

X chromosome BACs

| Pool | BAC | Start* | Finish | Genes |
|--------|---------------|------------|-----------|------------------------|
| 1 | RP11-102P23 | 80307324 | 80471160 | SH3BGRL |
| | RP11-73009 | 84981302 | 85164535 | CHM |
| | RP11-471G13 | 94723861 | 94900251 | None |
| | RP11-358K18 | 94891268 | 95033143 | None |
| | RP11-402K9 | 99210925 | 99386056 | None |
| | RP11-99E24 | 99373843 | 99554021 | PCDH19 |
| 2 | RP11-643F6 | 99643781 | 99890275 | TNMD, TSPAN6 + SYTL4 |
| | RP11-794023 | 100121535 | 100322783 | TRMT2B, TMEM35 + CENPI |
| | RP11-192G13 | 100579944 | 100766192 | ARMCX4, ARMCX1 + ARMC6 |
| | RP11-353J17 | 101454691 | 101642126 | NXF2B |
| | RP11-398N9 | 102643087 | 102806818 | MORF4L2 |
| | RP11-541I12 | 103373831 | 103557164 | ESX1 |
| | RP11-647M7 | 105695984 | 105872849 | CXORF57 + RNF128 |
| | RP11-150F24 | 105832705 | 106002508 | RNF128 + TBC1D8B |
| 3 | RP11-425O12 | 109203067 | 109404393 | AMMECR1 |
| | RP11-12D18 | ~109140001 | 109290000 | TMEM164 |
| | RP11-733H21 | 109968123 | 110073451 | PAK3 |
| | RP11-1066D24 | 111533817 | 111631587 | ZCCHC16 |
| | RP11-485D13 | 113641141 | 113824886 | HTR2C |
| 4 | RP11-268A15 | 114639781 | 114792452 | PLS3 |
| | 2272M5 | 114 | | DXZ4 |
| | RP11-761E20 | 114943586 | 115133311 | DXZ4 distal |
| | RP11-517D11 | 115089412 | 115231372 | AGTR2 |
| | RP11-27G9 | 115938027 | 116104434 | None |
| | RP11-197B12 | 117211298 | 117315606 | None |
| | ** RP11-423G5 | 117448177 | 117632053 | DOCK11 |
| 5 | RP11-166C19 | 119 | | CT47 |
| | RP11-100L19 | 122926111 | 123093111 | STAG2 |
| | RP11-13E5 | 124998007 | 125165297 | WDR40C |
| | RP11-388O17 | 129 | | SUHW3 |
| | RP11-754H22 | 130636029 | 130790513 | X130 array |
| 6 | RP11-274K13 | 134872885 | 135024305 | MMGT1 + SLC9A6 |
| | RP11-535K18 | 135051131 | 135233538 | FHL1, MAP7D3 + GPR112 |
| | 82404 | 153 | | H2A.Bbd & F8c |
| | 1047B3 | 153 | | H2A.Bbd & F8c |
| | 968D14 | 154 | | H2A.Bbd |
| | 999H8 | 154 | | H2A.Bbd |
| 905N17 | 154 | | H2A.Bbd | |
| 933C9 | 154 | | H2A.Bbd | |

*Coordinates based on hg18. Precise start and finish base pairing given for those BACs that are sequenced.

Coordinates of non-sequenced BACs are based on FISH data and gene content.

**DOCK11 containing BAC clone that frequently overlapped with WSTF enrichment at the Xi.