# CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|---|---|---|---|---|---|---|---|---|---|---|---|
| **MmagnB** | **TAC** | False | False | 115 | -1.871968 | 13 | 104 | False | 1 | 1 | 0.000 |
| **MmagnA** | **TAC** | False | False | 119 | -1.874893 | 13 | 108 | False | 1 | 1 | 0.000 |
| **MingeA** | **TAT** | False | False | 142 | -1.867855 | 13 | 124 | False | 1 | 1 | 0.000 |
| **MstarA** | **TAT** | False | False | 145 | -1.872307 | 13 | 129 | False | 1 | 1 | 0.000 |
| **CsalmA** | **TAC** | False | False | 2 | 0.519608 | 2 | 1 | False | 1 | 1 | 0.065 |
| **MtetrA** | **TAC** | False | False | 136 | -1.837884 | 13 | 115 | False | 1 | 1 | 0.000 |
| **SpombA** | **TAT** | False | False | 55 | -1.753374 | 6 | 41 | False | 1 | 1 | 0.000 |
| **CzempA** | **TAT** | False | False | 82 | -1.804701 | 6 | 64 | False | 1 | 1 | 0.000 |
| **SoctoA** | **TAC** | False | False | 17 | -1.677384 | 5 | 13 | False | 1 | 1 | 0.000 |
| **OanguA** | **TAT** | False | False | 124 | -1.896237 | 12 | 112 | False | 1 | 1 | 0.000 |
| **ScryoA** | **TAC** | False | False | 27 | -1.738701 | 4 | 19 | False | 1 | 1 | 0.000 |