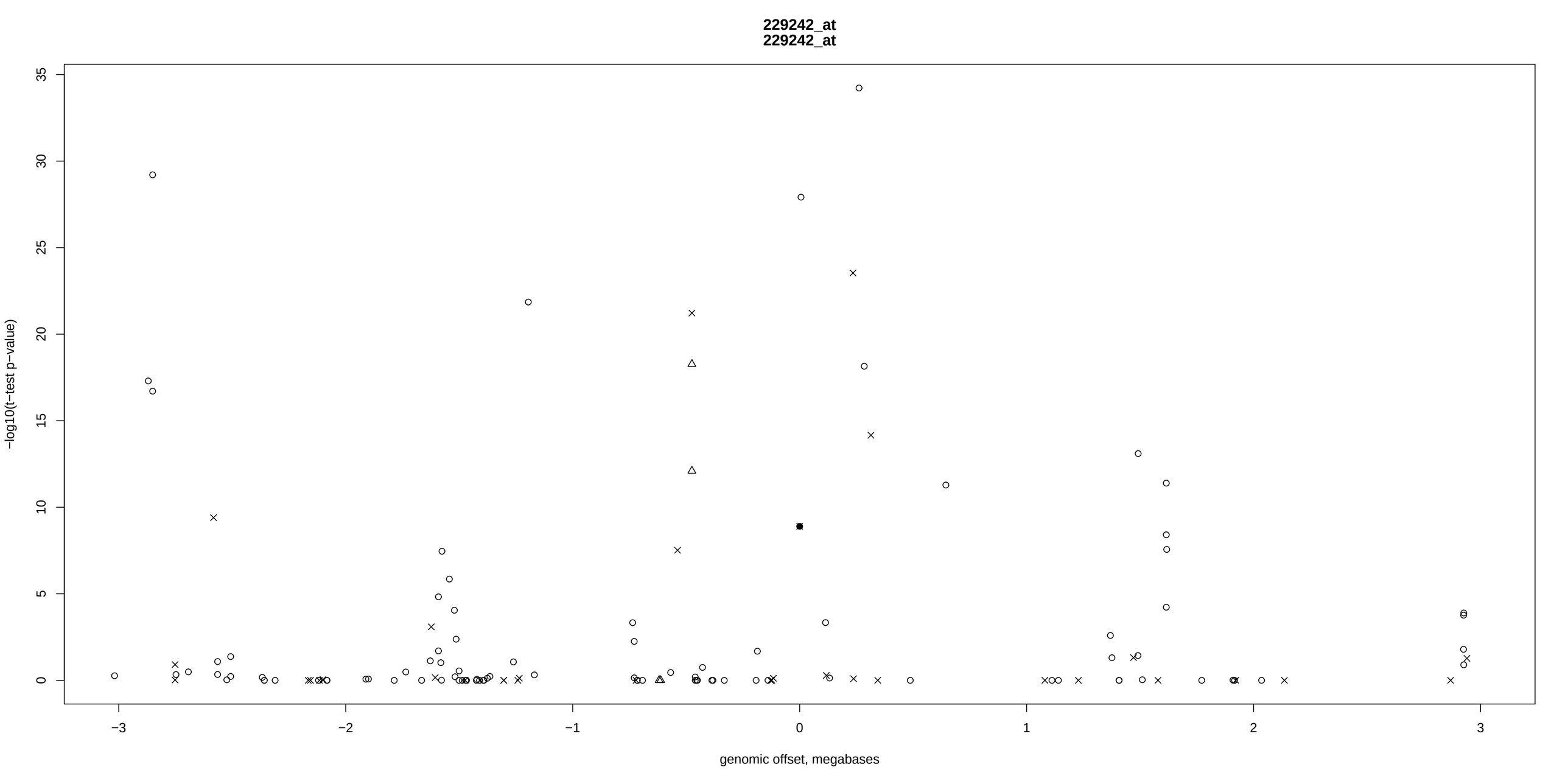
−2
−1
0
1
2
3
genomic offset, megabases


229242_at
229242_at
−log10(t−test p−value)
0
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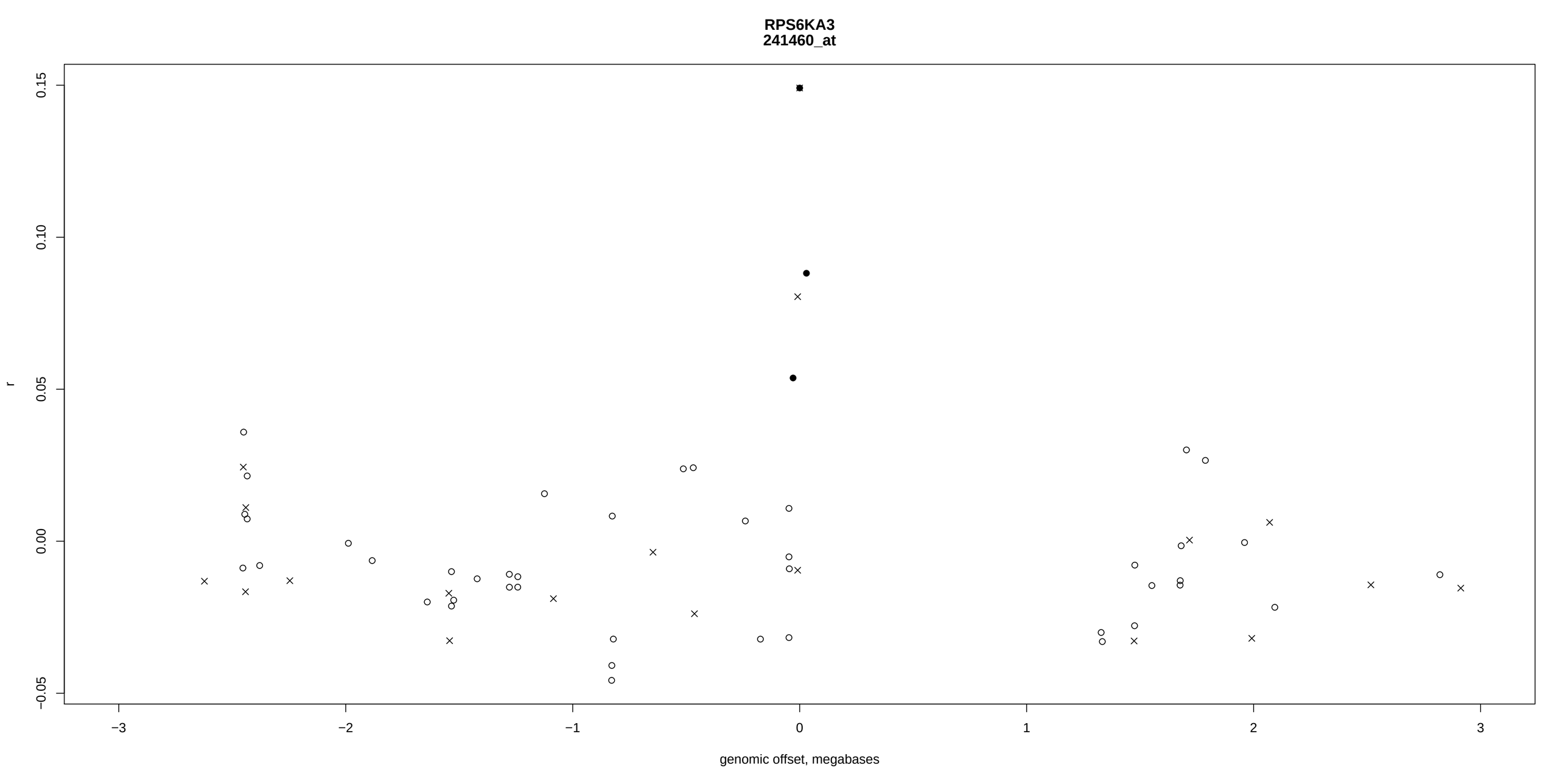
10
15
20
25
30
35
−3
−2
−1
0
1
2
3
genomic offset, megabases


RPS6KA3
241460_at
r
0.15
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0.05
0.00
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−3
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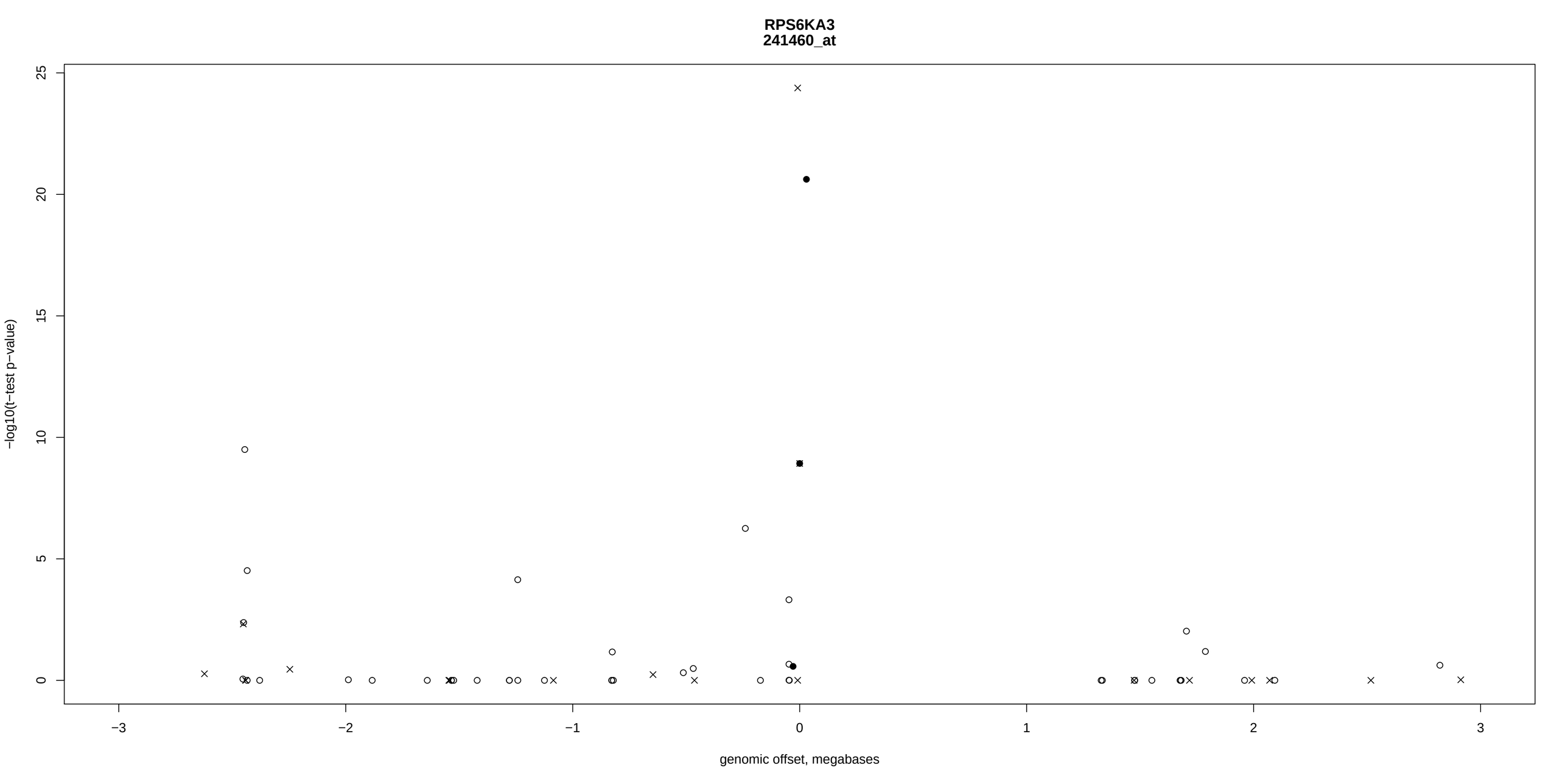
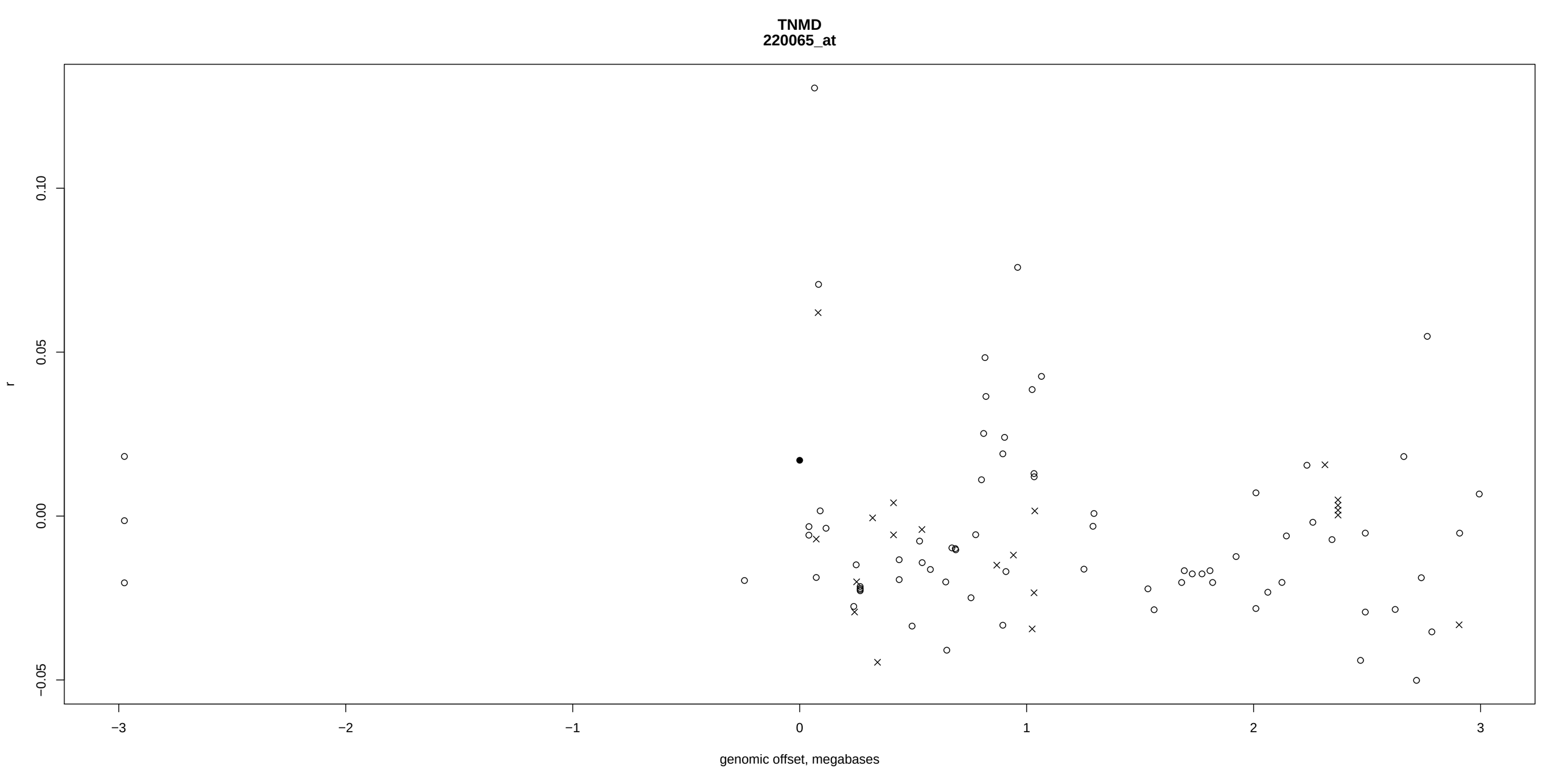
−1
0
1
2
3
genomic offset, megabases


RPS6KA3
241460_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


TNMD
220065_at
r
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0.05
0.00
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0
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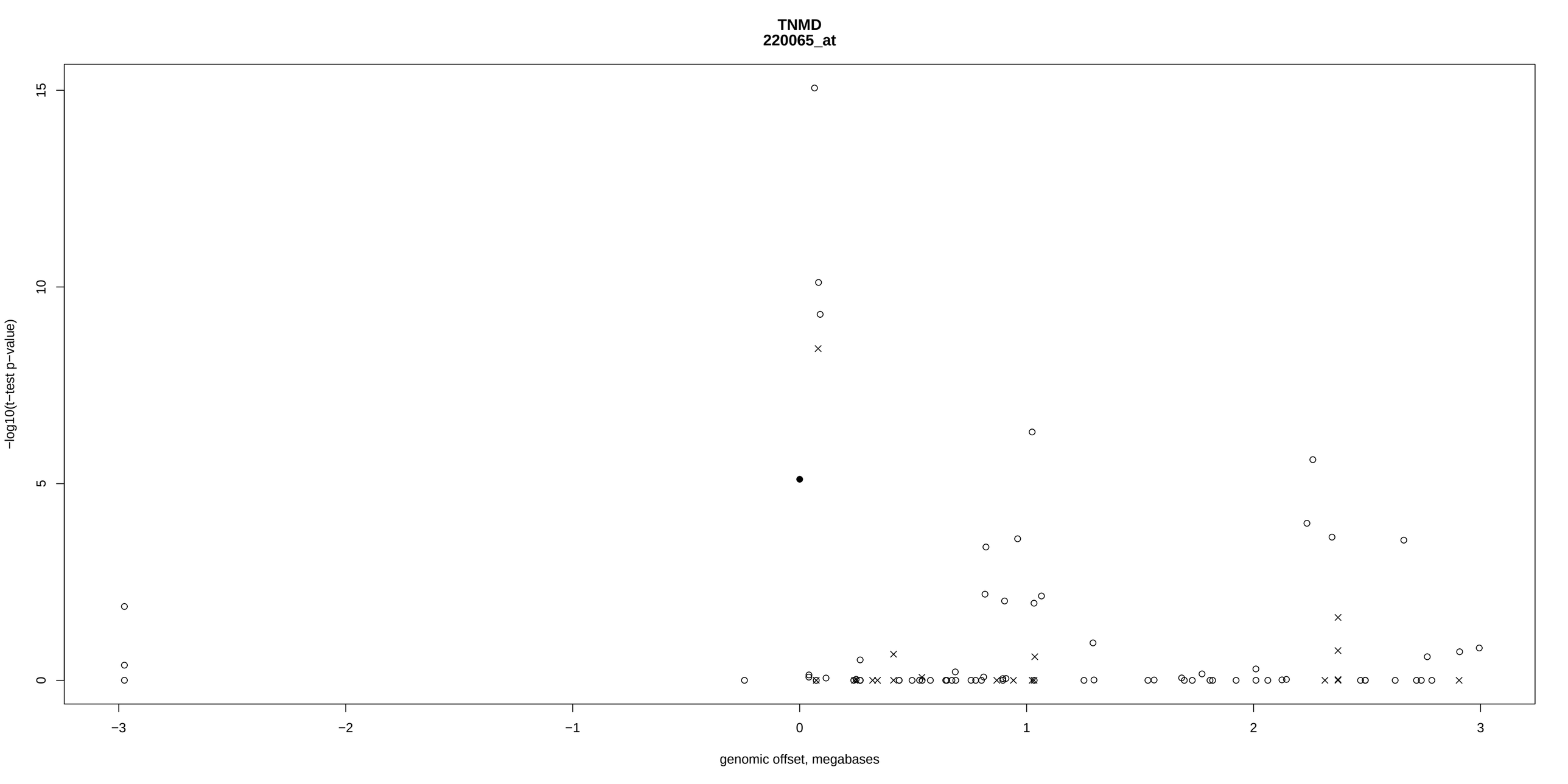
2
3
genomic offset, megabases


TNMD
220065_at
−log10(t−test p−value)
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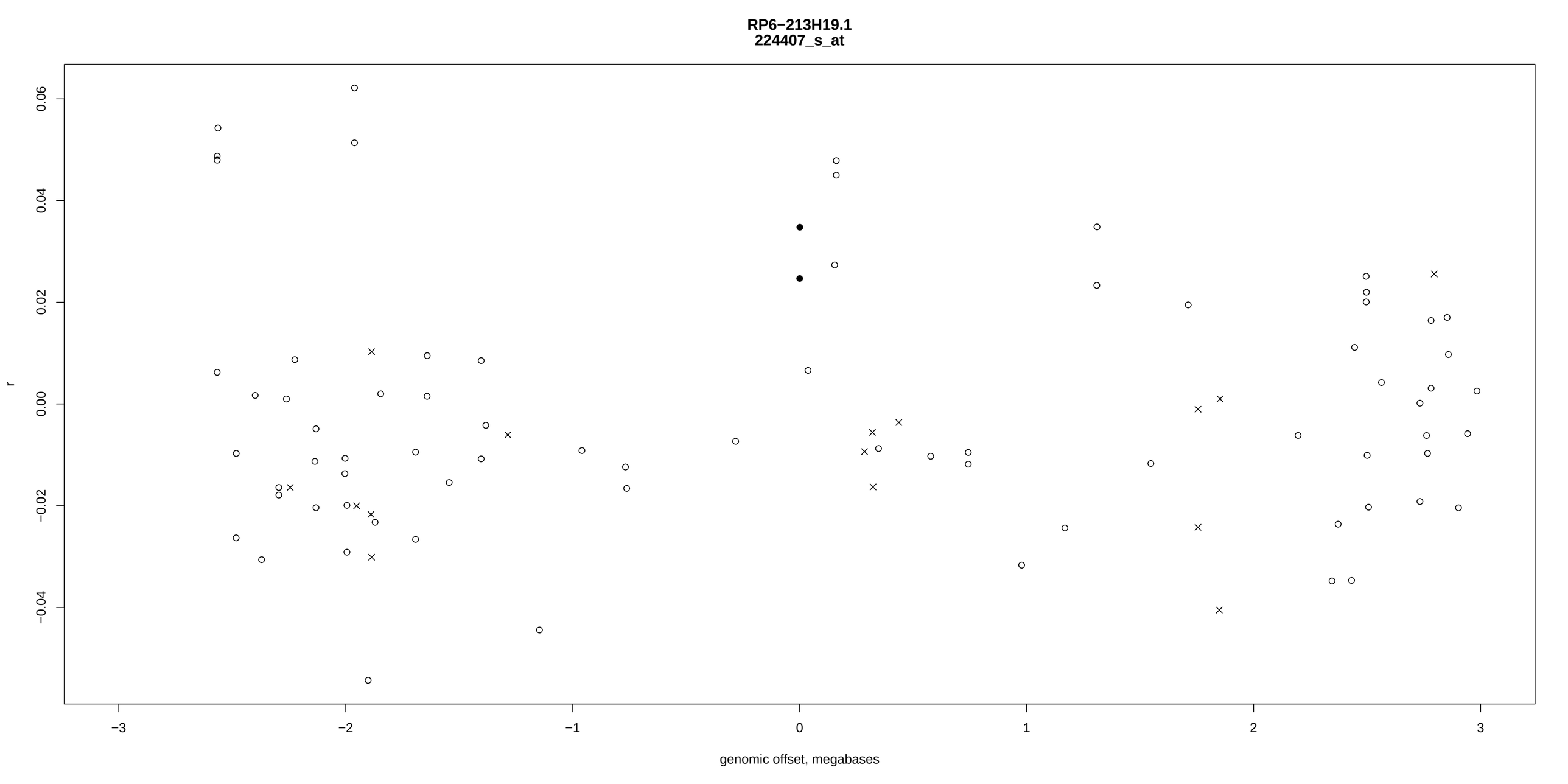
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


RP6−213H19.1
224407_s_at
r
0.06
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0.00
−0.02
−0.04
−3
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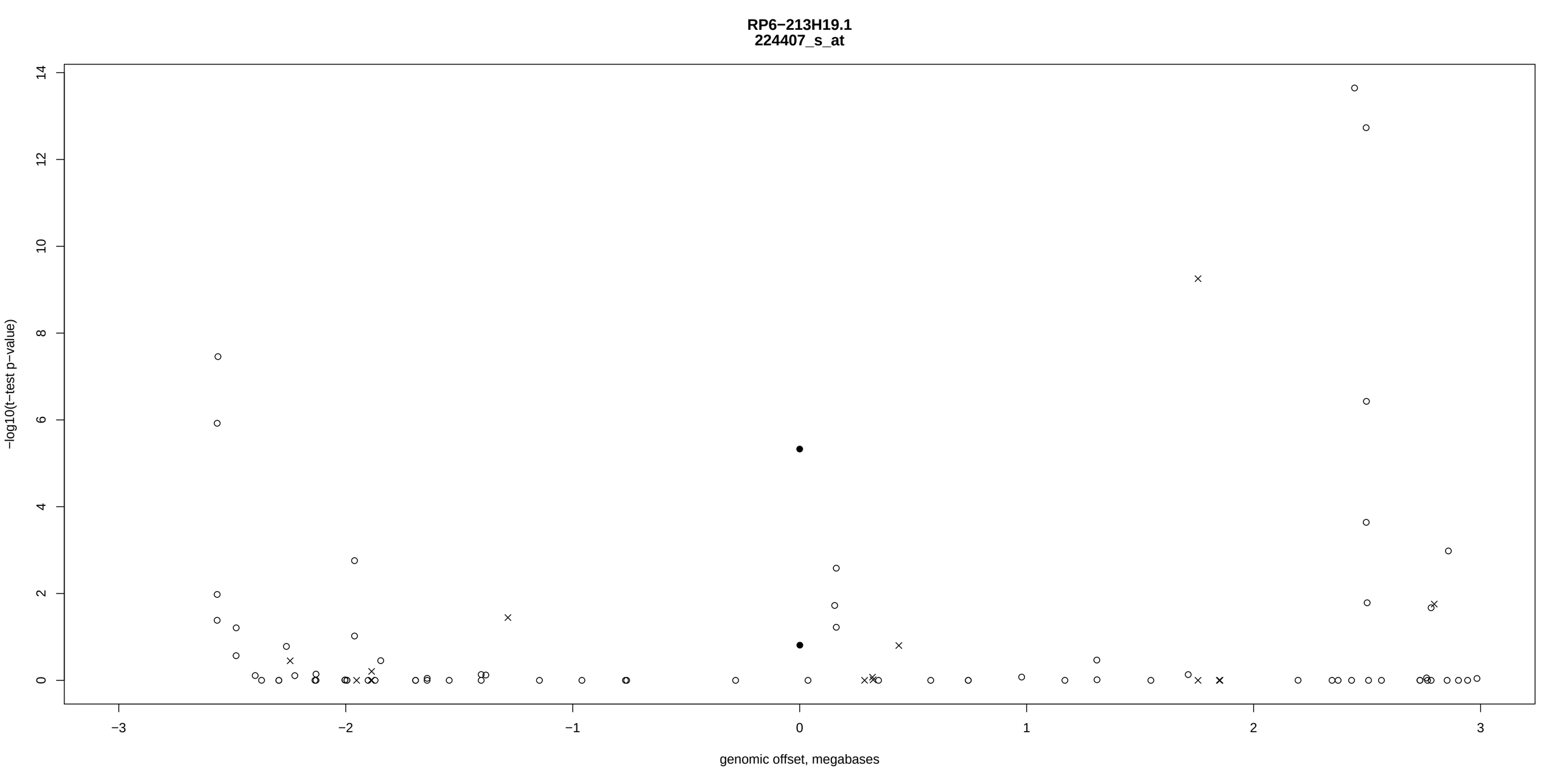
−1
0
1
2
3
genomic offset, megabases


RP6−213H19.1
224407_s_at
−log10(t−test p−value)
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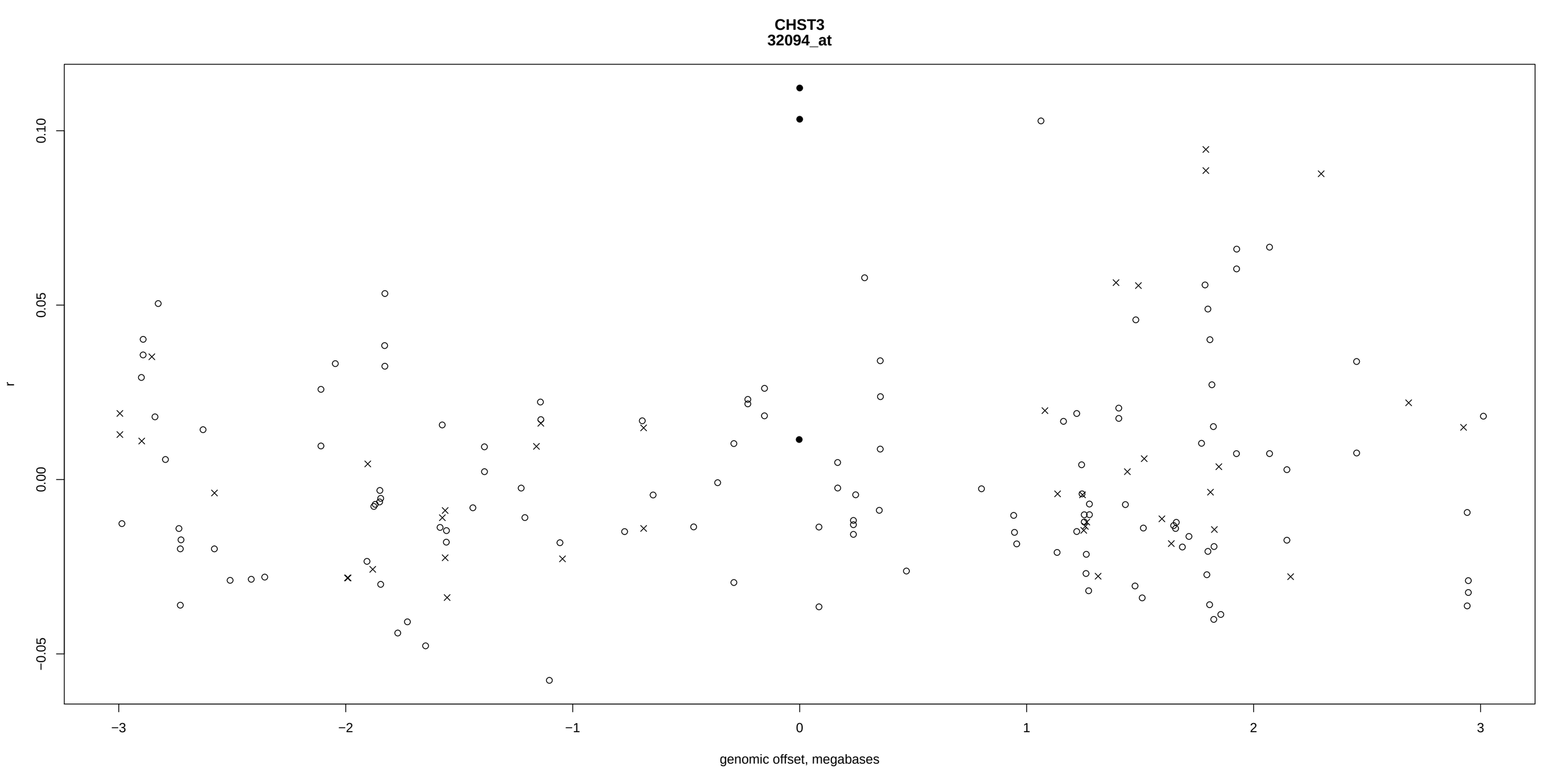
6
8
10
12
14
−3
−2
−1
0
1
2
3
genomic offset, megabases


CHST3
32094_at
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0.00
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−1
0
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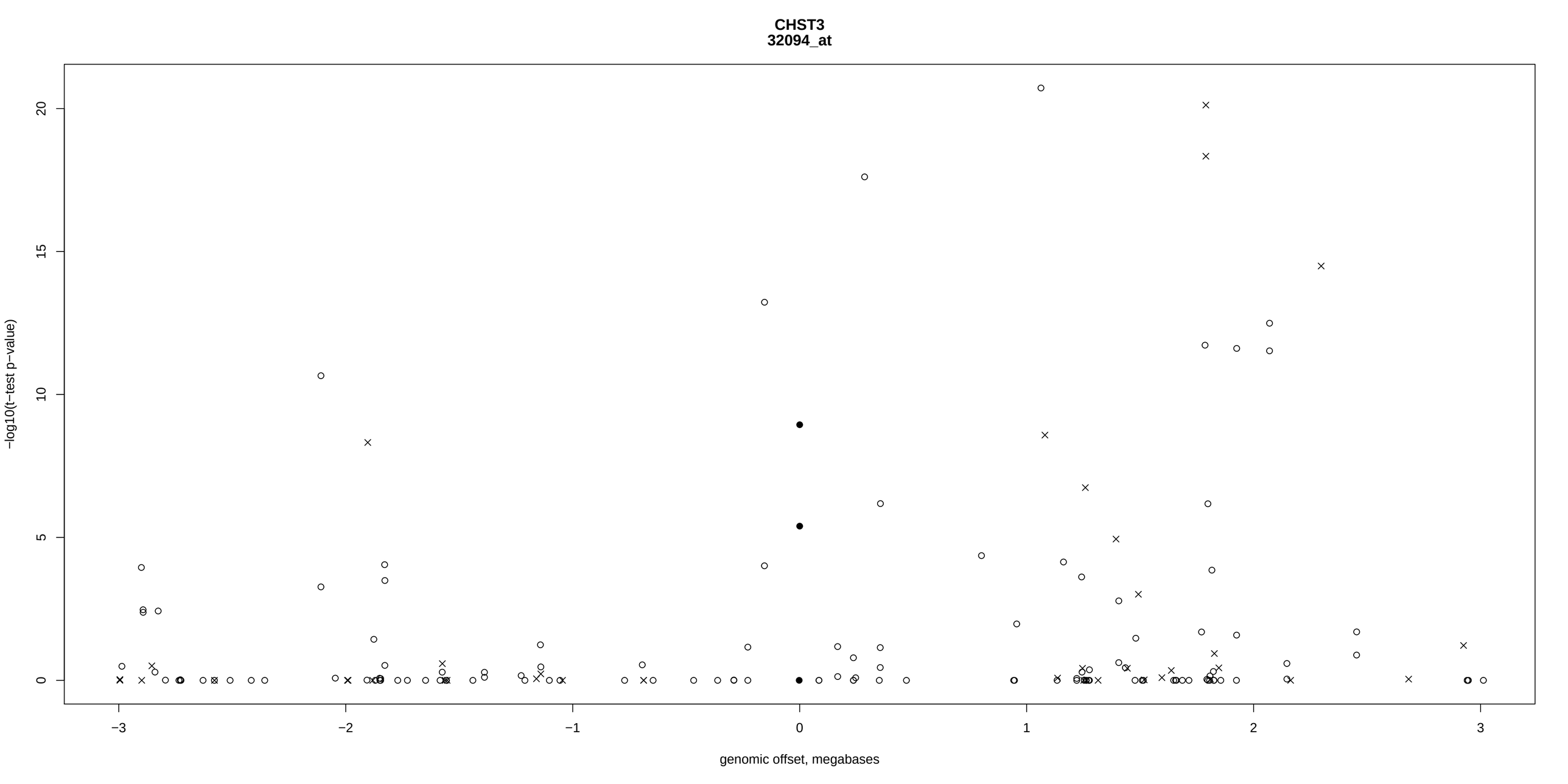
2
3
genomic offset, megabases


CHST3
32094_at
−log10(t−test p−value)
0
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15
20
−3
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−1
0
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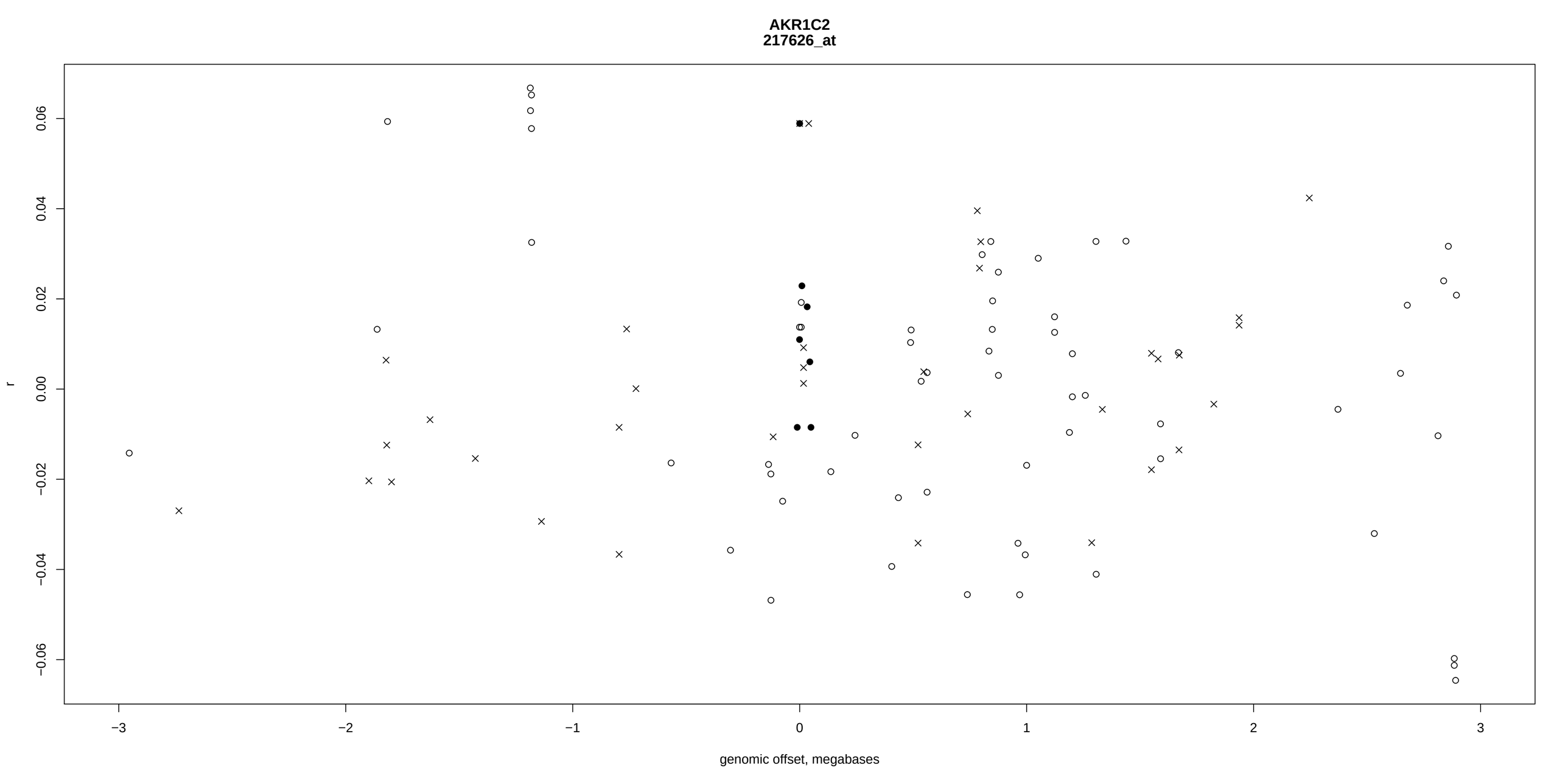
2
3
genomic offset, megabases


AKR1C2
217626_at
r
0.06
0.04
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−0.04
−0.06
−3
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−1
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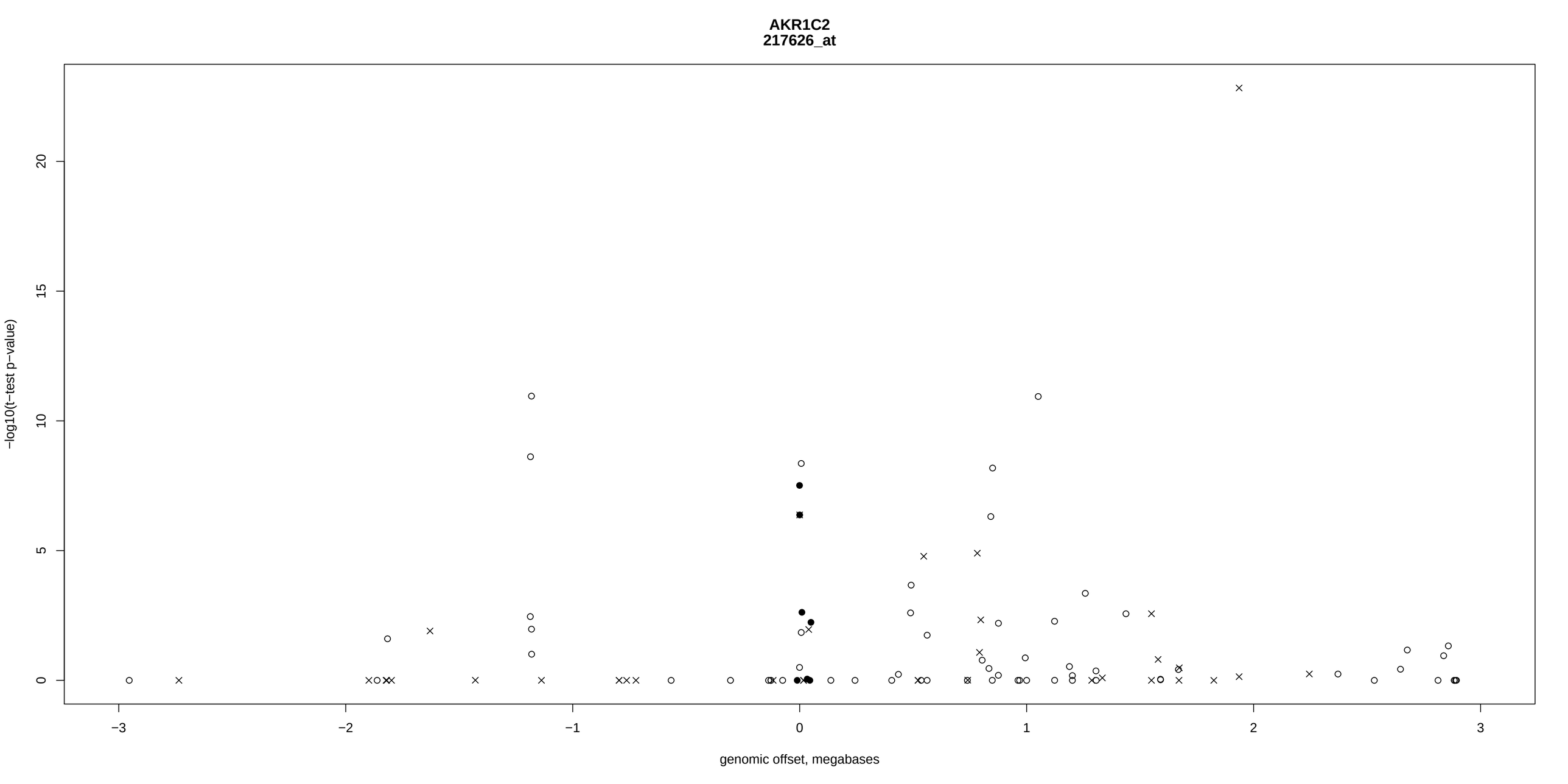
2
3
genomic offset, megabases


AKR1C2
217626_at
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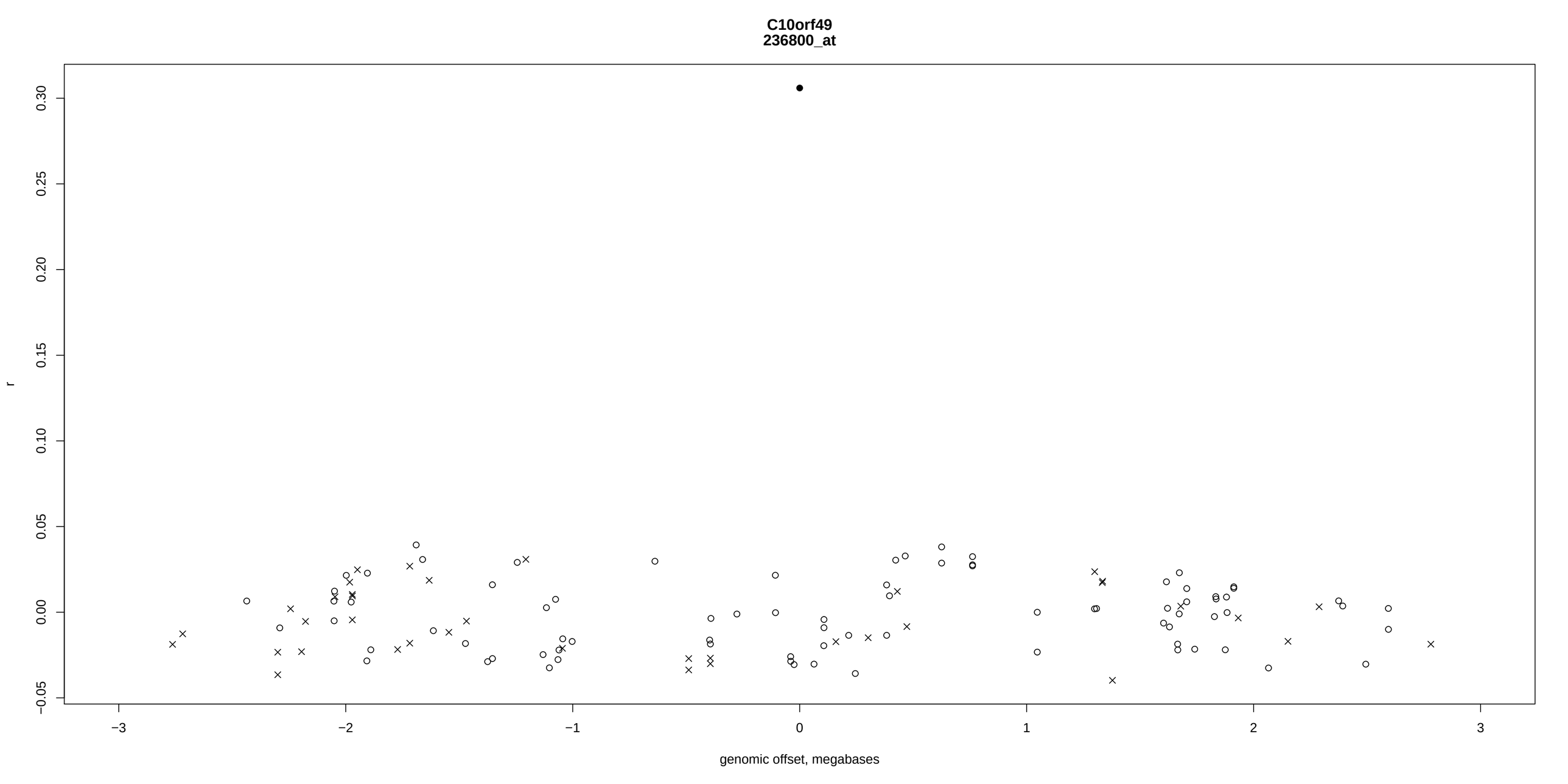
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


