

CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | |
|--------------------------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|---|
| Metaseiulus_occidentalis | AGA | False | False | 21 | -0.318456 | 9 | 0 | False | 1 | 1 | 0 |
| Halocynthia | CGC | False | False | 1 | 2.850000 | 1 | 0 | False | 1 | 1 | 0 |
| Schistosoma | AGG | False | False | 17 | -0.288922 | 9 | 0 | False | 1 | 1 | 0 |
| Pseudogarypus_banksi | AGA | False | False | 30 | -0.385204 | 9 | 0 | False | 1 | 1 | 0 |
| Ciona | AGG | False | False | 25 | -1.491803 | 10 | 0 | False | 1 | 1 | 0 |
| Tetrancistrum_nebulosi | AGA | False | False | 20 | -0.037179 | 6 | 0 | False | 1 | 1 | 0 |
| Versteria_mustelae | AGG | False | False | 7 | -0.007299 | 4 | 0 | False | 1 | 1 | 0 |
| Haplorchis_taichui | AGG | False | False | 22 | -0.168605 | 8 | 1 | False | 1 | 1 | 0 |