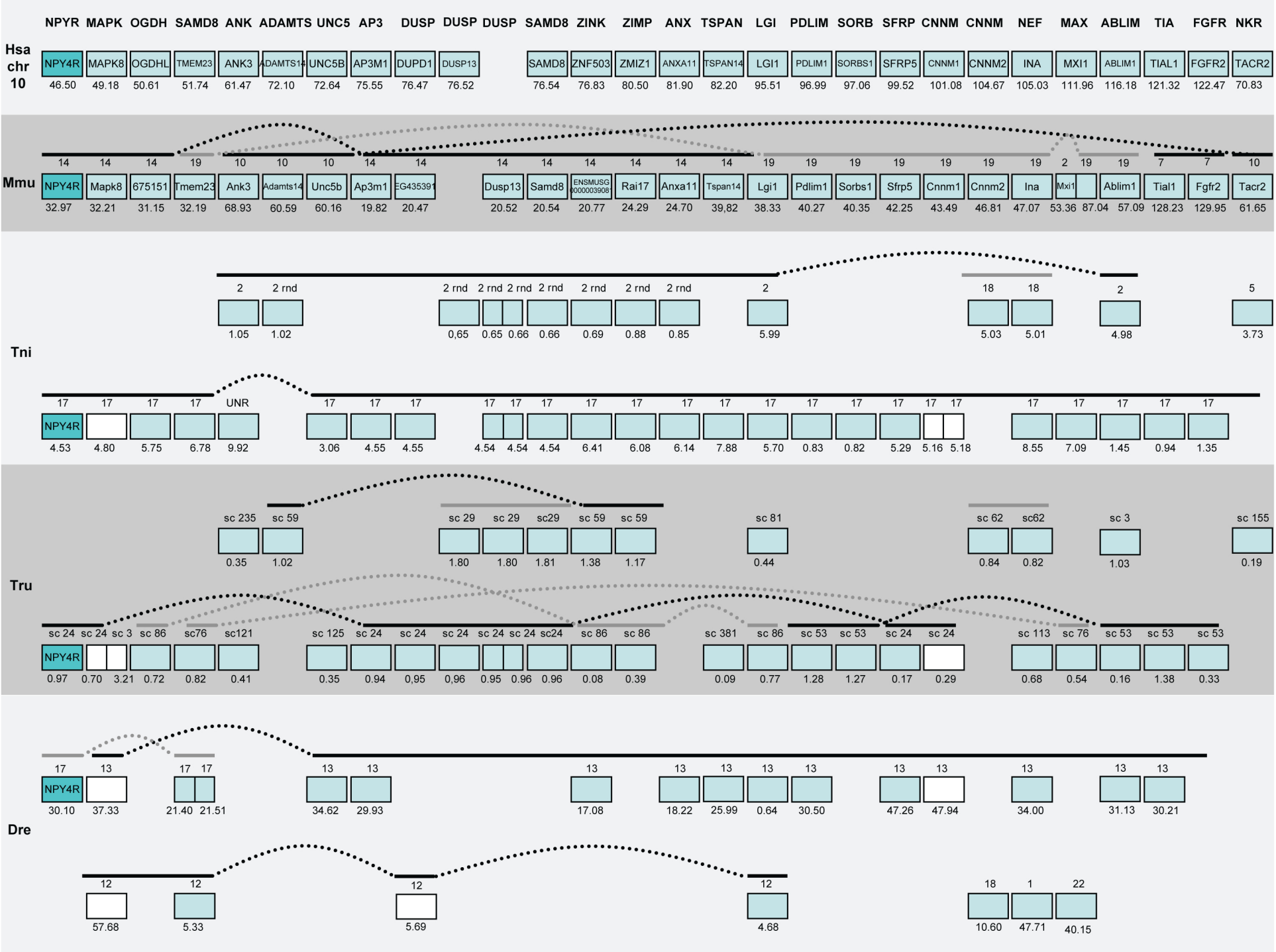
NPYR MAPK OGDH SAMD8 ANK ADAMTS UNC5 AP3 DUSP DUSP DUSP SAMD8 ZINK ZIMP ANX TSPAN LGI PDLIM SORB SFRP CNNM CNNM NEF MAX ABLIM TIA FGFR NKR
Hsa chr 10
NPY4R MAPK8 OGDHL TMEM23 ANK3 ADAMTS14 UNC5B AP3M1 DUPD1 DUSP13 SAMD8 ZNF503 ZMIZ1 ANXA11 TSPAN14 LGI1 PDLIM1 SORBS1 SFRP5 CNNM1 CNNM2 INA MXI1 ABLIM1 TIAL1 FGFR2 TACR2
46.50 49.18 50.61 51.74 61.47 72.10 72.64 75.55 76.47 76.52 76.54 76.83 80.50 81.90 82.20 95.51 96.99 97.06 99.52 101.08 104.67 105.03 111.96 116.18 121.32 122.47 70.83
Mmu
14 14 14 19 10 10 10 14 14 14 14 14 14 14 14 19 19 19 19 19 19 19 2 19 19 7 7 10
NPY4R Mapk8 675151 Tmem23 Ank3 Adamts14 Unc5b Ap3m1 EG435391 Dusp13 Samd8 ENSMUSG00000039081 Rai17 Anxa11 Tspan14 Lgi1 Pdlim1 Sorbs1 Sfrp5 Cnnm1 Cnnm2 Ina Mxi1 Ablim1 Tial1 Fgfr2 Tacr2
32.97 32.21 31.15 32.19 68.93 60.59 60.16 19.82 20.47 20.52 20.54 20.77 24.29 24.70 39,82 38.33 40.27 40.35 42.25 43.49 46.81 47.07 53.36 87.04 57.09 128.23 129.95 61.65
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1.05 1.02 0,65 0.65 0.66 0.66 0.69 0.88 0.85 5.99 5.03 5.01 4.98 3.73
17 17 17 17 UNR 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
NPY4R
4.53 4.80 5.75 6.78 9.92 3.06 4.55 4.55 4.54 4.54 4.54 6.41 6.08 6.14 7.88 5.70 0.83 0.82 5.29 5.16 5.18 8.55 7.09 1.45 0.94 1.35
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sc 235 sc 59 sc 29 sc 29 sc29 sc 59 sc 59 sc 81 sc 62 sc62 sc 3 sc 155
0.35 1.02 1.80 1.80 1.81 1.38 1.17 0.44 0.84 0.82 1.03 0.19
sc 24 sc 24 sc 3 sc 86 sc76 sc121 sc 125 sc 24 sc 24 sc 24 sc 24 sc 24 sc24 sc 86 sc 86 sc 381 sc 86 sc 53 sc 53 sc 24 sc 24 sc 113 sc 76 sc 53 sc 53 sc 53
NPY4R
0.97 0.70 3.21 0.72 0.82 0.41 0.35 0.94 0,95 0,96 0.95 0.96 0.96 0.08 0.39 0.09 0.77 1.28 1.27 0.17 0.29 0.68 0.54 0.16 1.38 0.33
Dre
17 13 17 17 13 13 13 13 13 13 13 13 13 13 13 13 13 13
NPY4R
30.10 37.33 21.40 21.51 34.62 29.93 17.08 18.22 25.99 0.64 30.50 47.26 47.94 34.00 31.13 30.21
12 12 12 12 18 1 22
57.68 5.33 5.69 4.68 10.60 47.71 40.15