C10orf49
236800_at
r
0.30
0.25
0.20
0.15
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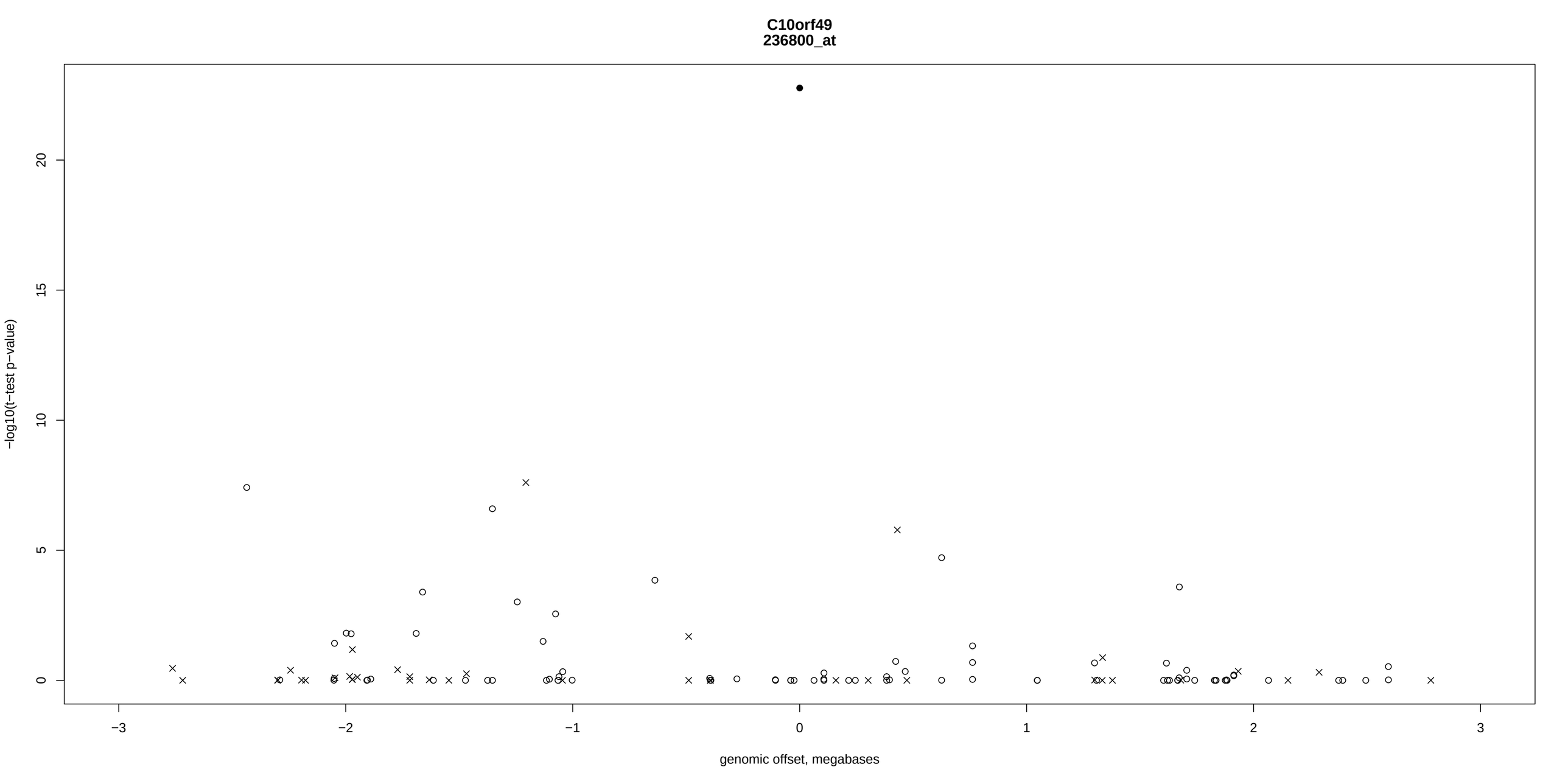
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


C10orf49
236800_at
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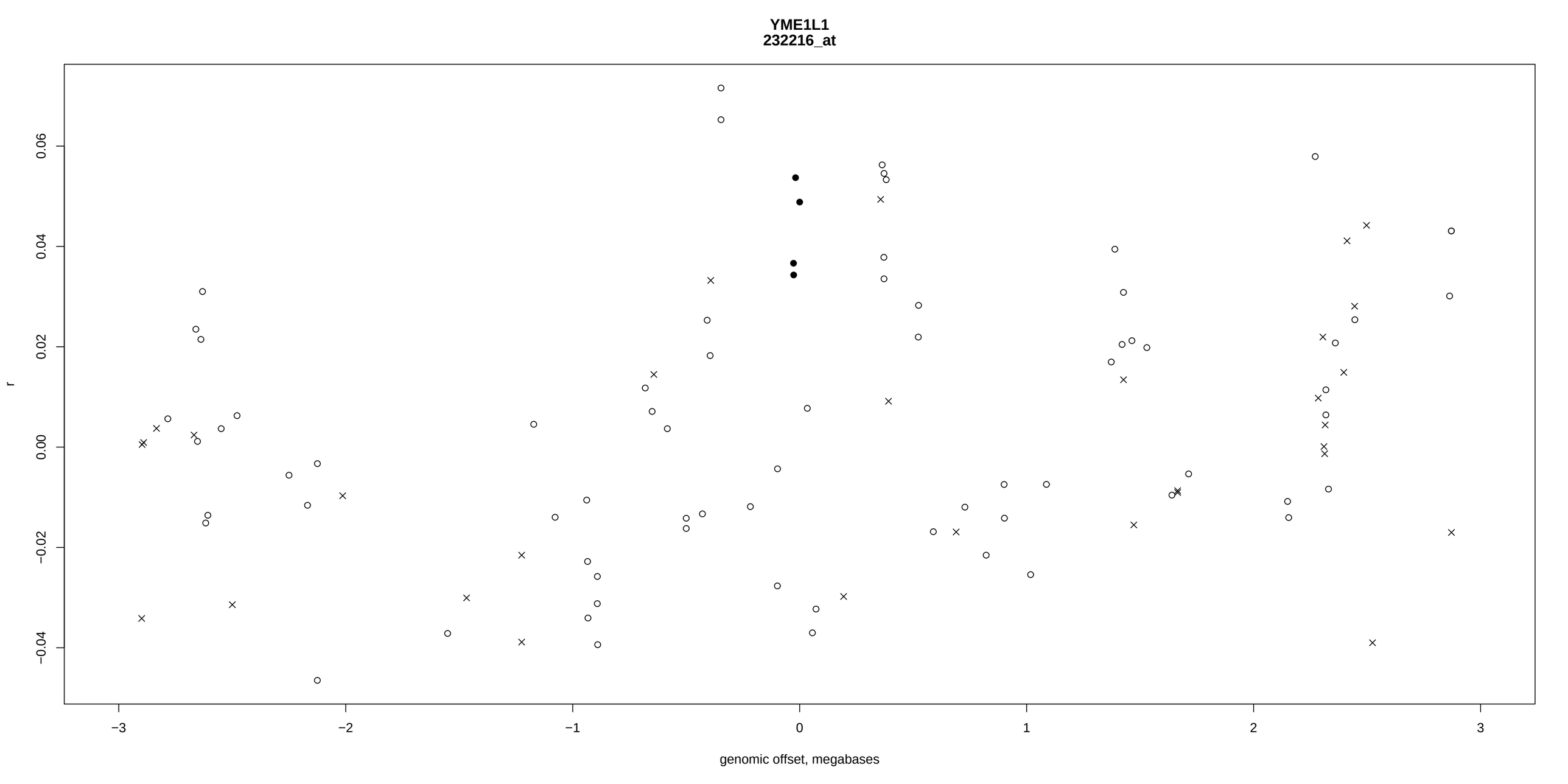
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


YME1L1
232216_at
r
0.06
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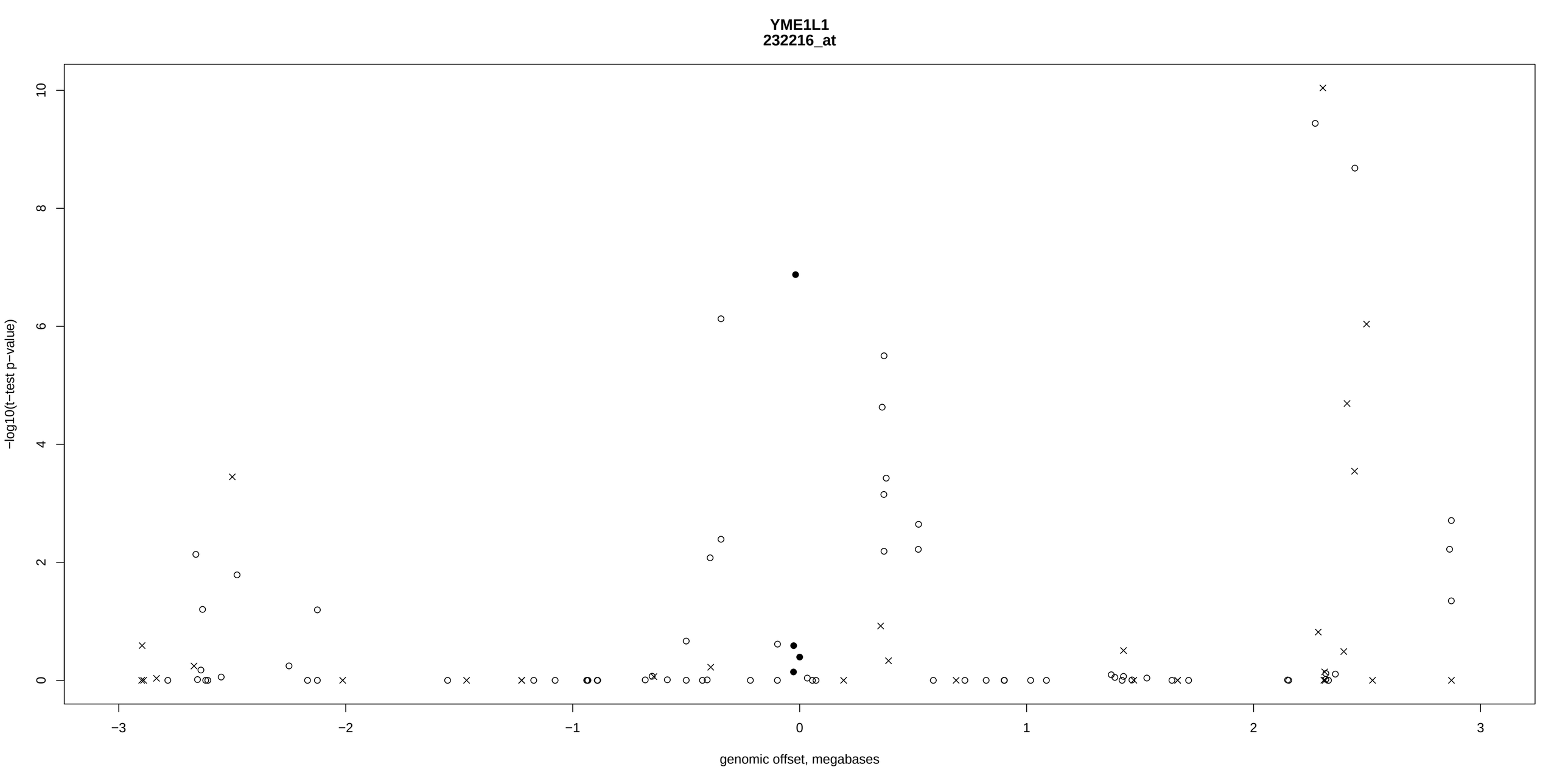
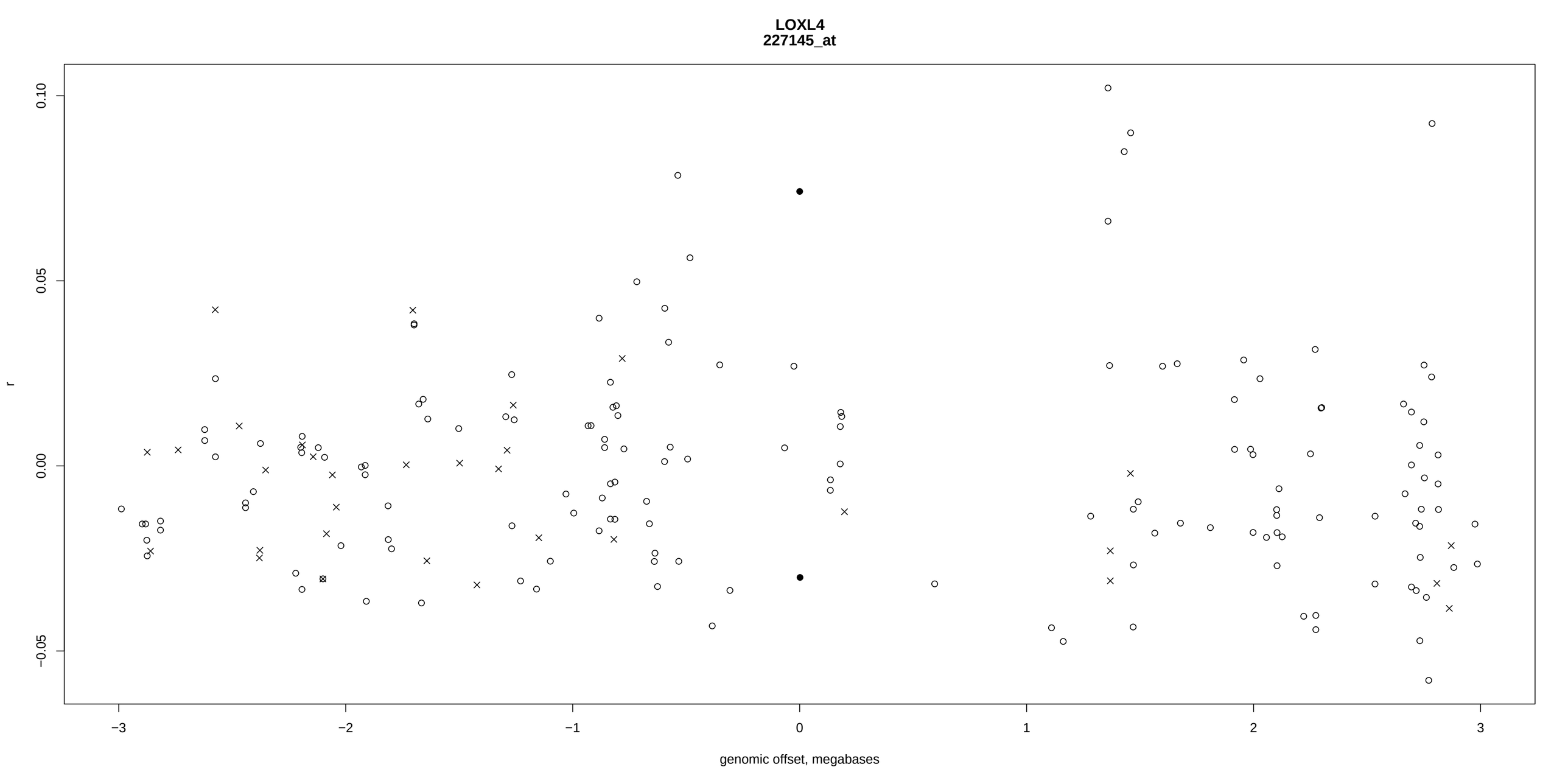
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases


YME1L1
232216_at
−log10(t−test p−value)
0
2
4
6
8
10
−3
−2
−1
0
1
2
3
genomic offset, megabases


LOXL4
227145_at
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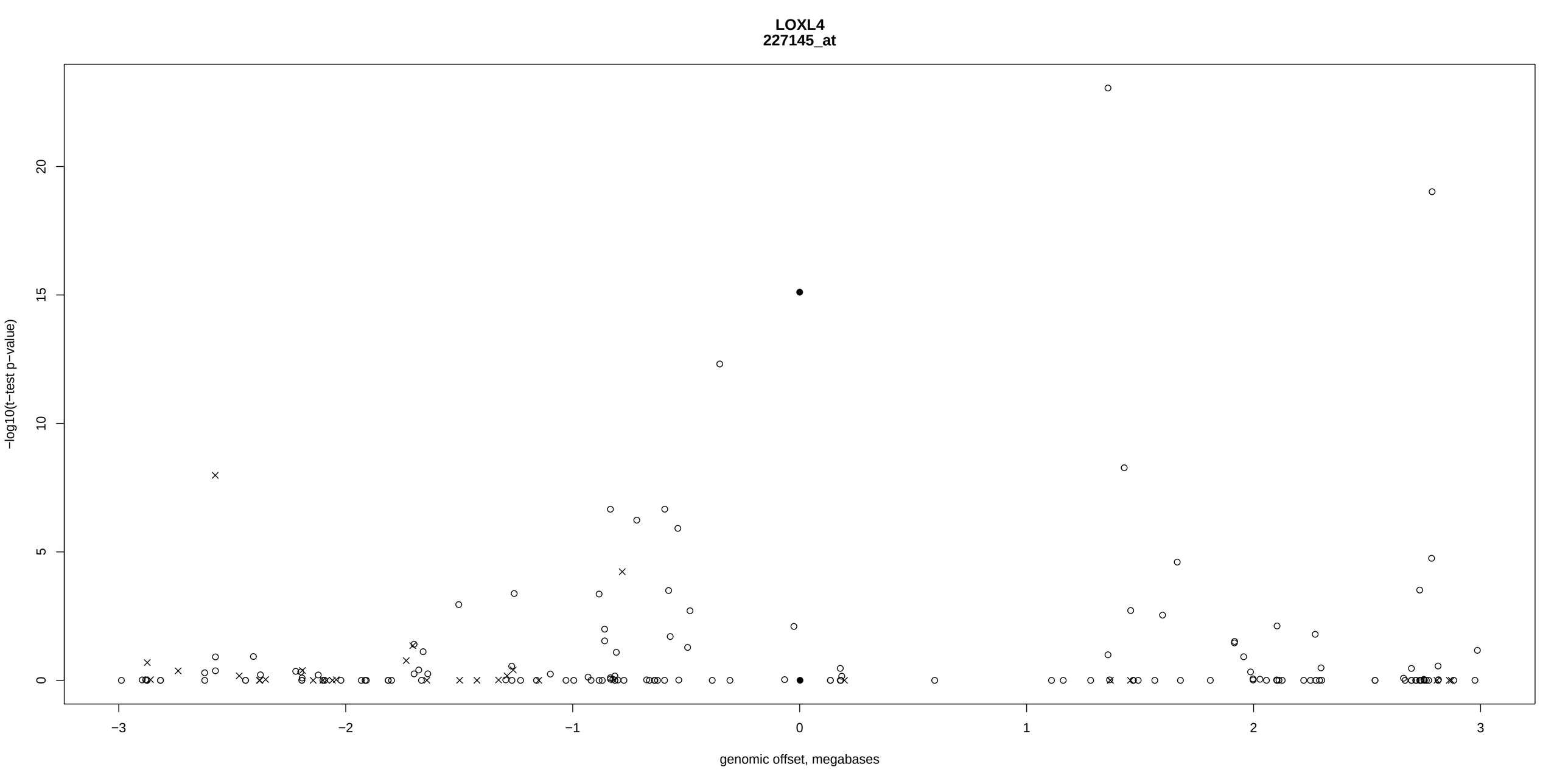
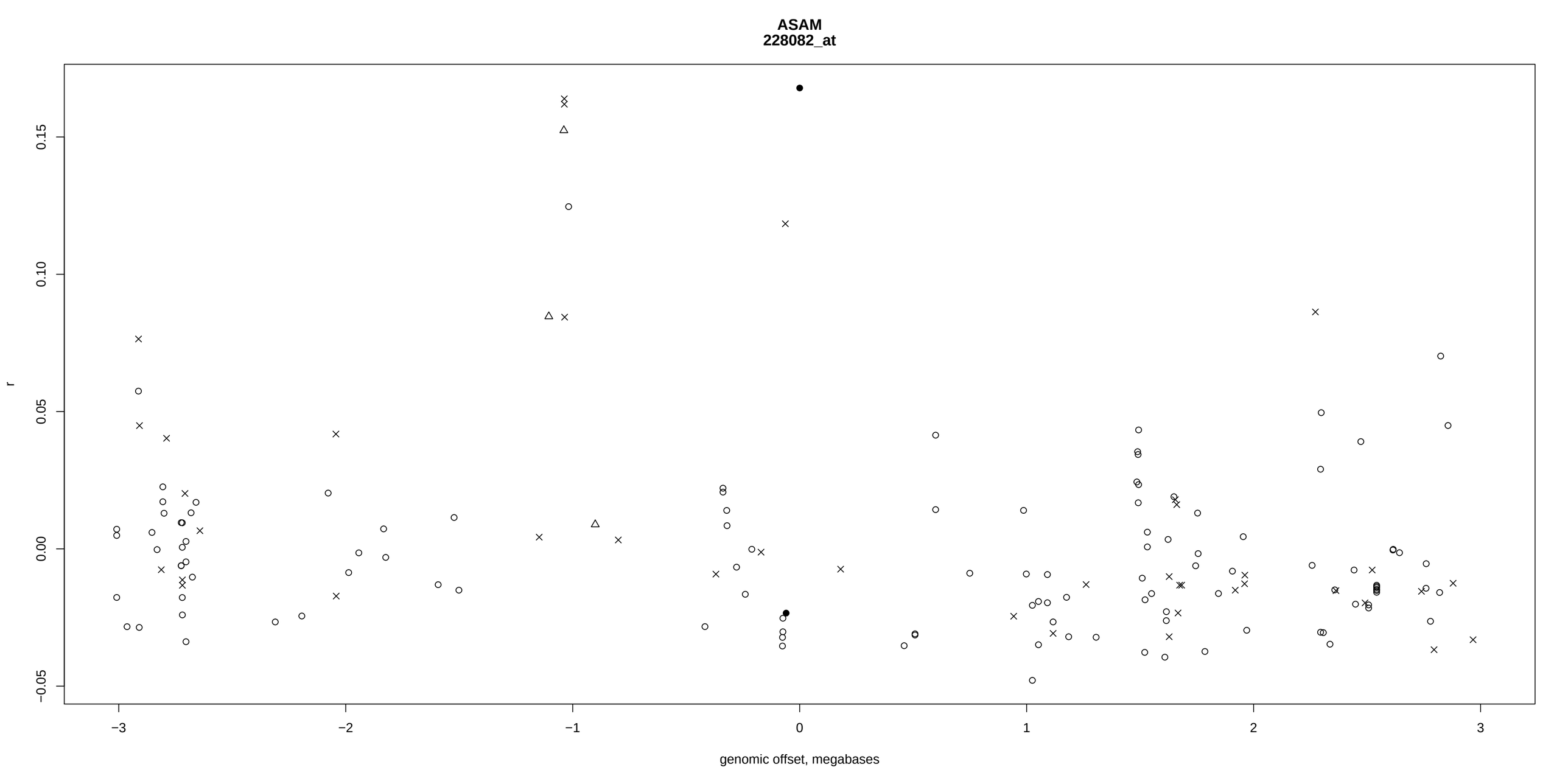
0.00
−0.05
r
−3
−2
−1
0
1
2
3
genomic offset, megabases


LOXL4
227145_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


ASAM
228082_at
r
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0.05
0.00
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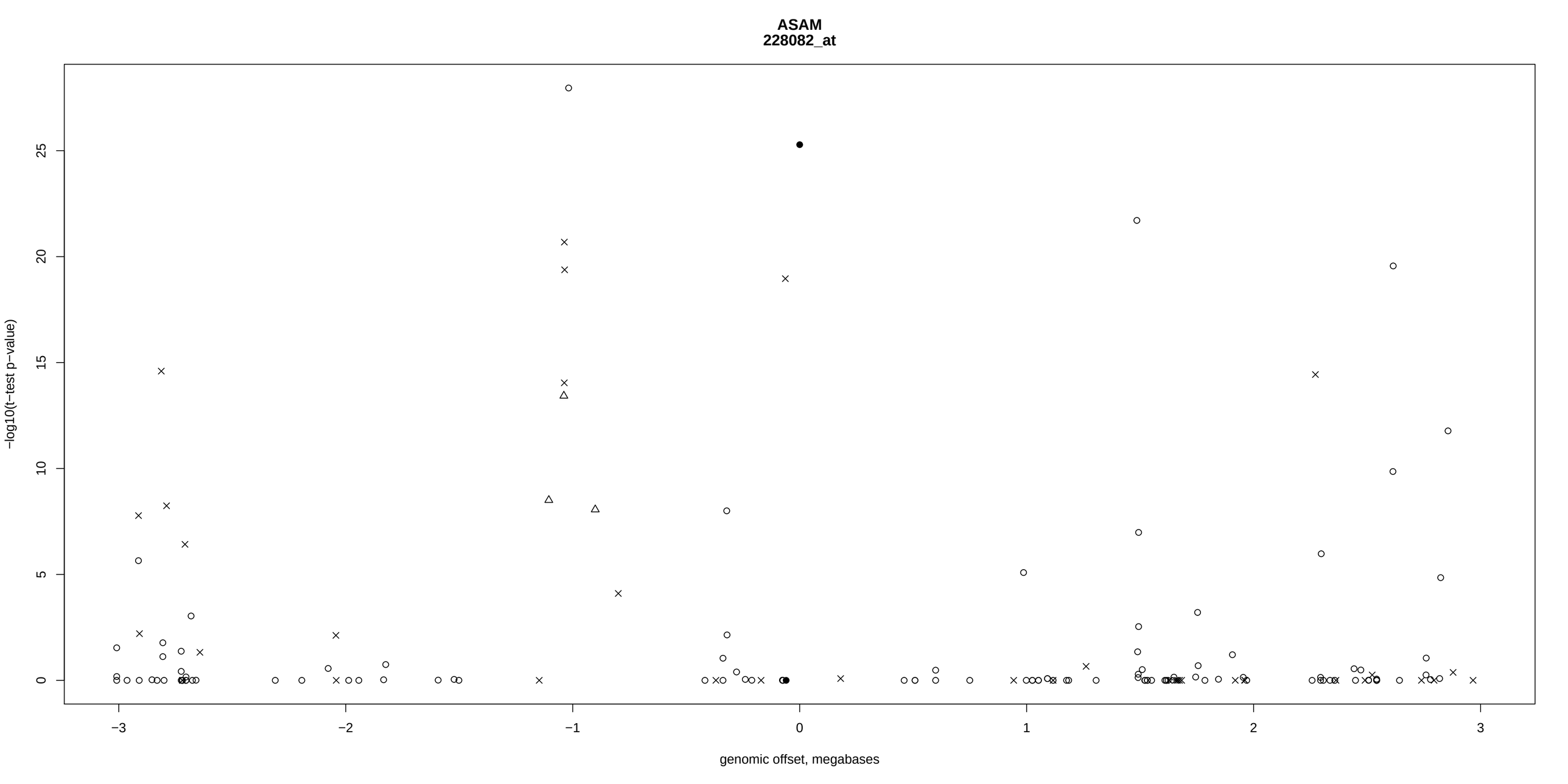
−2
−1
0
1
2
3
genomic offset, megabases


ASAM
228082_at
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0
5
10
15
20
25
−3
−2
−1
0
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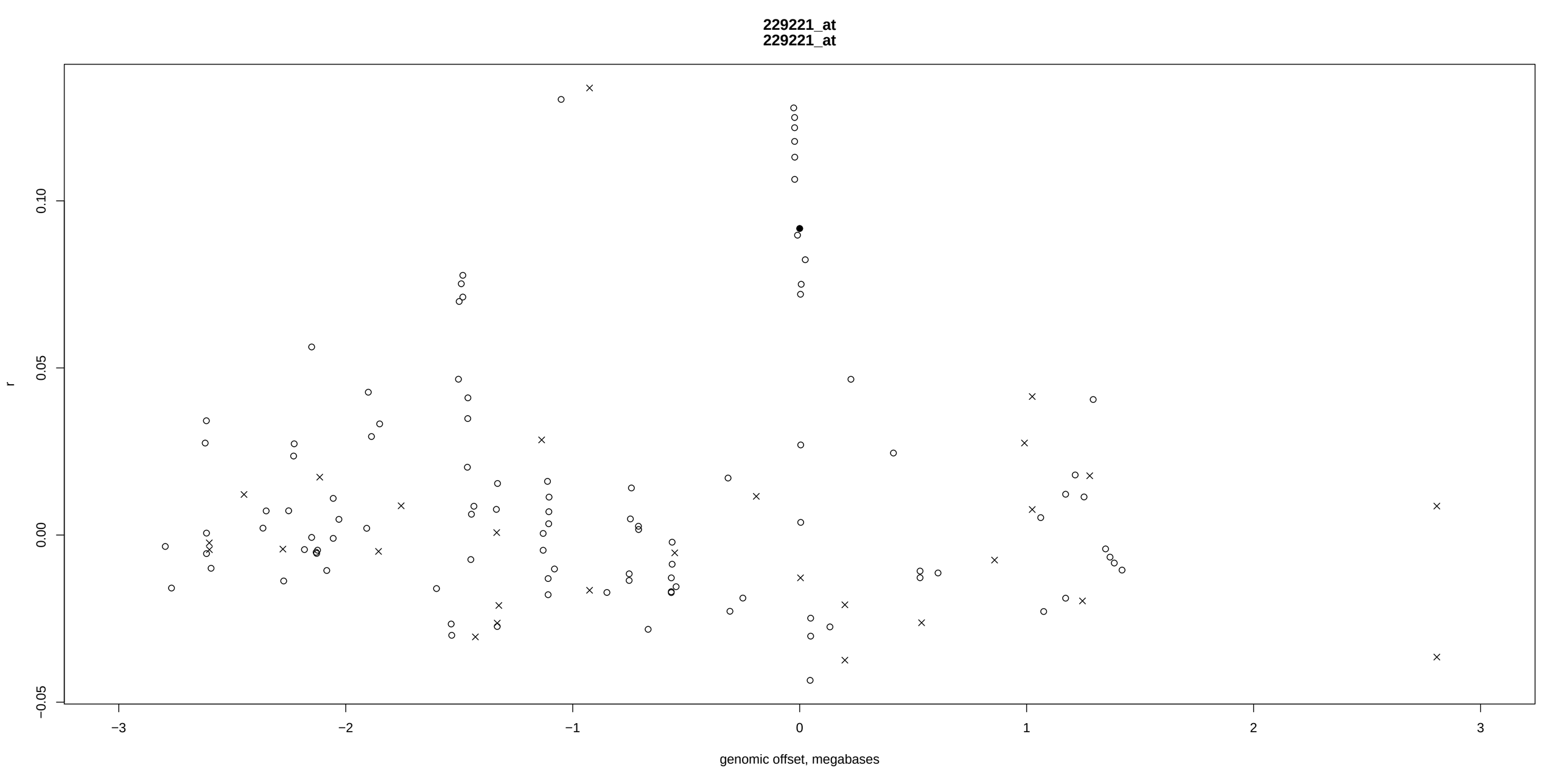
2
3
genomic offset, megabases


229221_at
229221_at
r
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0.05
0.00
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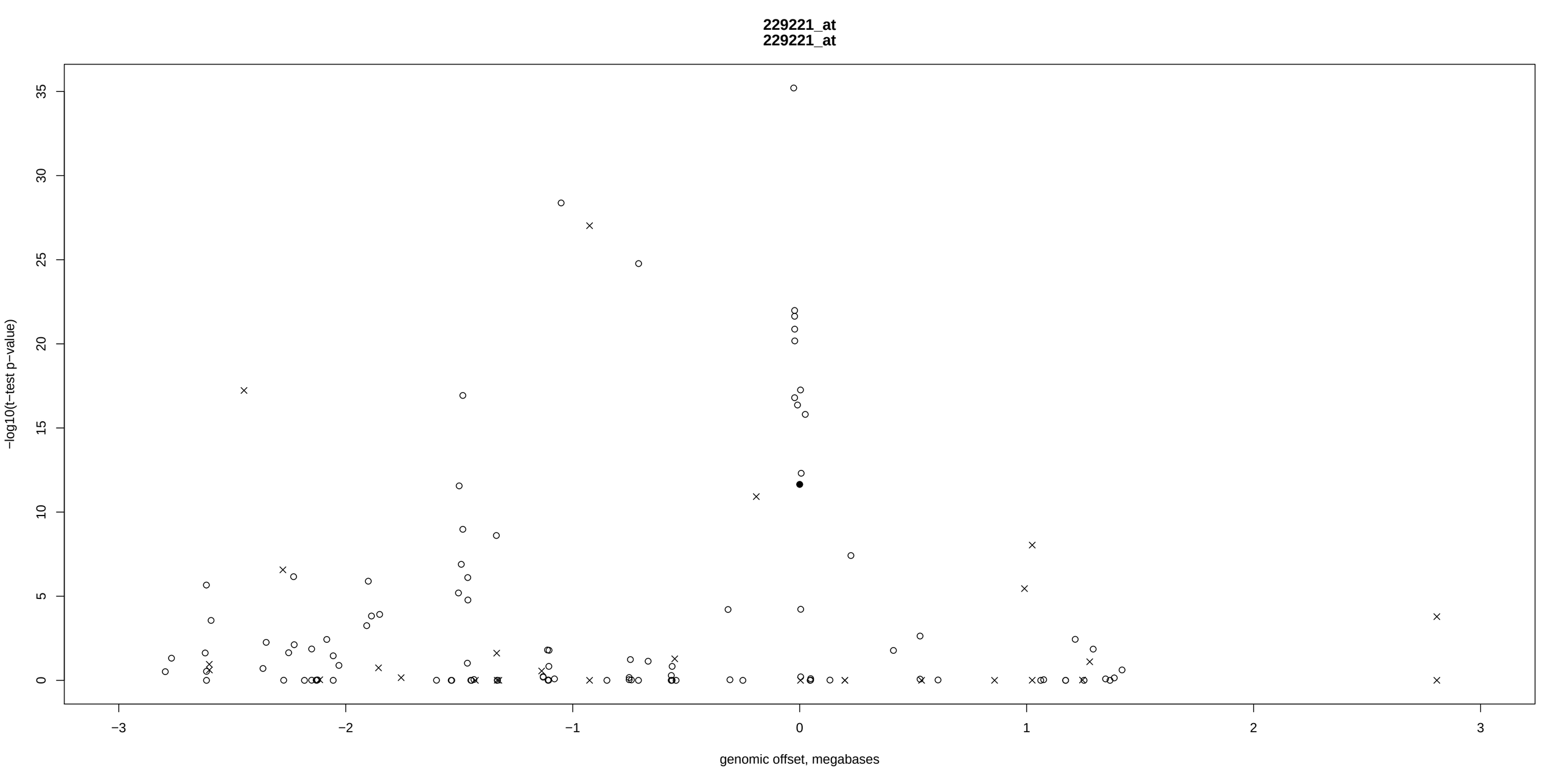
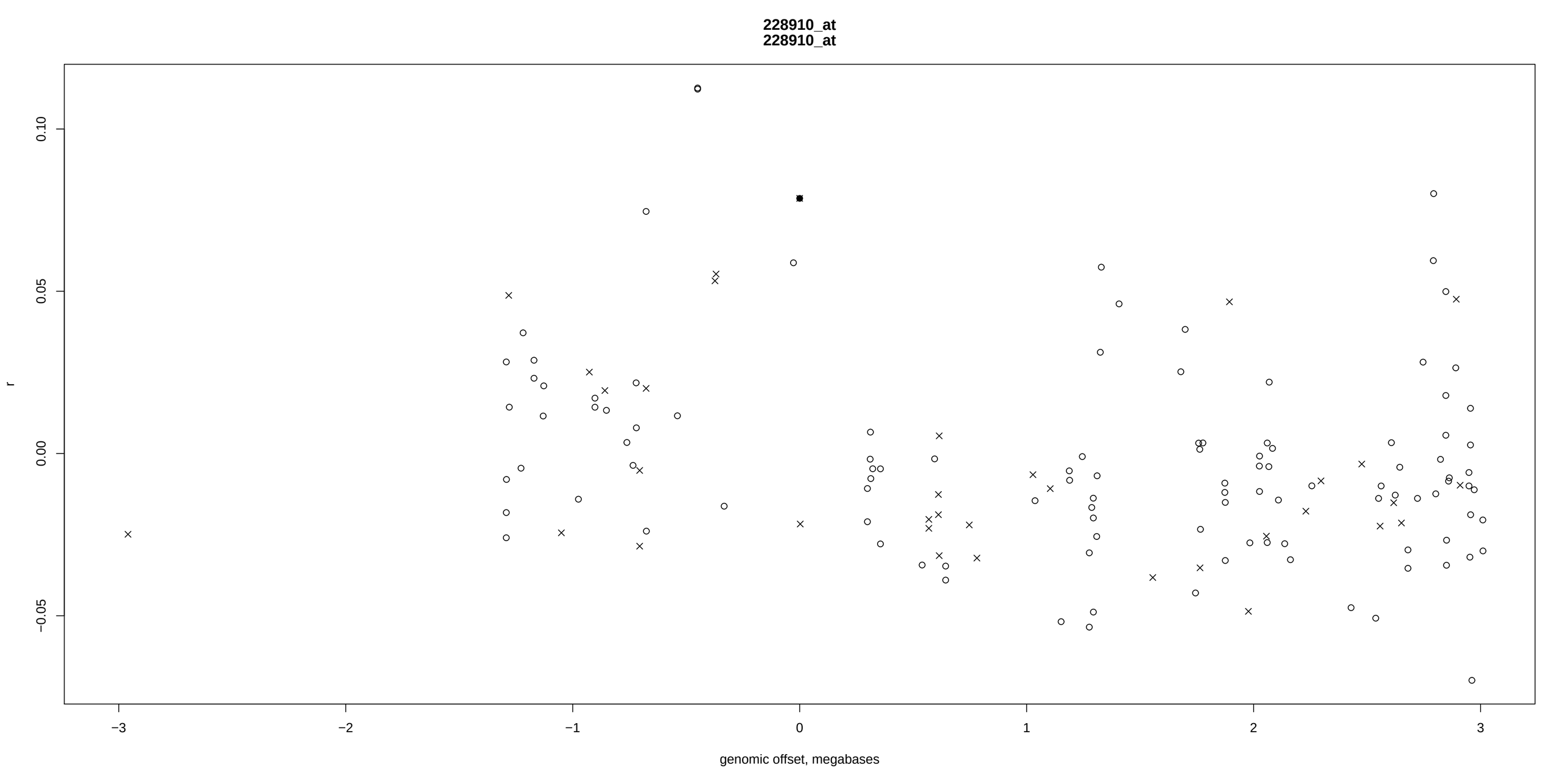
−3
−2
−1
0
1
2
3
genomic offset, megabases


229221_at
229221_at
−log10(t−test p−value)
0
5
10
15
20
25
30
35
−3
−2
−1
0
1
2
3
genomic offset, megabases


228910_at
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−1
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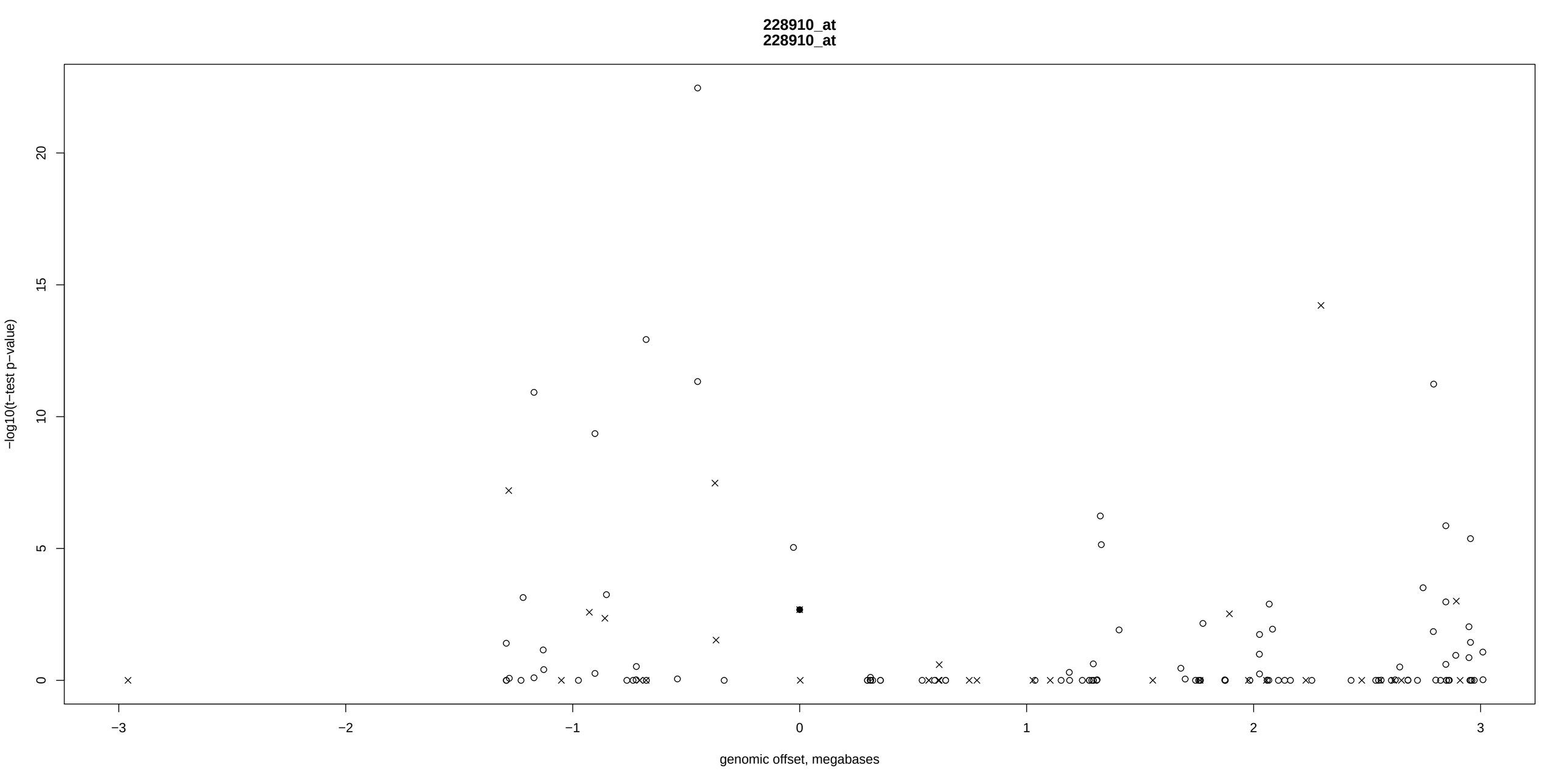
3
genomic offset, megabases


228910_at
228910_at
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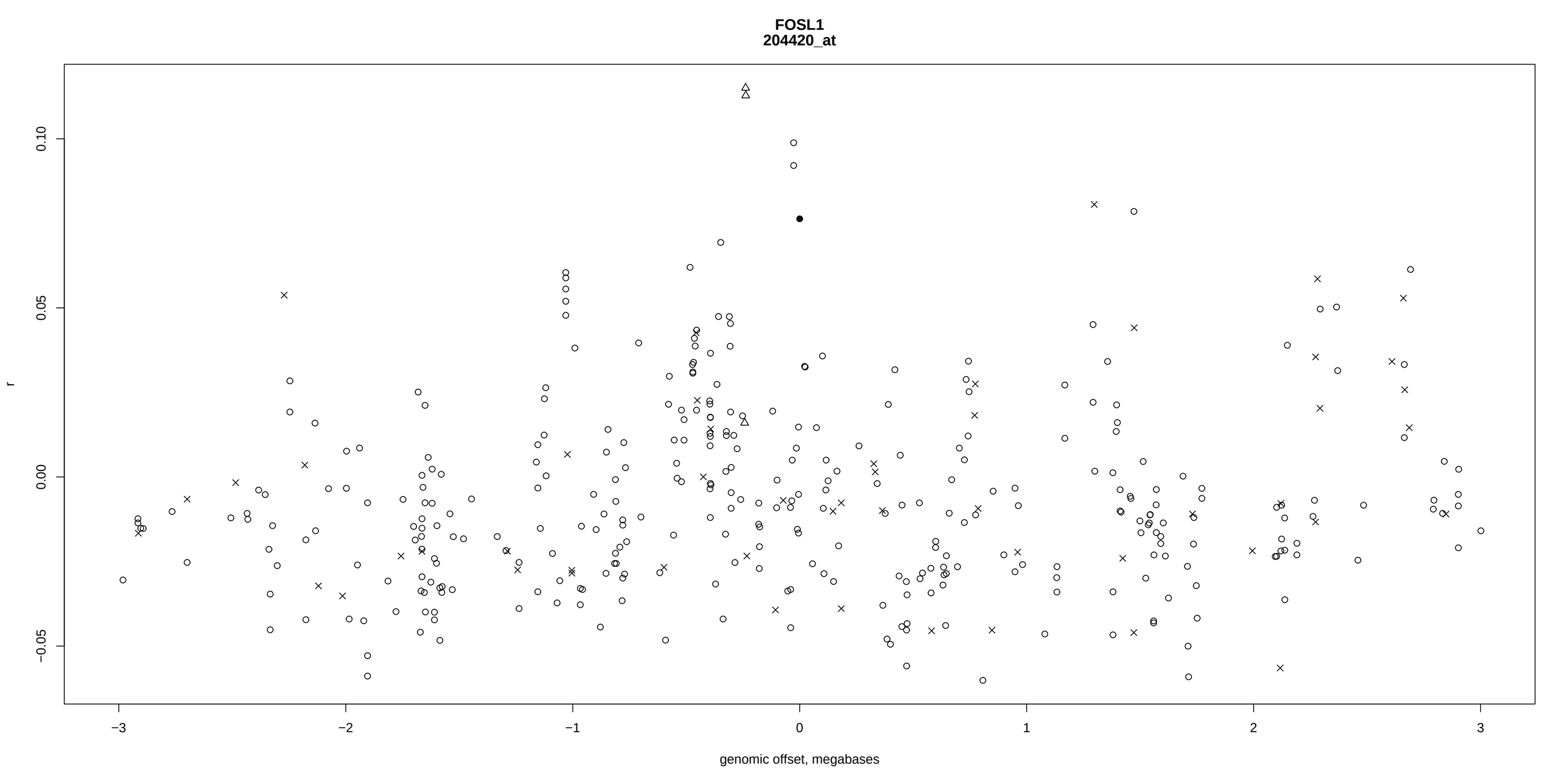
3
genomic offset, megabases


