**Table S5. Frequency of recurrent missegregation events.**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Chromosome | Whole Chr aneuploidy | | | Whole Chr LOH | |
|  | 1N | 3N | 4N | AA | BB |
| 1 | 6 | 5 | 1 | 9 | 0 |
| 2 | 6 | 4 | 0 | 7 | 0 |
| 3 | 6 | 1 | 0 | 6 | 8 |
| 4 | 5 | 6 | 3 | 6 | 0 |
| 5 | 6 | 8 | 1 | 6 | 4 |
| 6 | 4 | 9 | 3 | 6 | 0 |
| 7 | 4 | 8 | 2 | 5 | 0 |
| R | 6 | 5 | 0 | 7 | 6 |

Data summary for mice with N ≥ 12 (9 mice). Number of recurrent genotypes are shown separately for whole chromosome aneuploidy (1N, monosomic, 3N, trisomic, 4N, tetrasomic) and whole chromosome LOH (AA, homozygous allele A, BB, homozygous allele B)