

CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|----------------|-----|-----------|------------|------------|----------|------------|------------|------------|------------|------------|-------|
| Halocynthia | AGT | False | False | 27 | 0.517405 | 5 | 19 | False | 1 | 1 | 0.005 |
| Schistosoma | AGT | False | False | 12 | 0.670213 | 5 | 4 | False | 1 | 1 | 0.051 |
| | TCT | False | False | 27 | 0.620472 | 4 | 10 | False | 2 | 4 | 0.050 |
| Terebratulina | TCT | False | False | 17 | 0.381313 | 5 | 10 | False | 1 | 1 | 0.005 |
| Caenorhabditis | TCT | False | False | 16 | 0.379404 | 4 | 8 | False | 1 | 1 | 0.013 |
| Myxine | AGC | False | False | 8 | 1.173184 | 3 | 5 | False | 1 | 1 | 0.051 |
| Ciona | AGT | False | False | 21 | 0.727642 | 5 | 14 | False | 1 | 2 | 0.005 |
| | AGT | False | False | 19 | 0.392377 | 5 | 9 | False | 2 | 2 | 0.009 |
| Fasciola | TCC | False | False | 4 | 0.156250 | 3 | 1 | False | 1 | 1 | 0.061 |
| | TCT | False | False | 27 | 0.689548 | 5 | 13 | False | 1 | 2 | 0.009 |
| Acropora | AGT | False | False | 13 | 0.735294 | 4 | 10 | False | 1 | 1 | 0.009 |
| Homo | AGT | False | False | 2 | 2.458333 | 2 | 1 | False | 1 | 1 | 0.153 |
| Trichinella | TCA | False | False | 22 | 0.165029 | 4 | 11 | False | 1 | 1 | 0.018 |
| | TCC | False | False | 11 | 0.681992 | 4 | 5 | False | 1 | 1 | 0.025 |