FOSL1
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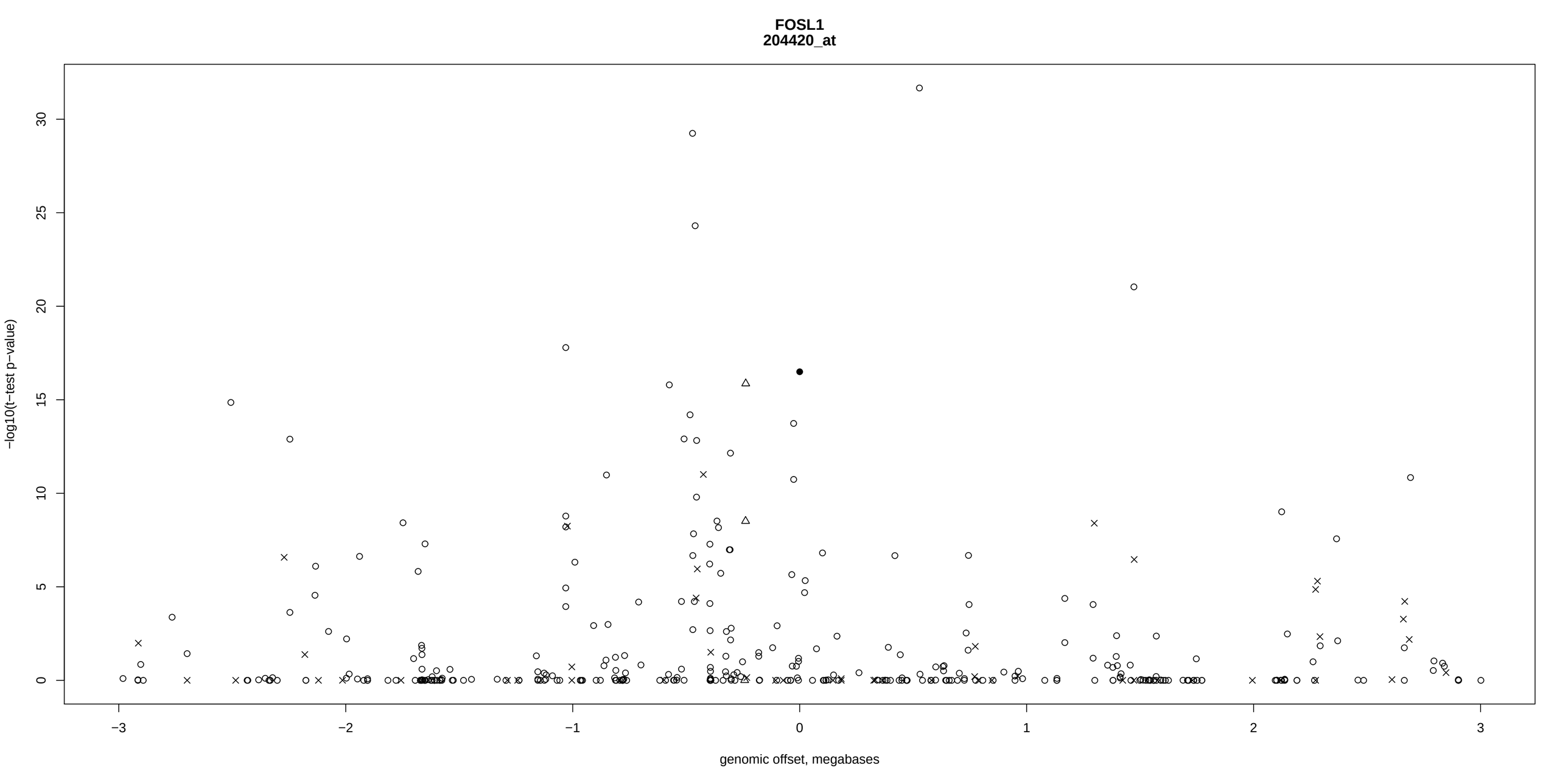
genomic offset, megabases


FOSL1
204420_at
−log10(t−test p−value)
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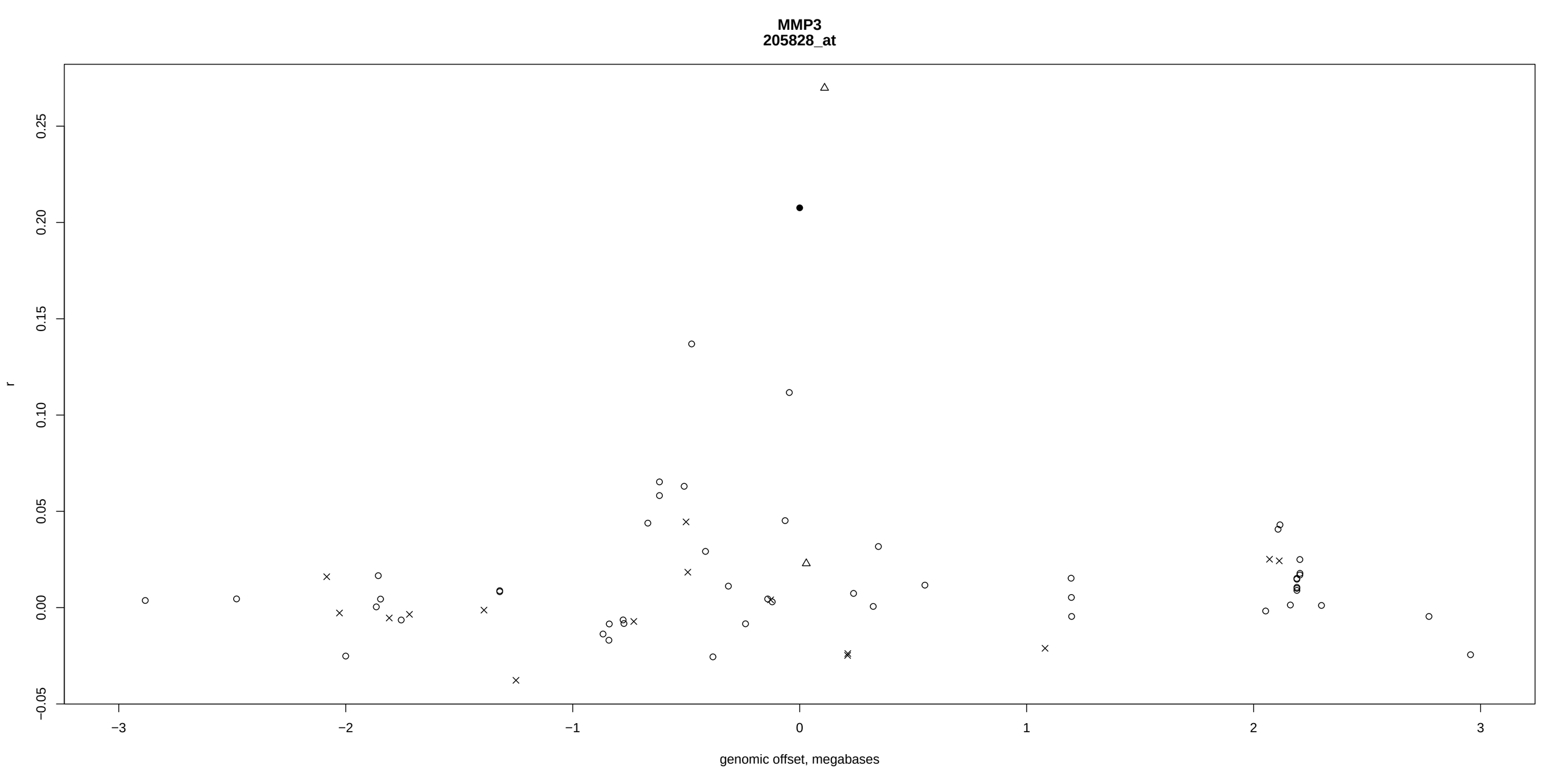
2
3
genomic offset, megabases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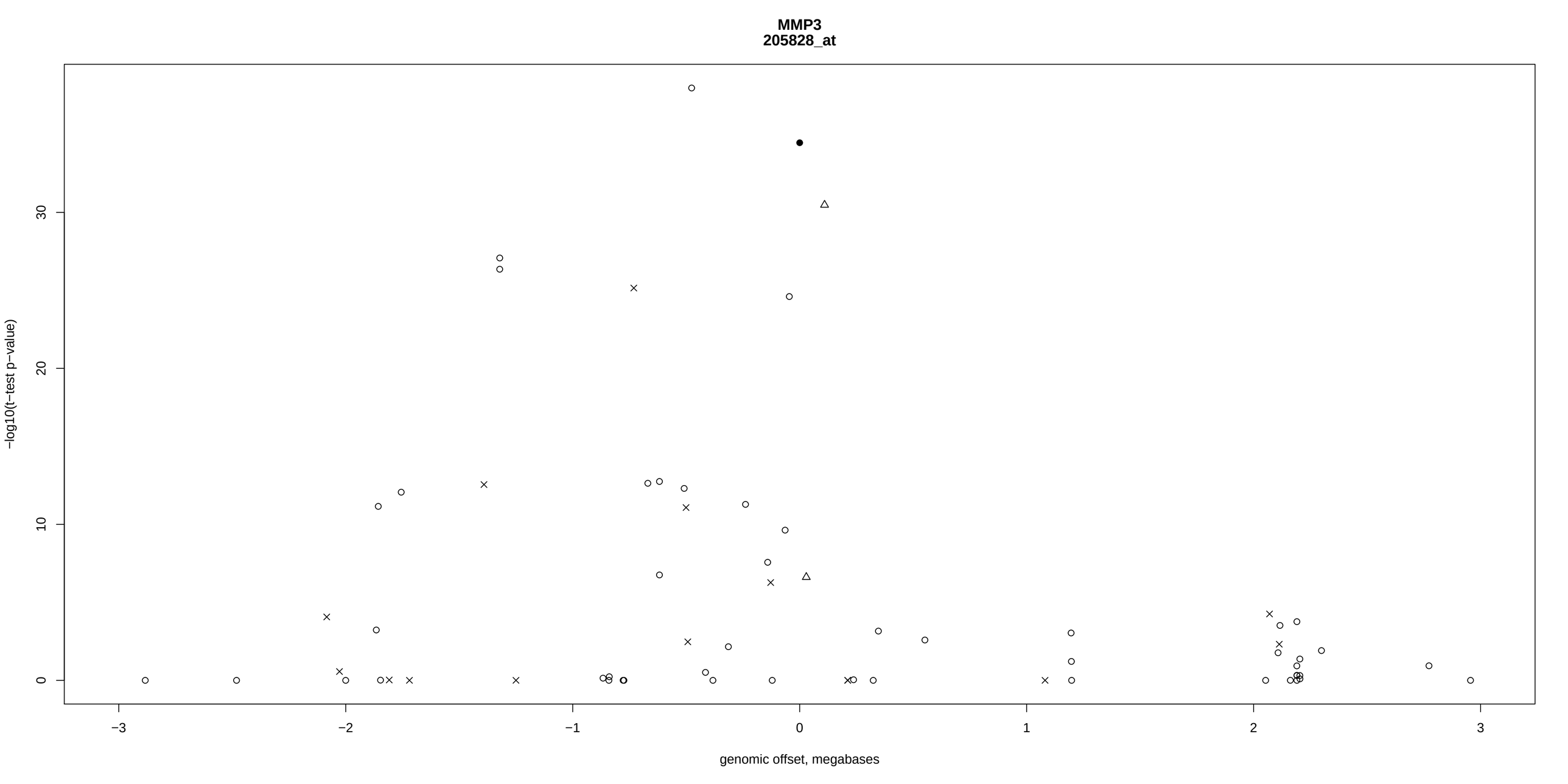
genomic offset, megabases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205828_at
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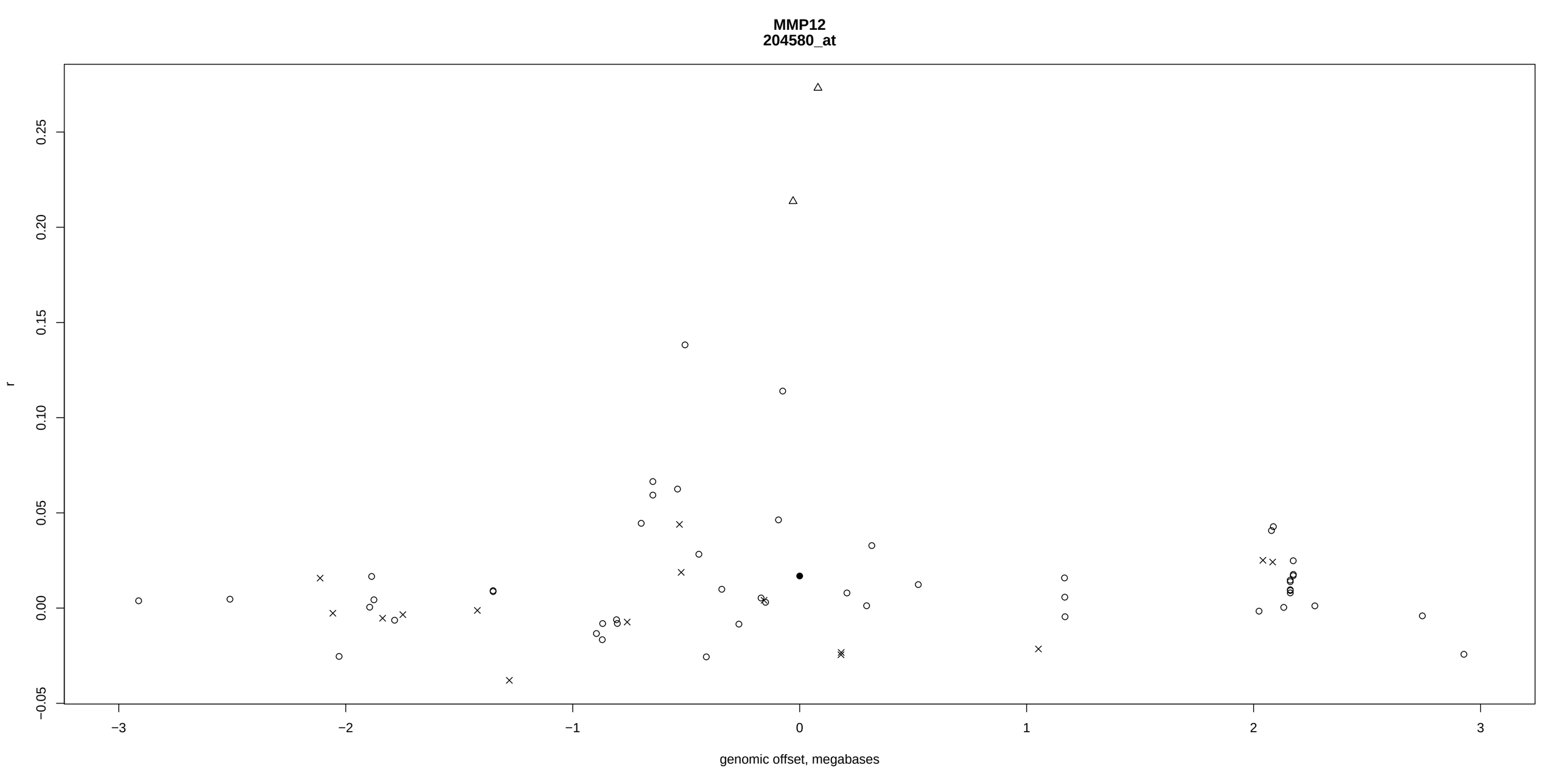
genomic offset, megabases


MMP12
204580_at
r
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0.05
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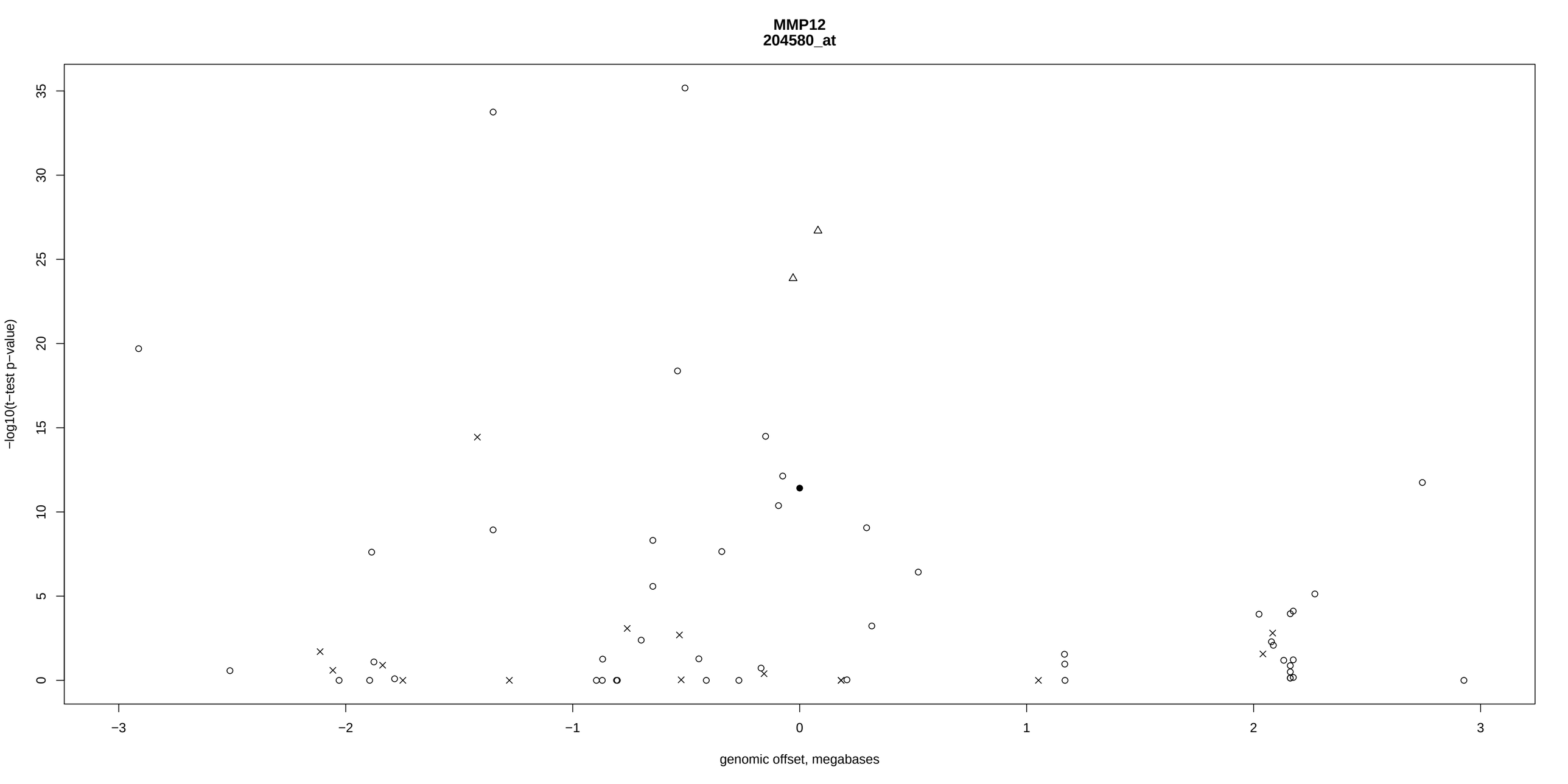
genomic offset, megabases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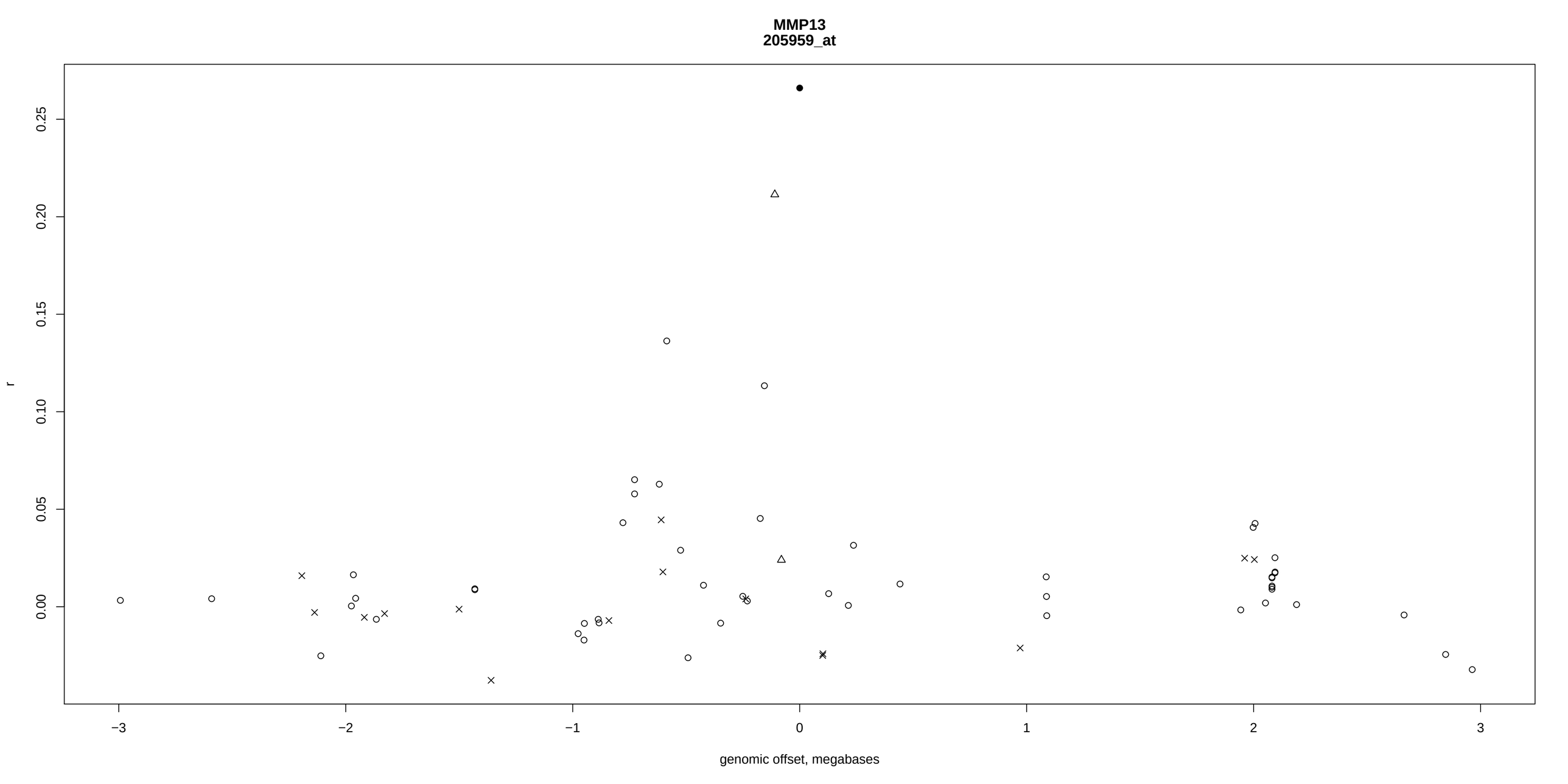
3
genomic offset, megabases


MMP13
205959_at
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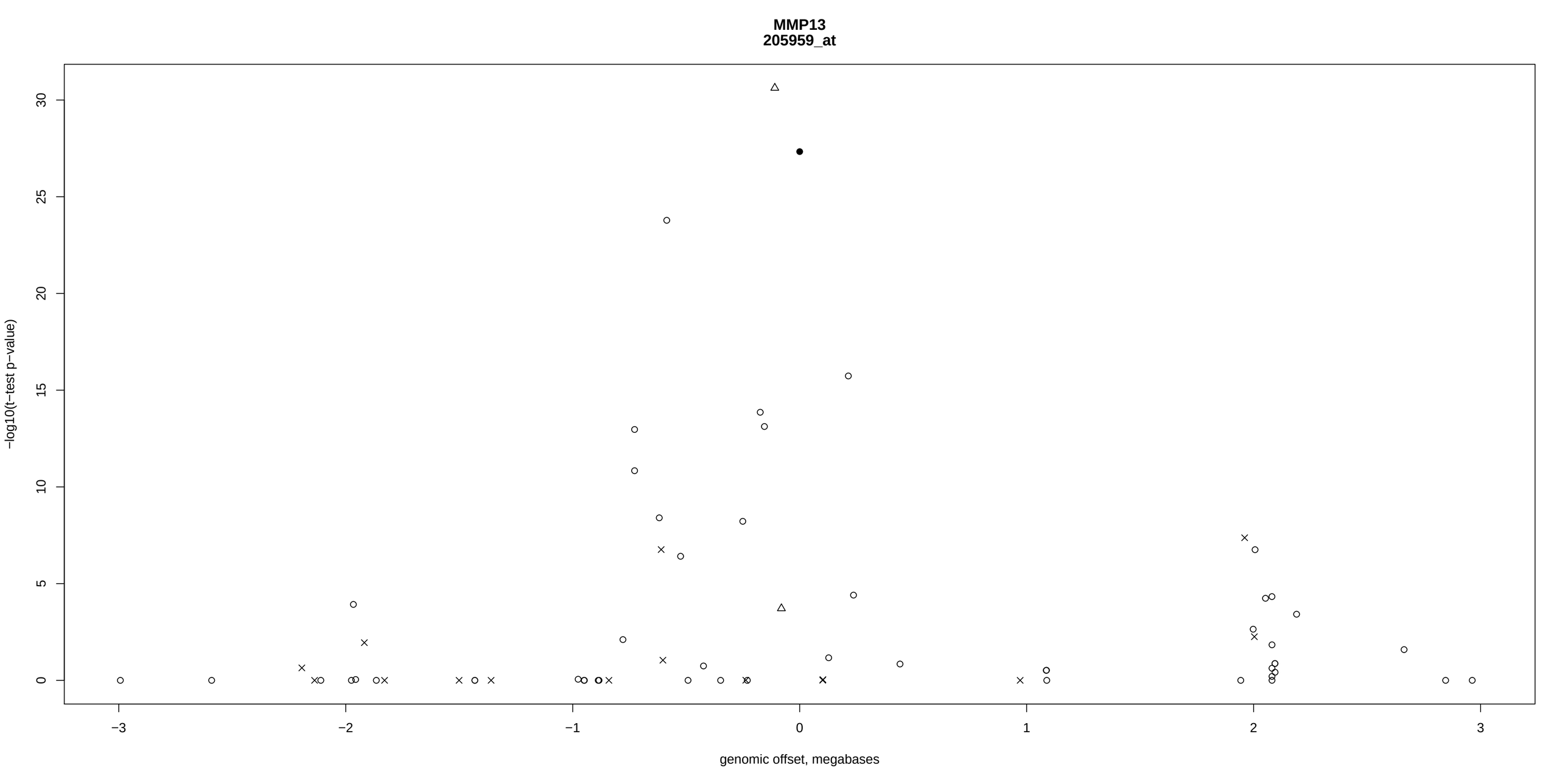
3
genomic offset, megabases


MMP13
205959_at
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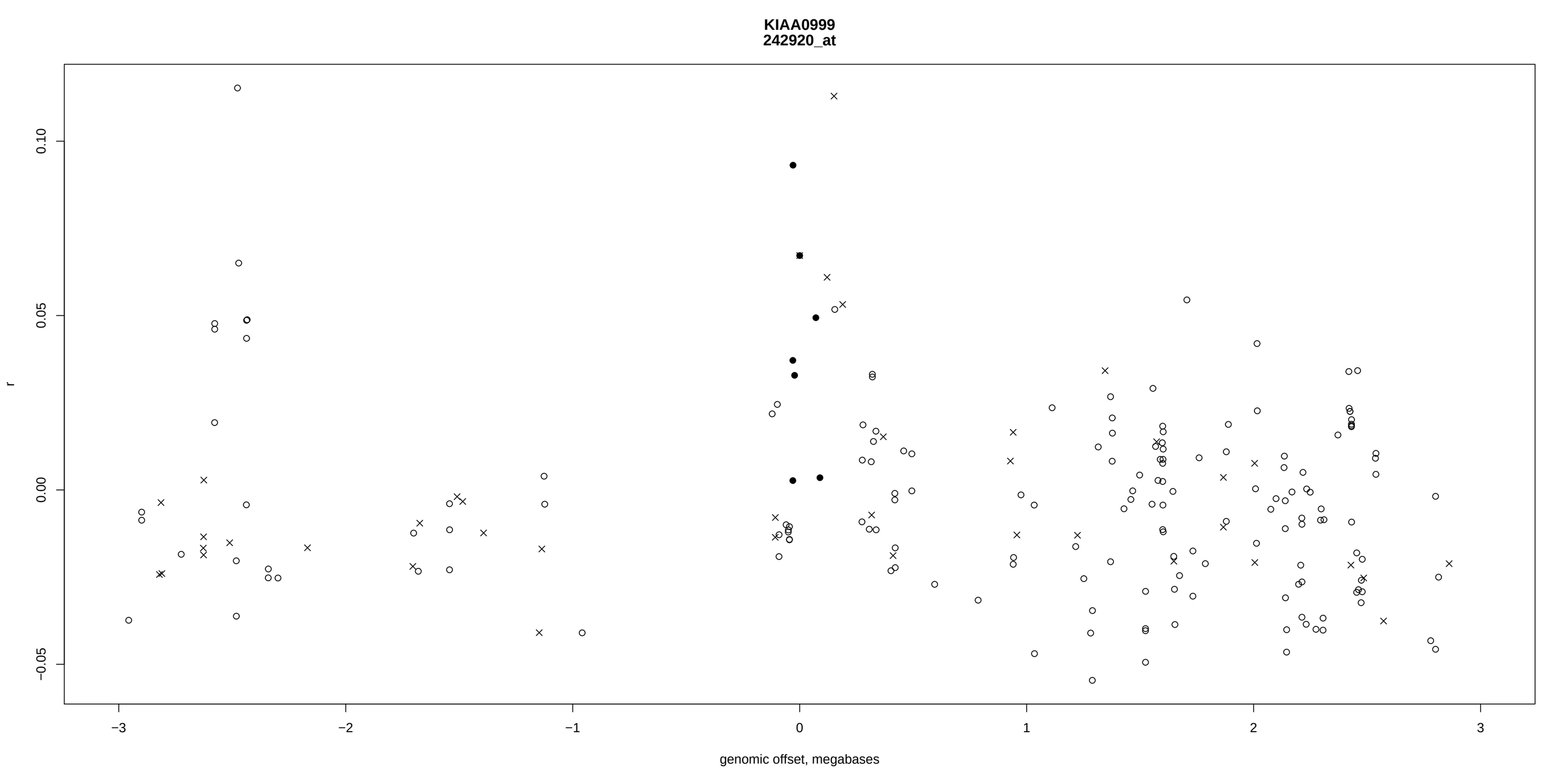
genomic offset, megabases


KIAA0999
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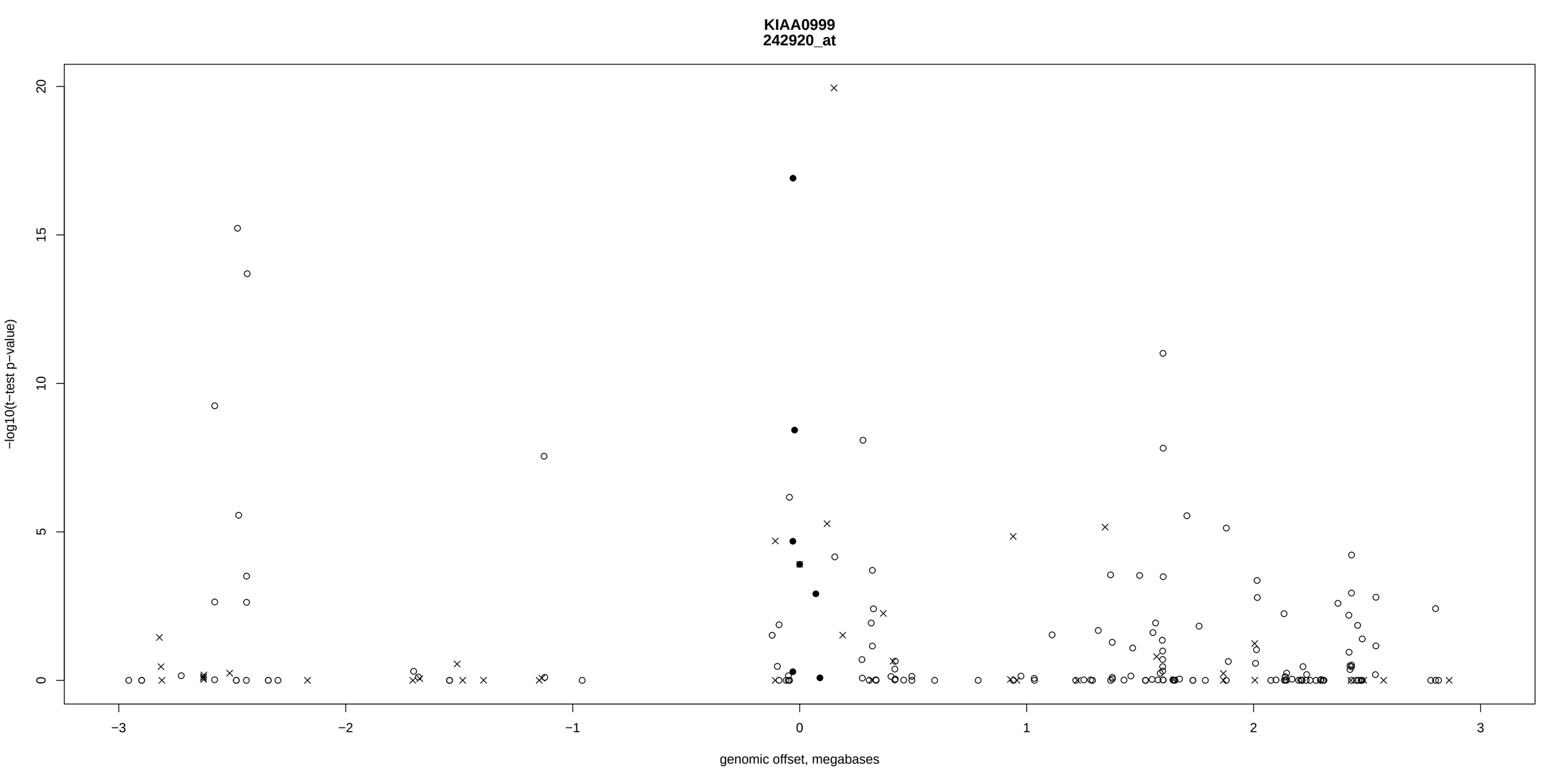
3
genomic offset, megabases


KIAA0999
242920_at
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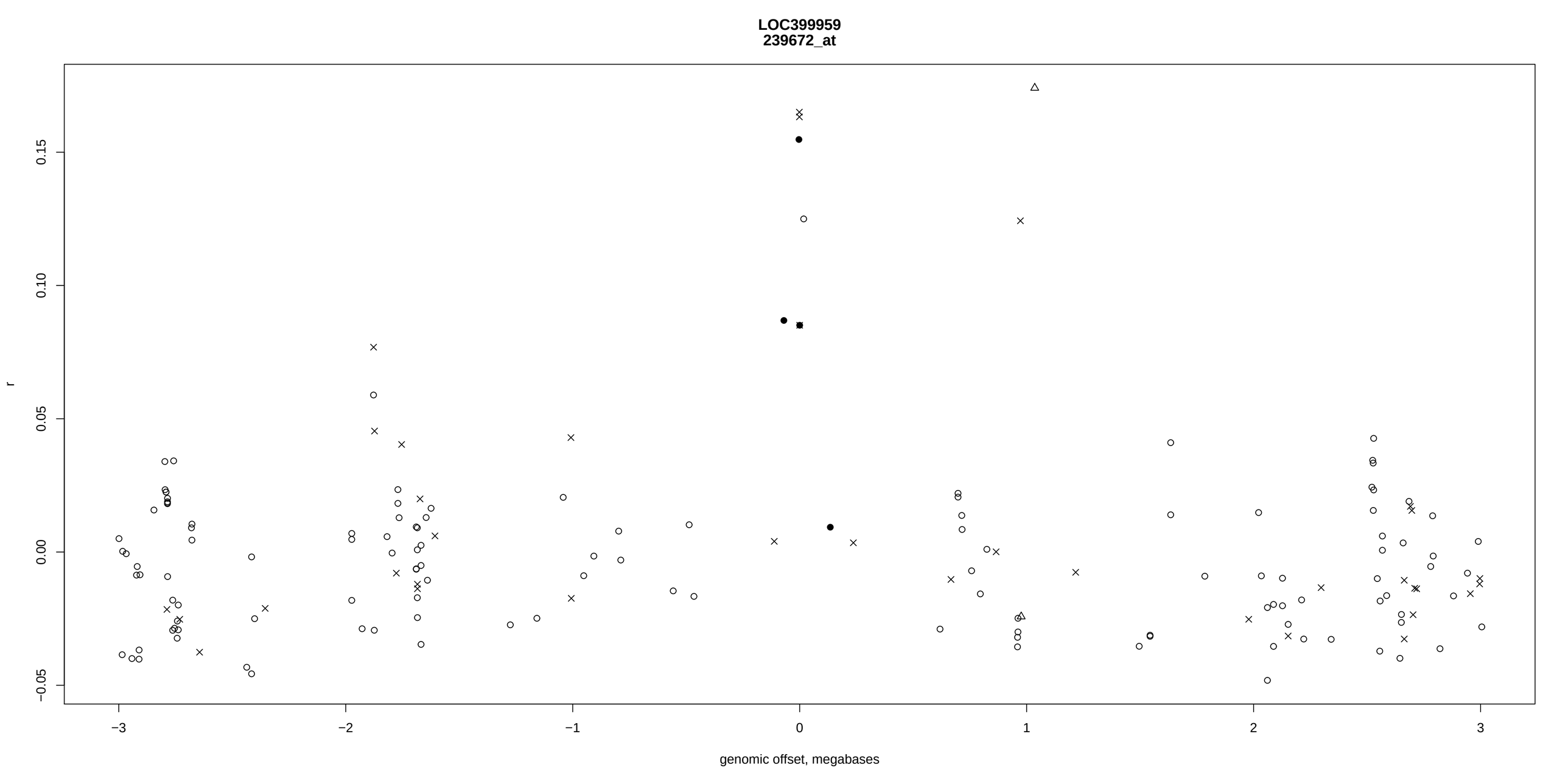
3
genomic offset, megabases


LOC399959
239672_at
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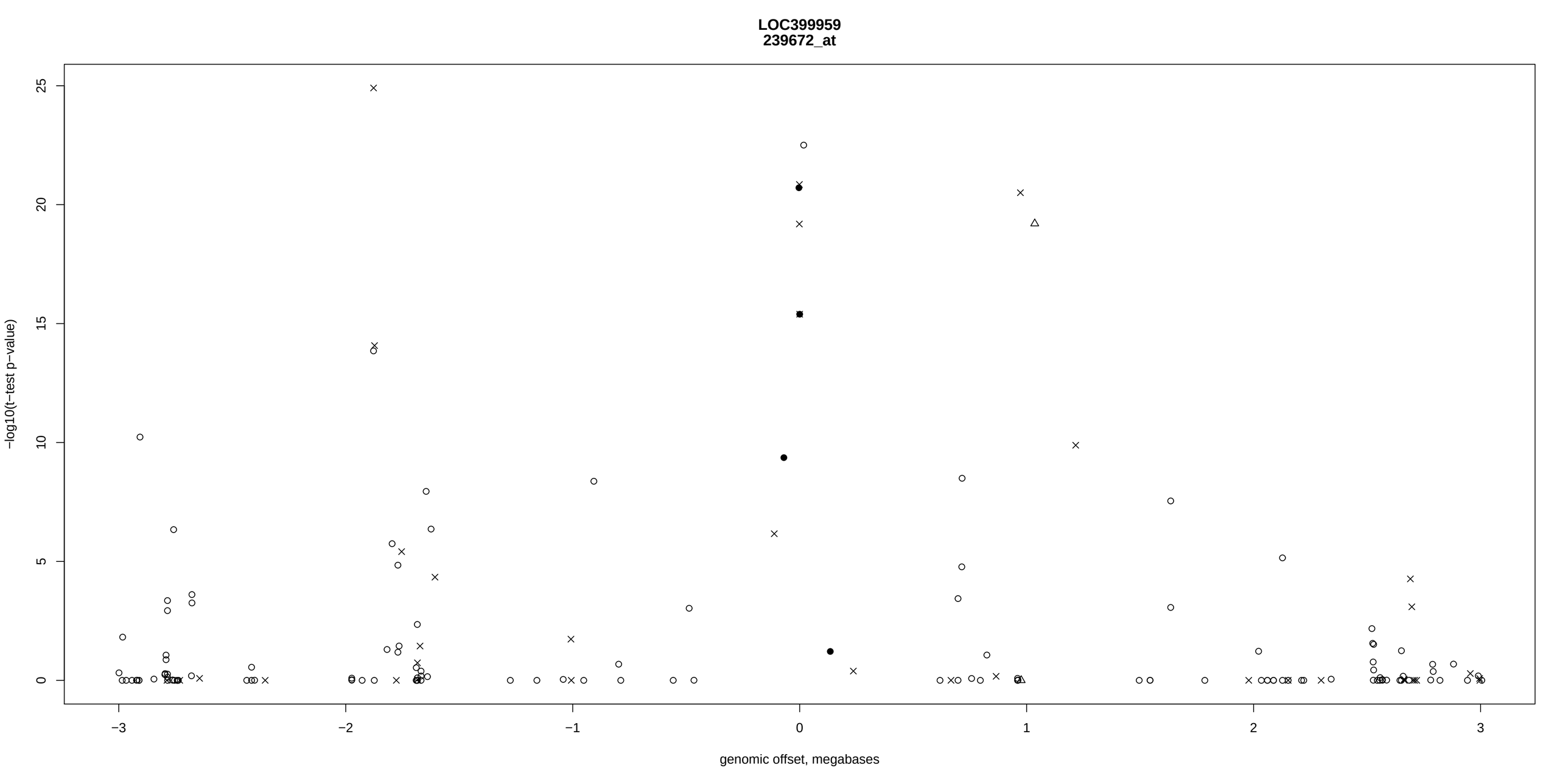
3
genomic offset, megabases


LOC399959
239672_at
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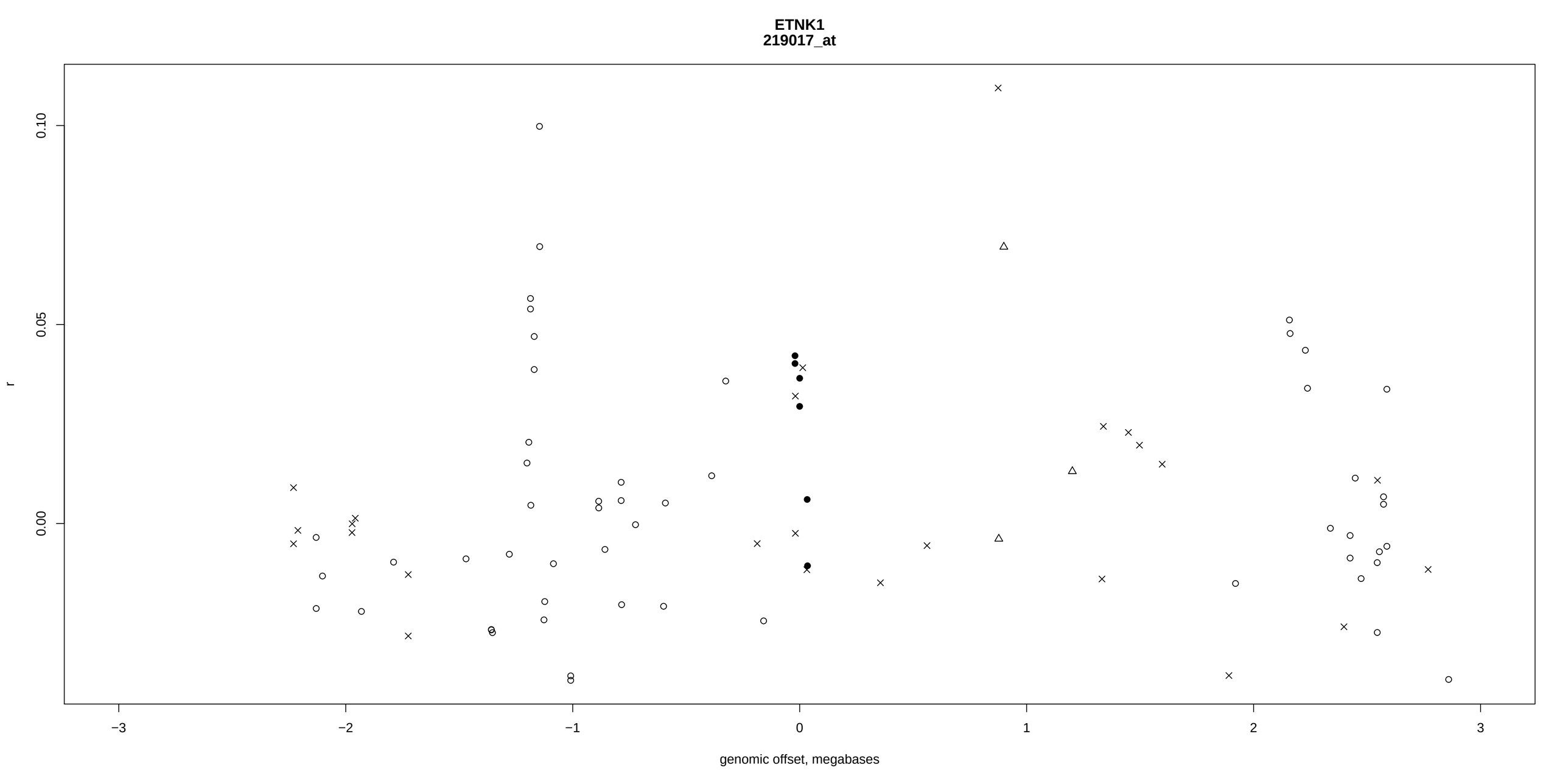
3
genomic offset, megabases


