

Subtel-only blocks

block	query sequence			len	aligned sequences	%ID	Named Transcripts
1	1p	1	25292	25292	17q: 19570 - 43429 5q: 15084 - 41588 6q: 9260 - 33611 16q: 12682 - 13816	97.61 98.10 98.42 97.15	sim to protein phosphatase 1 inhibitor subunit 2
2	15q	29000	116788	87789	19p: 20386 - 109078 6p: 1 - 58403 11p: 1 - 39455 6q: 66845 - 117940 8p: 32210 - 81762 1p: 67656 - 112213 5q: 77269 - 122537	98.33 97.97 98.00 97.86 97.84 97.92 97.86	OR4F3, OR4F4, OR4F5, OR4F29, OR4F21, OR4F16, OR4F17, C6orf88
3	1p	245037	280290	35254	8p: 194471 - 231807	97	
3a	1p	237884	242735	4852	8p: 134360 - 139200	98	Znf596
3b	1p	231380	233742	2363	8p: 279211 - 281570	97	FAM87B
4	1q	12455	15423	2969	6p: 95196 - 98868 5p: 2433 - 5242 4q: 8814 - 11635 13q: 2258 - 5129 4p: 17171 - 22773 2p: 6592 - 10454	93.21 92.86 90.73 91.40 91.14 91.24	sim to RPL23AP7 (multiple transcripts)
5	2p	6601	23214	16614	13q: 2258 - 15153 21q: 19455 - 21054 5p: 2449 - 16995 4p: 17171 - 27400 6p: 95211 - 98868 1q: 12594 - 15423 4q: 8814 - 14548	91.23 92.00 91.02 90.49 91.07 91.22 90.94	
6	3q	60958	99281	38324	9q: 25703 - 62989 7p: 75073 - 141501 16q: 105940 - 131364 11p: 115754 - 129425 17p: 1 - 21439	97.89 97.71 97.15 97.35 97.69	sim to RPL23AP7 (multiple transcripts)
6'	11p	125961	136823	10863	17p: 21439 - 29146	97.98	SCGB1C1
7	4q	15654	43535	27882	10q: 14673 - 88676	*	DUX4
8	4q	111450	125571	14122	1p: 286892 - 291767 10p: 30149 - 56397 18p: 34540 - 42205 9q: 73305 - 75051 3q: 111837 - 116459 16q: 131398 - 136973	91.33 94.60 93.00 93.47 92.00 93.47	TUBB4q
9	5p	4451	16125	11675	4q: 8933 - 14656 4p: 19173 - 24814 13q: 5070 - 11775 2p: 9585 - 16953	92 92 92.42 91.26	
10	2q	42832	91779	48948	8p: 290588 - 328066	97.14	FBOXO25
11	9q	88452	124137	35686	18p: 30838 - 69538 10p: 39691 - 74445 Xq+Yq: 18812 - 30108 16p: 17757 - 26950 16q: 138869 - 143475 4q: 100856 - 109294	95.68 95.37 92.18 91.88 94.63 91.26	IL9R
12	12p	27892	42389	14498	20q: 3777 - 18388	97.89	IQSEC3