Supplementary Table S6. Mapping proportion of the *Hypsibius dujardini* Trinity assembled transcriptome

|  |  |  |  |
| --- | --- | --- | --- |
| ***Hypsibius dujardini*** |  |  |  |
| **10k Individuals** | Total number of transcripts assembled | Number (proportion) mapped to the *H. dujardini* genome | Number (proportion) mapped to the *H. dujardini* transcriptome |
| act-1 | 66886 | 59283 (88.63%) | 43608 (65.20%) |
| act-2 | 68941 | 60998 (88.48%) | 45168 (65.52%) |
| act-3 | 73982 | 68767 (99.74%) | 52217 (70.58%) |
| tun-1 | 63670 | 59503 (93.46%) | 43667 (68.58%) |
| tun-2 | 77919 | 74304 (95.37%) | 54768 (61.59%) |
| tun-3 | 149853 | 117883 (78.67%) | 72078 (48.10%) |