ETNK1
219017_at
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0.00
r
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−1
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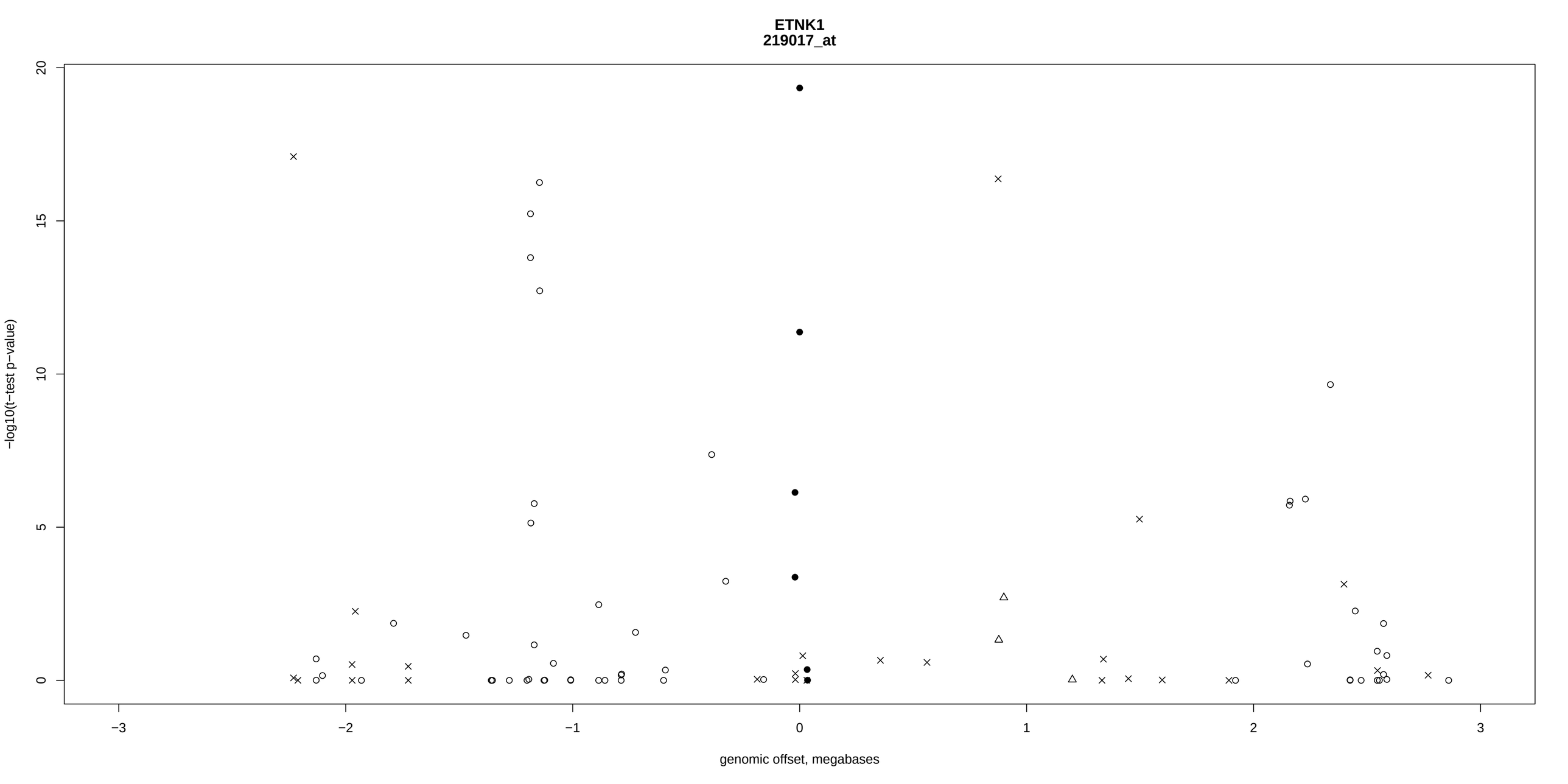
1
2
3
genomic offset, megabases


ETNK1
219017_at
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10
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20
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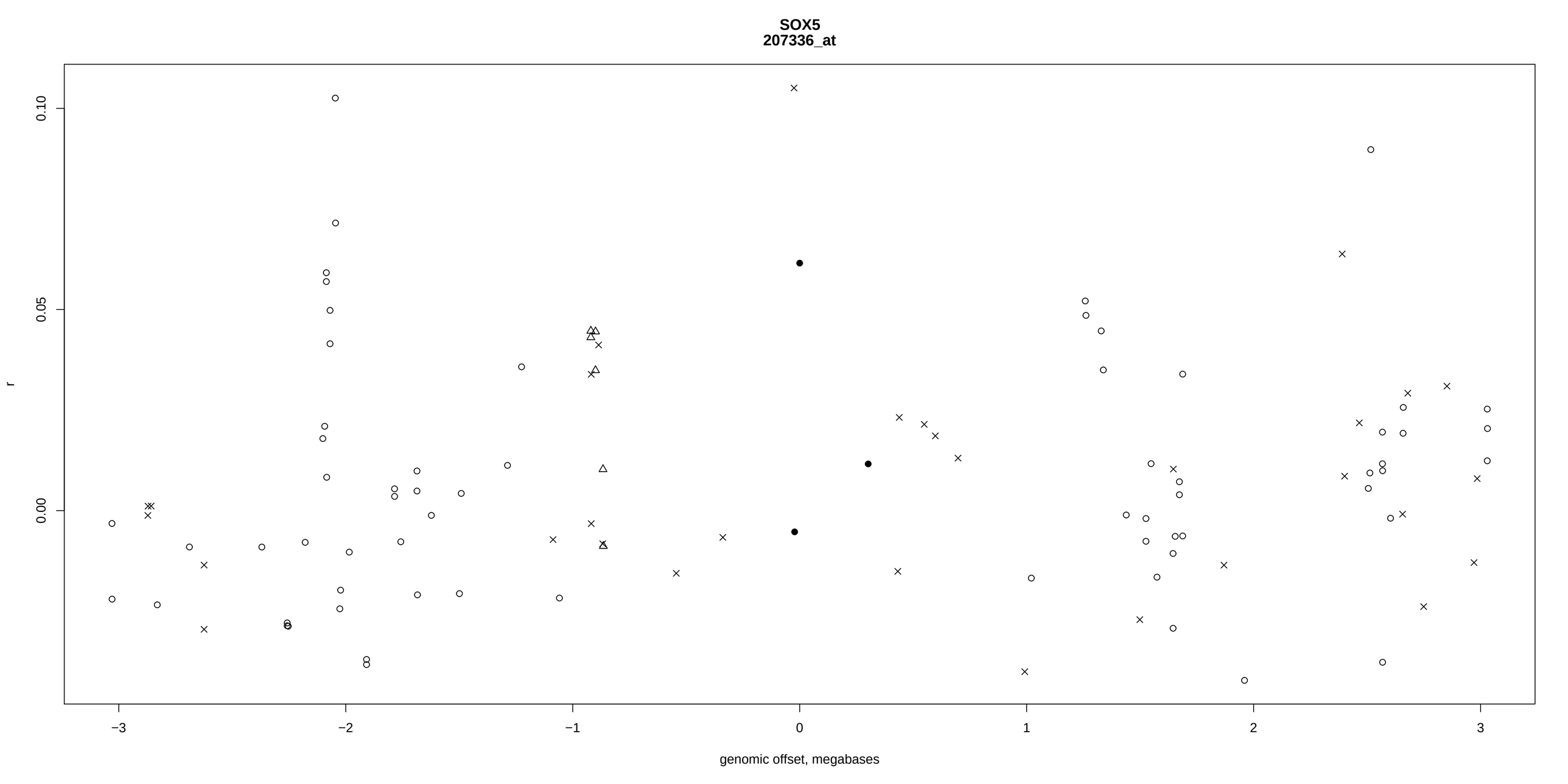
3
genomic offset, megabases


SOX5
207336_at
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0.10
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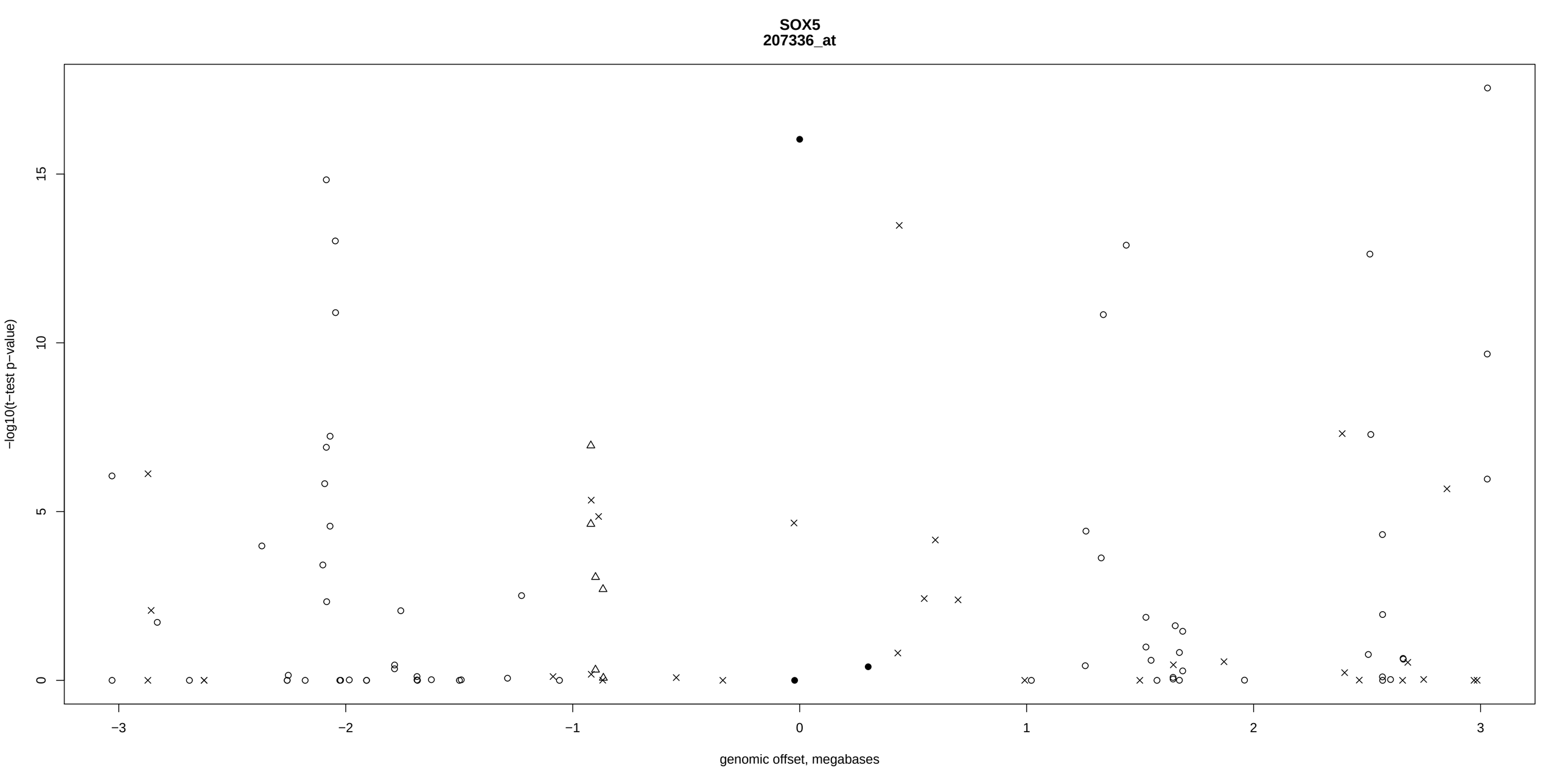
2
3
genomic offset, megabases


SOX5
207336_at
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−3
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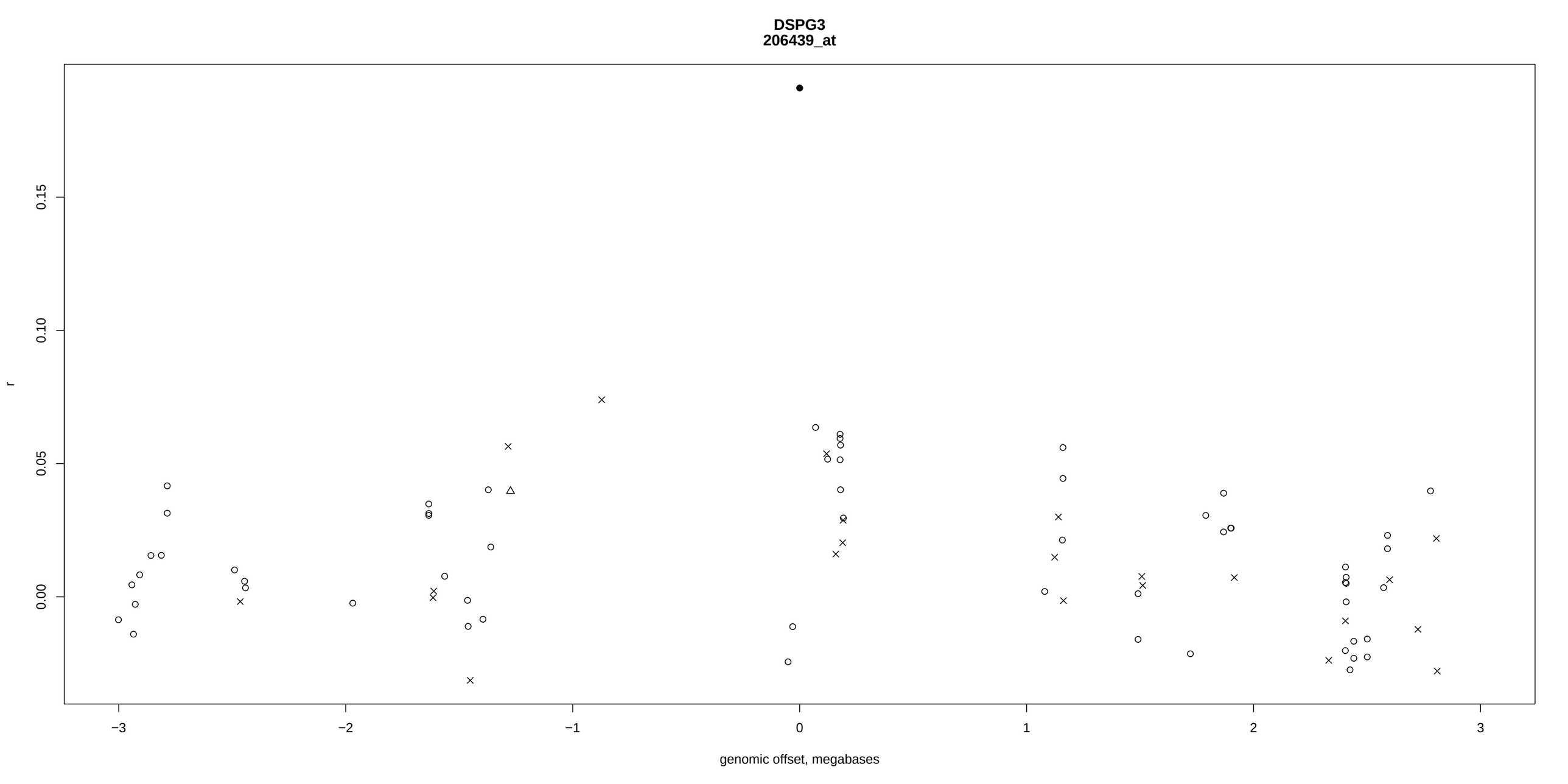
2
3
genomic offset, megabases


DSPG3
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r
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0.10
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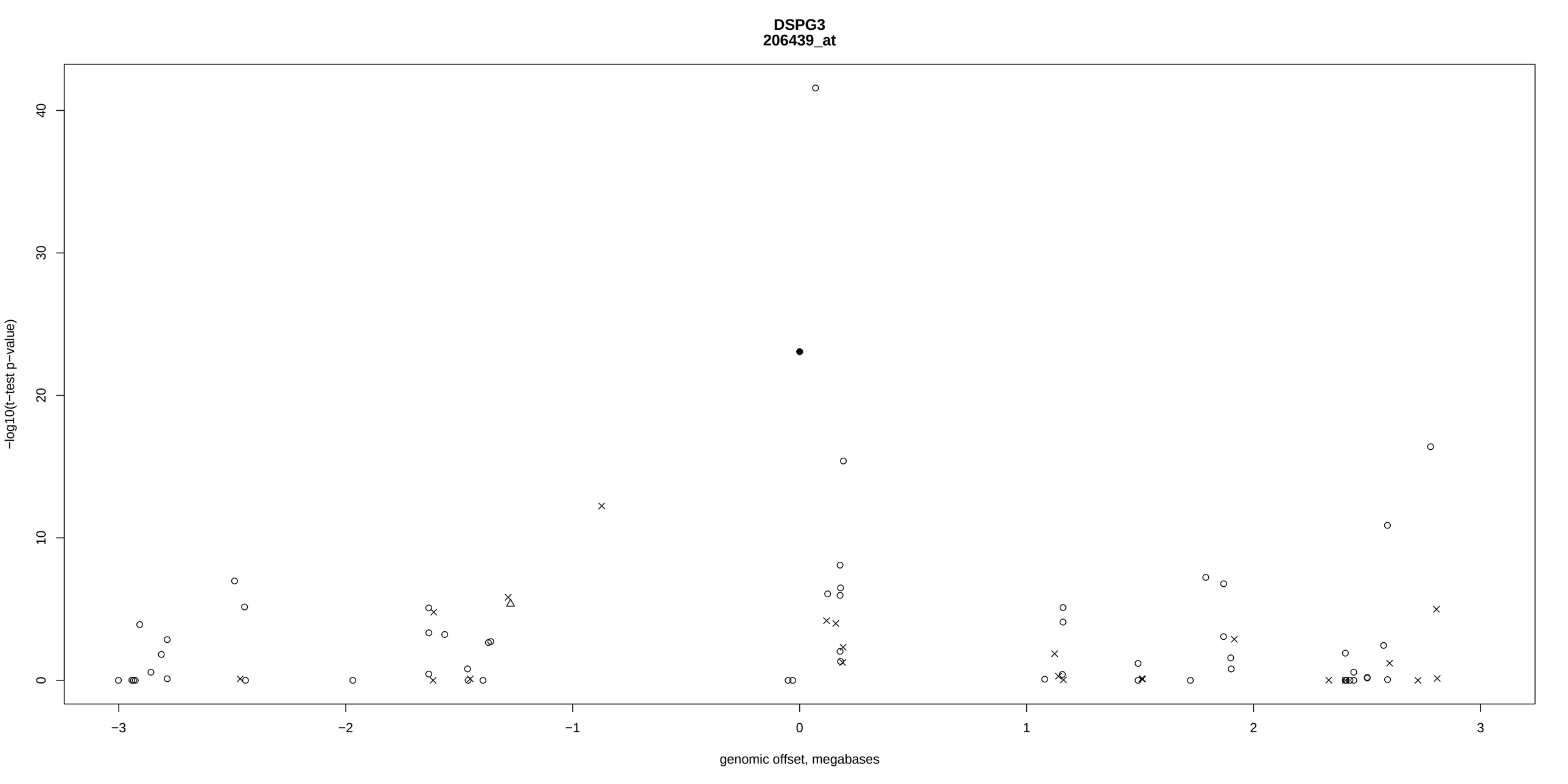
3
genomic offset, megabases


DSPG3
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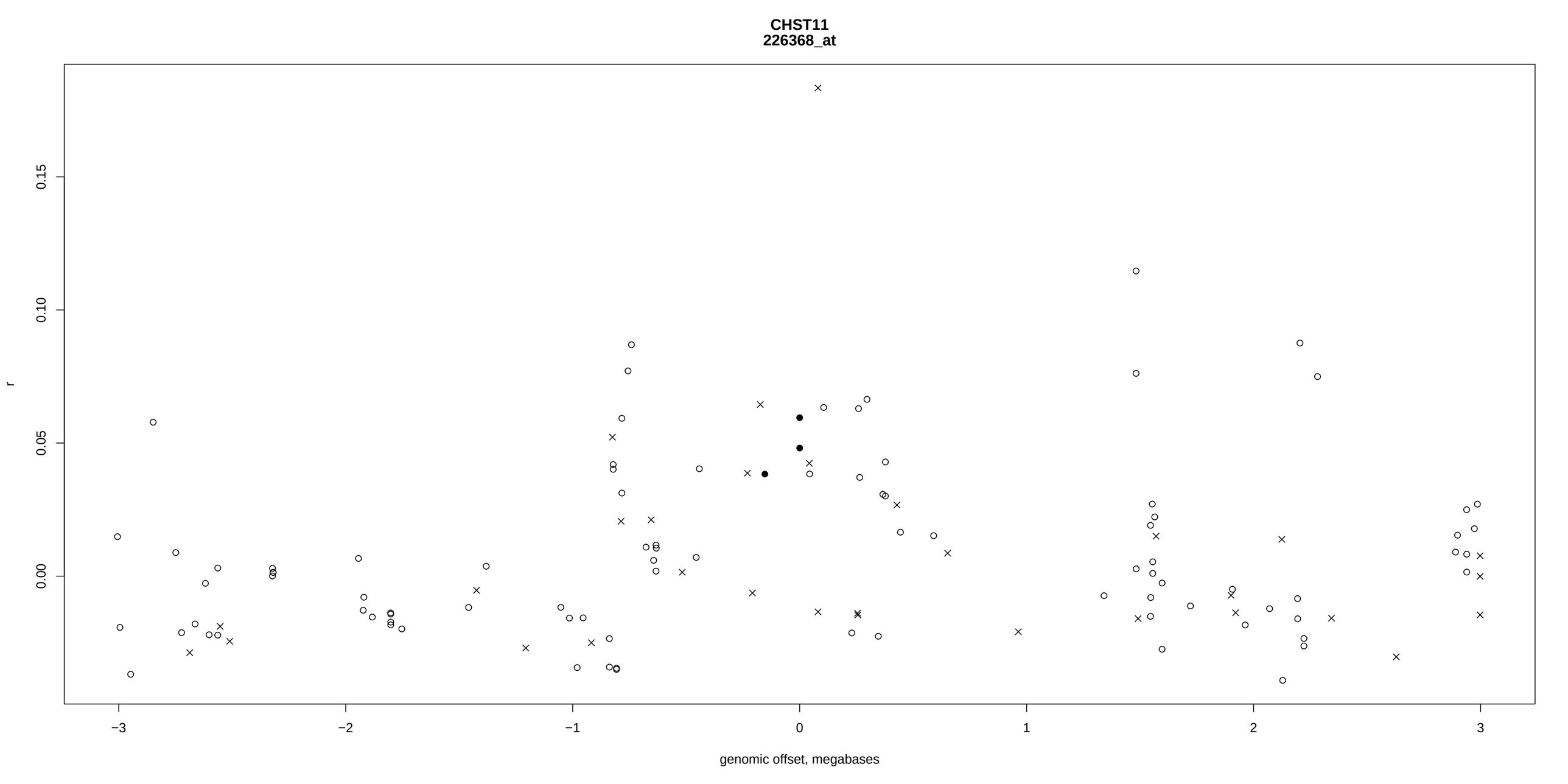
−1
0
1
2
3
genomic offset, megabases


CHST11
226368_at
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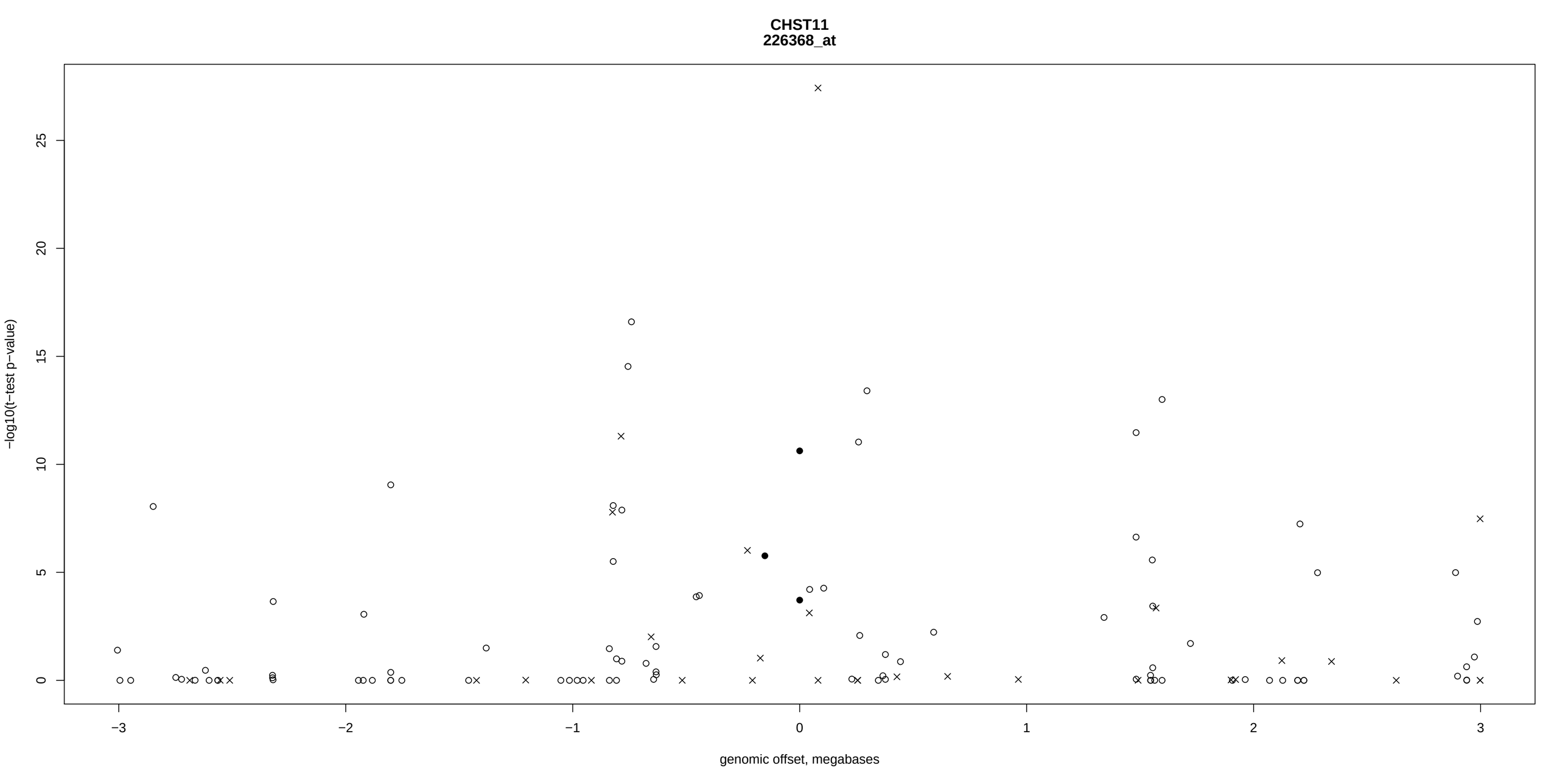
3
genomic offset, megabases


CHST11
226368_at
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20
25
−3
−2
−1
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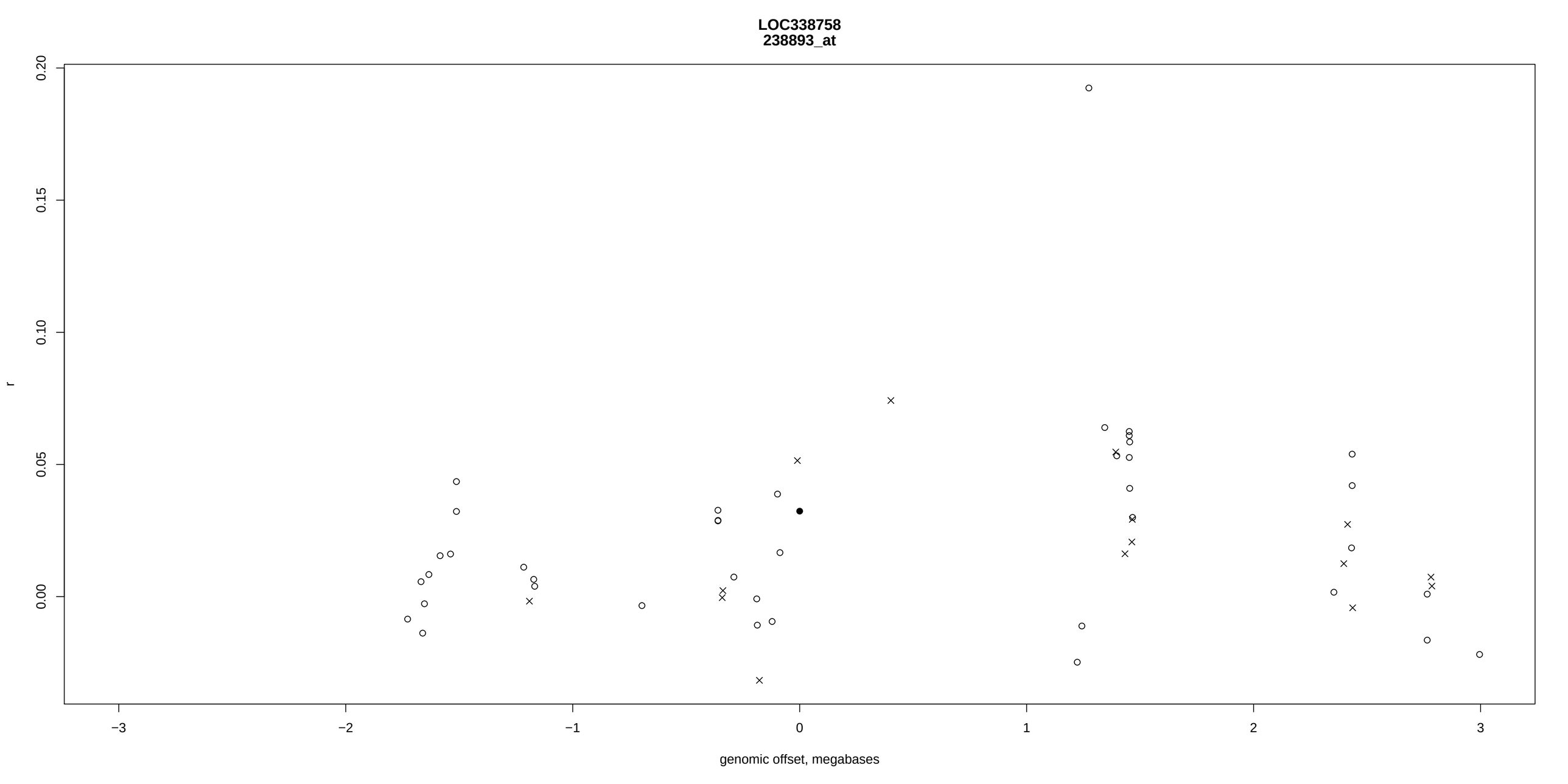
2
3
genomic offset, megabases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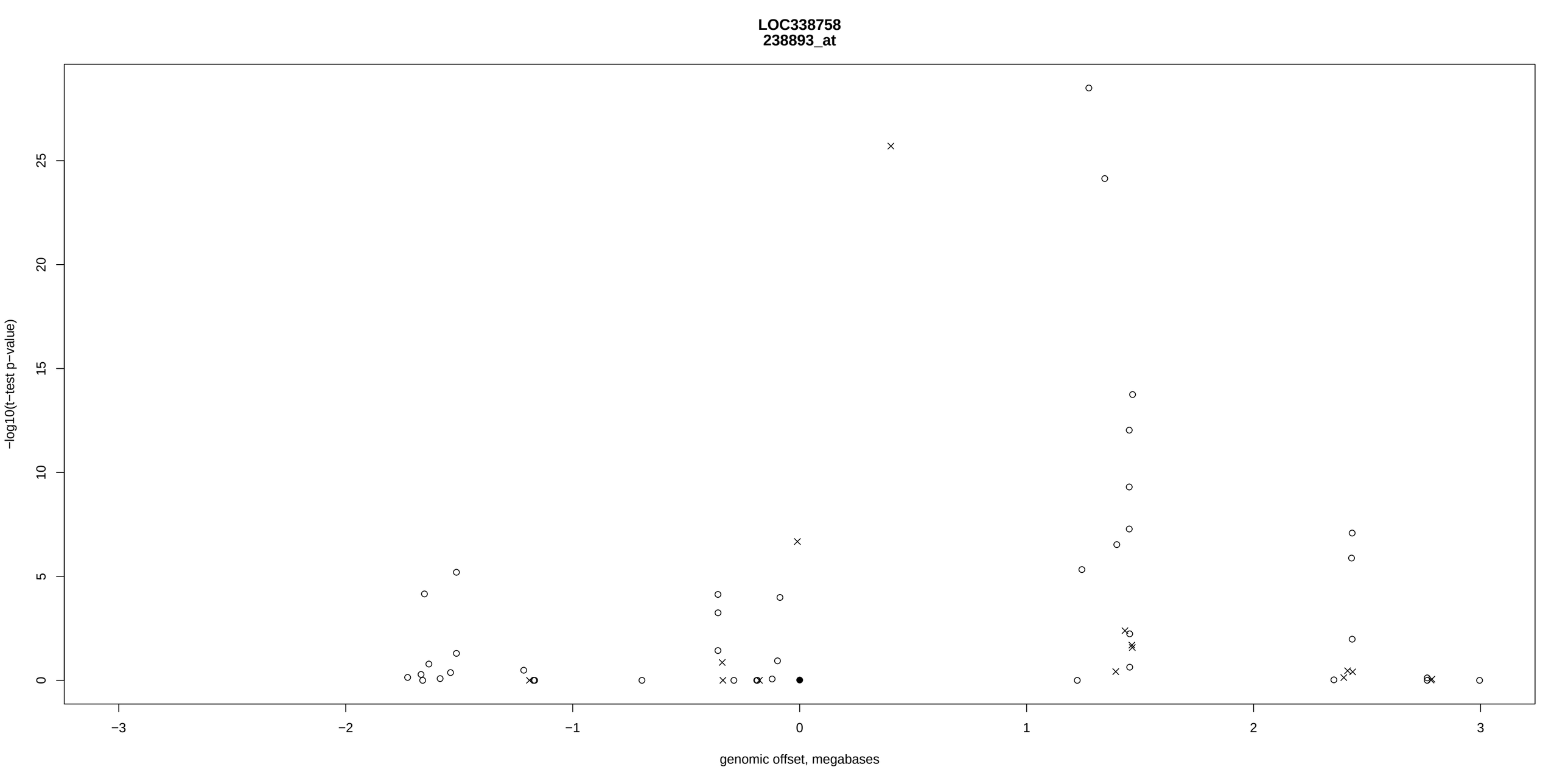
genomic offset, megabases


LOC338758
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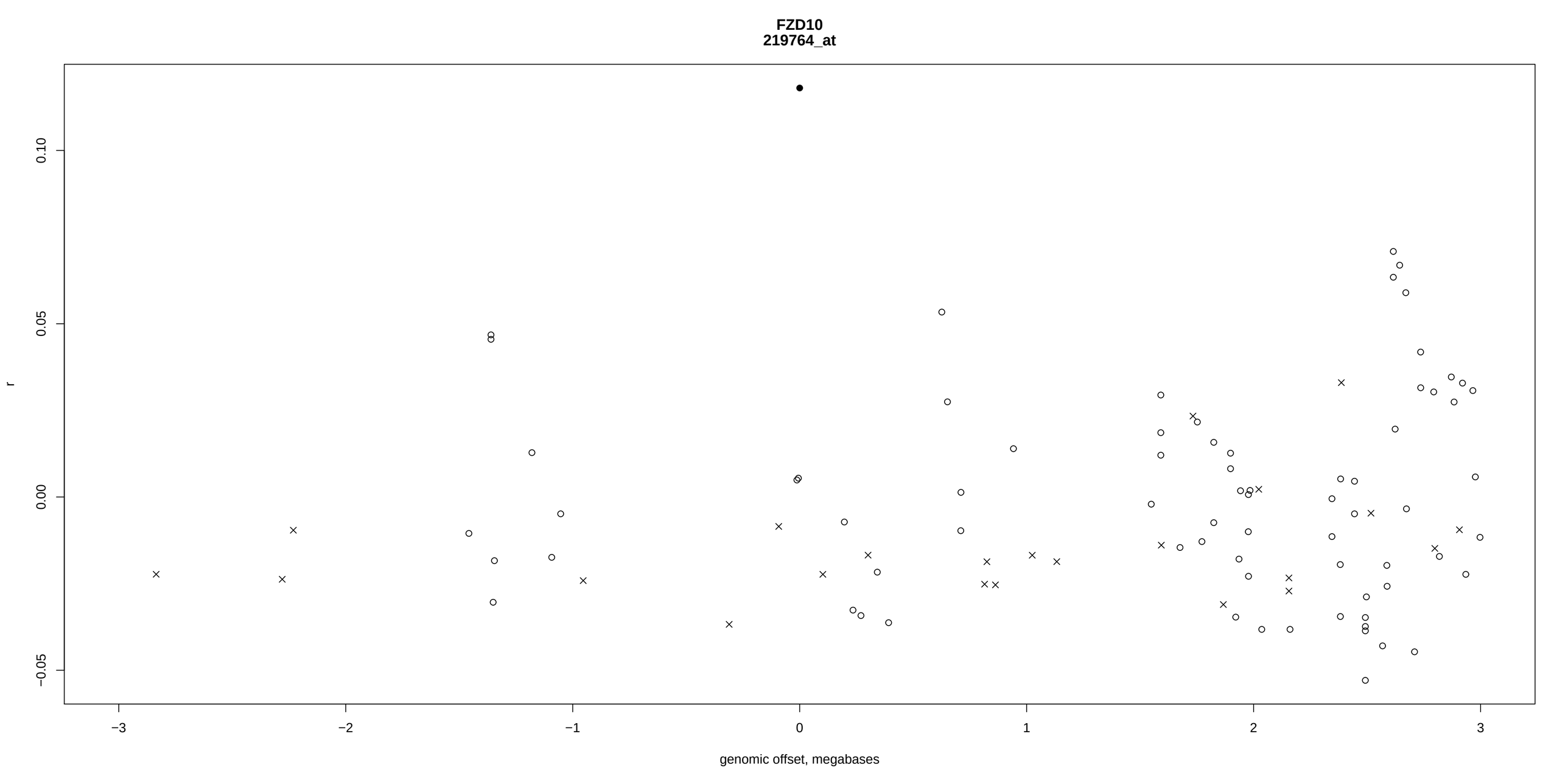
3
genomic offset, megabases


FZD10
219764_at
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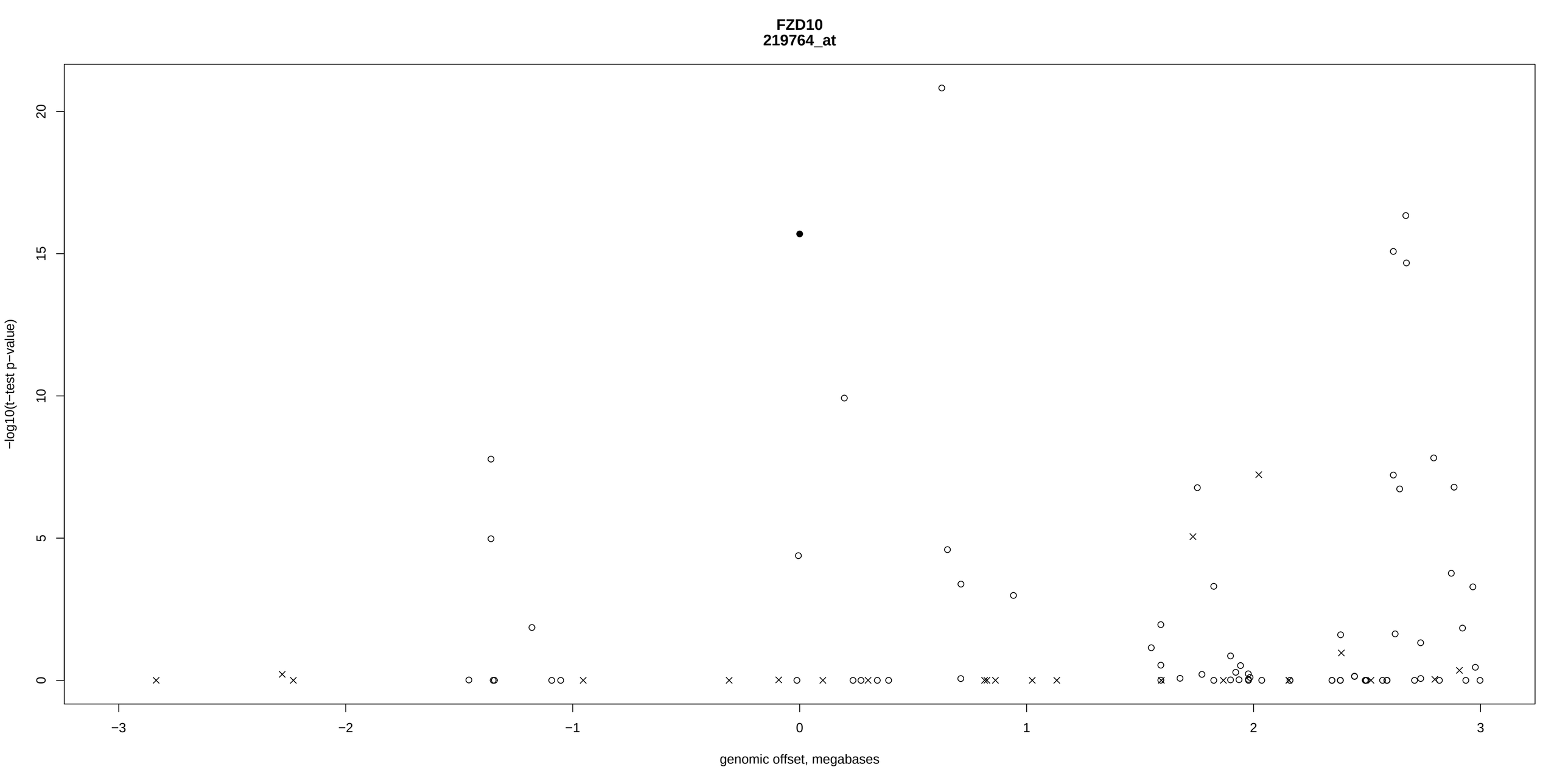
genomic offset, megabases


FZD10
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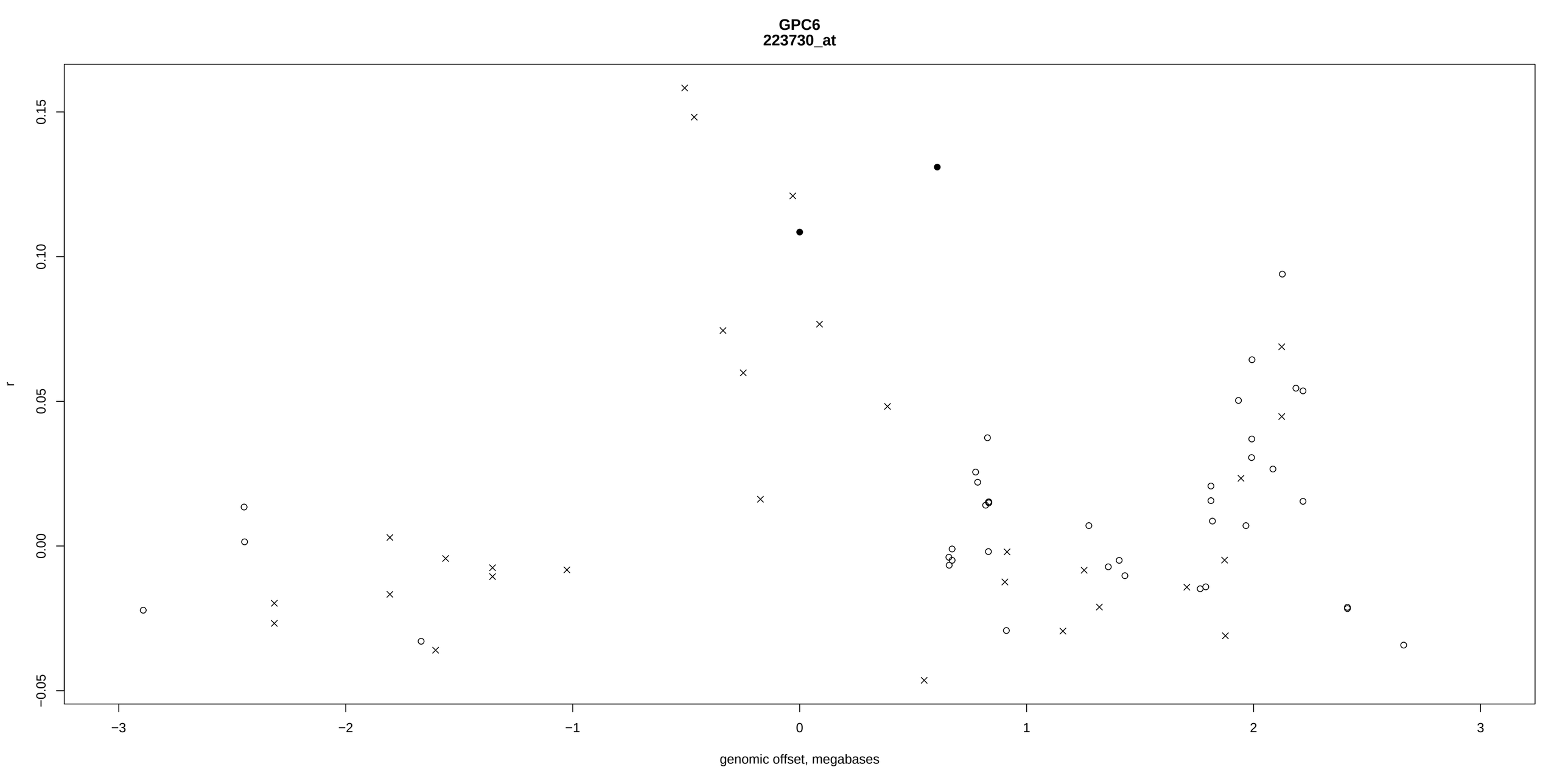
3
genomic offset, megabases


GPC6
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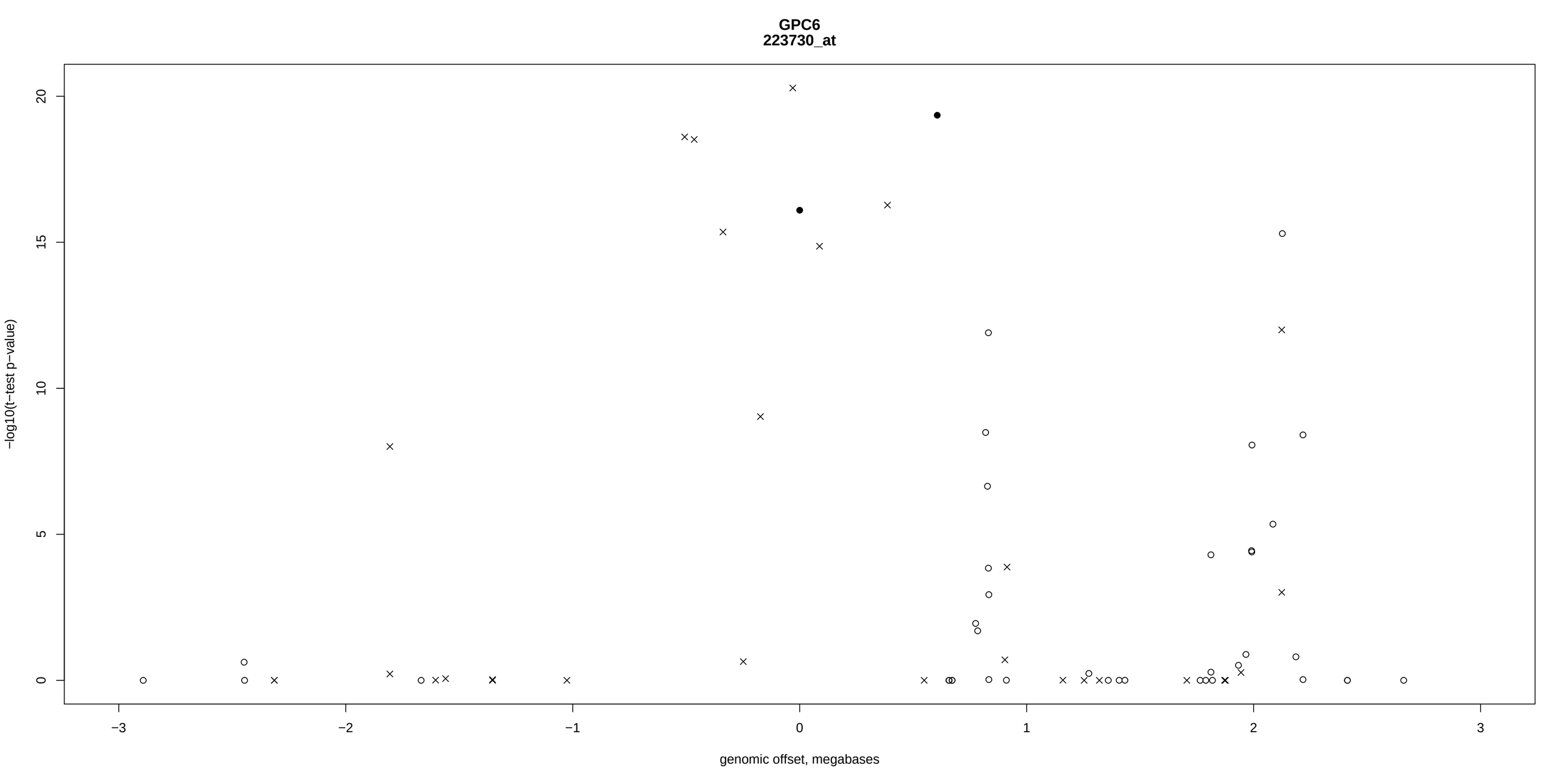
3
genomic offset, megabases


