# CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|---|---|---|---|---|---|---|---|---|---|---|---|
| **KmarxA** | **GCC** | False | False | 4 | -0.476082 | 1 | 2 | False | 1 | 1 | 0.306 |
| **CbohiA** | **GCT** | False | False | 108 | -1.896798 | 13 | 93 | False | 1 | 1 | 0.000 |
| **MmagnB** | **GCC** | False | False | 31 | -1.756180 | 9 | 25 | False | 1 | 1 | 0.000 |
| **CphanA** | **GCT** | False | False | 108 | -1.890998 | 12 | 71 | False | 1 | 1 | 0.000 |
| **CgallA** | **GCA** | False | False | 58 | -1.818653 | 13 | 38 | False | 1 | 1 | 0.000 |
| **MmagnA** | **GCC** | False | False | 31 | -1.756180 | 9 | 25 | False | 1 | 1 | 0.000 |
| **CdefoA** | **GCA** | False | False | 61 | -1.842908 | 13 | 38 | False | 1 | 1 | 0.000 |
| **YlipoB** | **GCT** | False | False | 98 | -1.876068 | 12 | 63 | False | 1 | 1 | 0.000 |
| **YlipoA** | **GCT** | False | False | 98 | -1.876068 | 12 | 63 | False | 1 | 1 | 0.000 |
| **CalimA** | **GCA** | False | False | 42 | -1.731532 | 9 | 27 | False | 1 | 1 | 0.000 |
| **CglabA** | **GCA** | False | False | 37 | -1.694060 | 6 | 30 | False | 1 | 1 | 0.000 |
| **SoctoA** | **GCA** | False | False | 23 | -1.529876 | 5 | 12 | False | 1 | 1 | 0.000 |
| **MtetrA** | **GCC** | False | False | 60 | -1.855773 | 10 | 43 | False | 1 | 1 | 0.000 |
| **CcastA** | **GCC** | False | False | 4 | -0.432367 | 4 | 3 | False | 1 | 1 | 0.000 |
| **NdelpA** | **GCA** | False | False | 35 | -1.811768 | 6 | 30 | False | 1 | 1 | 0.000 |
| **ScryoA** | **GCT** | False | False | 54 | -1.506043 | 6 | 27 | False | 1 | 1 | 0.000 |