Supplementary Table S3. Repeat content of the genomes of *Hypsibius dujardini* and *Ramazzottius varieornatus*

|  |  |  |  |
| --- | --- | --- | --- |
| Category | Term | *H. dujardini* | *R. varieornatus* |
| Simple | #elements | 65,638 | 3,301 |
| #length (bp) | 5,391,682 | 137,297 |
| %genome | 5.18 | 0.25 |
| Unclassified | #elements | 158,522 | 65,730 |
| #length (bp) | 24,232,698 | 11,020,138 |
| %genome | 23.27 | 19.74 |