GPC6
223730_at
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20
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−2
−1
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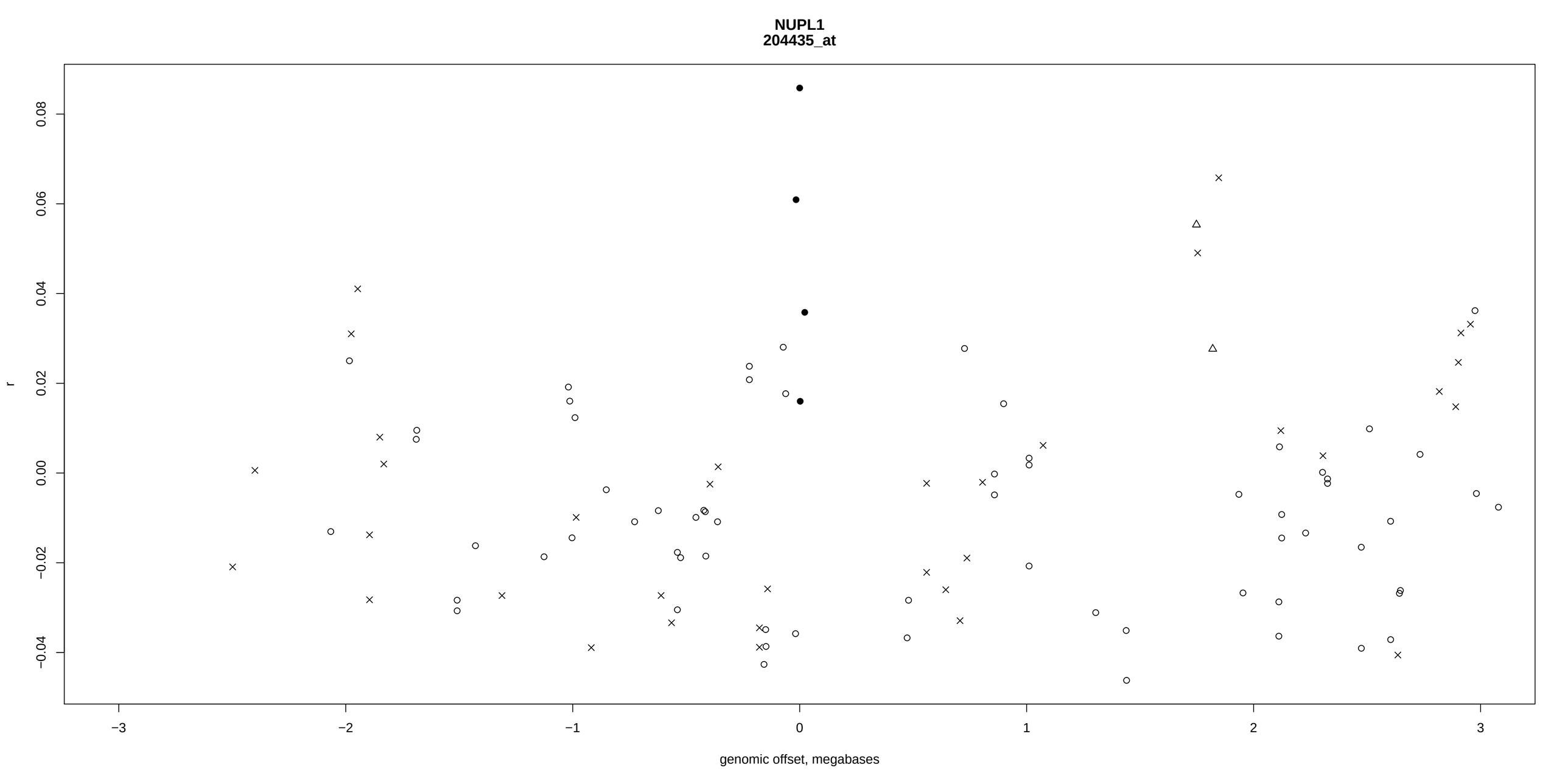
2
3
genomic offset, megabases


NUPL1
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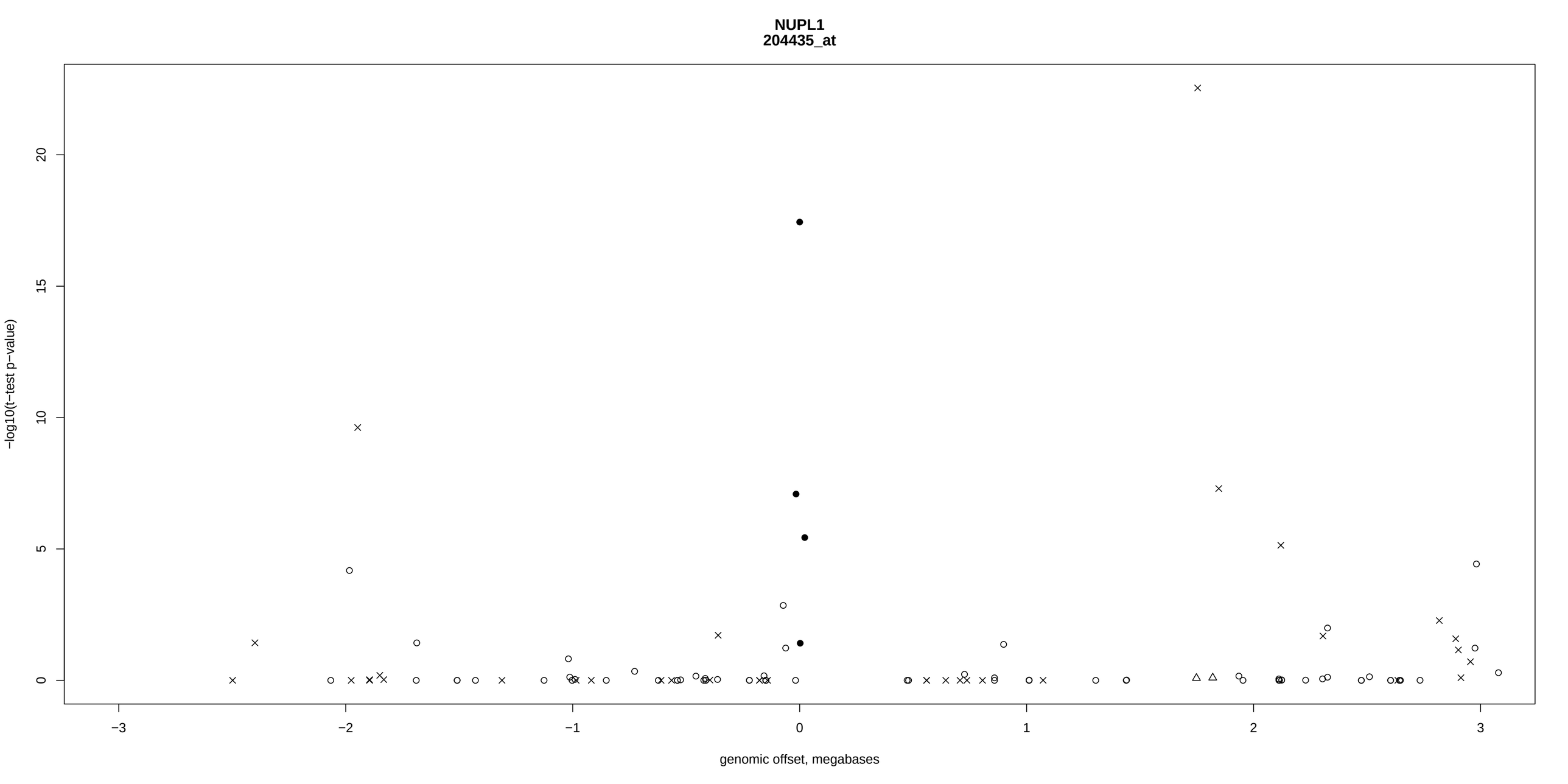
3
genomic offset, megabases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204435_at
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−1
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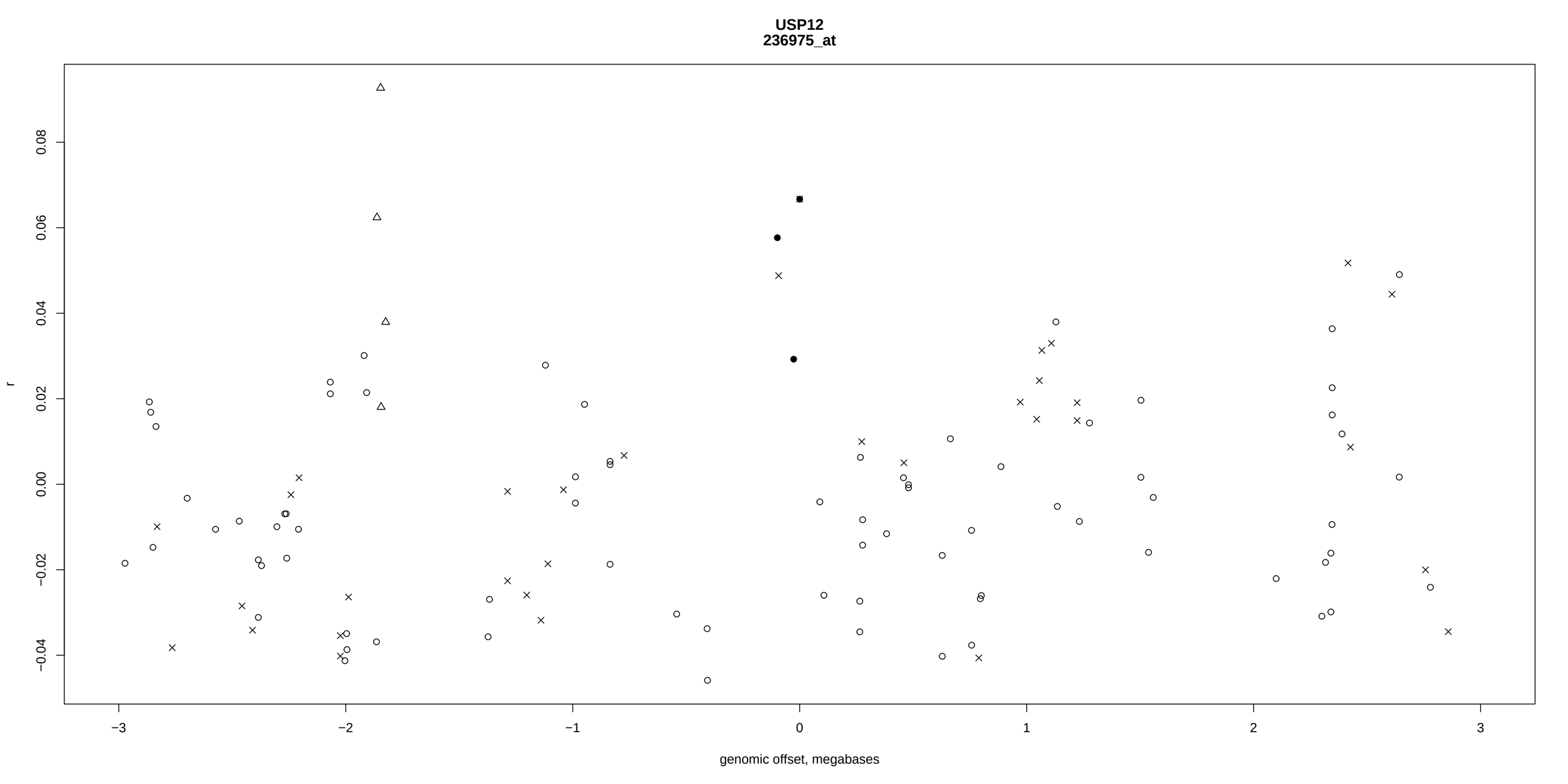
2
3
genomic offset, megabases


USP12
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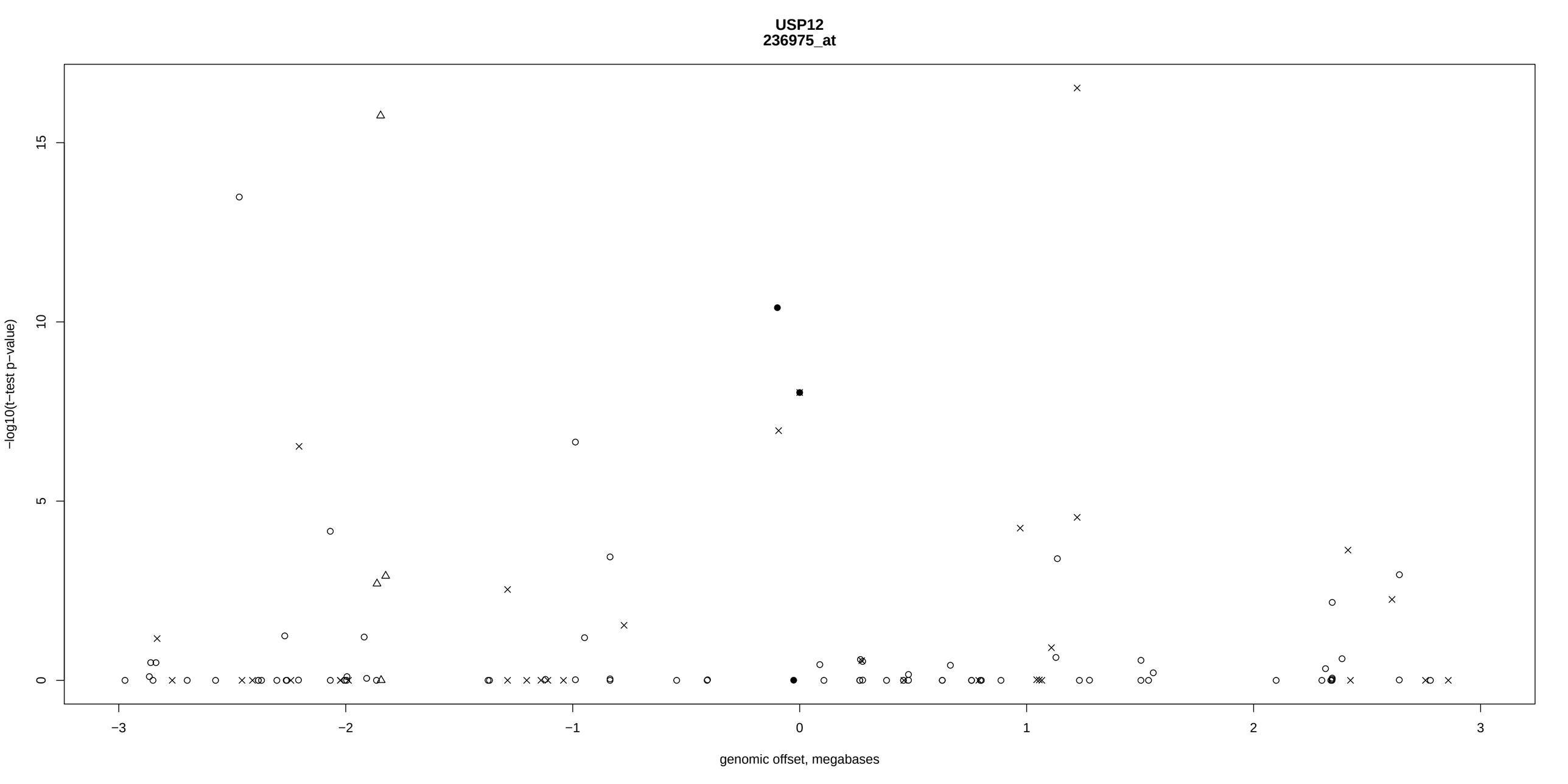
3
genomic offset, megabases


USP12
236975_at
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−2
−1
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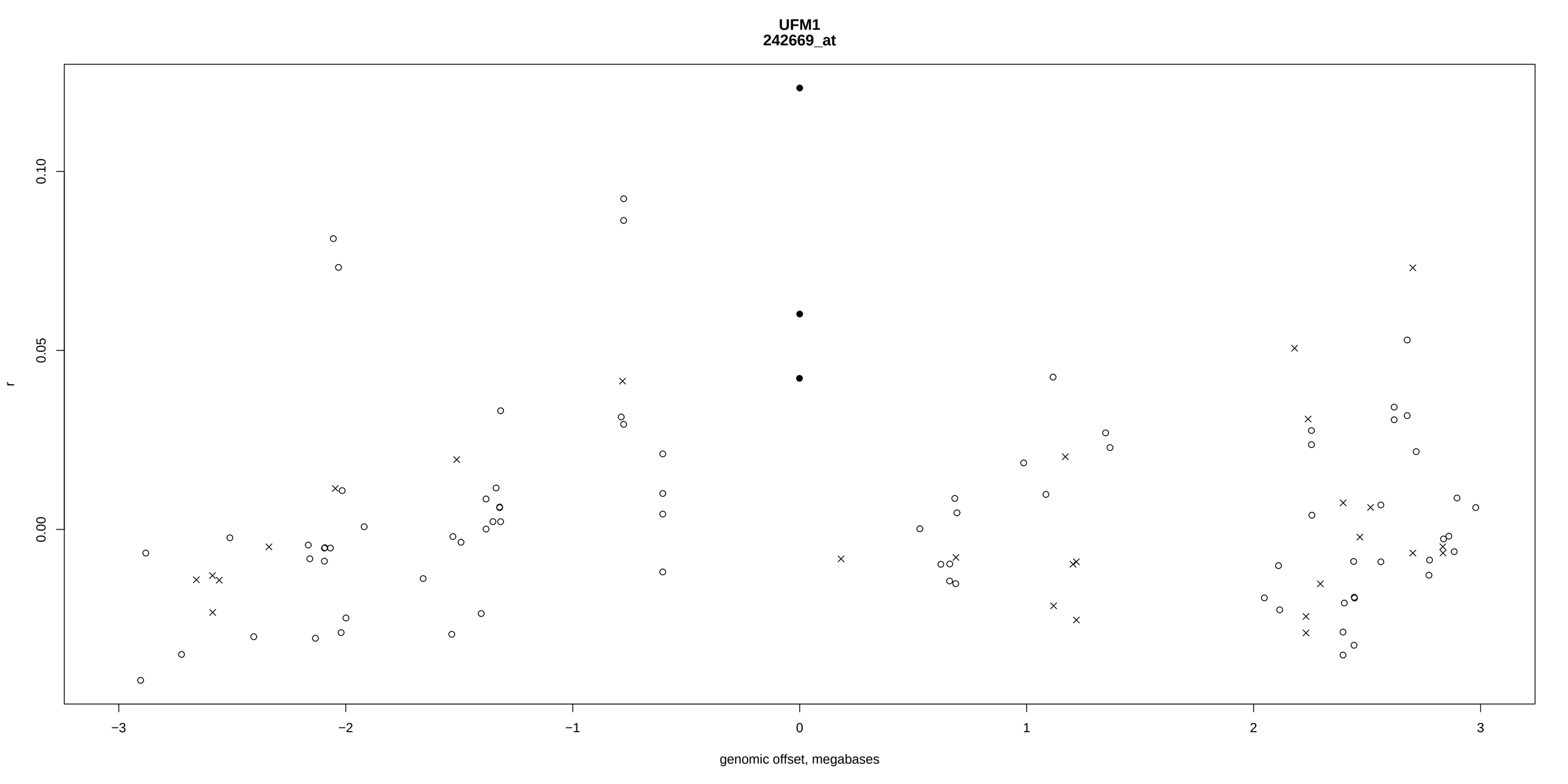
2
3
genomic offset, megabases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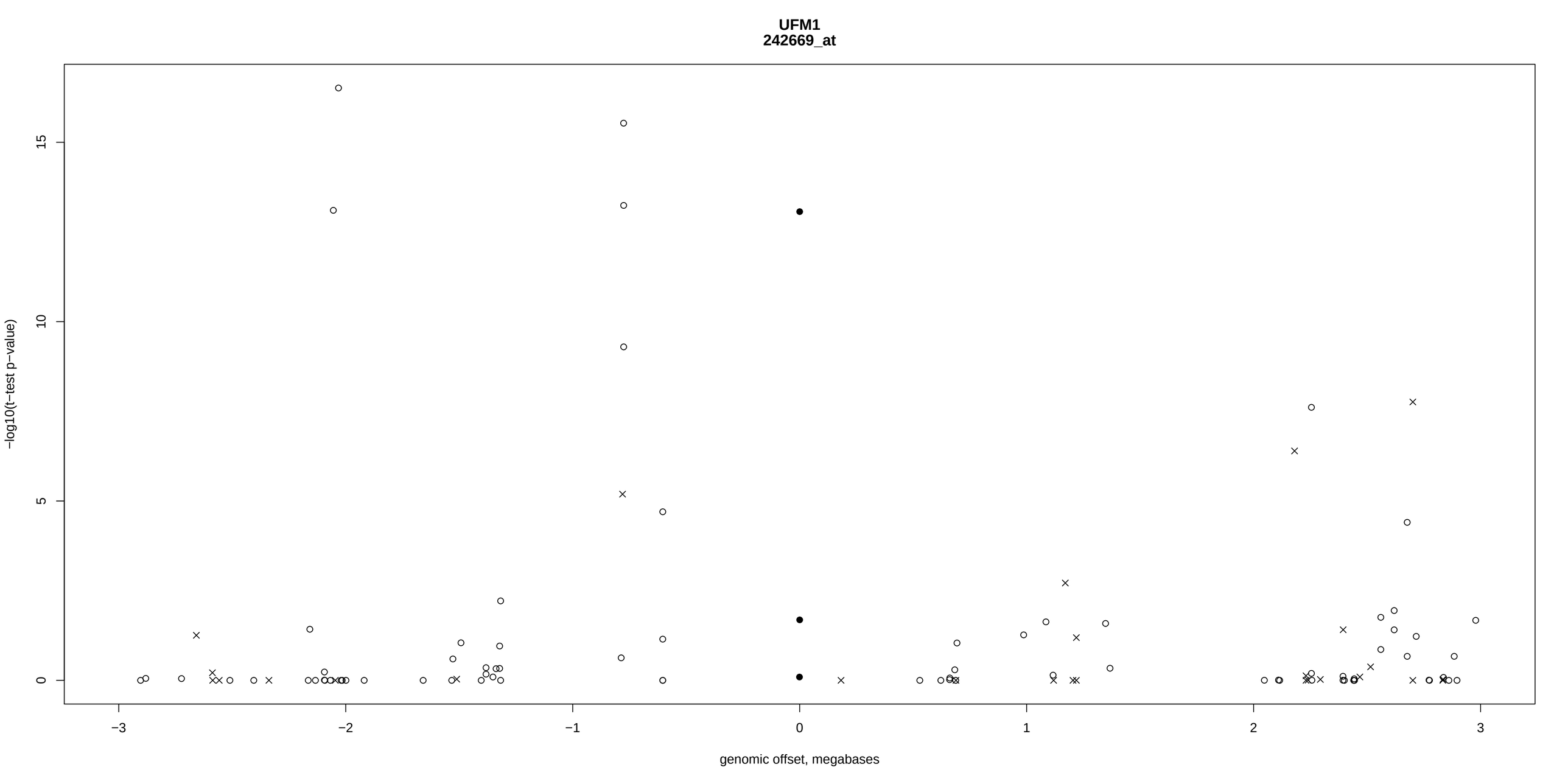
3
genomic offset, megabases


UFM1
242669_at
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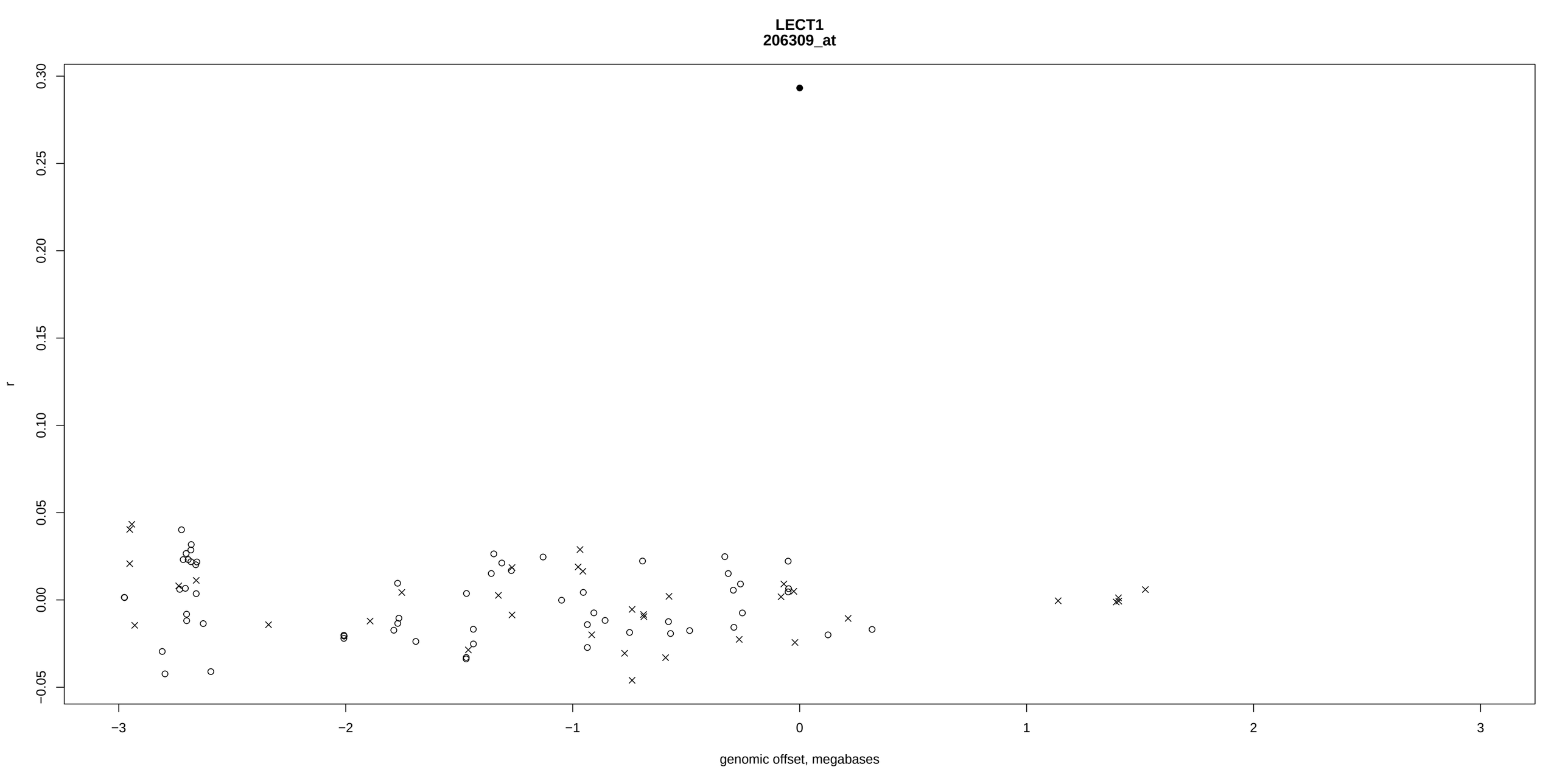
2
3
genomic offset, megabases


LECT1
206309_at
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0.00
−0.05
−3
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−1
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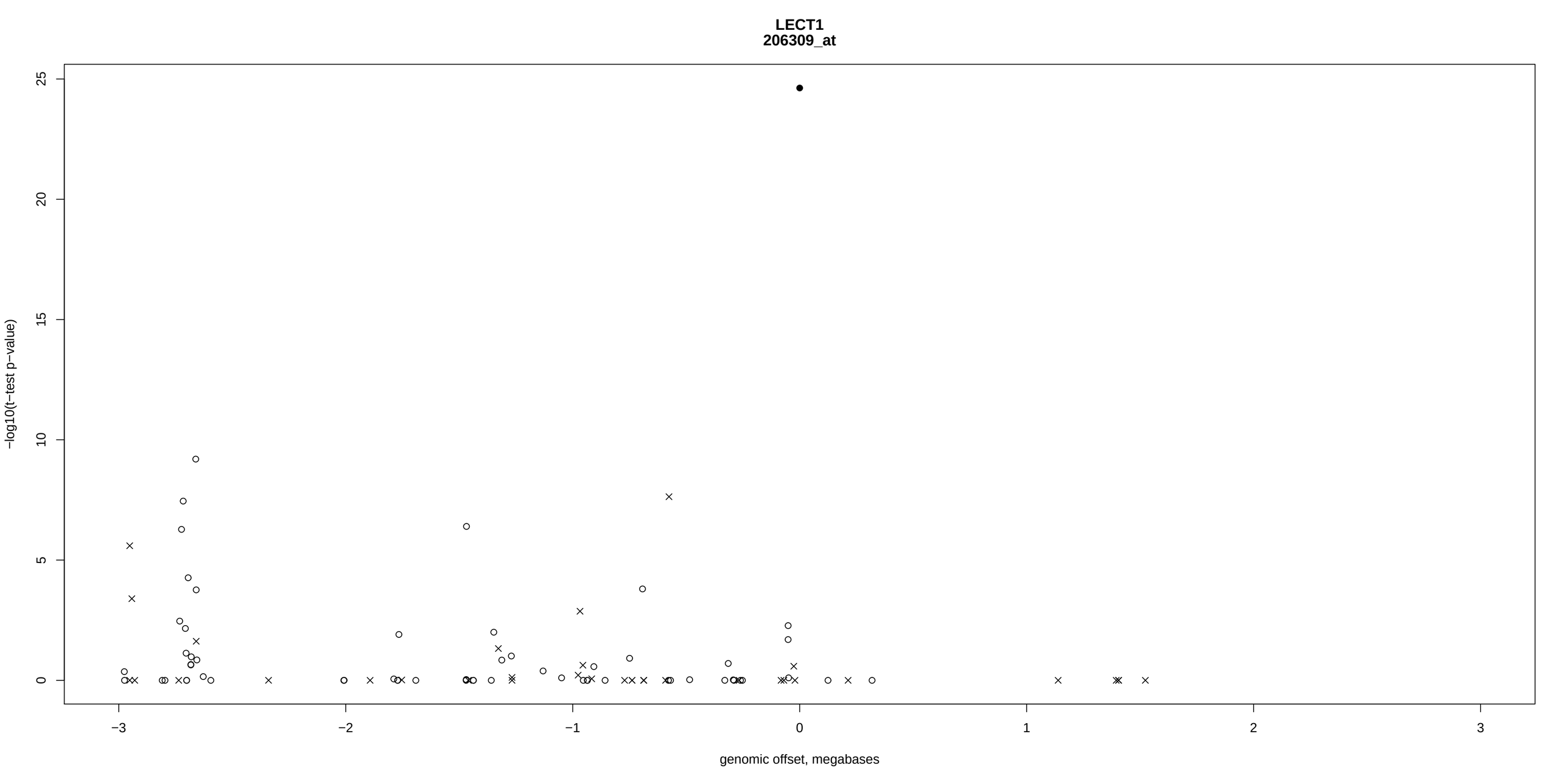
2
3
genomic offset, megabases


LECT1
206309_at
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−3
−2
−1
0
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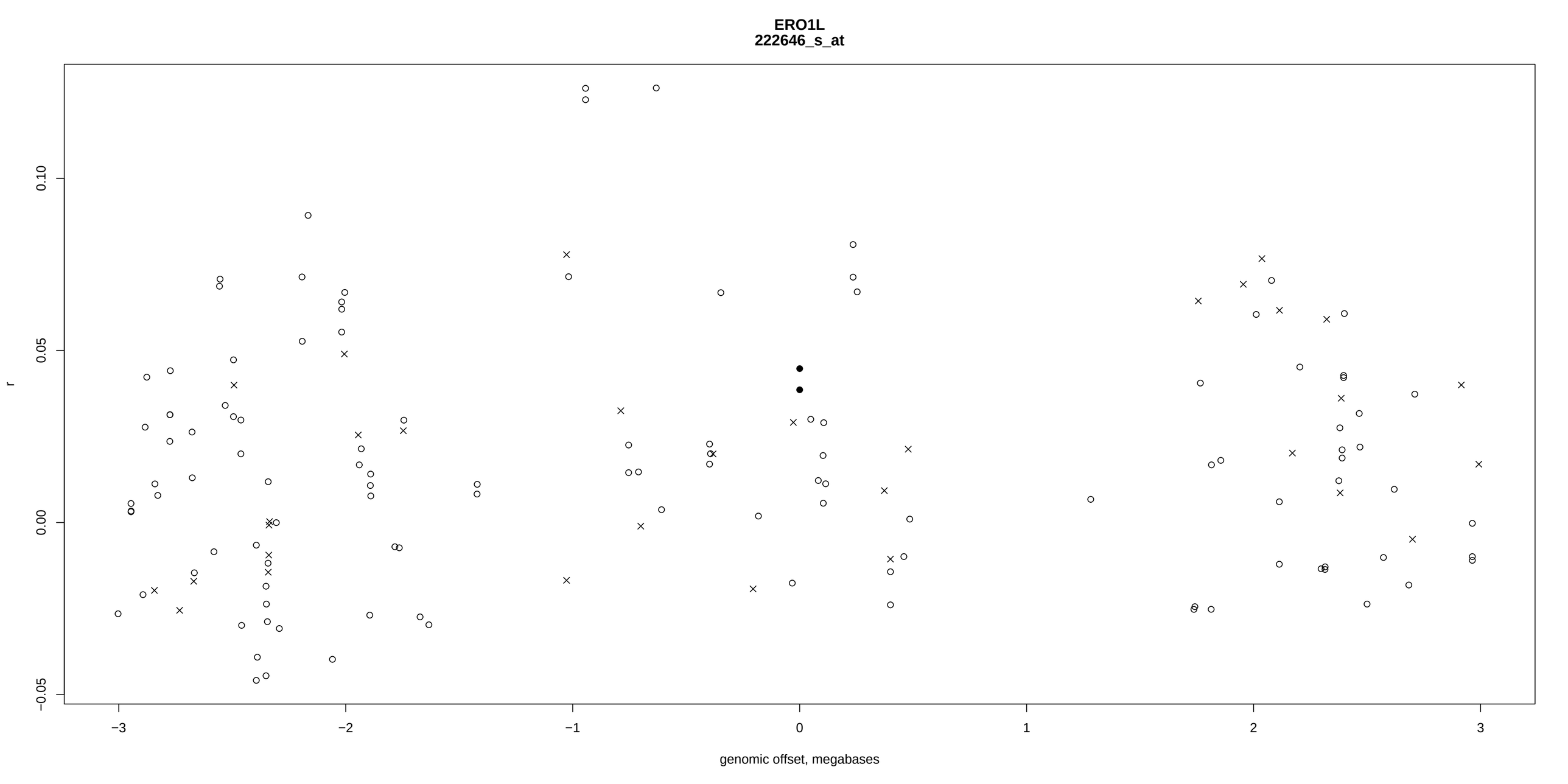
2
3
genomic offset, megabases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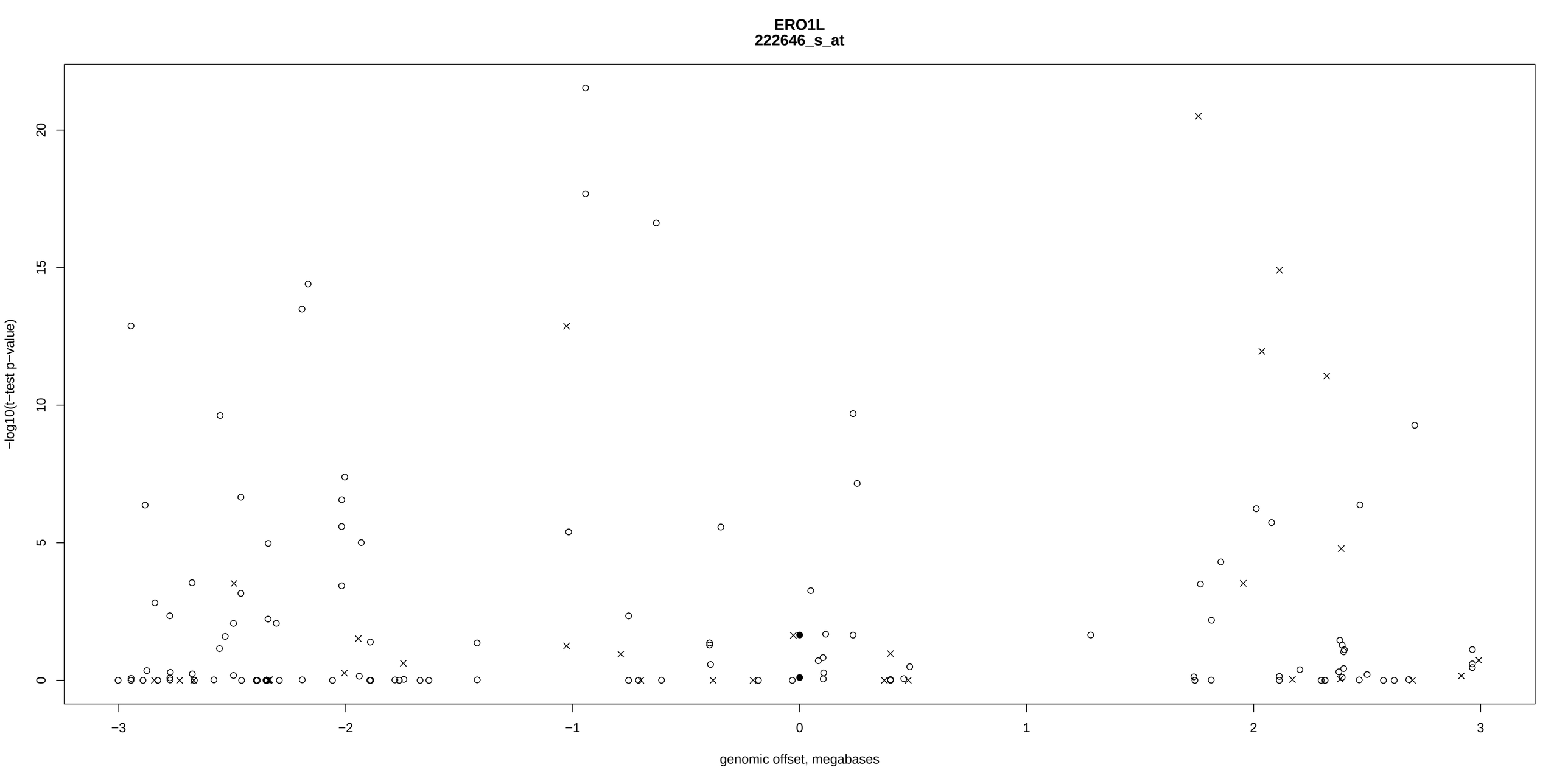
3
genomic offset, megabases


ERO1L
222646_s_at
−log10(t−test p−value)
0
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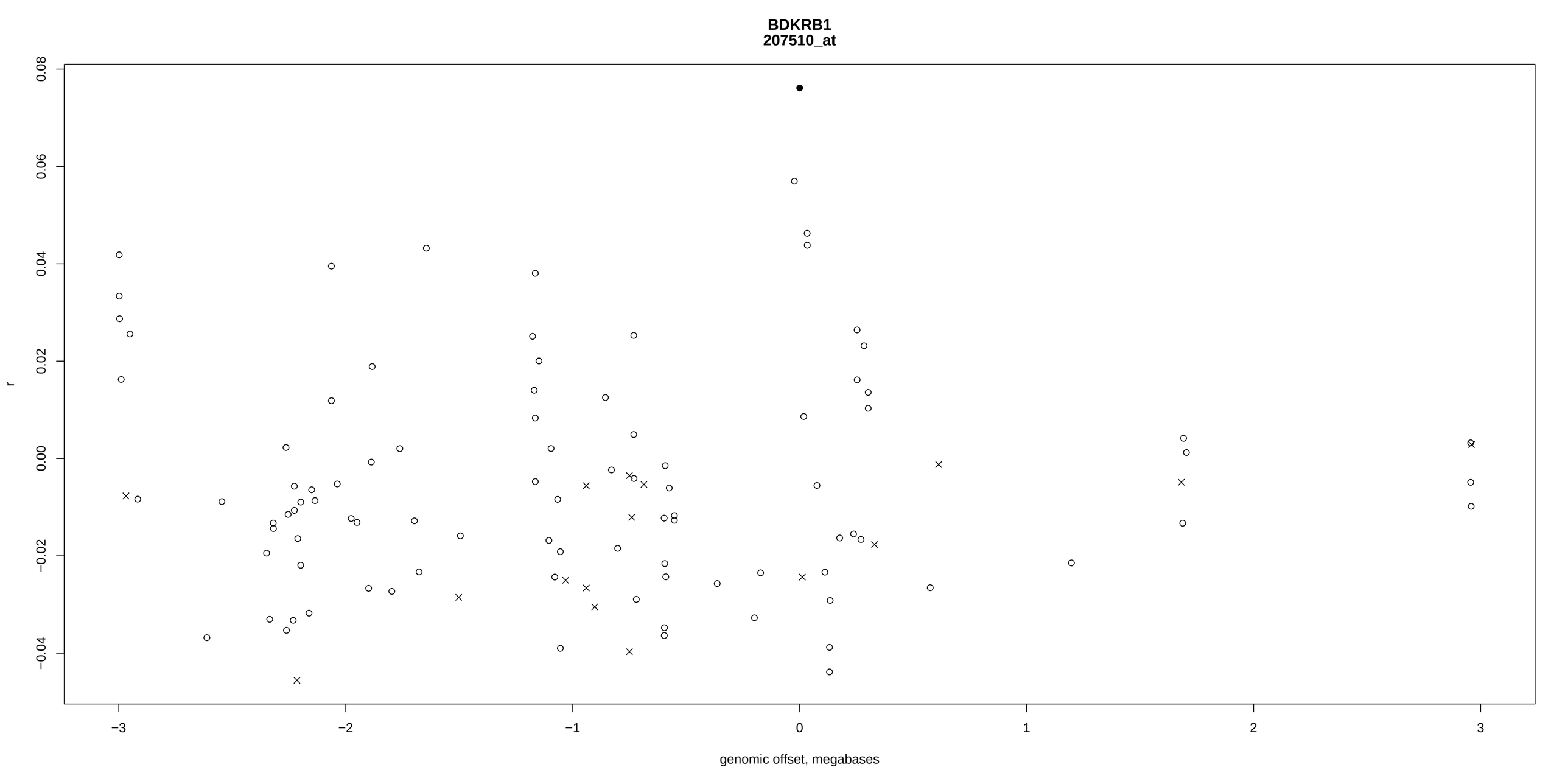
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


BDKRB1
207510_at
r
0.08
0.06
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0.00
−0.02
−0.04
−3
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−1
0
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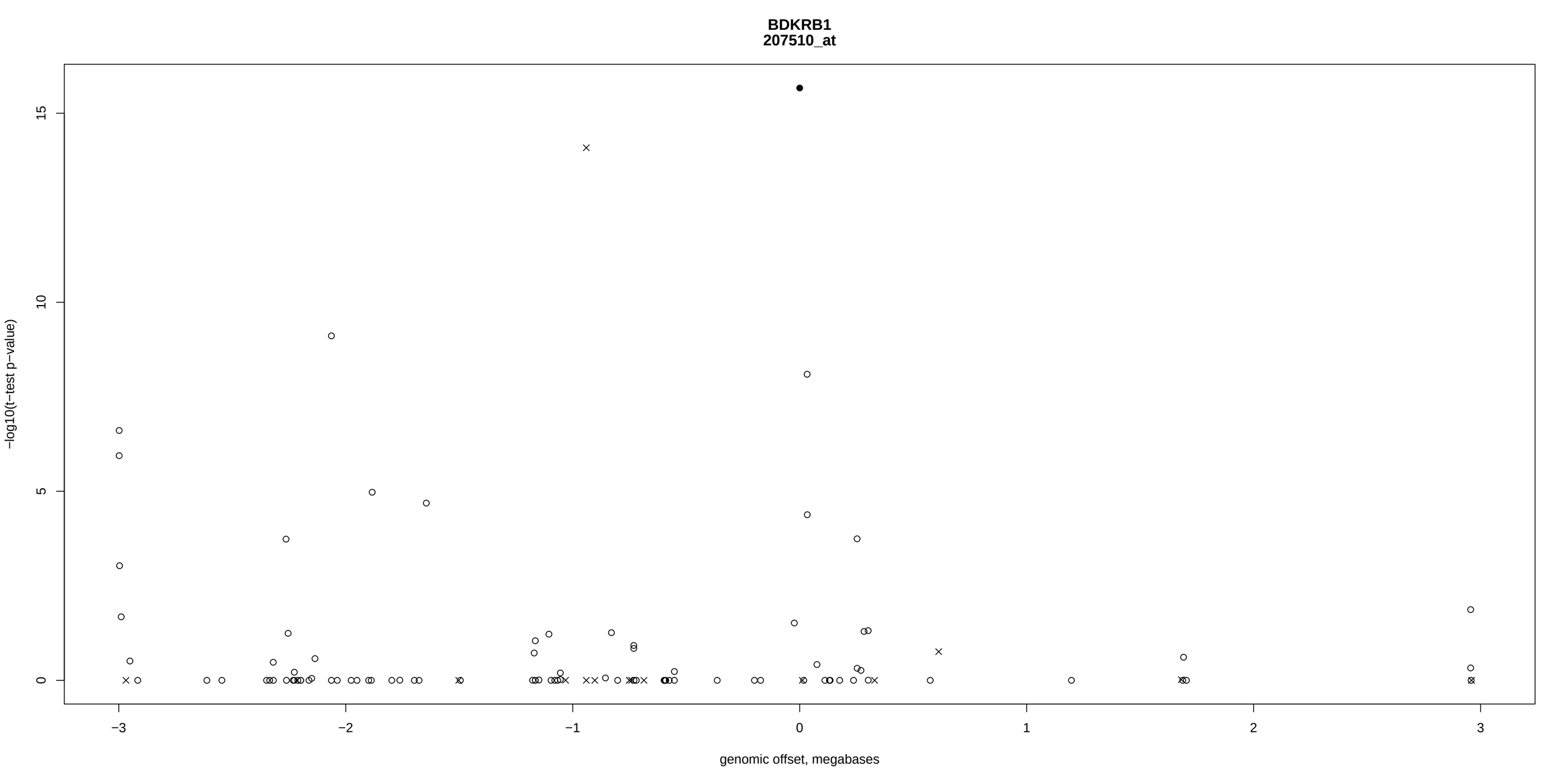
3
genomic offset, megabases


