

CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|----------------|-----|-----------|------------|------------|----------|------------|------------|------------|------------|------------|-------|
| Fasciola | AAT | False | False | 19 | 0.620767 | 5 | 9 | False | 1 | 1 | 0.120 |
| Halocynthia | AAT | False | False | 25 | 0.972789 | 5 | 23 | False | 1 | 1 | 0.005 |
| Schistosoma | AAC | False | False | 2 | 2.369565 | 2 | 1 | False | 1 | 1 | 0.165 |
| Caenorhabditis | AAT | False | False | 31 | 0.632823 | 5 | 21 | False | 1 | 1 | 0.005 |
| Ciona | AAT | False | False | 35 | 0.964806 | 5 | 25 | False | 2 | 2 | 0.001 |