|  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Recovery factor | $x\_{1}$ [codons] | $x\_{2}$ [codons] | $$x\_{3}$$[codons] | mean($v\_{1}$) [codons/second] | mean($v\_{2}$) [codons/second] | KS-test P value | Mean($v\_{1}$,$ v\_{2}$) [codons/second] | Median of $v\_{2}$/$v\_{1}$ [codons/second] | Median of |$v\_{1}-v2|/min⁡(v\_{1},v\_{2})$ | Number of genes $x\_{1}<x\_{2}<x\_{3}$ |
| 0.4 | 142+/-64 | 267+/-65 | 445+/-87 | 4.1+/-2.2 | 5.9+/-2.4 | ‎<1.85\*10-36 | 5.0+/-1.3 | 1.45 | 0.81 | 692 |
| 0.5 | ‎177+/-74 | ‎308+/-72 | ‎489+/-85 | ‎4.3+/-2.6 | ‎6.0+/-2.5 | ‎<1.78\*10-24 | 5.2+/-1.2 | ‎1.37 | 0.82 | ‎692‎ |
| 0.6 | 207+/-81 | 347+/-72 | 525+/-88 | 4.7+/-2.7 | 5.9+/-2.6 | ‎<5.96\*10-15 | 5.3+/-0.9 | 1.24 | 0.81 | 691 |
| 0.7 | 237+/-88 | 389+/-77 | 561+/-93 | 5.0+/-2.9 | 5.7+/-2.7 | ‎<2.52\*10-07 | 5.4+/-0.5 | 1.14 | 0.86 | 677 |
| 0.8 | 270+/-96 | 431+/-84 | 598+/-92 | 5.4+/-3.2 | 5.6+/-2.8 | ‎<0.0296 | 5.5+/-0.1 | 1.06 | 0.91 | 651 |