BDKRB1
207510_at
−log10(t−test p−value)
0
5
10
15
−3
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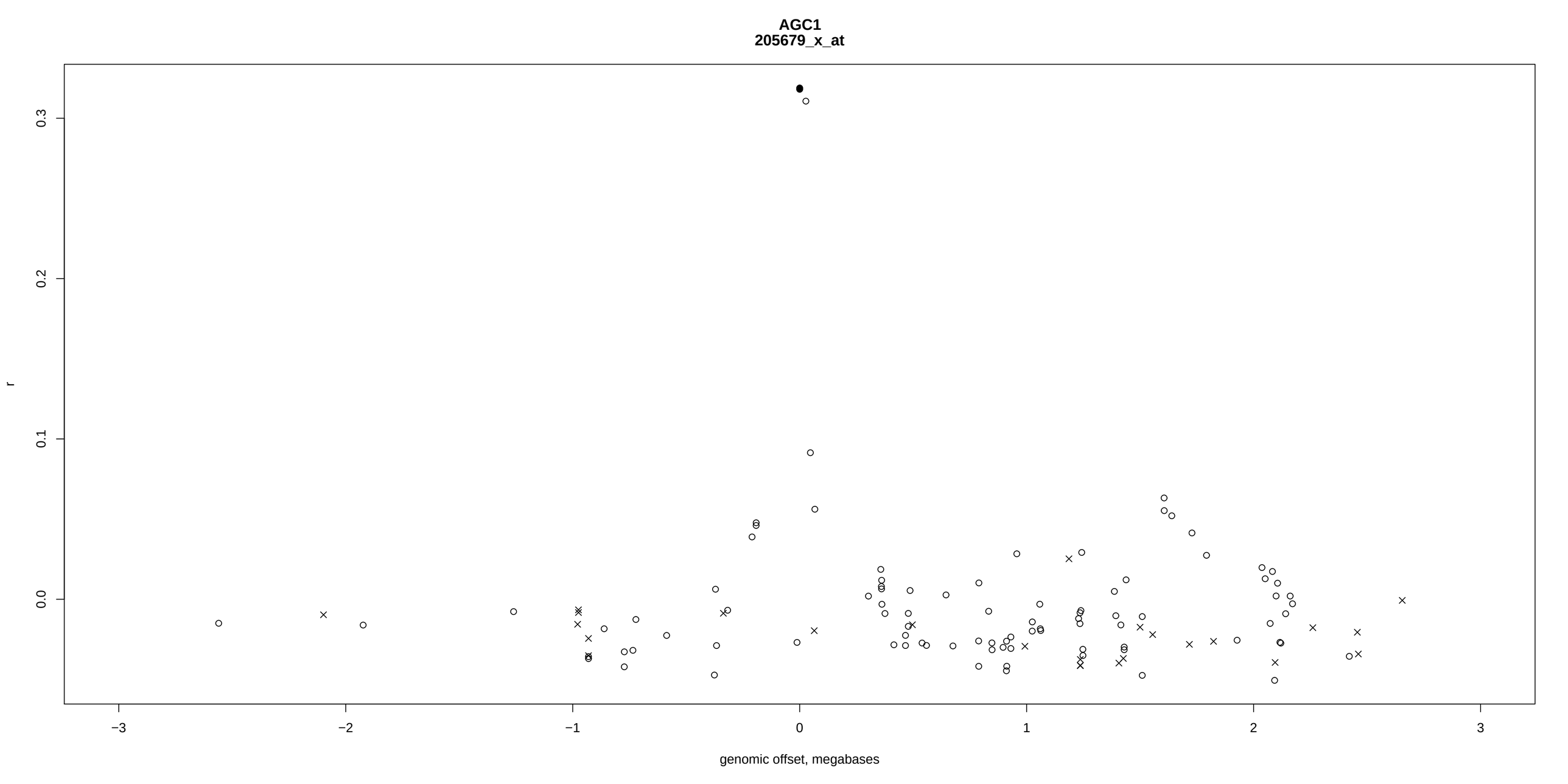
−1
0
1
2
3
genomic offset, megabases


AGC1
205679_x_at
r
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0.2
0.3
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−1
0
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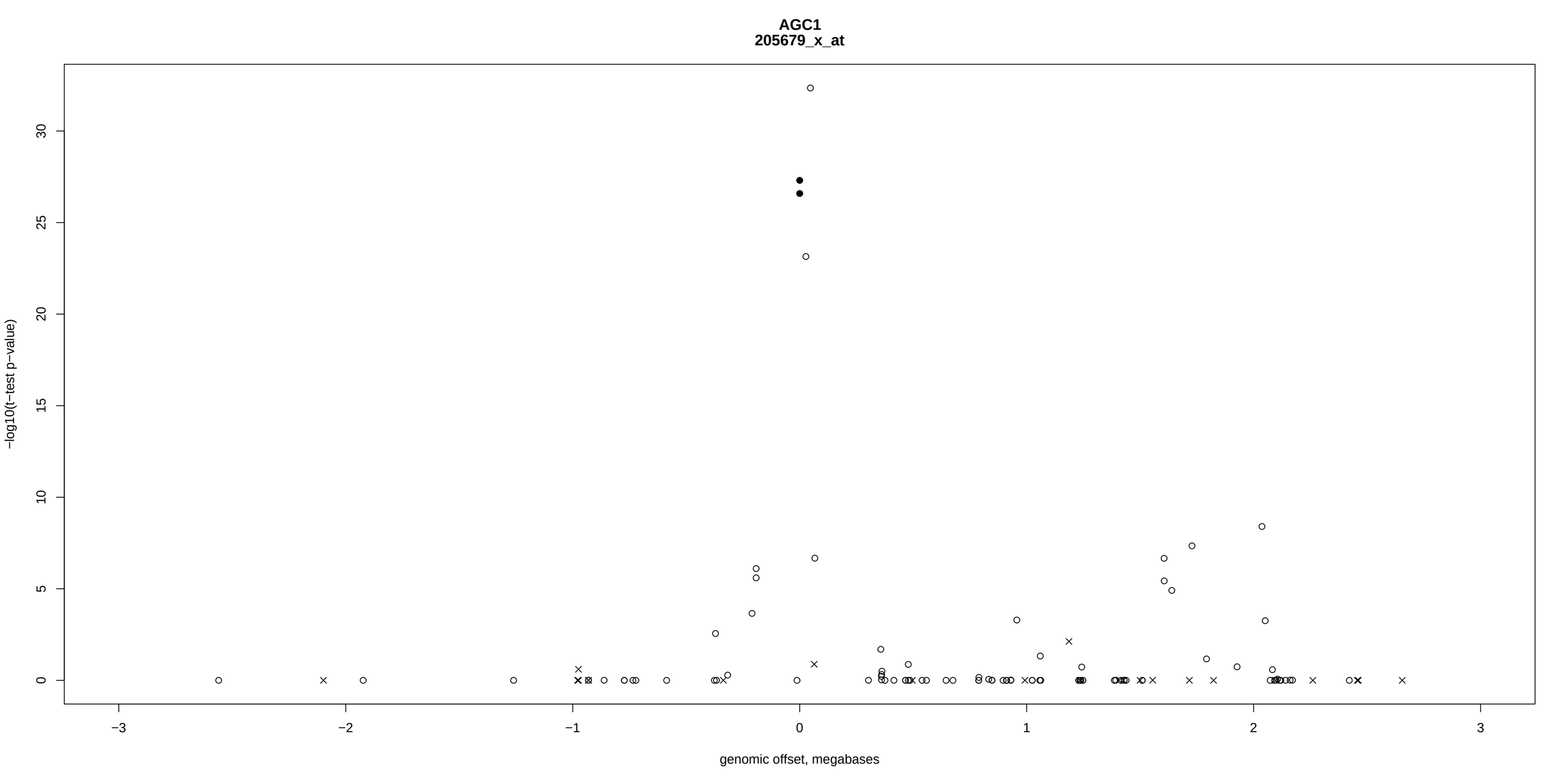
3
genomic offset, megabases


AGC1
205679_x_at
−log10(t−test p−value)
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15
20
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30
−3
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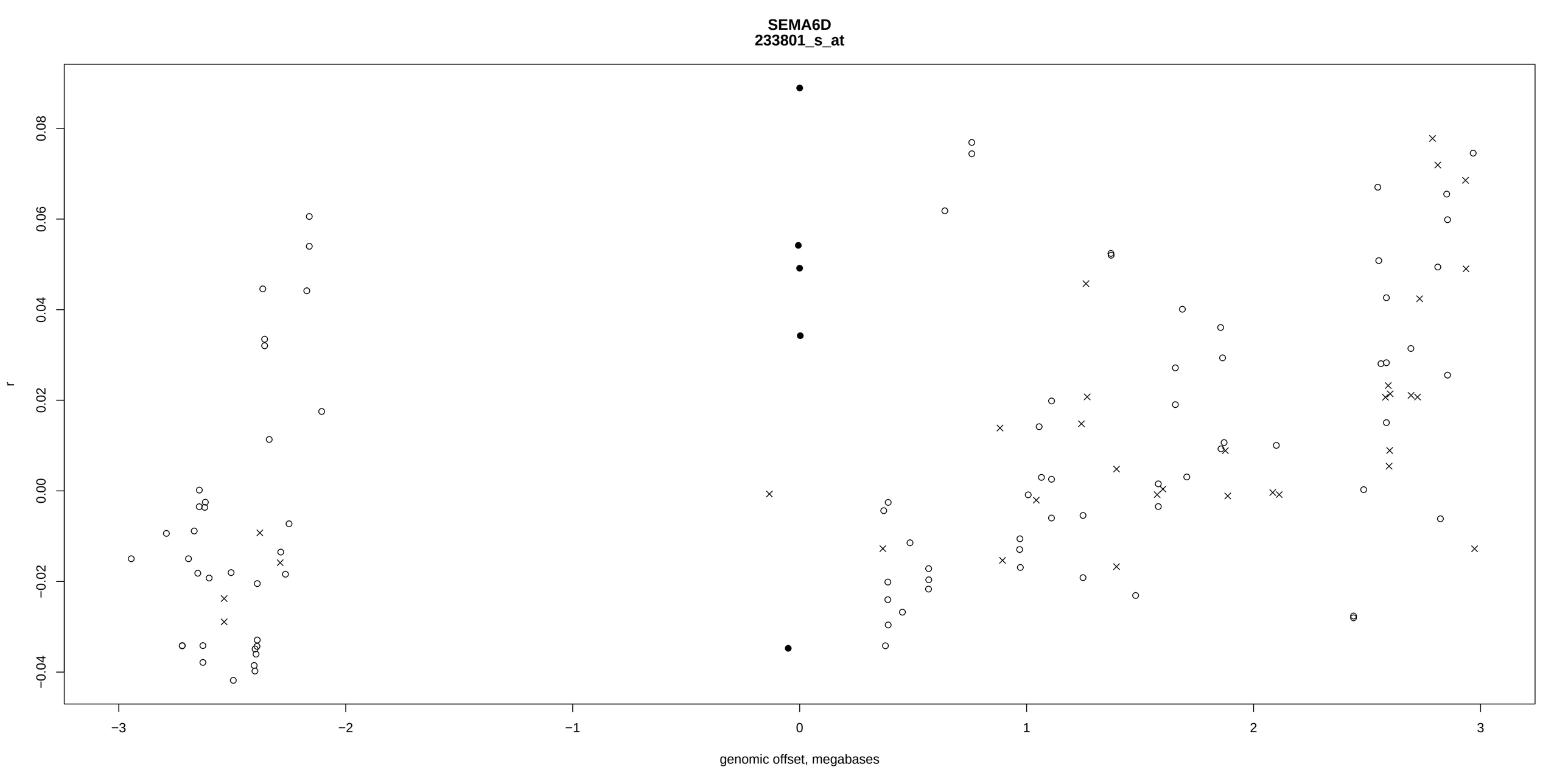
−1
0
1
2
3
genomic offset, megabases


SEMA6D
233801_s_at
r
0.08
0.06
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−0.04
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−1
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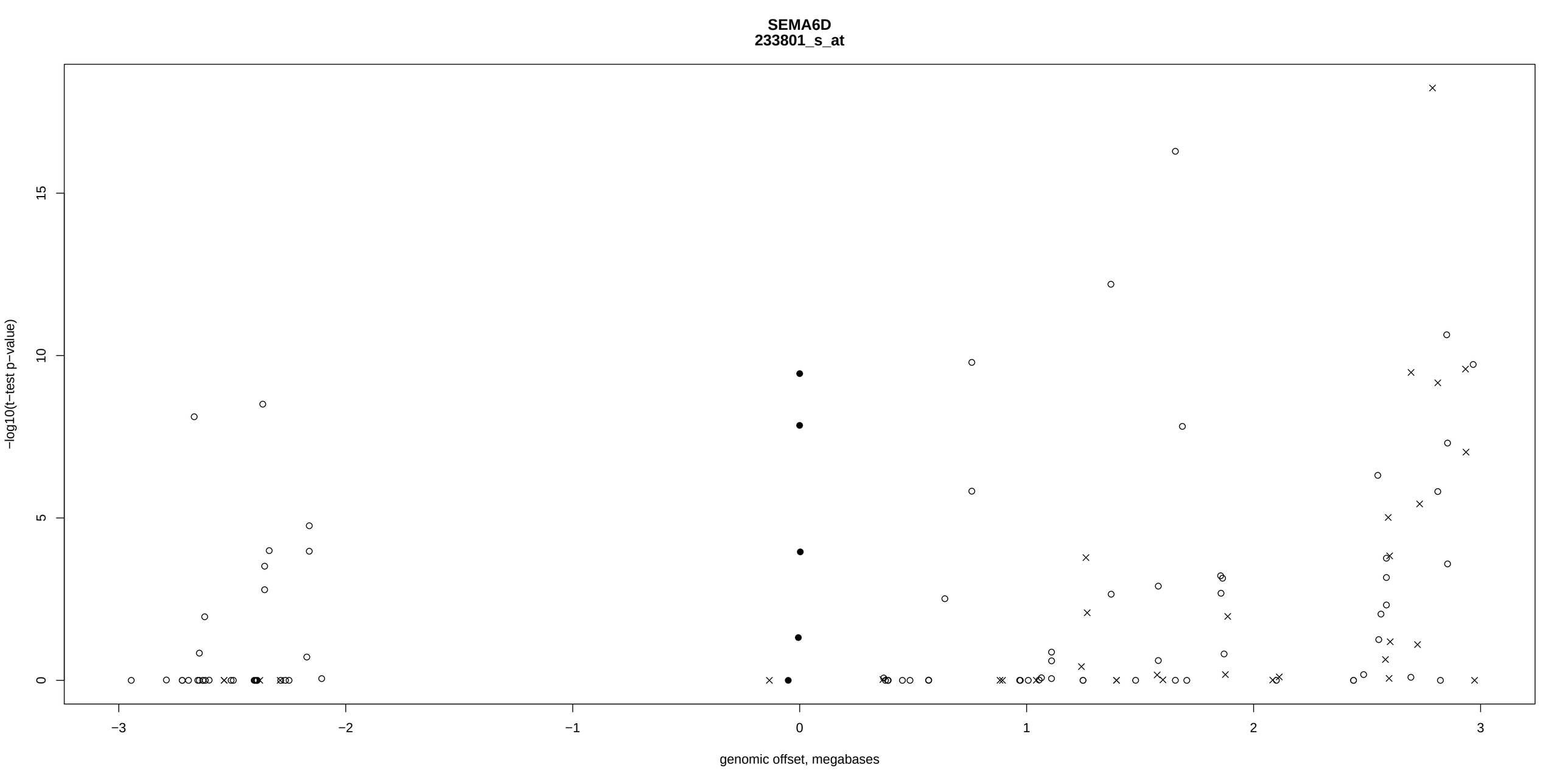
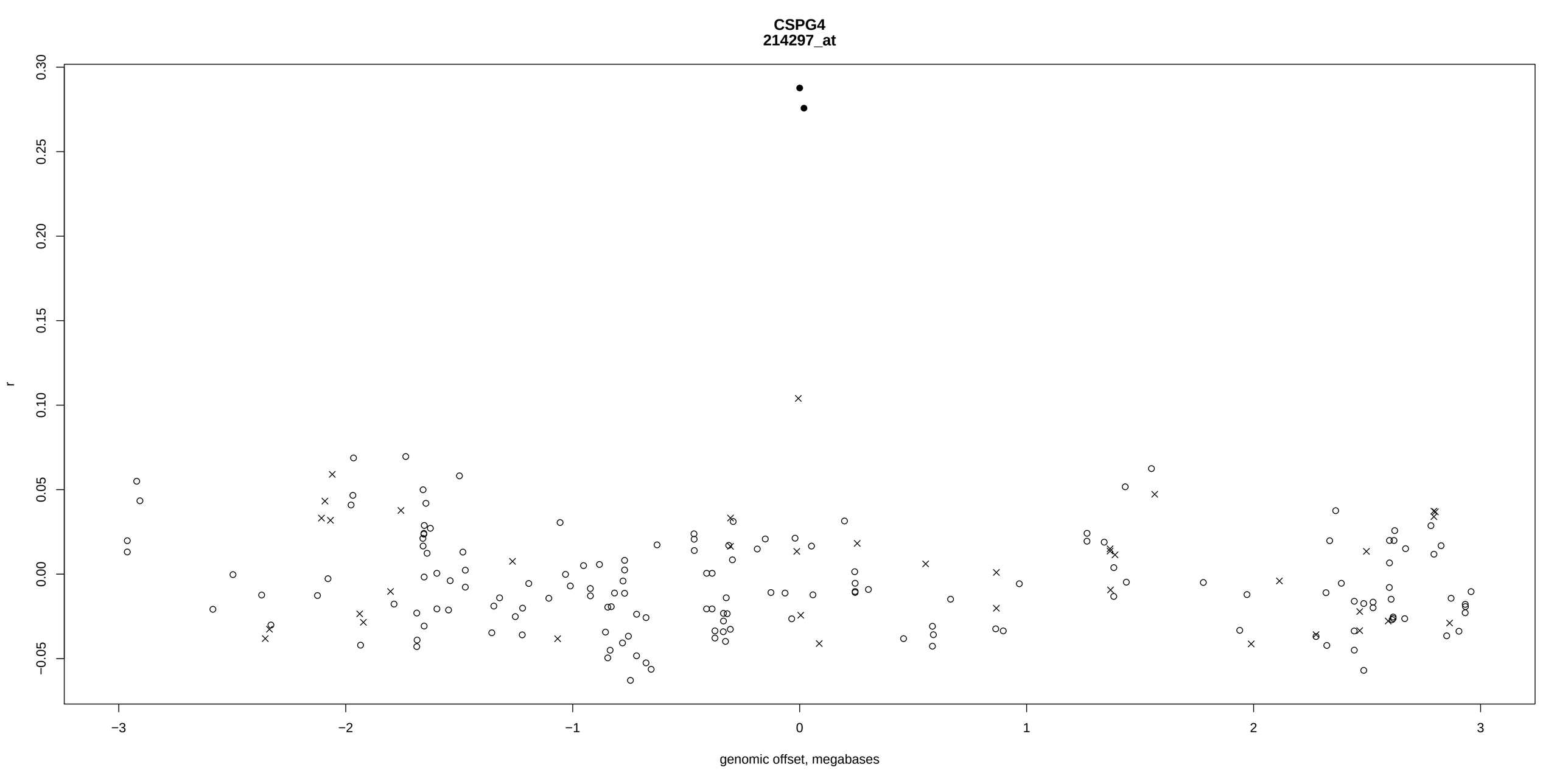
3
genomic offset, megabases


SEMA6D
233801_s_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


CSPG4
214297_at
r
0.30
0.25
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0.05
0.00
−0.05
−3
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−1
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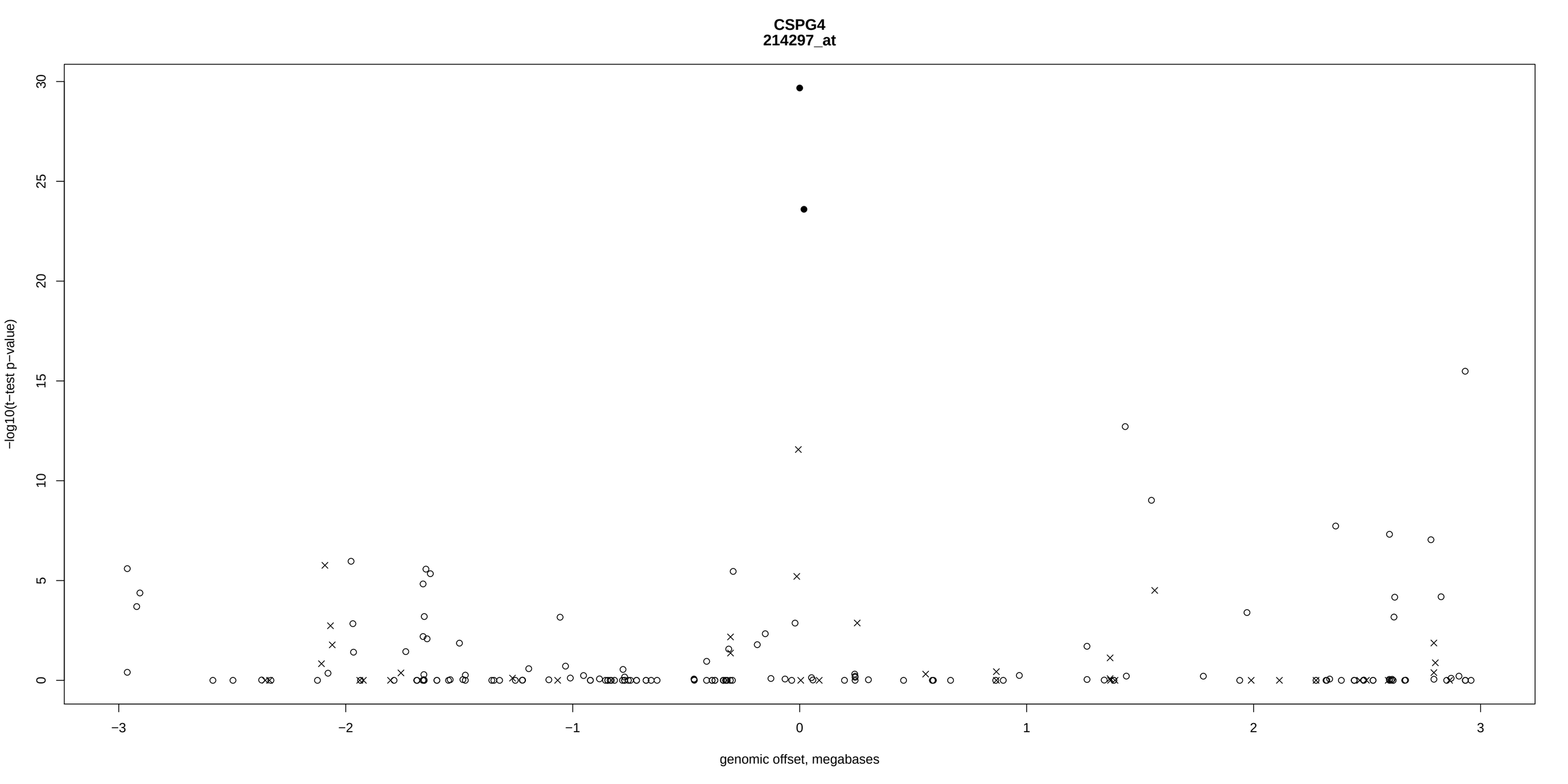
3
genomic offset, megabases


CSPG4
214297_at
−log10(t−test p−value)
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10
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20
25
30
−3
−2
−1
0
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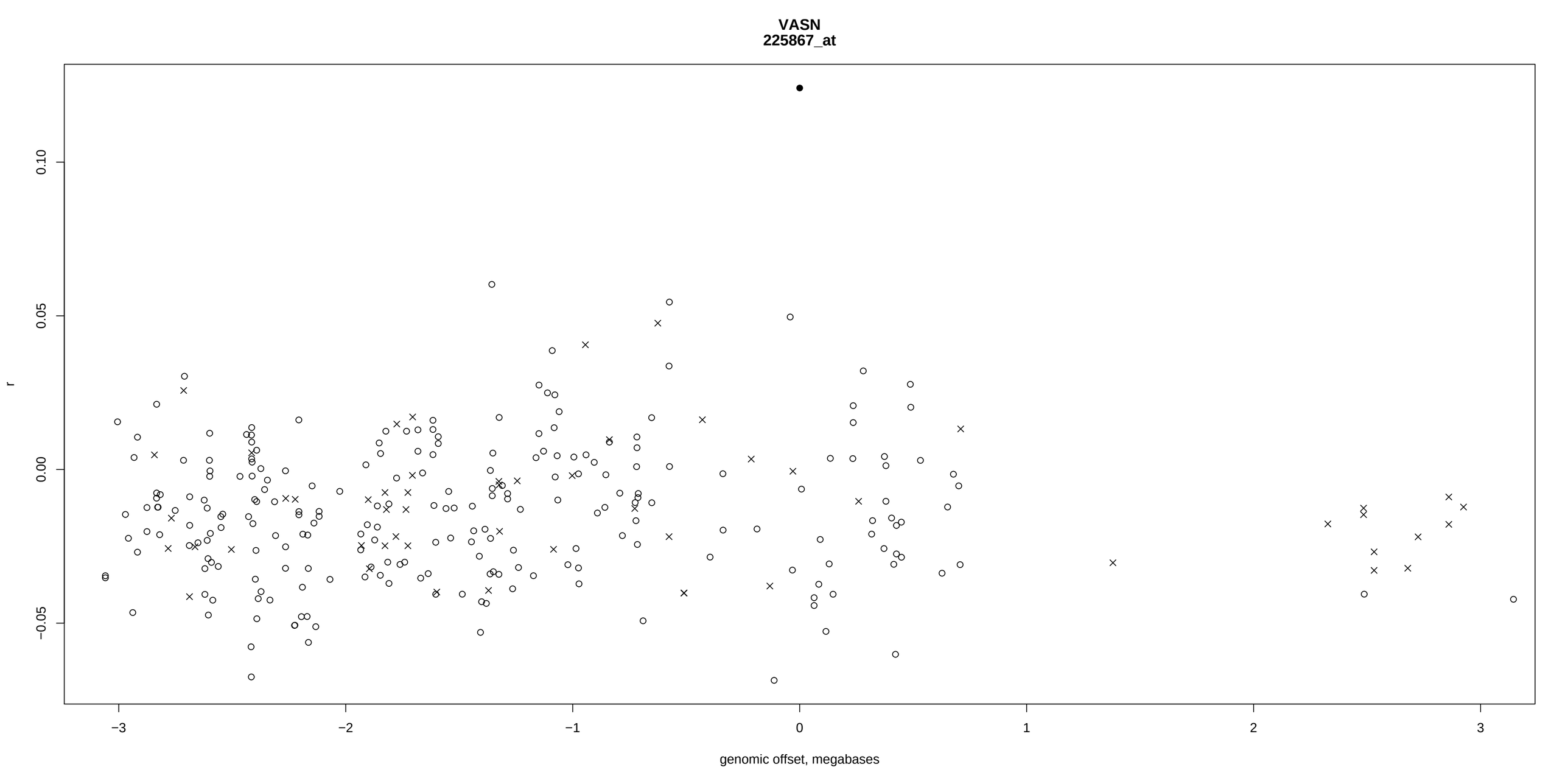
2
3
genomic offset, megabases


VASN
225867_at
r
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0.00
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−1
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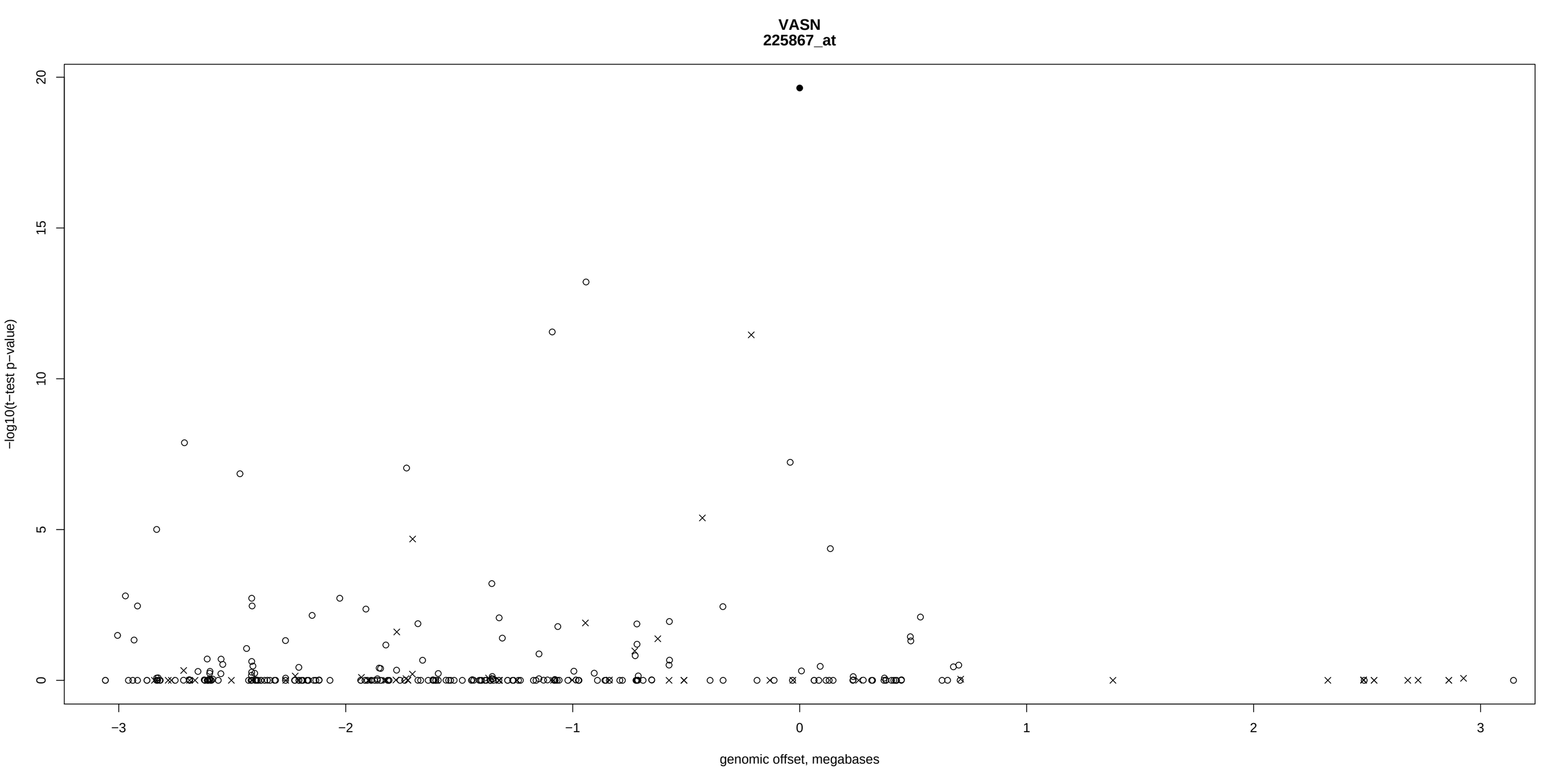
3
genomic offset, megabases


VASN
225867_at
−log10(t−test p−value)
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20
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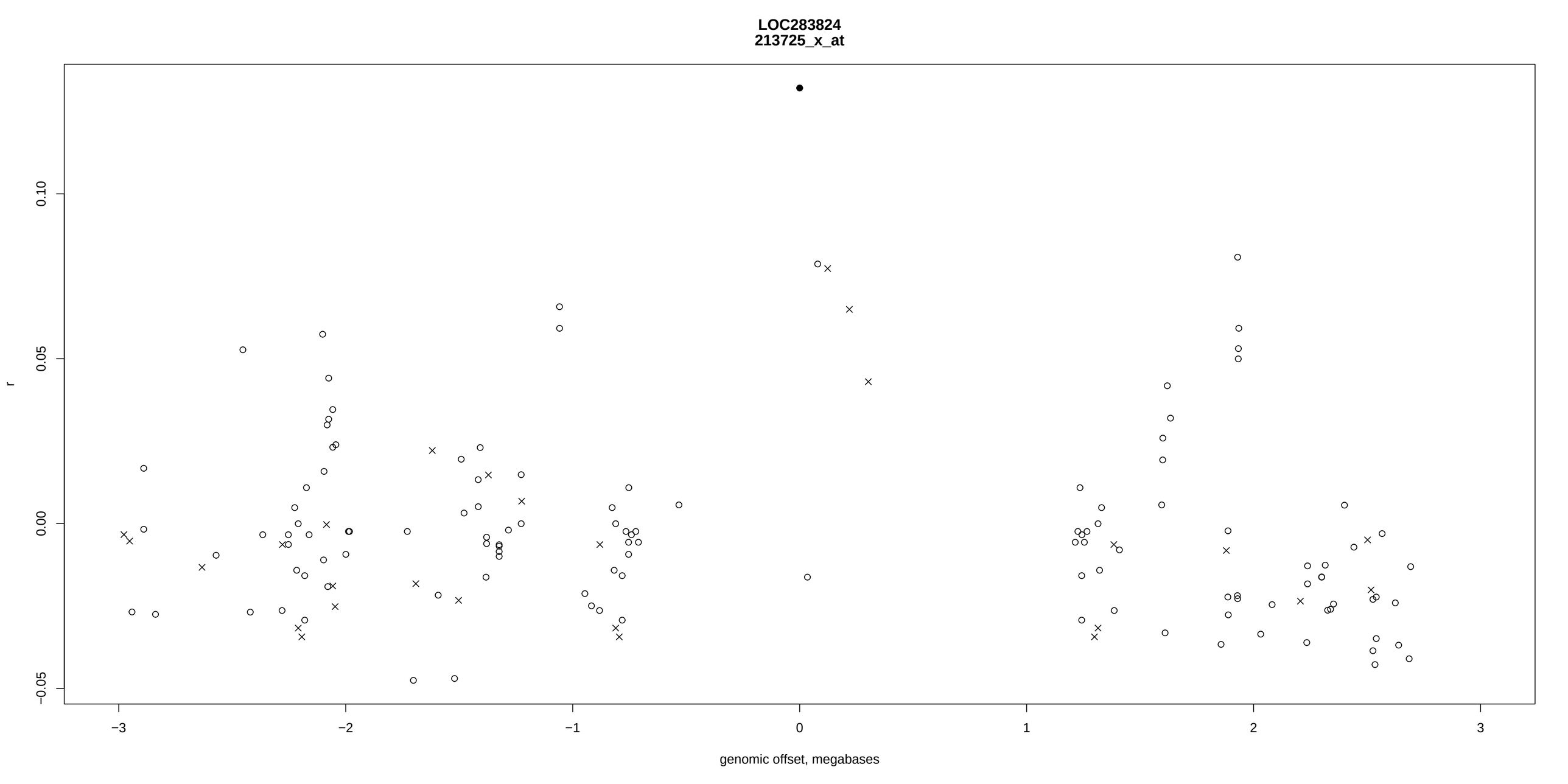
−2
−1
0
1
2
3
genomic offset, megabases


LOC283824
213725_x_at
r
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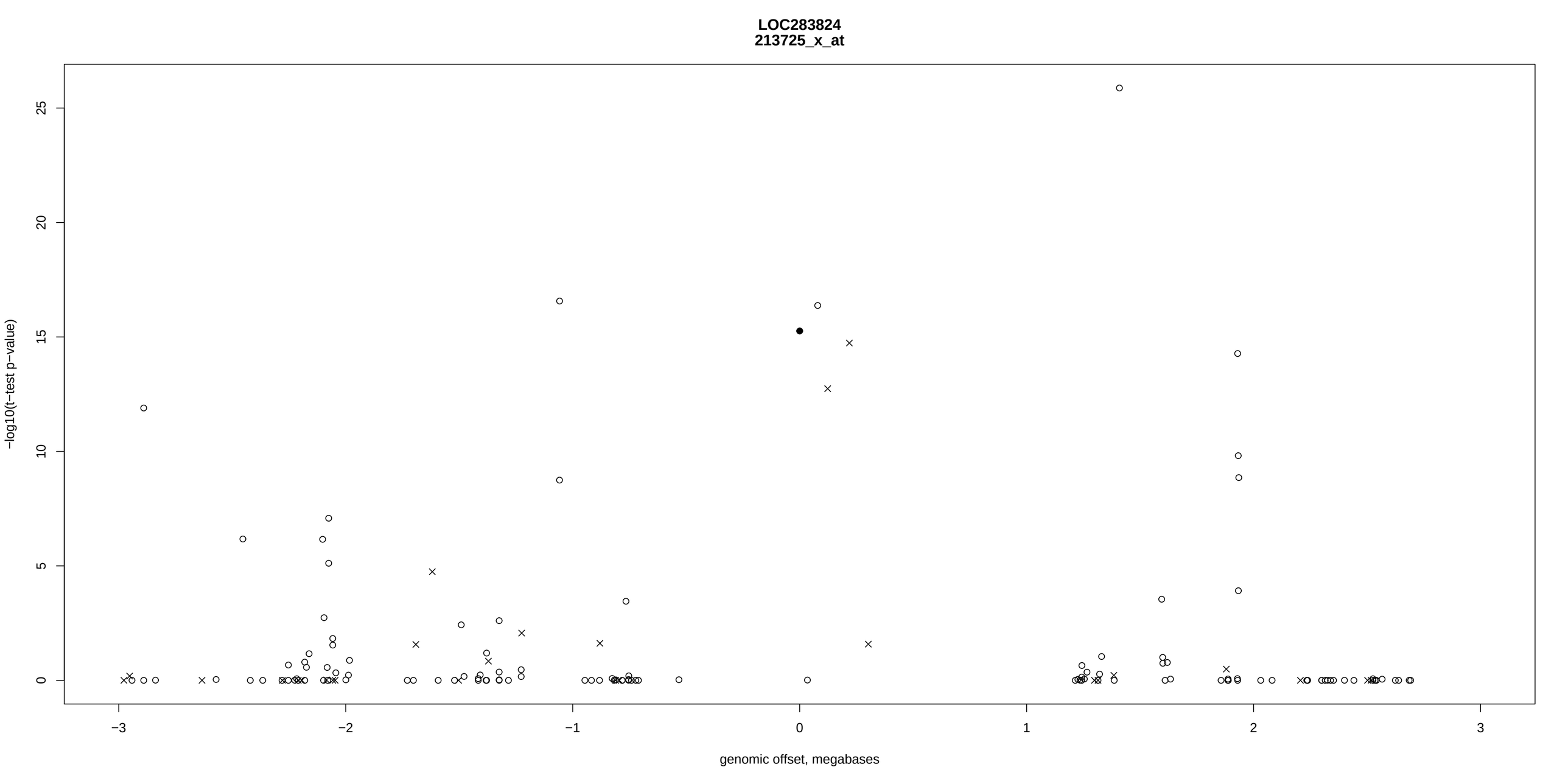
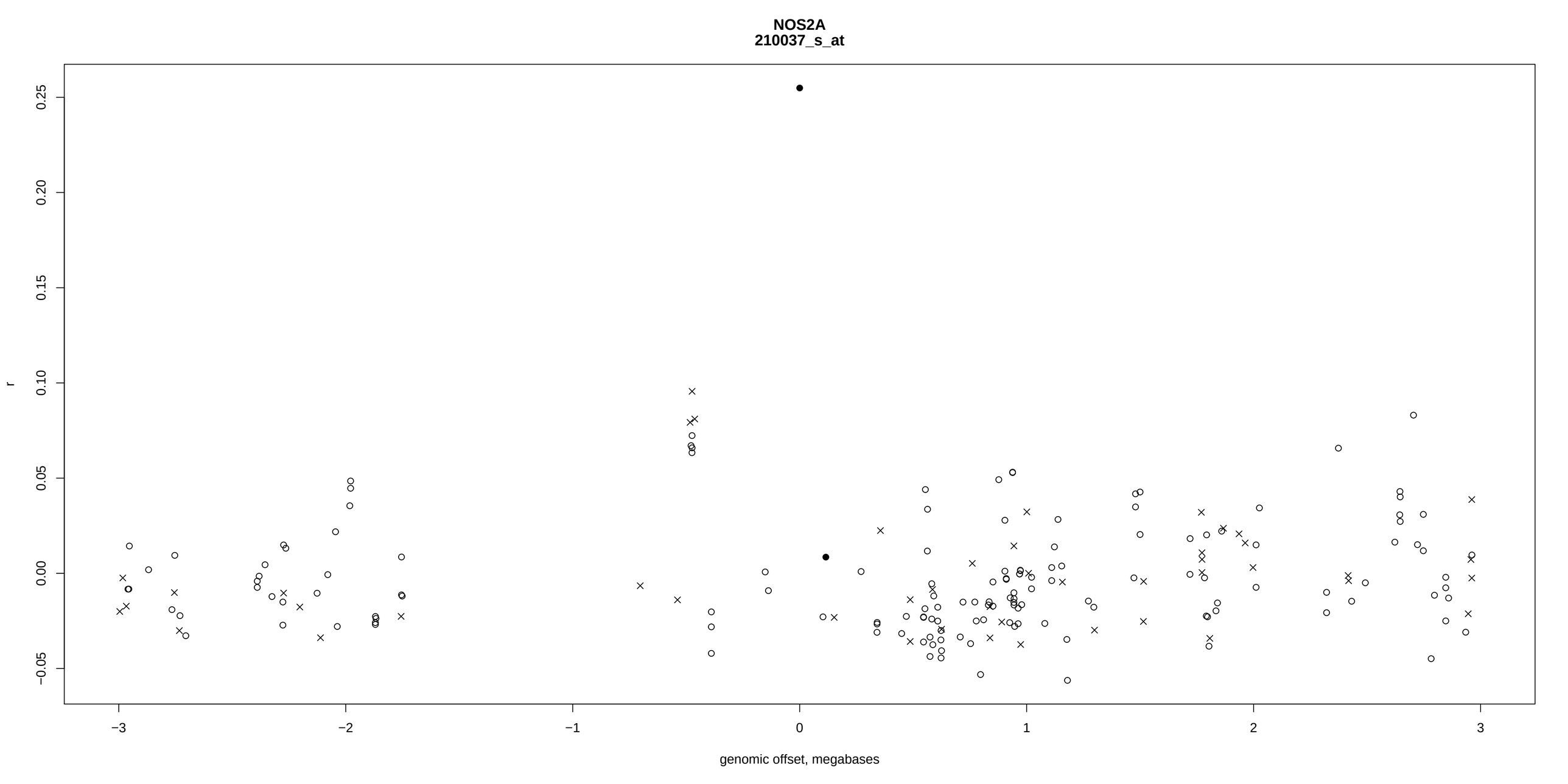
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


