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|---|---------------------------------------|----------------------------------|---------------------------|-------------------------|-----------------------|
| | by clustering | recovered in <i>G. raimondii</i> | recovered in sister genus | | |
| G. kirkii gains | 211 | 4 (2%) | 12 (6%) | | |
| G. kirkii gains with "good" translations | | 0 (0%) | 2 (1%) | | |
| | | | | | |
| K. drynarioides gains | 394 | 6 (2%) | 64 (16%) | | |
| K. drynarioides gains with "good" translations | | 0 (0%) | 33 (8%) | | |
| | | | | | |
| | by clustering | recovered in itself | | | |
| G. kirkii losses | 2144 | 1,465 (68%) | | | |
| G. kirkii losses with "good" translations | | 988 (46%) | | | |
| | | | | | |
| K. drynarioides losses | 1458 | 872 (60%) | | | |
| K. drynarioides losses with "good" translations | | 514 (35%) | | | |
| | | | | | |
| | | | | | |
| | <i>G. raimondii</i> unclustered genes | recovered in itself | "good" translations | overlapping annotations | valid and overlapping |
| G. kirkii unclustered verification | 10408 | 6229 | 3330 | 1983 | 1512 |
| K. drynarioides unclustered verification | 10408 | 6035 | 3513 | 2009 | 1527 |