**Table S3.** Statistics of the repeat sequences in the genome assembly of the Japanese eel.

|  |  |  |
| --- | --- | --- |
| **Type** | **Repeat Size (bp)** | **Percentage of genome (%)** |
| Trf | 74,462,905 | 6.57 |
| Repeatmasker | 85,570,095 | 7.55 |
| Proteinmask | 27,406,207 | 2.42 |
| De novo | 200,071,462 | 17.66 |
| Total | 259,849,065 | 22.94 |