LOC283824
213725_x_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


NOS2A
210037_s_at
r
0.25
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0.15
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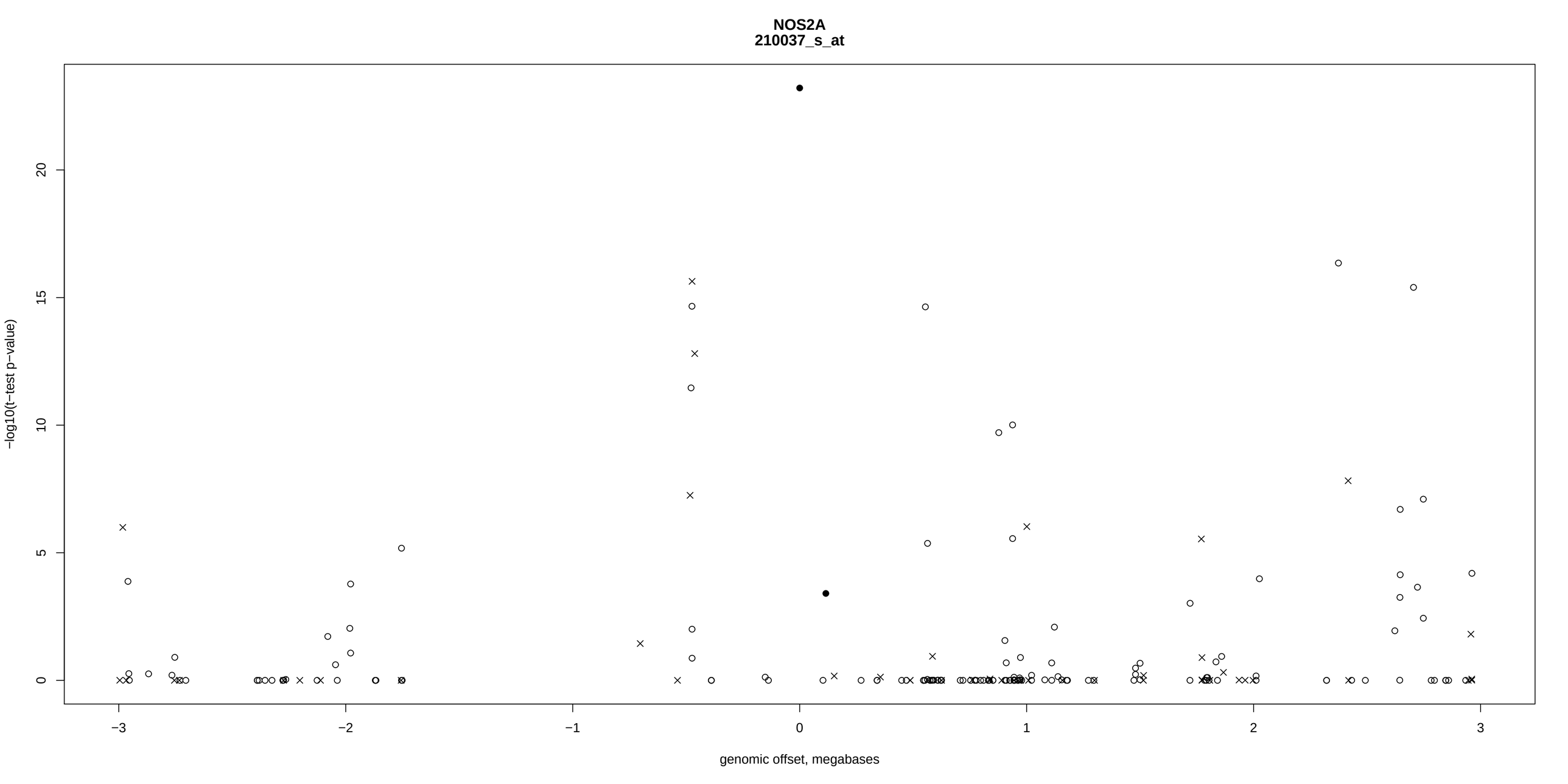
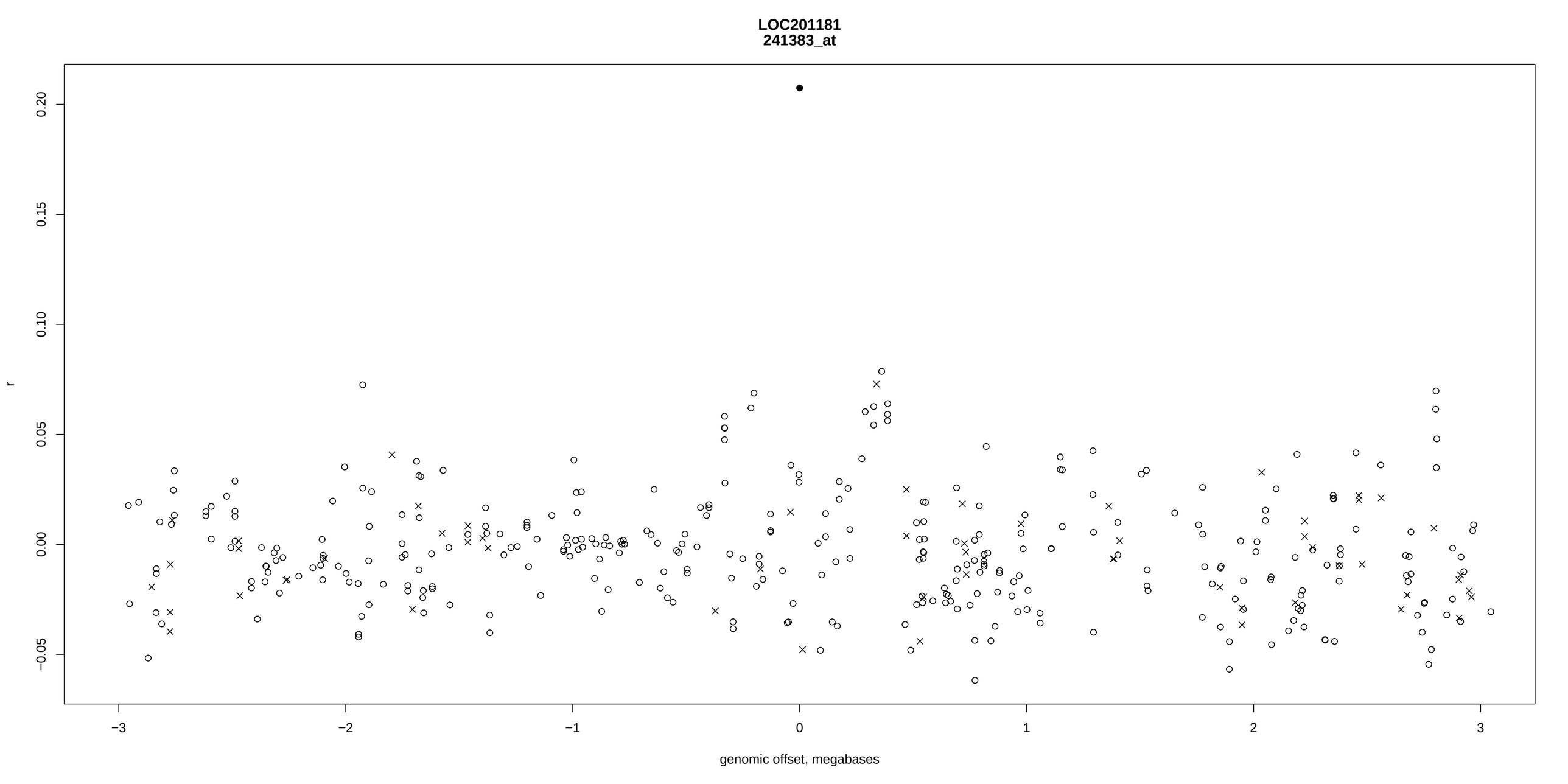
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


NOS2A
210037_s_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


LOC201181
241383_at
r
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0.15
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0.05
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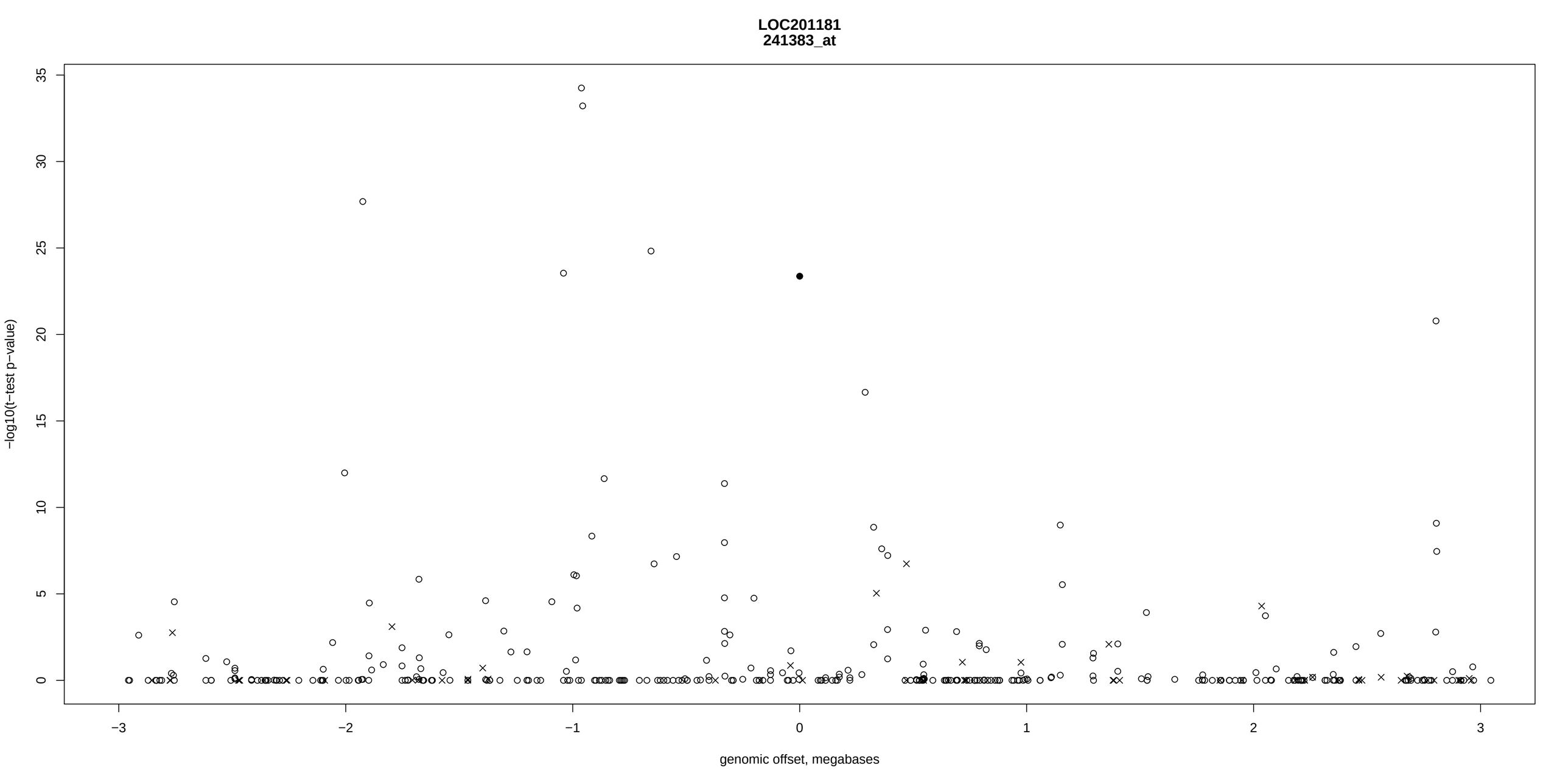
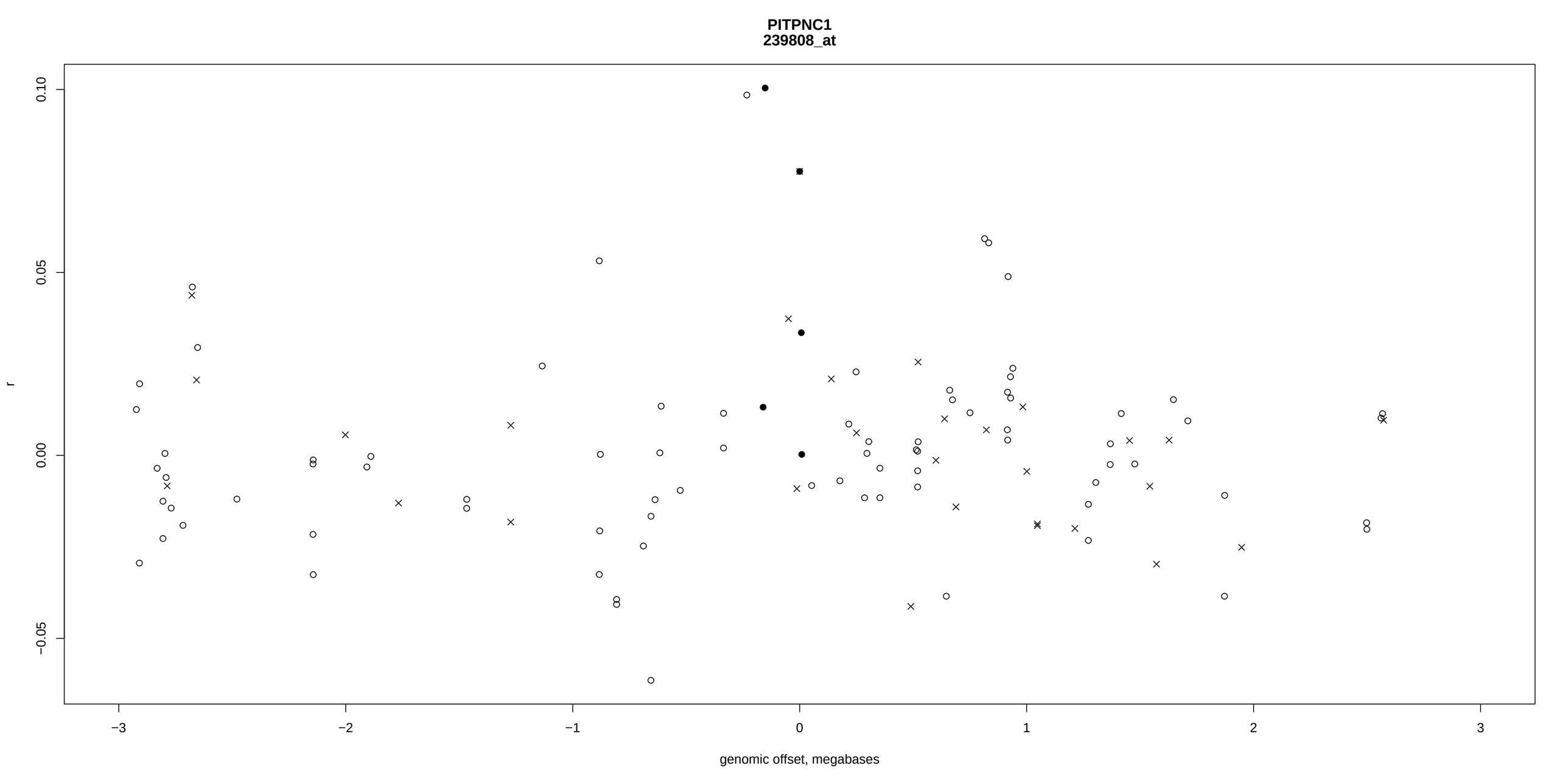
−3
−2
−1
0
1
2
3
genomic offset, megabases


LOC201181
241383_at
−log10(t−test p−value)
0
5
10
15
20
25
30
35
−3
−2
−1
0
1
2
3
genomic offset, megabases


PITPNC1
239808_at
r
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0.05
0.00
−0.05
−3
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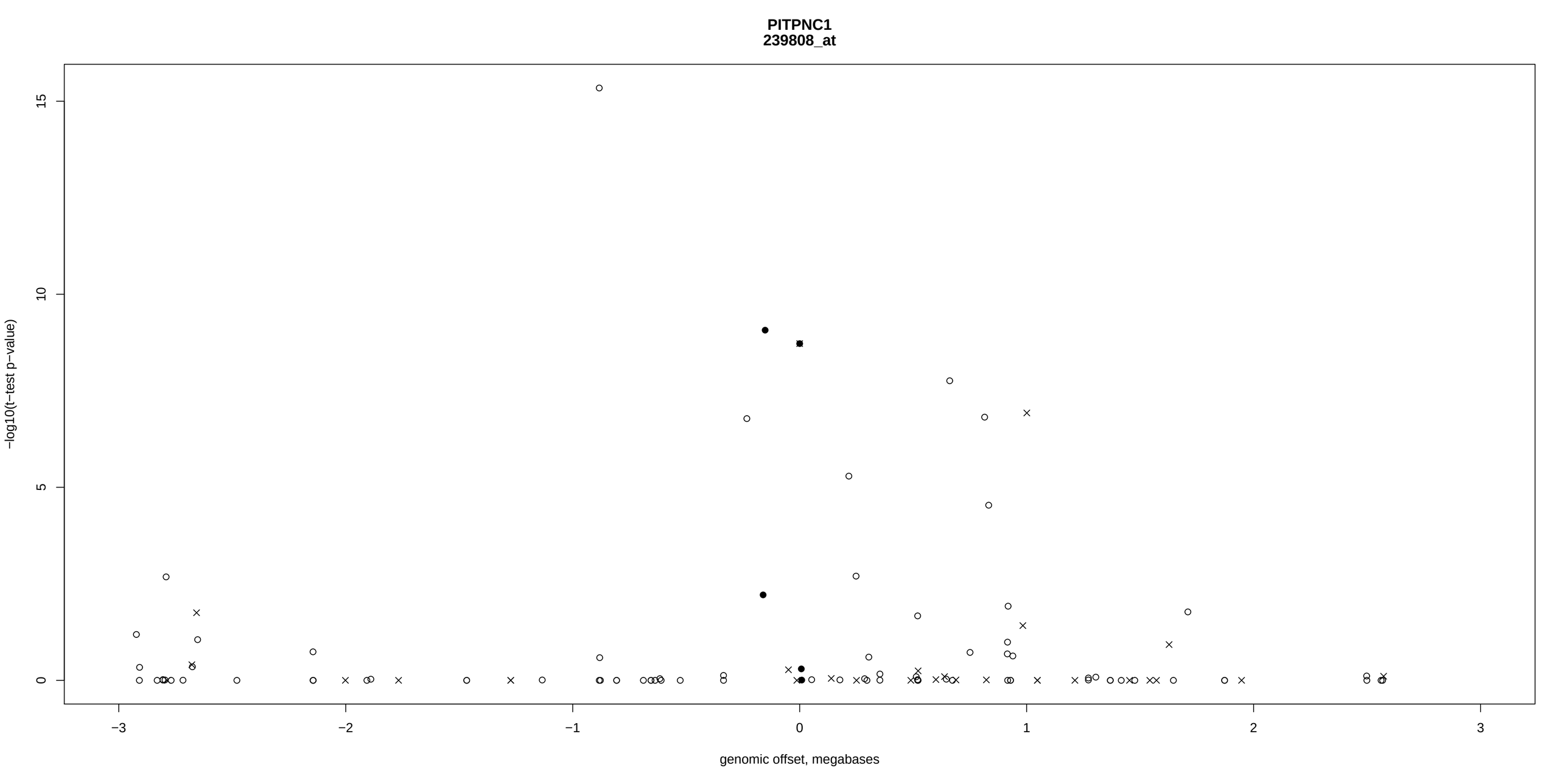
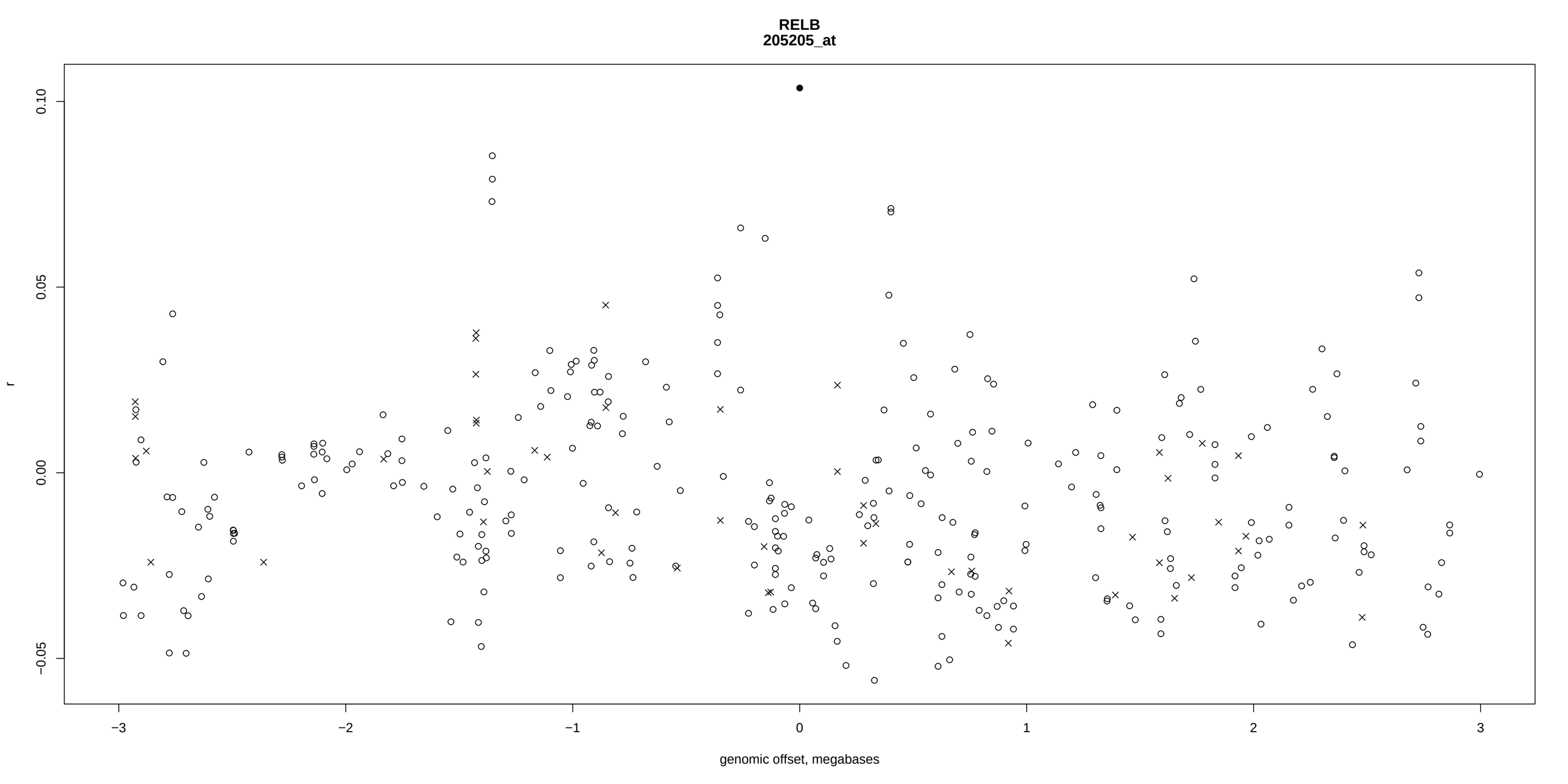
−1
0
1
2
3
genomic offset, megabases


PITPNC1
239808_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


RELB
205205_at
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0.00
−0.05
r
−3
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−1
0
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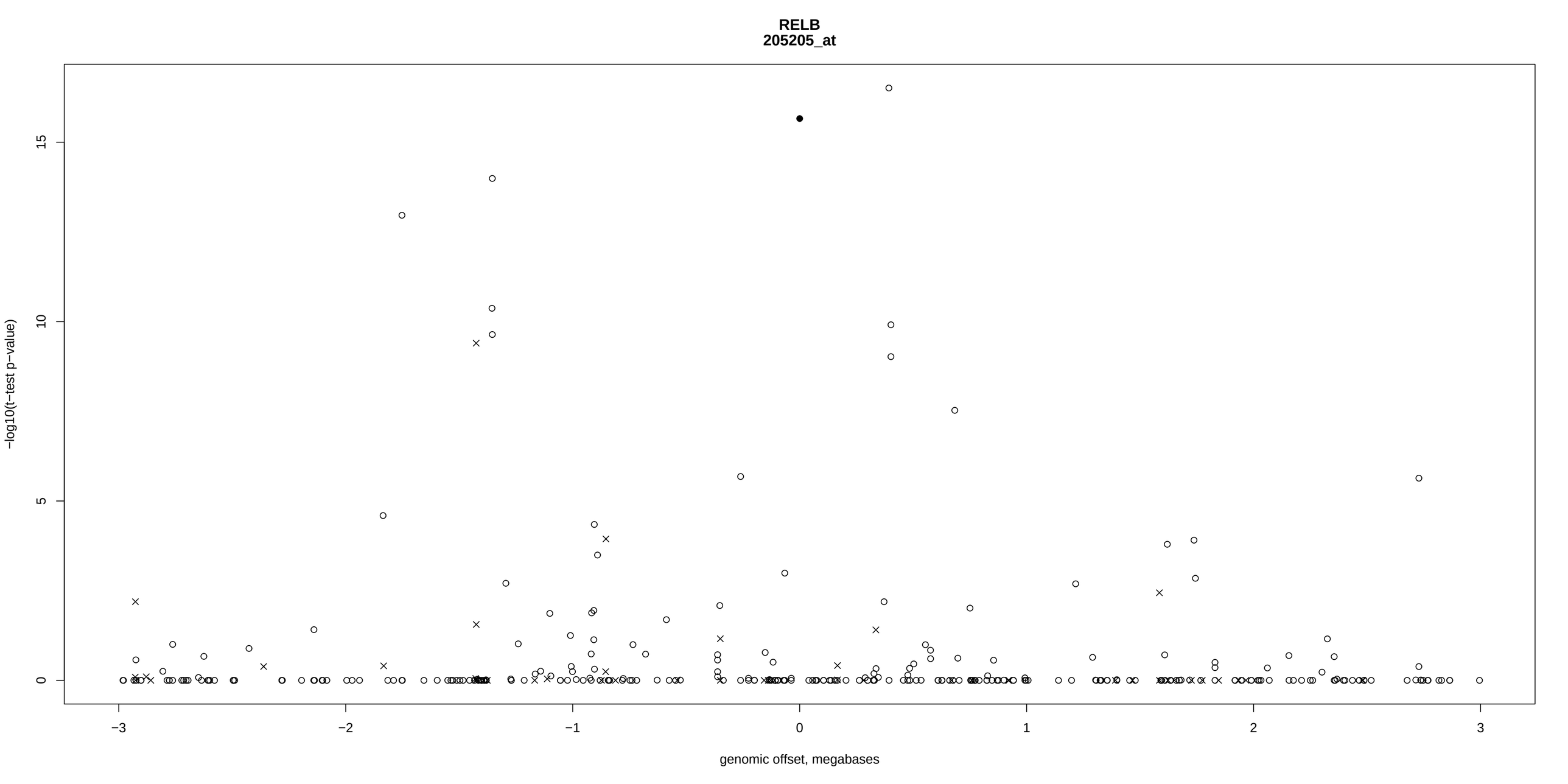
2
3
genomic offset, megabases


RELB
205205_at
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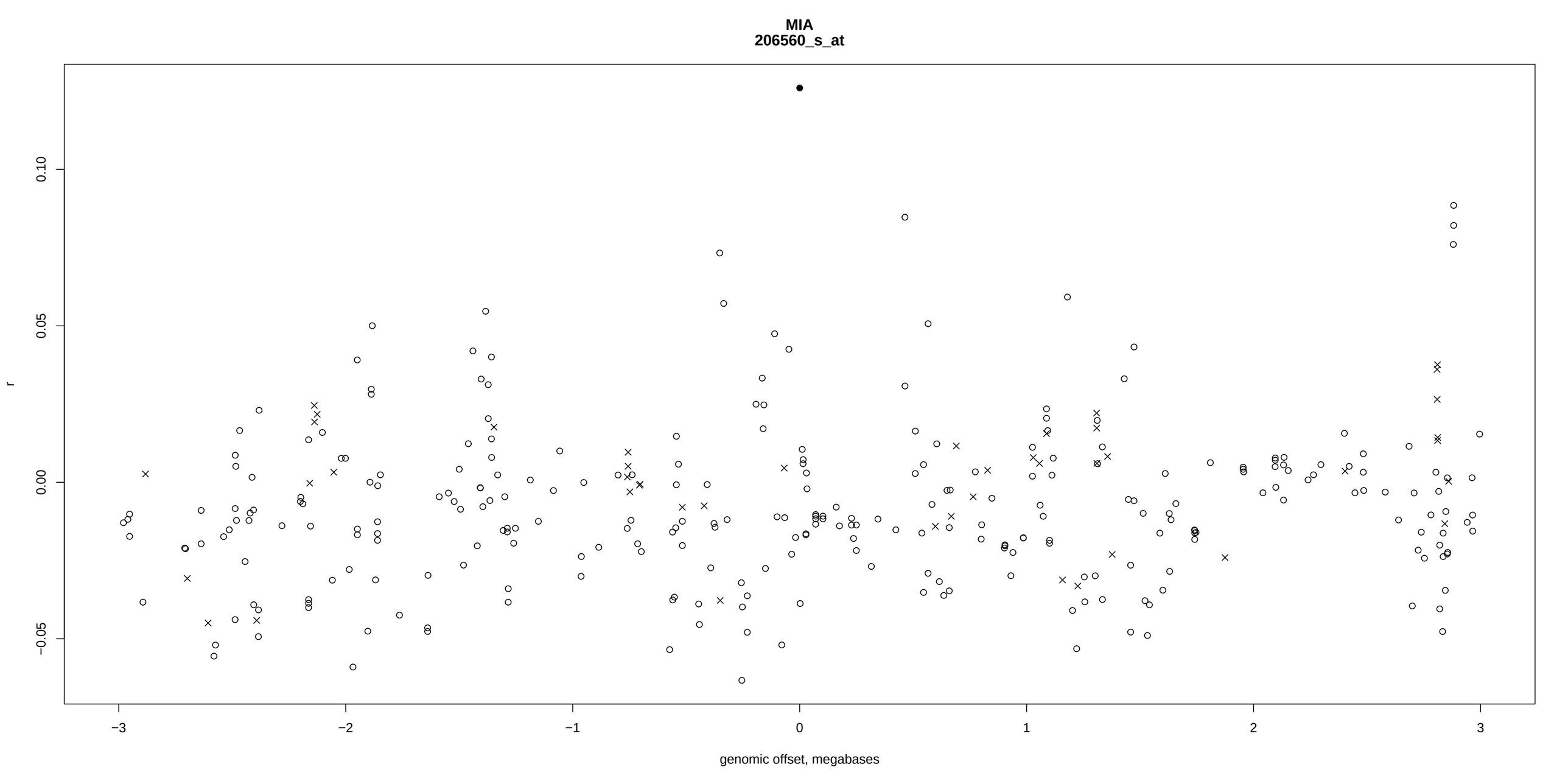
−3
−2
−1
0
1
2
3
genomic offset, megabases


MIA
206560_s_at
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0.00
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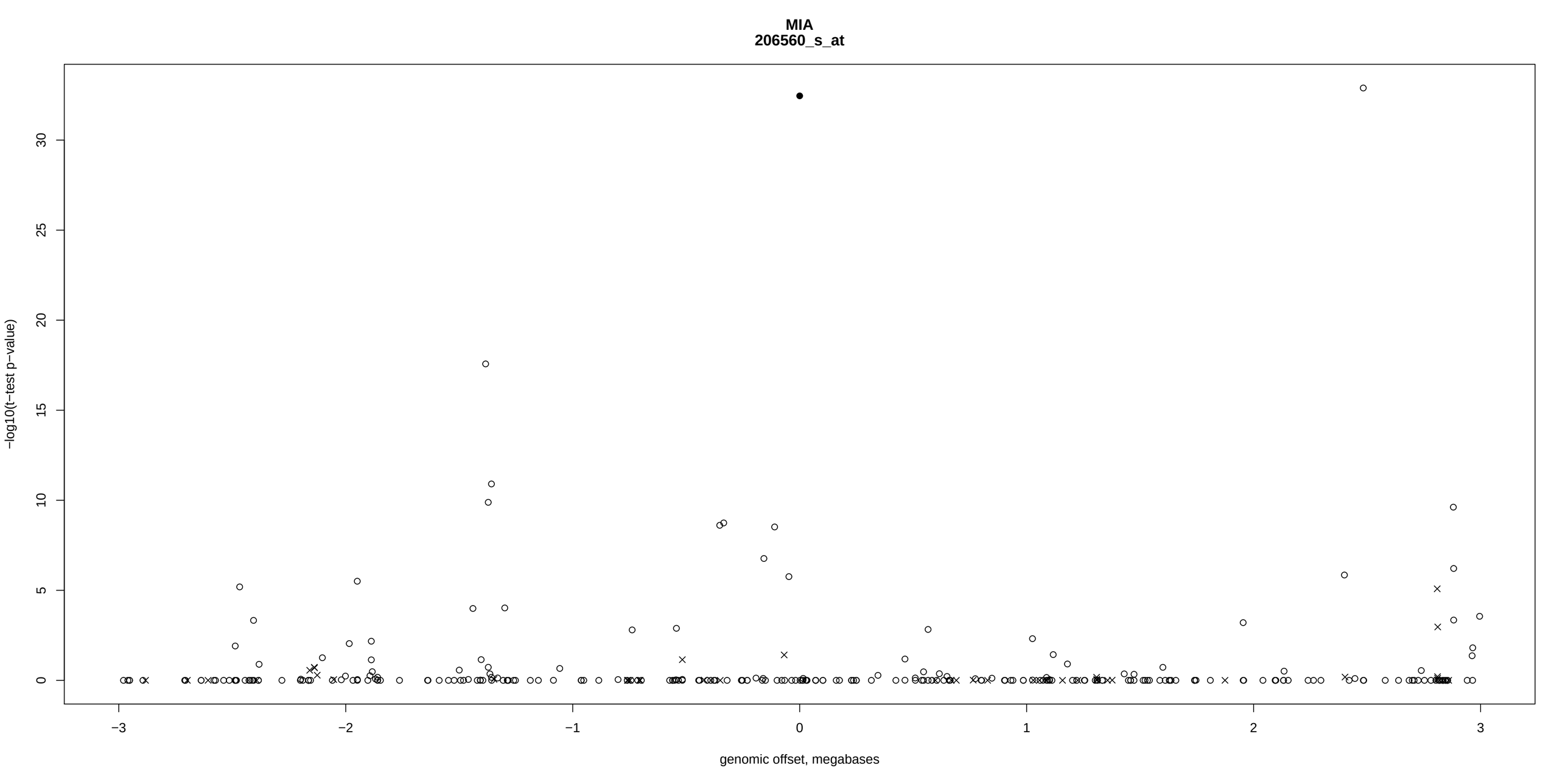
−3
−2
−1
0
1
2
3
genomic offset, megabases


MIA
206560_s_at
−log10(t−test p−value)
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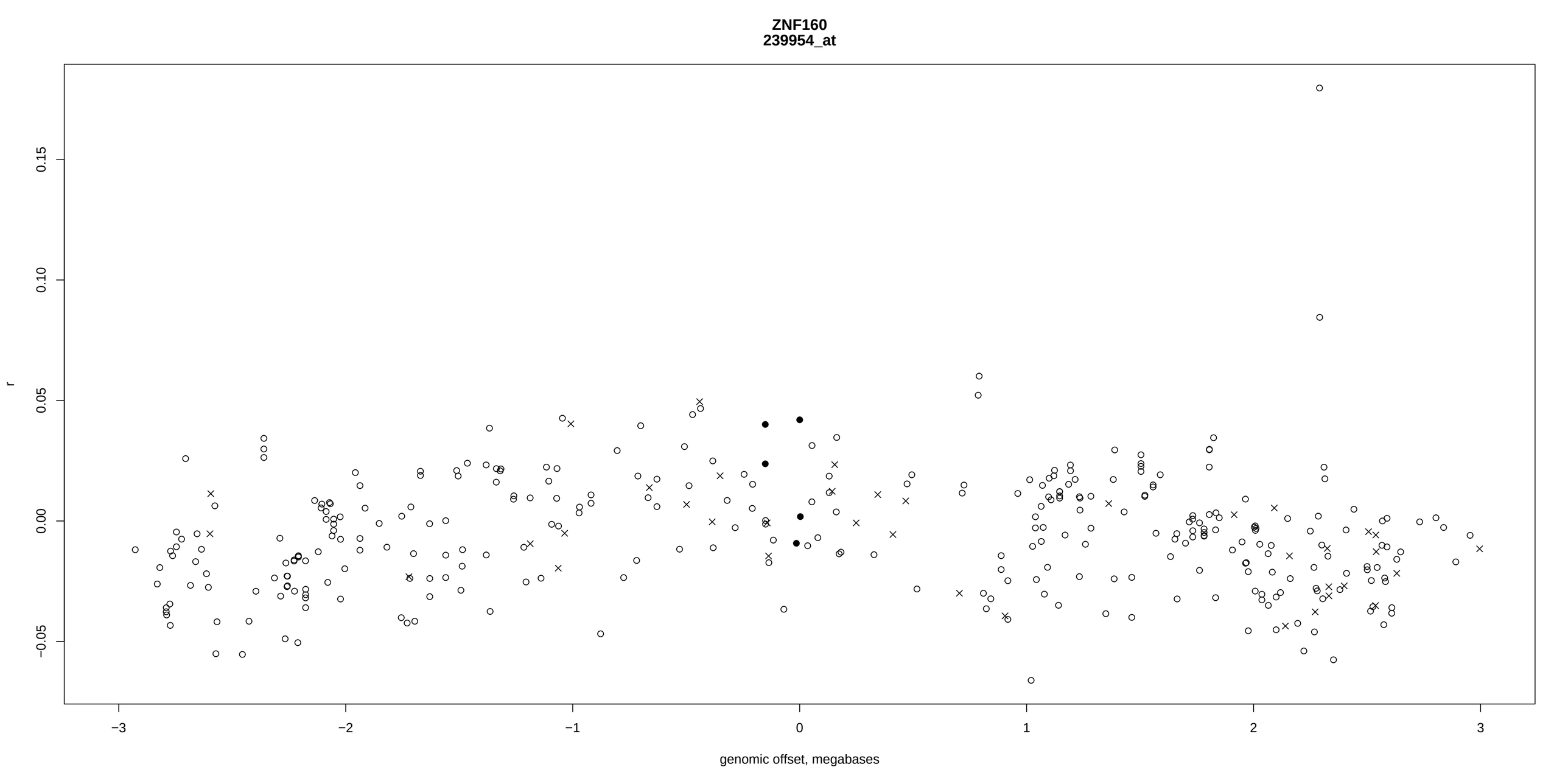
5
10
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


ZNF160
239954_at
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0.05
0.00
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−2
−1
0
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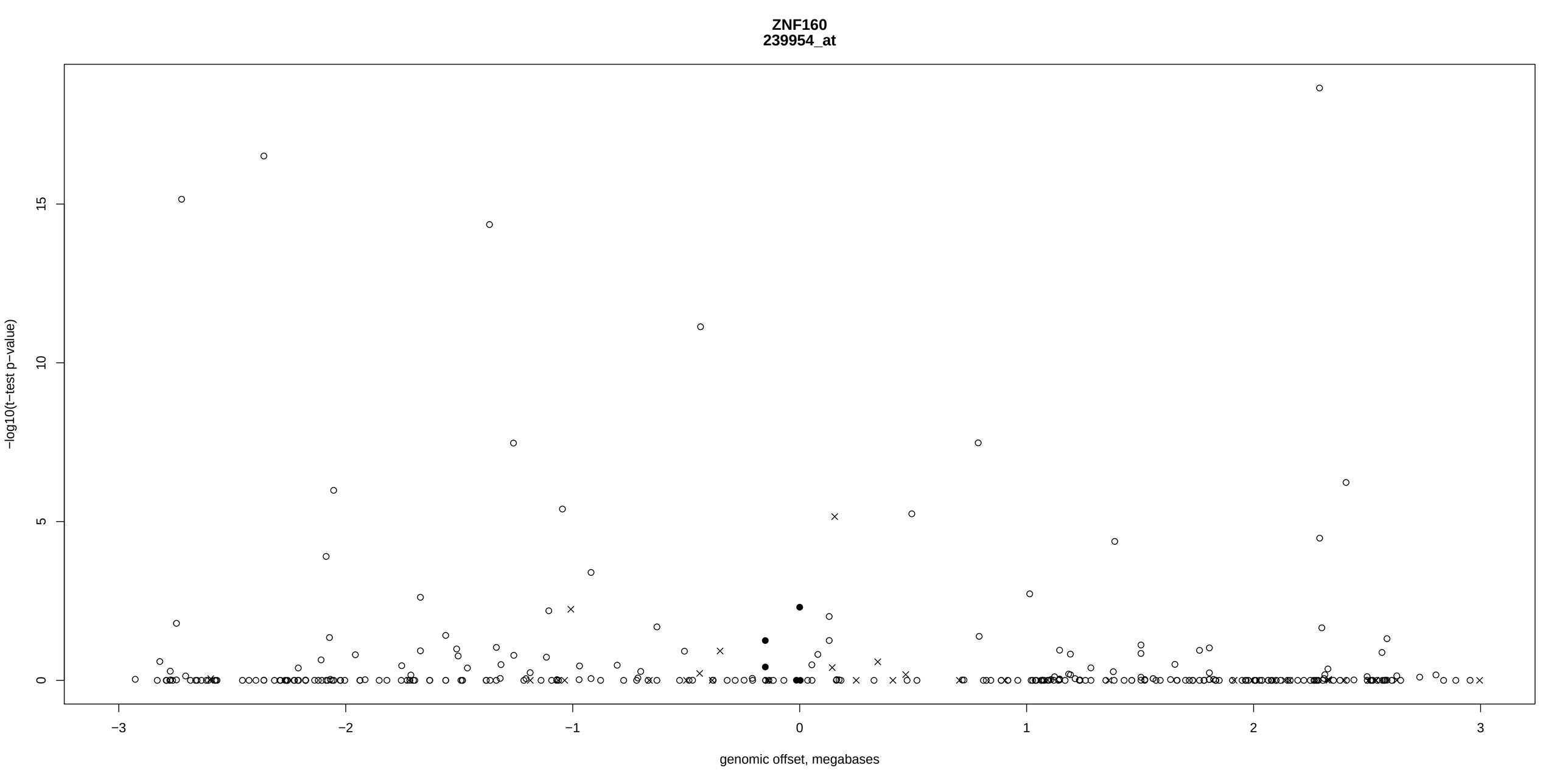
2
3
genomic offset, megabases


ZNF160
239954_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
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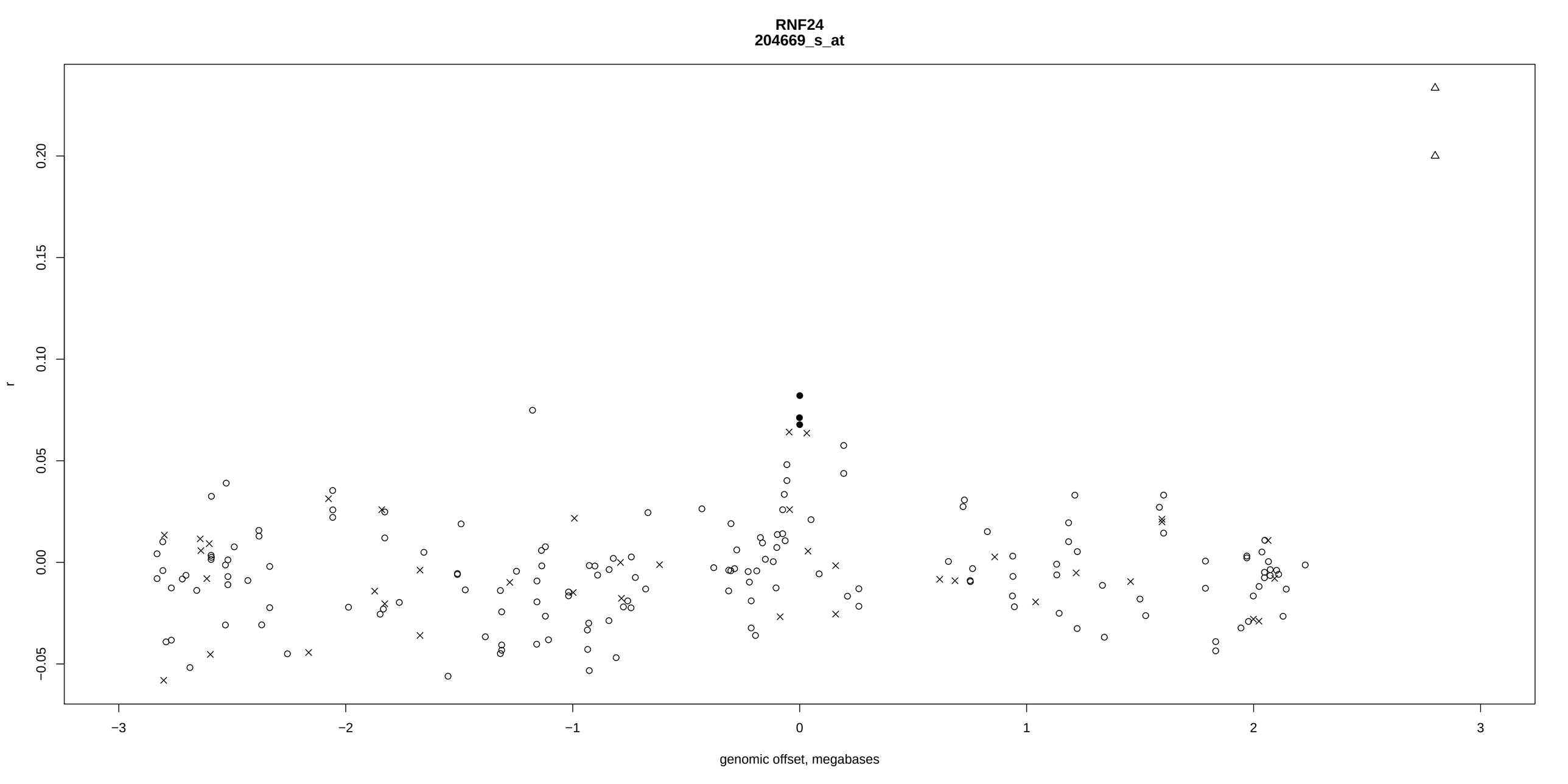
2
3
genomic offset, megabases


RNF24
204669_s_at
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−3
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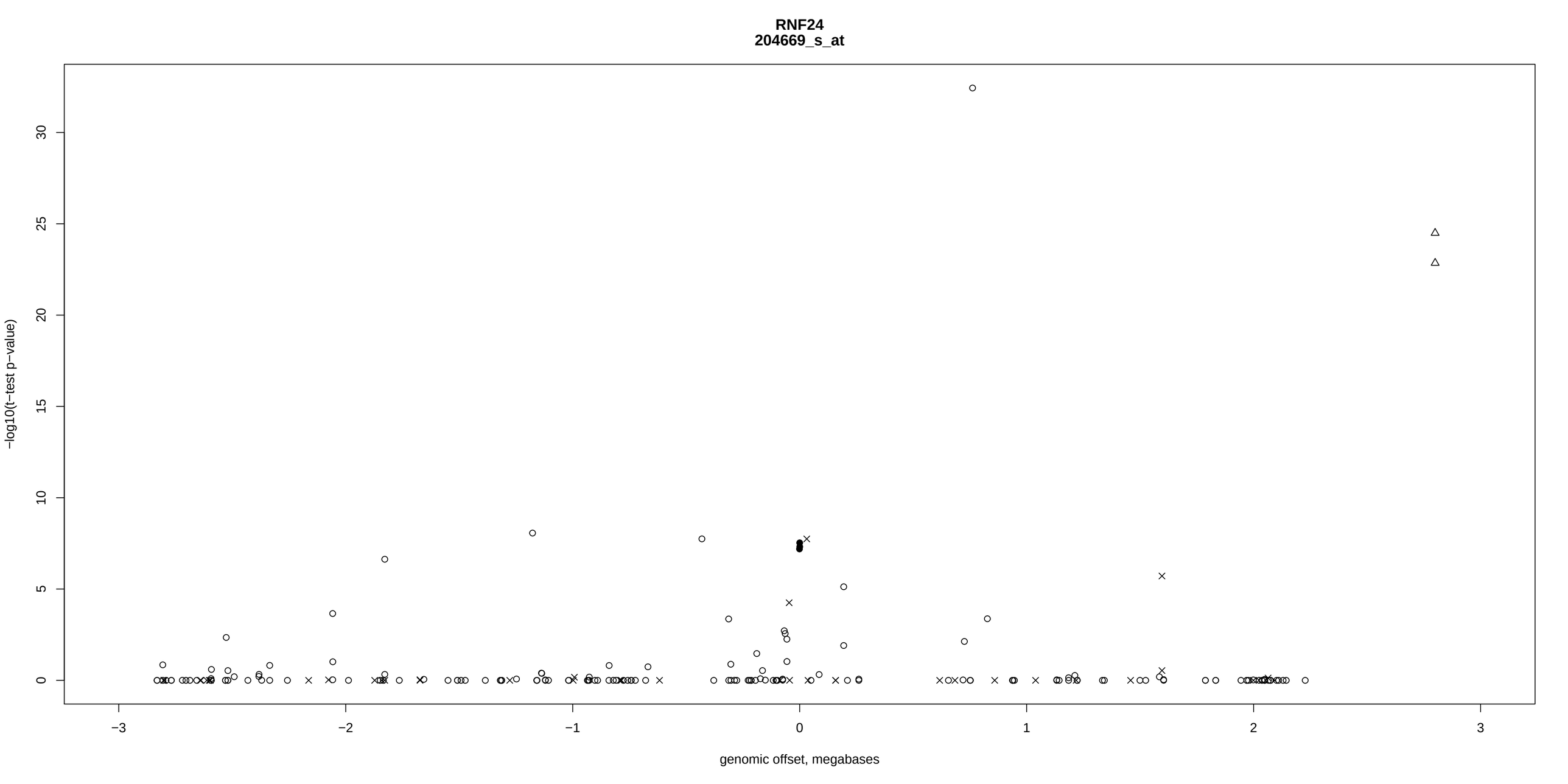
2
3
genomic offset, megabases


