**Supplementary Table 3.** *1.688* satDNA euchromatic insertion distribution of BLAST hits using species-specific sequences as query in the annotated genomes of *D. melanogaster, D. simulans* and *D. yakuba*.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | **X** | **2L** | **2R** | **3L** | **3R** | **4** | **Y** |
| ***D. melanogaster*** | 561 | 29 | 26 | 46 | 13 | - | - |
| ***D. simulans*** | 366 | 23 | 18 | 13 | 36 | - | - |
| ***D. yakuba*** | 807 | 144 | 72 | 189 | 111 | - | - |