

Table S3. Primers

| Case | Name | Sequence (5'->3') | Annealing temperature [°C] | Used for |
|------|--------------|---------------------------|----------------------------|--------------------------------------|
| #4 | del(17)_FOR1 | AGGGAAAACCCAGATACCAAAC | 60.7 | primer walking |
| | del(17)_FOR2 | CCCACCACTACTGCTATACTC | 62.1 | primer walking, breakpoint spanning |
| | del(17)_FOR3 | TGCATCTCTCAACTCAGACC | 60.3 | primer walking |
| | del(17)_REV1 | TTGCCATTGATAGGGTAGGGC | 59.8 | primer walking, breakpoint spanning |
| | del(17)_REV2 | TGCCTACAGTATATTAGGTGAAAGC | 59.7 | primer walking |
| | del(17)_REV3 | TGTGGGAAGGGTTTAGACGC | 59.8 | primer walking |
| #9 | del(19)_FOR1 | GACCTGACTCAAGCCCCTC | 61.4 | primer walking, break point spanning |
| | del(19)_FOR2 | GAGTGTCTCAGGGCTTCCC | 61.4 | primer walking |
| | del(19)_REV1 | CAGGACCCATCGACCACAA | 59.4 | primer walking |
| | del(19)_REV2 | GAGAGGTGGGAAGGAGGGTC | 63.5 | primer walking |
| | del(19)_REV3 | GGGTGCCAAGAAAGGGAACT | 59.4 | break point spanning |
| #15 | t(2;12)_FOR1 | ATCAGCTTCACCTTGGGTCG | 59.4 | primer walking |
| | t(2;12)_FOR2 | CCTATTCTGAGGCTGAATTATGGC | 61. | break point spanning |
| | t(2;12)_REV1 | CAGTGCCCTTCATCCATT | 59.4 | primer walking |
| | t(2;12)_REV2 | ACAAGAGGCAACCCCATCAG | 59.4 | break point spanning |
| #22 | t(8;12)_FOR1 | TTGAGGGCAGTTGTCAAAGC | 57.3 | primer walking |
| | t(8;12)_FOR2 | CTCCTCTCCCTGCTCTCAT | 61.4 | primer walking |
| | t(8;12)_FOR3 | TAATGAGAGACGGCAGCCG | 58.8 | primer walking/sequencing |
| | t(8;12)_FOR4 | AAATTCCCTAAATGCTAACAA | 51.7 | primer walking/sequencing |
| | t(8;12)_FOR5 | ATACAGGTCTAGGCATGCTTTGT | 59.3 | primer walking/sequencing |
| | t(8;12)_FOR6 | CCAATTCTGAAAACACTTTGTGTG | 58.5 | primer walking/sequencing |
| | t(8;12)_FOR7 | TGCACATACATCATCTCGCGT | 57.90 | break point spanning |
| | t(8;12)_REV1 | CCTTCCTGCGTGTGGATGT | 59.4 | primer walking |
| | t(8;12)_REV2 | TCTCCAAAGCCATCAGCCAG | 59.4 | primer walking |
| | t(8;12)_REV3 | AATGTACAATATGTGCTAACT | 50.6 | primer walking/sequencing |
| | t(8;12)_REV4 | GAGGGCAGGGACTCTATGAC | 61.40 | primer walking/sequencing |
| | t(8;12)_REV5 | CCTCAGATACTGGCAGC | 60.99 | primer walking/sequencing |
| | t(8;12)_REV6 | CACCCCATTAACGGAGGACA | 59.40 | break point spanning |