RNF24
204669_s_at
−log10(t−test p−value)
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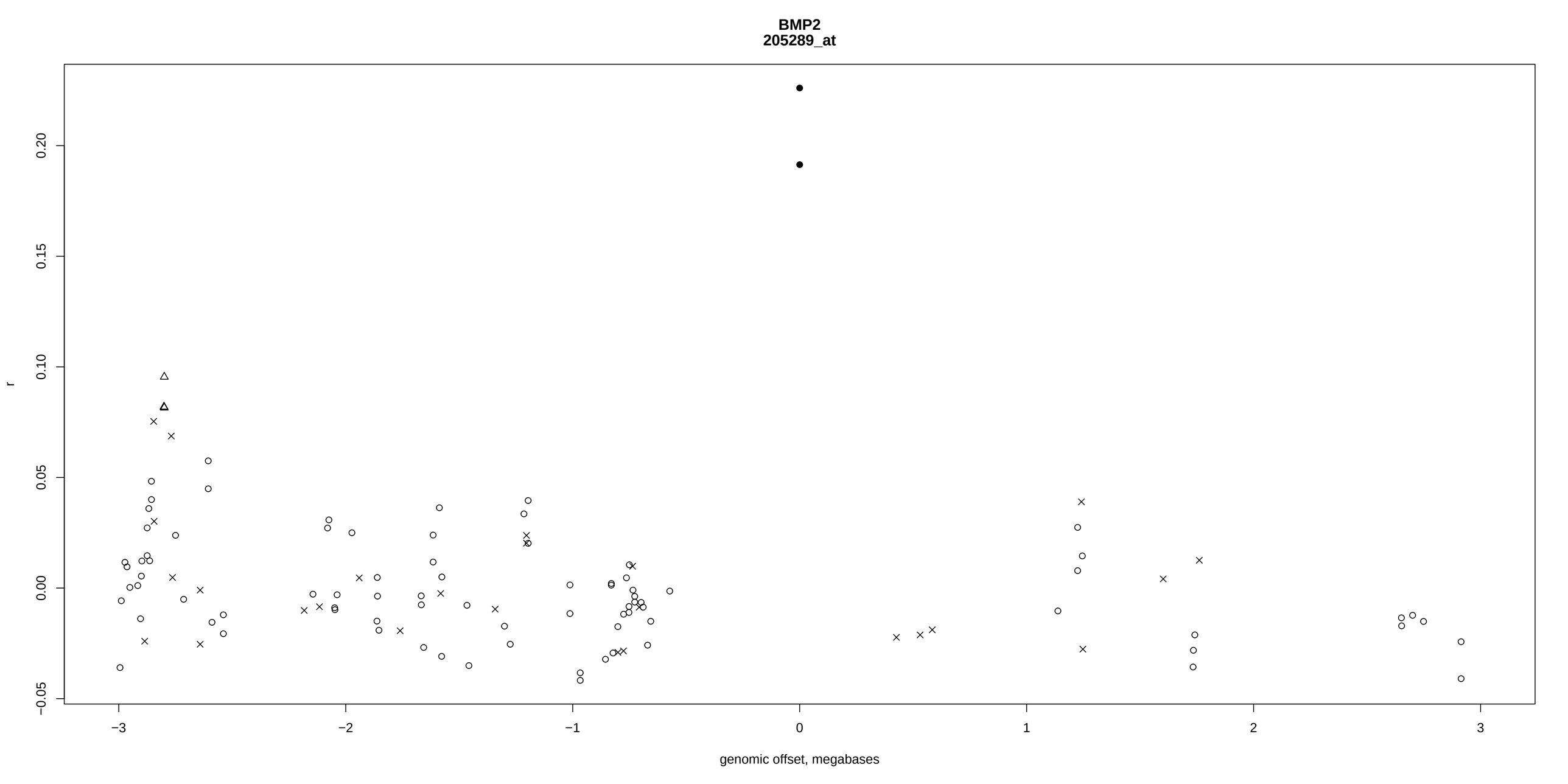
10
15
20
25
30
−3
−2
−1
0
1
2
3
genomic offset, megabases


BMP2
205289_at
r
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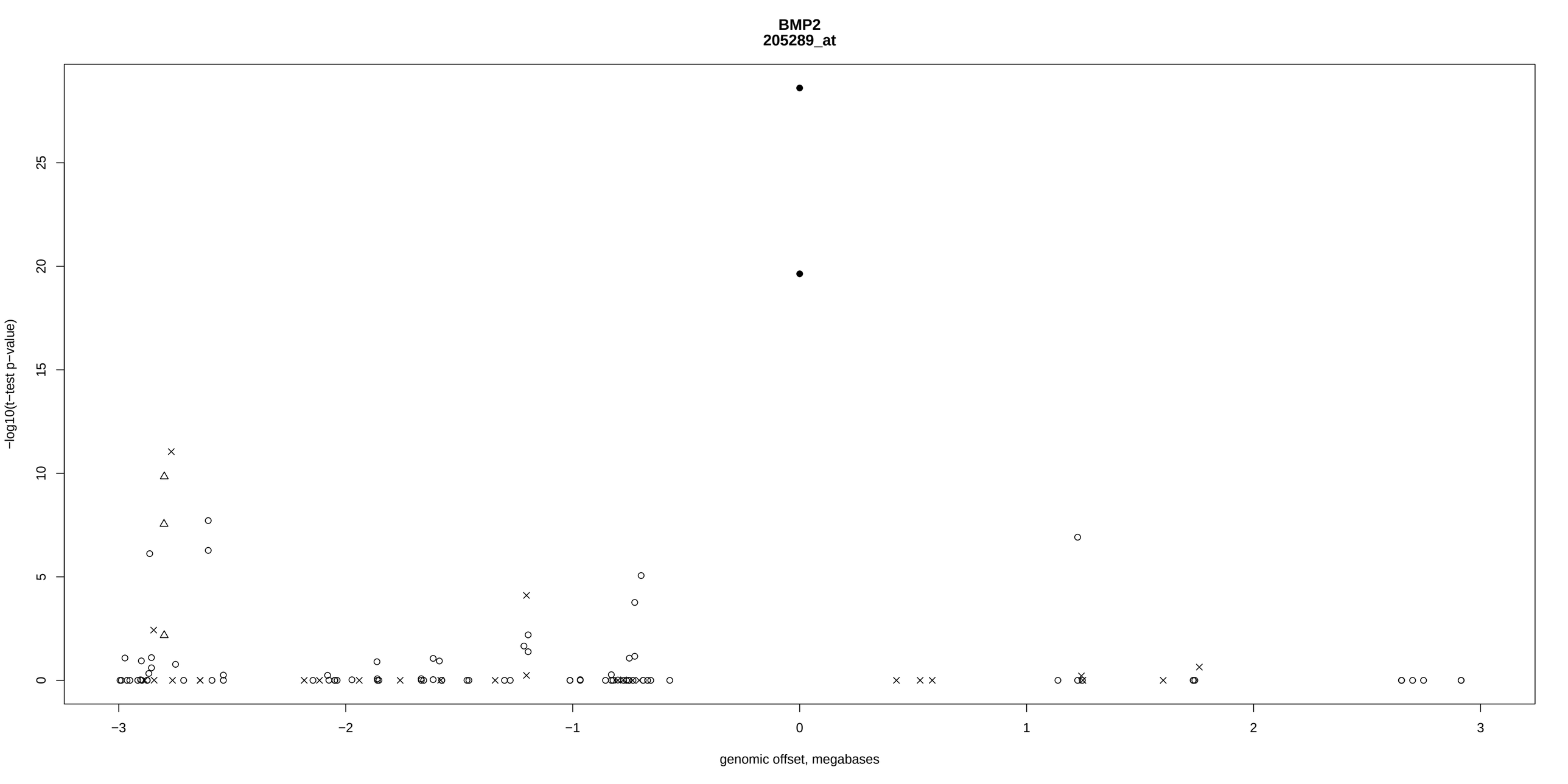
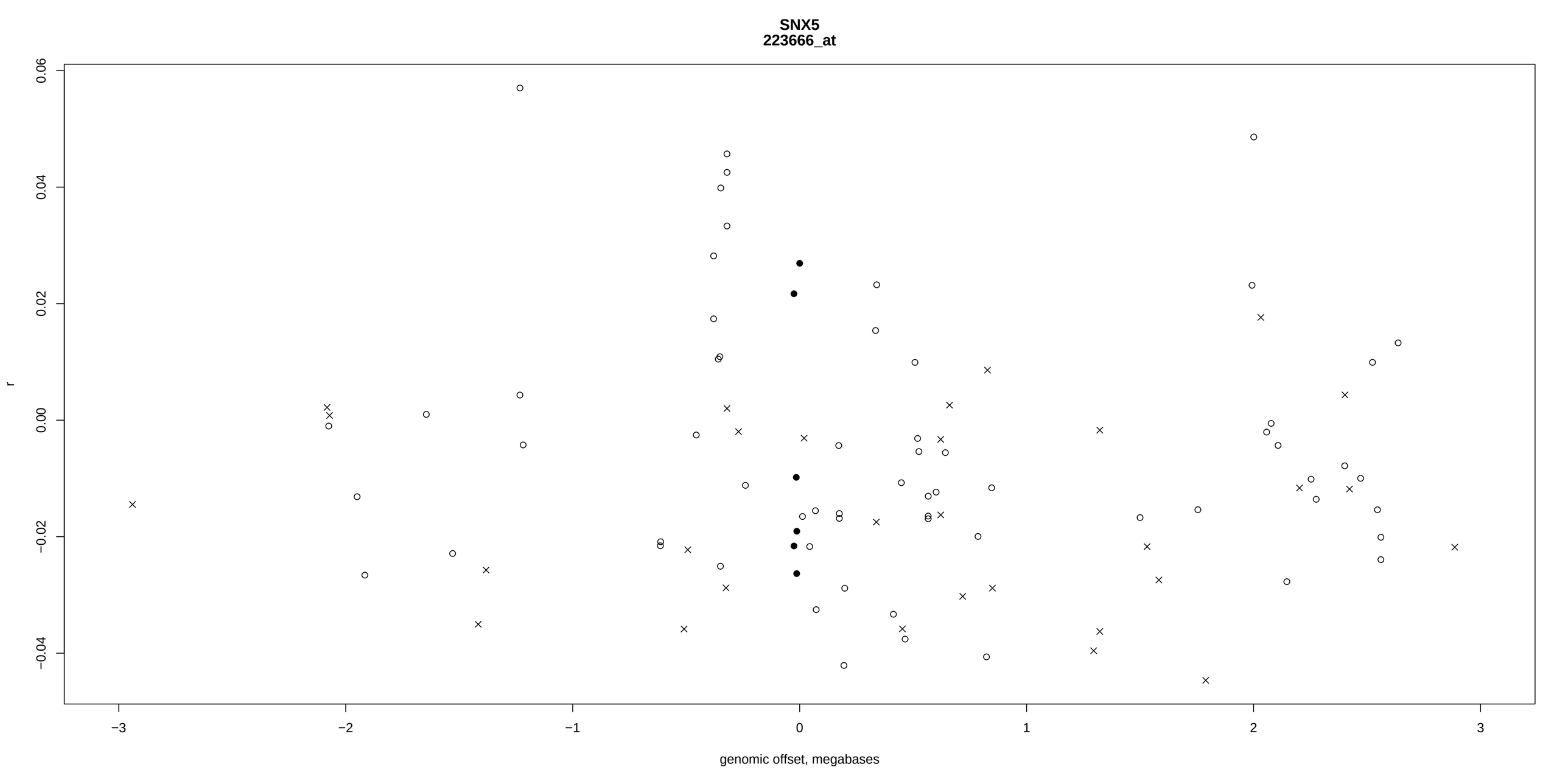
0.05
0.00
−0.05
−3
−2
−1
0
1
2
3
genomic offset, megabases


BMP2
205289_at
−log10(t−test p−value)
0
5
10
15
20
25
−3
−2
−1
0
1
2
3
genomic offset, megabases


SNX5
223666_at
0.06
0.04
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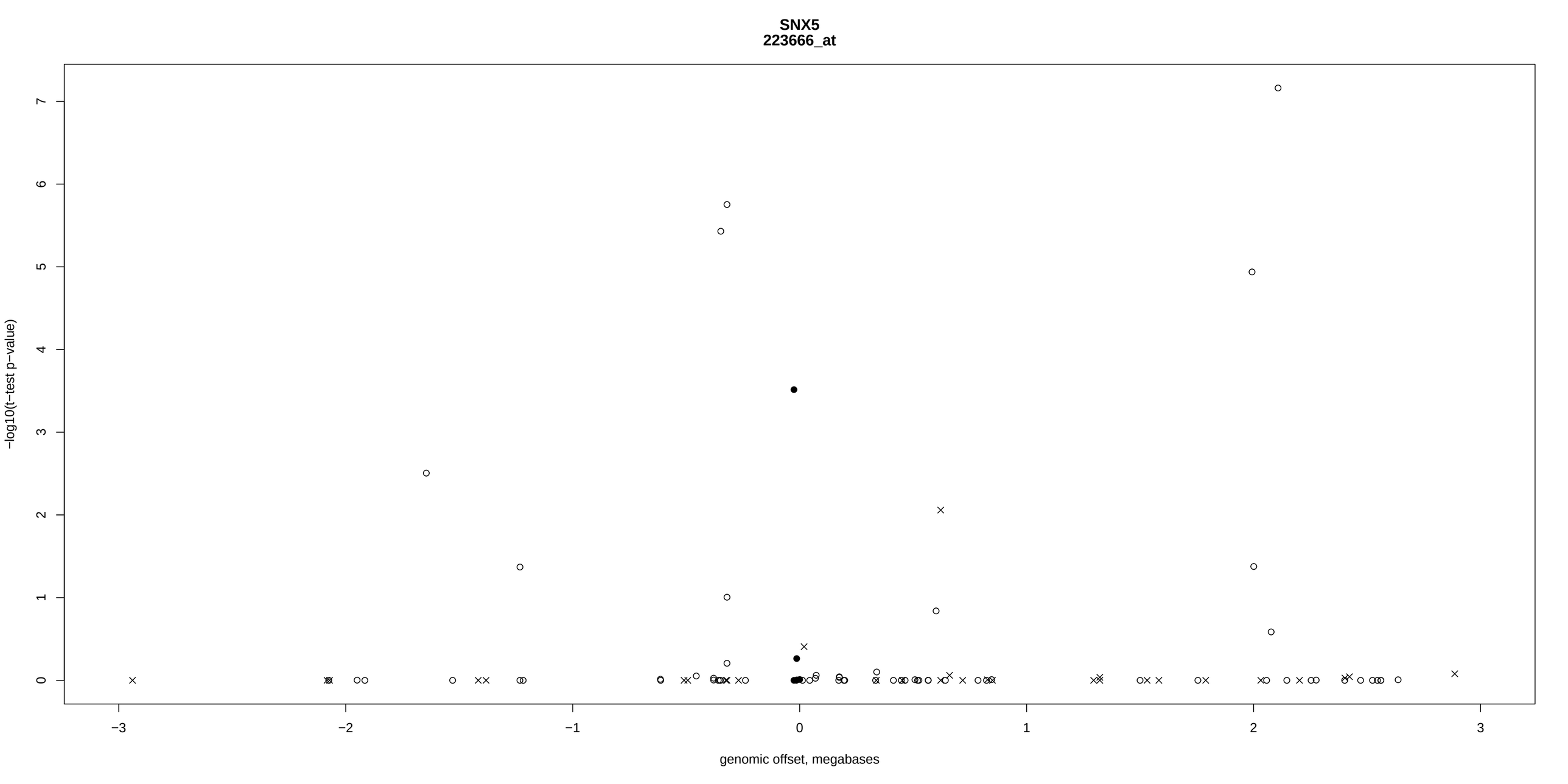
−0.02
−0.04
r
−3
−2
−1
0
1
2
3
genomic offset, megabases


SNX5
223666_at
−log10(t−test p−value)
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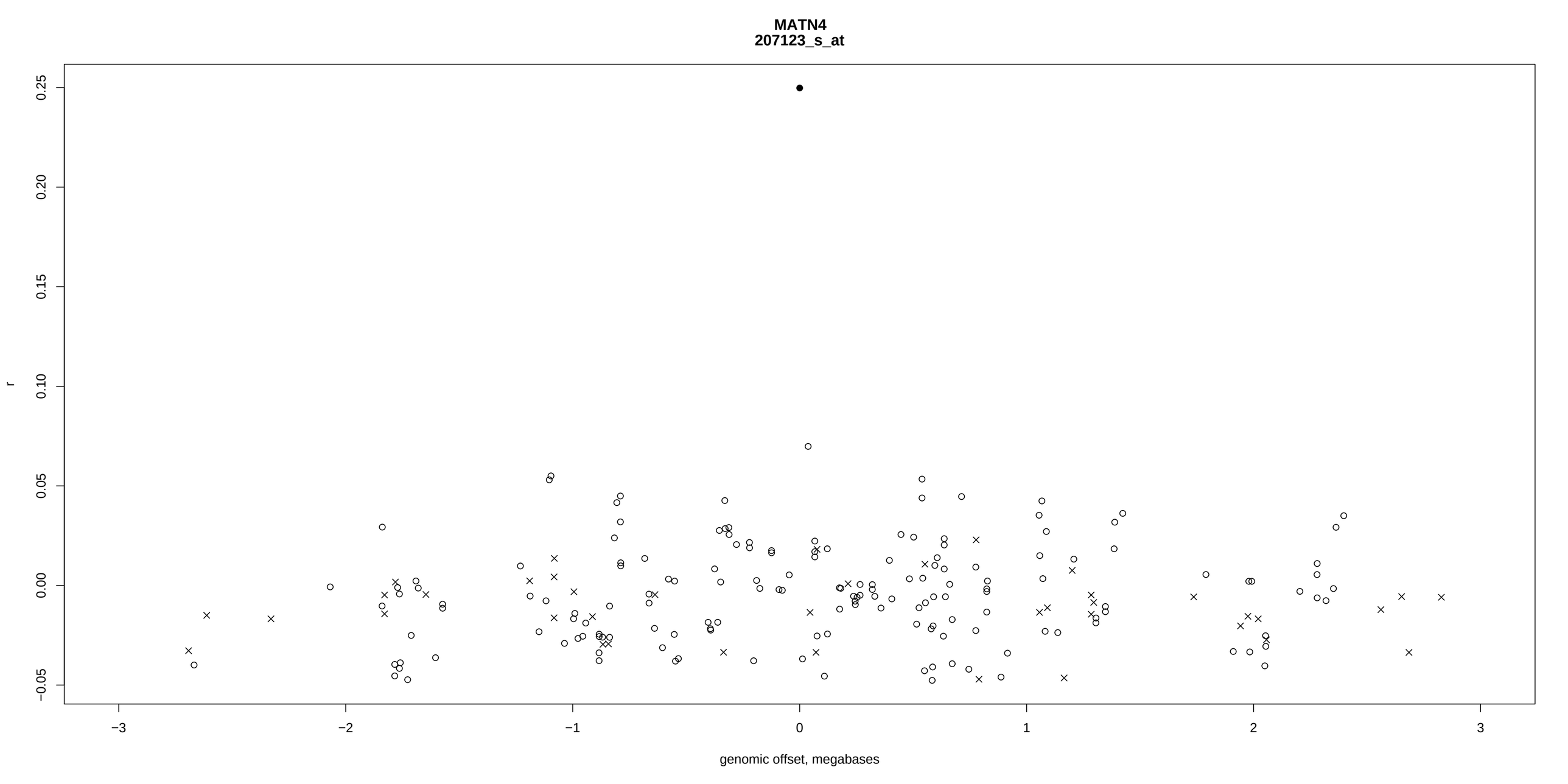
2
3
4
5
6
7
−3
−2
−1
0
1
2
3
genomic offset, megabases


MATN4
207123_s_at
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0.05
0.00
−0.05
−3
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−1
0
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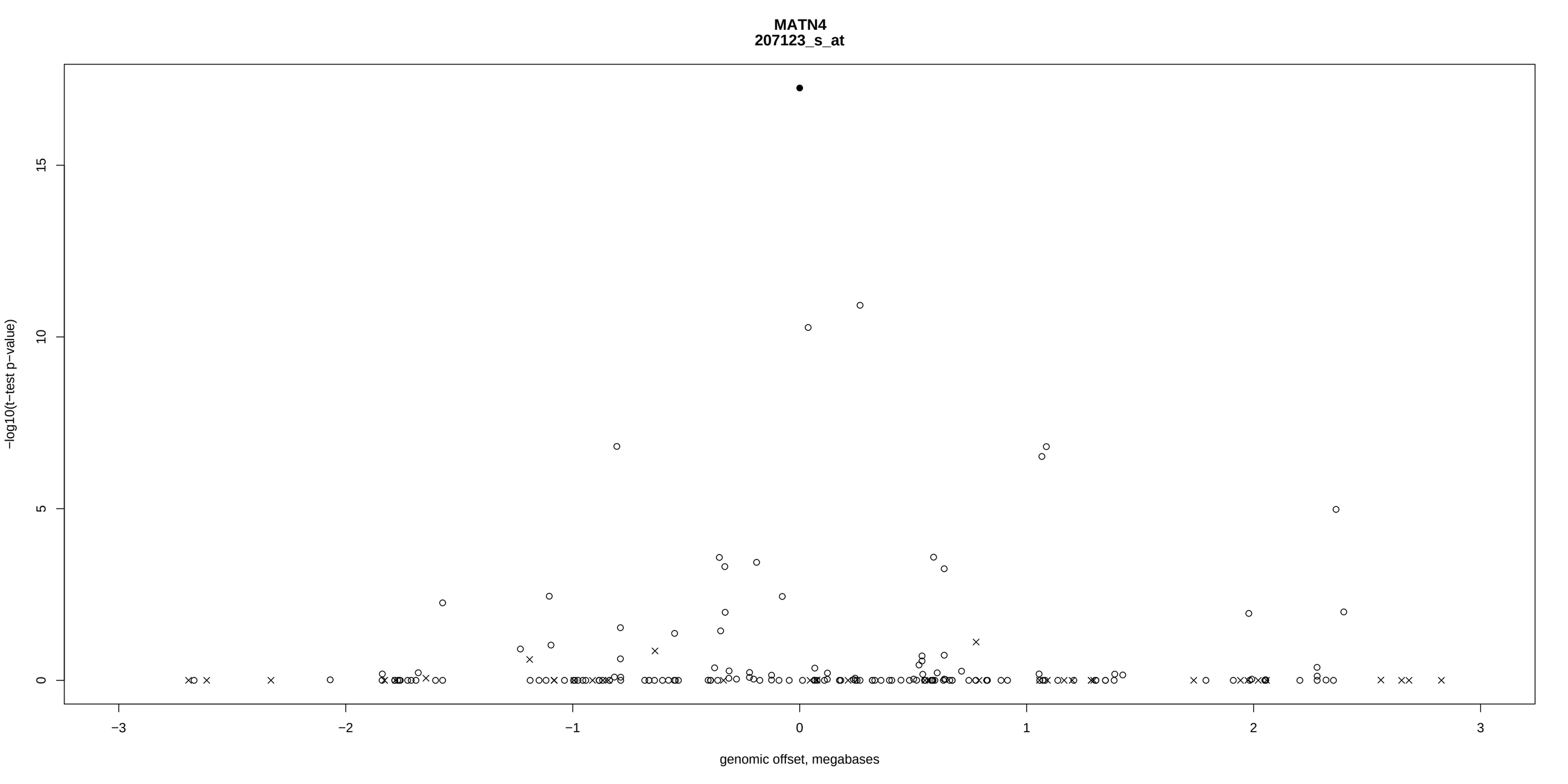
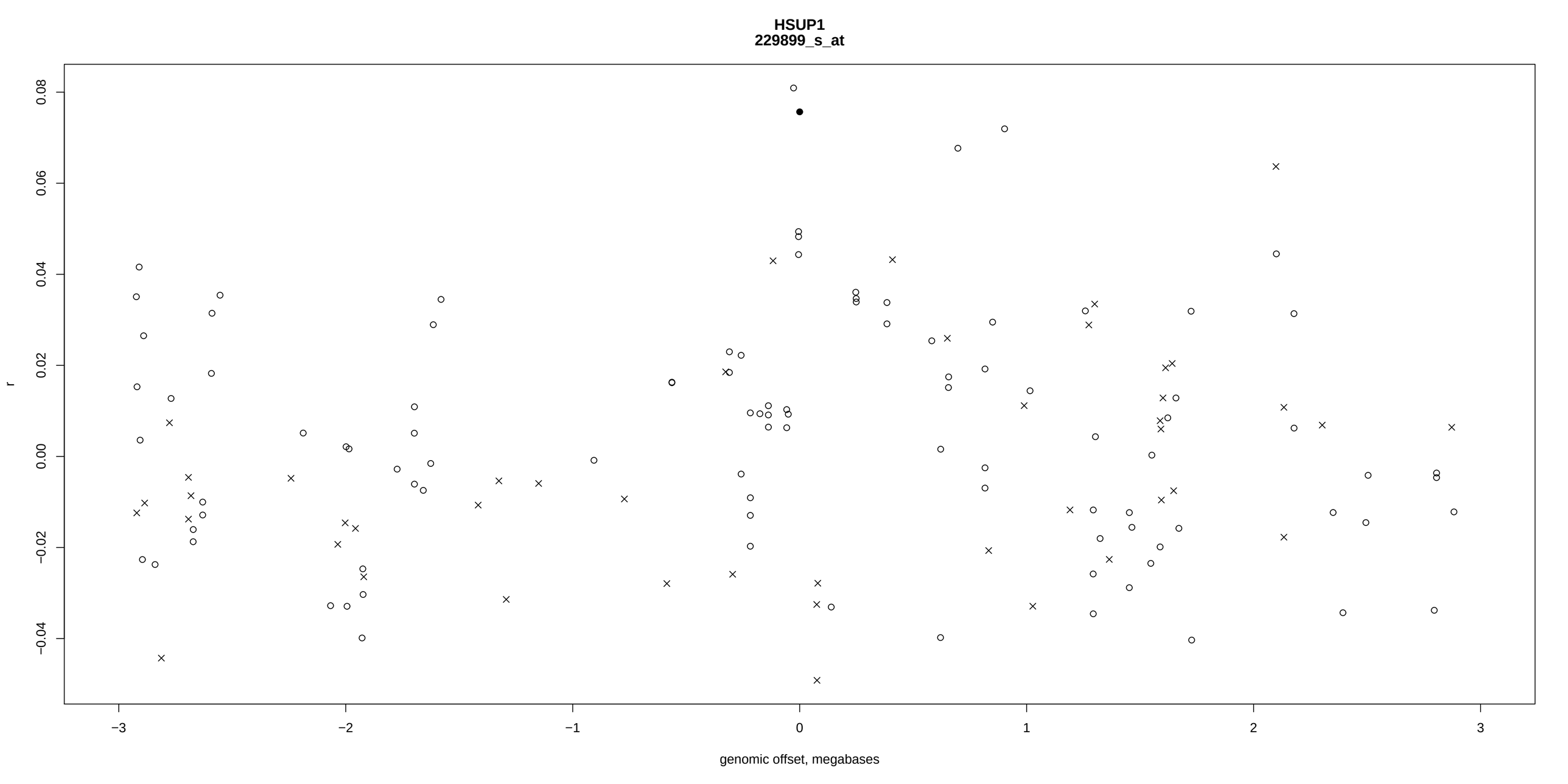
2
3
genomic offset, megabases


MATN4
207123_s_at
−log10(t−test p−value)
0
5
10
15
−3
−2
−1
0
1
2
3
genomic offset, megabases


HSUP1
229899_s_at
r
0.08
0.06
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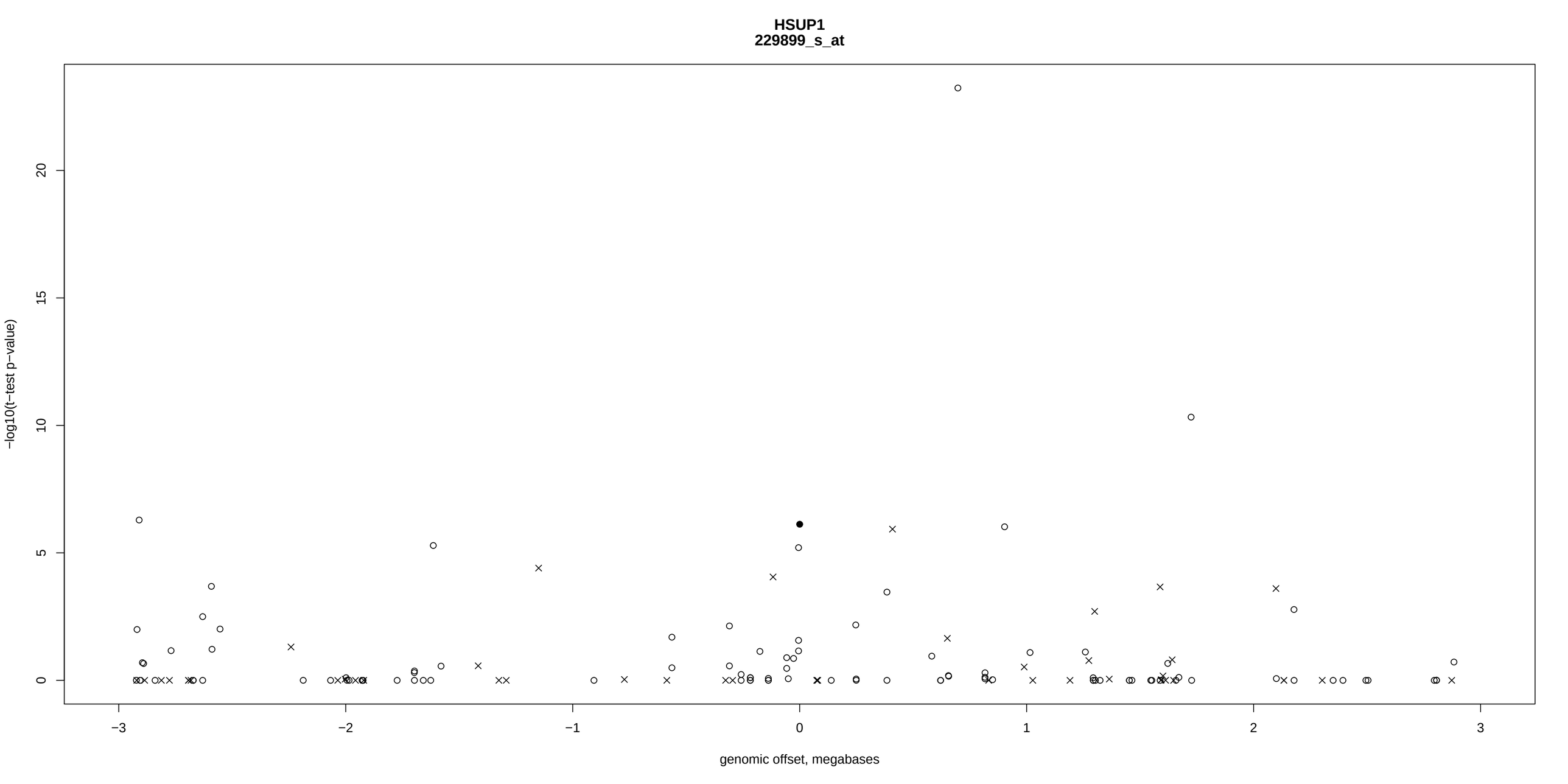
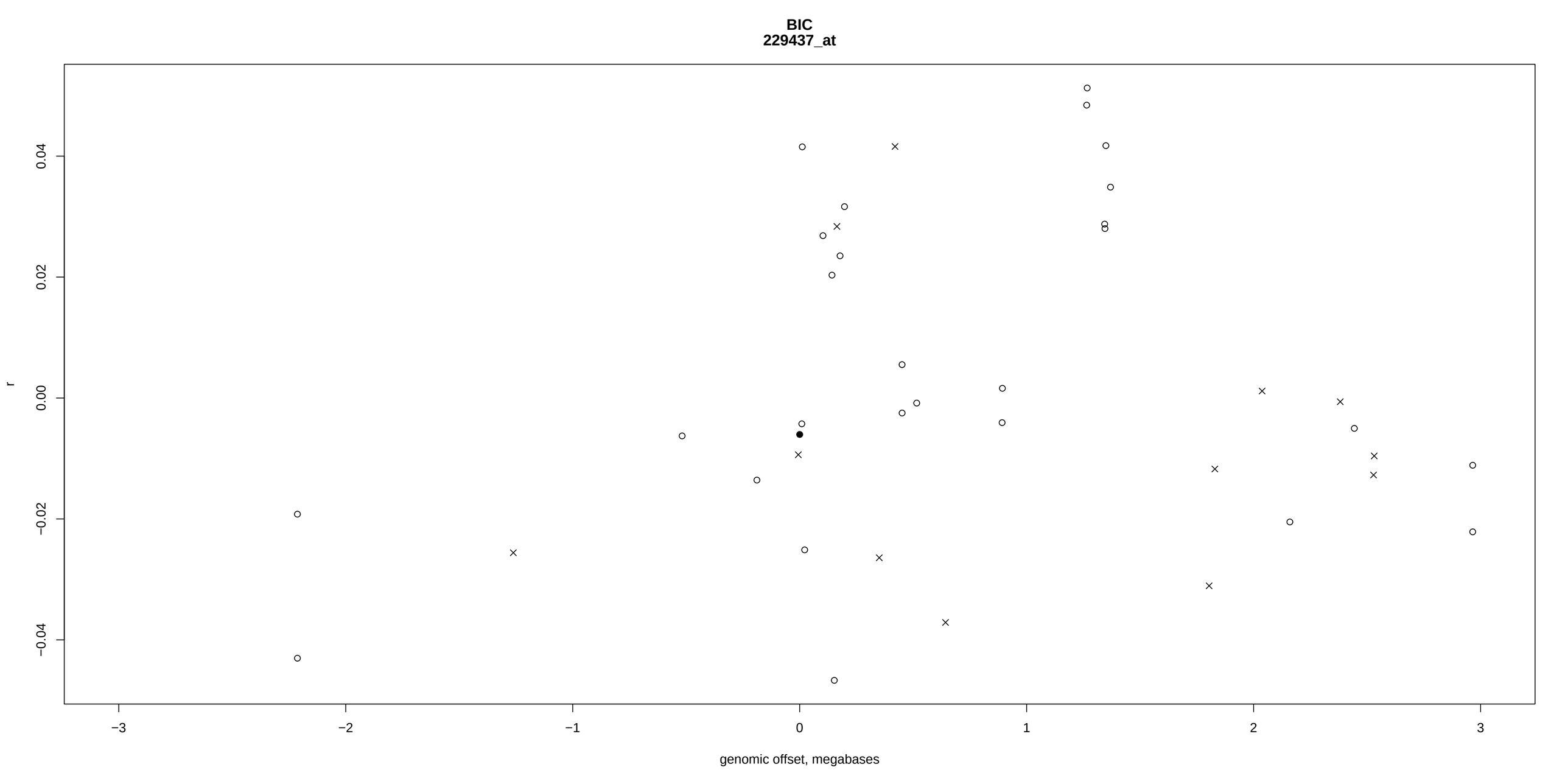
0.00
−0.02
−0.04
−3
−2
−1
0
1
2
3
genomic offset, megabases


HSUP1
229899_s_at
−log10(t−test p−value)
0
5
10
15
20
−3
−2
−1
0
1
2
3
genomic offset, megabases


BIC
229437_at
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0.00
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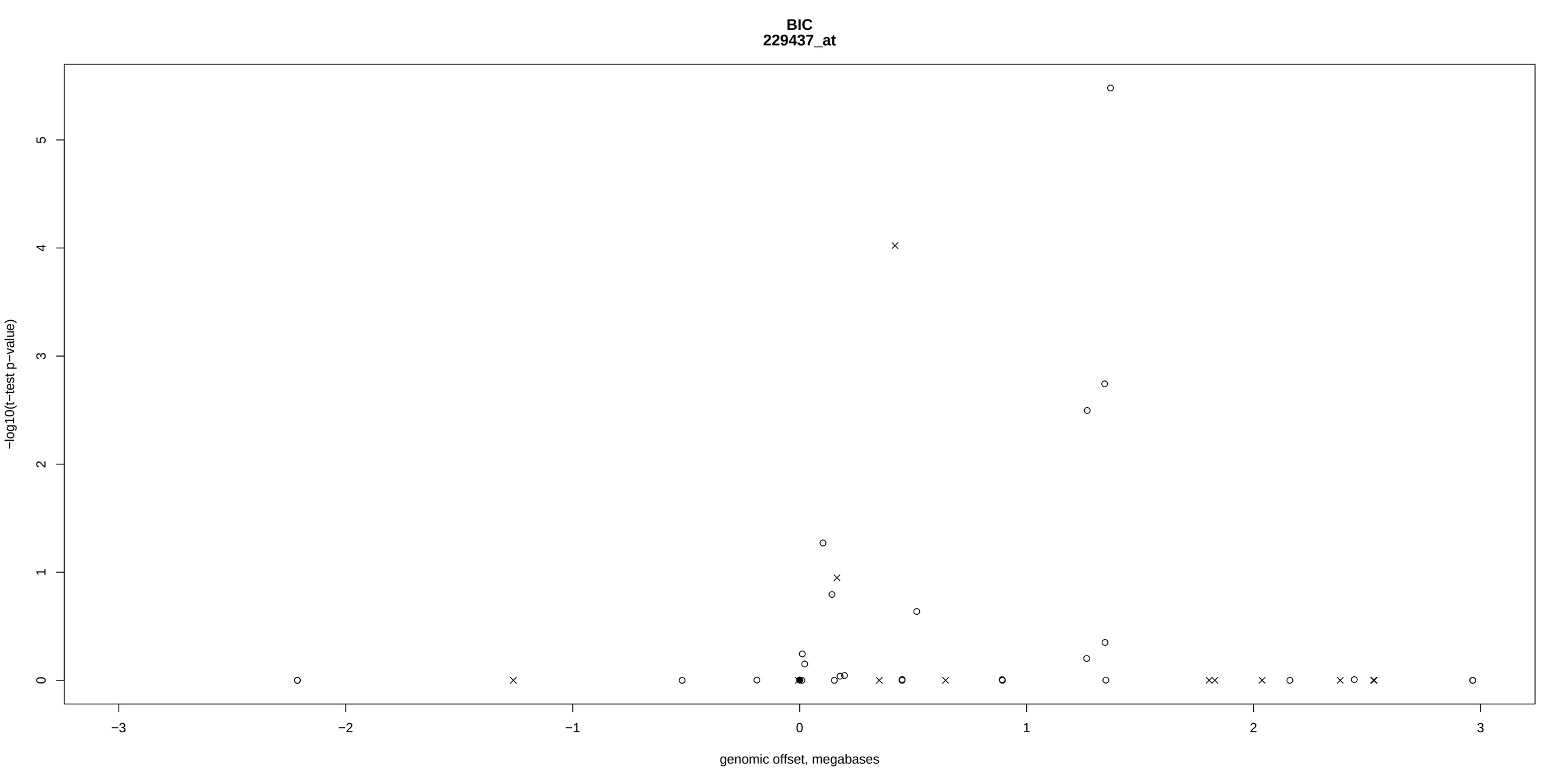
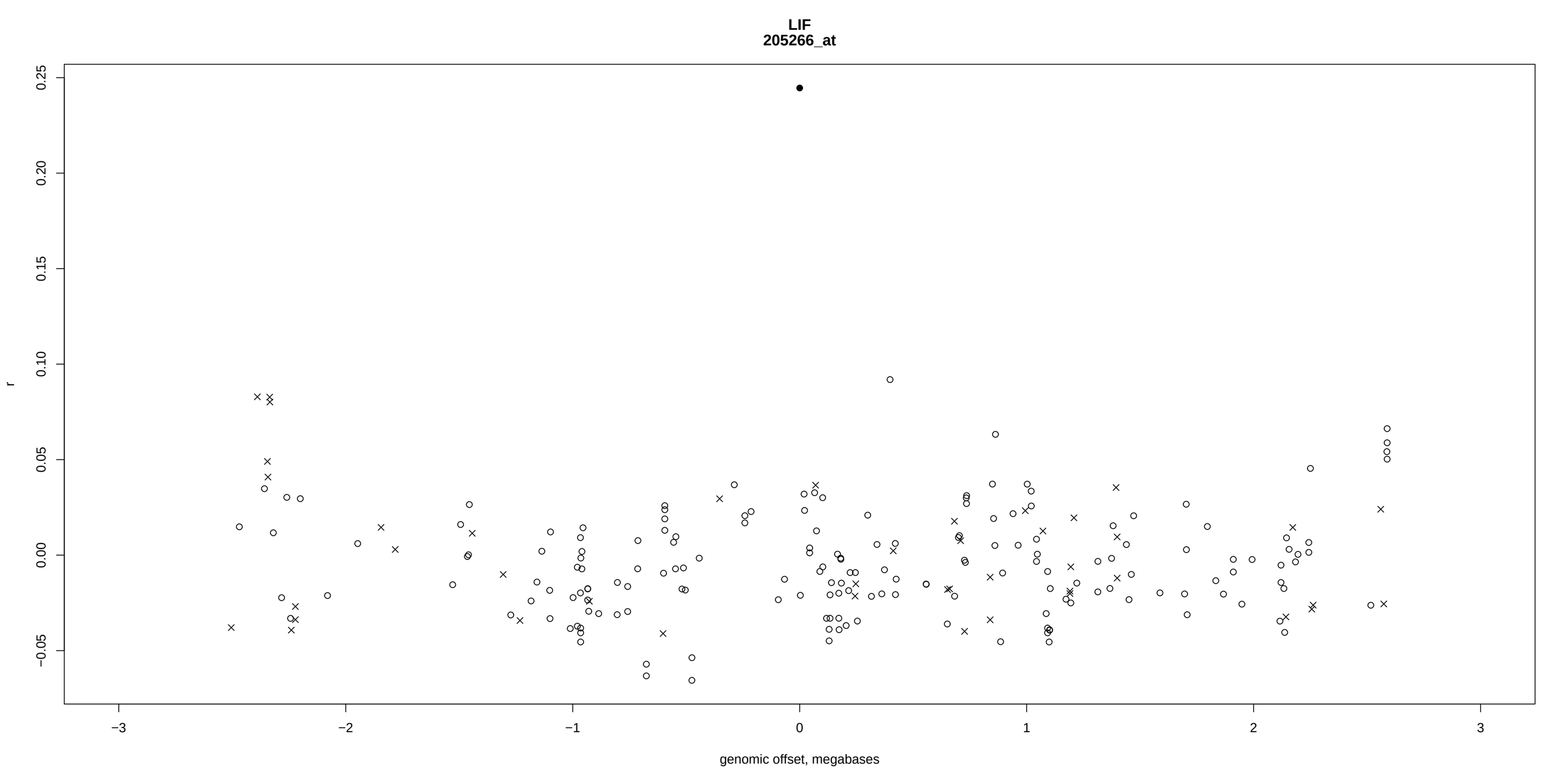
−3
−2
−1
0
1
2
3
genomic offset, megabases


BIC
229437_at
−log10(t−test p−value)
0
1
2
3
4
5
−3
−2
−1
0
1
2
3
genomic offset, megabases


LIF
205266_at
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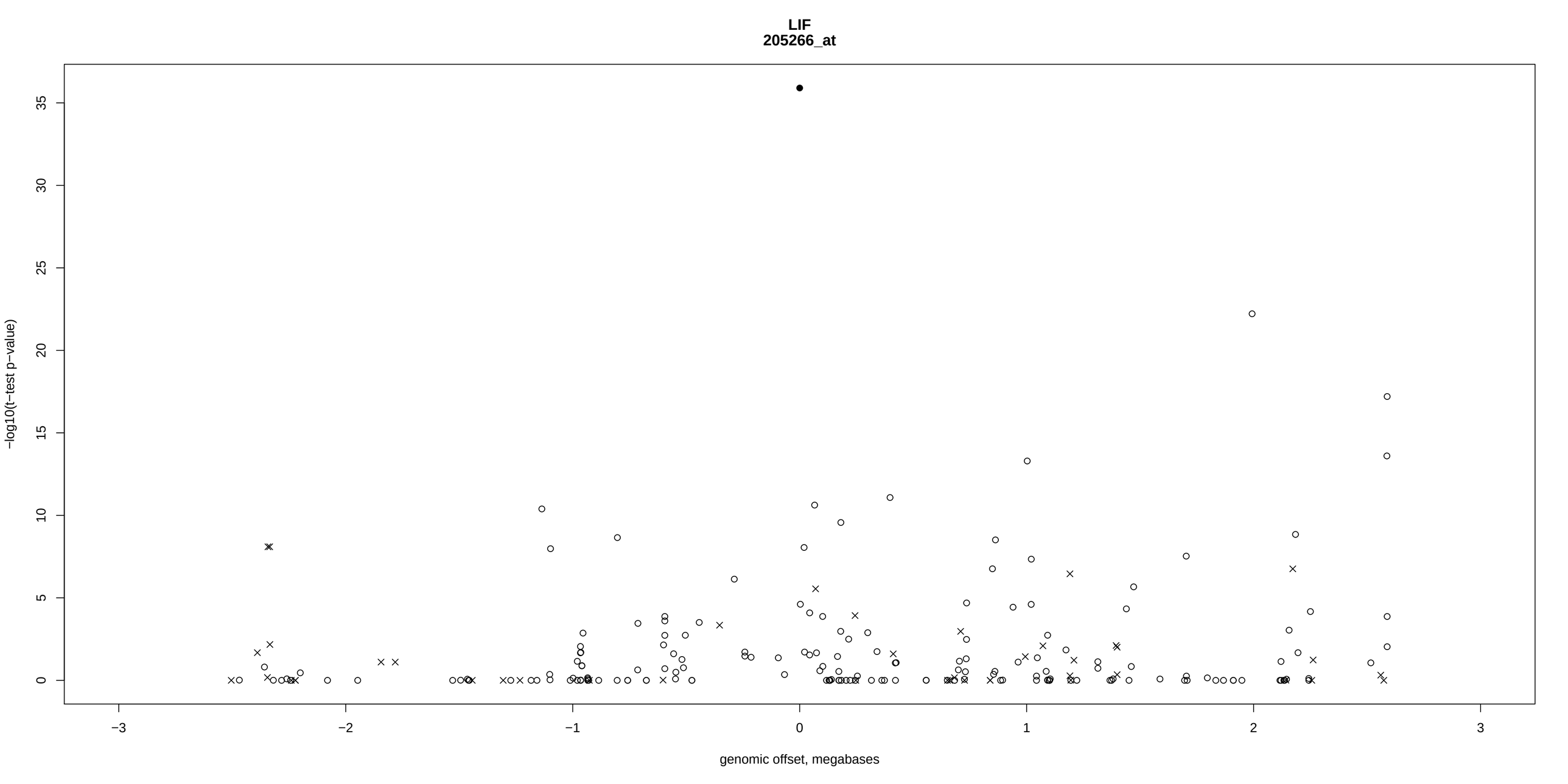
−3
−2
−1
0
1
2
3
genomic offset, megabases


LIF
205266_at
−log10(t−test p−value)
0
5
10
15
20
25
30
35
−3
−2
−1
0
1
2
3
genomic offset, megabases