CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|-------------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|-------|
| Fasciola | AGT | False | False | 19 | -0.500000 | 5 | 9 | False | 1 | 1 | 0.013 |
| | TCT | False | False | 27 | -0.611544 | 5 | 13 | False | 1 | 1 | 0.013 |
| Halocynthia | AGT | False | False | 27 | -0.386076 | 5 | 19 | False | 1 | 1 | 0.005 |
| Schistosoma | TCA | False | False | 11 | -0.192157 | 5 | 7 | False | 1 | 1 | 0.069 |
| Acropora | TCC | False | False | 7 | -0.190476 | 4 | 3 | False | 1 | 1 | 0.017 |