

CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|--------|-----|-----------|------------|------------|-----------|------------|------------|------------|------------|------------|-------|
| CnorvA | TGT | False | False | 23 | -0.508318 | 10 | 14 | False | 2 | 2 | 0.038 |
| LmeyeA | TGT | False | False | 13 | -0.700447 | 5 | 8 | False | 1 | 1 | 0.000 |
| CfrijA | TGT | False | False | 14 | -0.573209 | 8 | 13 | False | 1 | 1 | 0.000 |
| CgallA | TGT | False | False | 16 | -0.144881 | 8 | 8 | False | 2 | 2 | 0.096 |
| CviswA | TGT | False | False | 15 | -0.651045 | 8 | 13 | False | 1 | 1 | 0.000 |
| CdefoA | TGT | False | False | 16 | -0.144881 | 8 | 8 | False | 2 | 2 | 0.096 |
| PpastA | TGT | False | False | 19 | -0.356188 | 10 | 14 | False | 2 | 2 | 0.035 |
| MstarA | TGT | False | False | 19 | -0.159838 | 10 | 9 | False | 2 | 3 | 0.149 |
| AgossA | TGT | False | False | 13 | -0.316928 | 6 | 9 | False | 2 | 2 | 0.101 |
| LkluyF | TGT | False | False | 13 | -0.709918 | 5 | 9 | False | 1 | 1 | 0.000 |
| LkluyE | TGT | False | False | 13 | -0.709918 | 5 | 9 | False | 1 | 1 | 0.000 |
| LkluyD | TGT | False | False | 12 | -0.686795 | 5 | 8 | False | 1 | 1 | 0.000 |
| SpombA | TGT | False | False | 13 | -0.348221 | 5 | 4 | False | 1 | 1 | 0.000 |
| CvartA | TGT | False | False | 25 | -0.520753 | 10 | 14 | False | 2 | 2 | 0.044 |
| LkluyA | TGT | False | False | 13 | -0.709918 | 5 | 9 | False | 1 | 1 | 0.000 |
| CtetrA | TGT | False | False | 14 | -0.573209 | 8 | 13 | False | 1 | 1 | 0.000 |
| ClusiA | TGT | False | False | 16 | -0.344151 | 10 | 14 | False | 2 | 2 | 0.088 |
| AacerA | TGT | False | False | 5 | -0.208812 | 2 | 4 | False | 1 | 1 | 0.244 |
| MmagnB | TGT | False | False | 19 | -0.349794 | 10 | 10 | False | 2 | 2 | 0.025 |
| CgigaA | TGC | False | False | 6 | -0.133333 | 5 | 5 | False | 1 | 1 | 0.000 |
| MmagnA | TGT | False | False | 19 | -0.349794 | 10 | 10 | False | 2 | 2 | 0.025 |
| KmarxA | TGT | False | False | 2 | 0.726027 | 1 | 1 | False | 1 | 1 | 0.334 |
| LstarA | TGT | False | False | 11 | 0.956092 | 7 | 4 | False | 4 | 4 | 0.304 |
| MguilA | TGT | False | False | 15 | -0.595203 | 10 | 14 | False | 1 | 1 | 0.000 |
| WmucoA | TGT | False | False | 19 | -0.635702 | 10 | 14 | False | 1 | 1 | 0.000 |
| CalimA | TGT | False | False | 16 | -0.144881 | 8 | 8 | False | 2 | 2 | 0.096 |
| CsantA | TGT | False | False | 21 | -0.050857 | 10 | 6 | False | 3 | 3 | 0.000 |
| MtetrA | TGT | False | False | 18 | -0.270964 | 10 | 10 | False | 2 | 2 | 0.022 |
| CsaraA | TGT | False | False | 12 | -0.484230 | 8 | 11 | False | 1 | 1 | 0.000 |
| CtropB | TGT | False | False | 13 | -0.536833 | 8 | 12 | False | 1 | 1 | 0.000 |
| CsuavA | TGT | False | False | 24 | -0.498881 | 10 | 13 | False | 2 | 2 | 0.000 |
| CtropA | TGT | False | False | 12 | -0.476809 | 8 | 11 | False | 1 | 1 | 0.000 |
| MfariA | TGT | False | False | 16 | -0.380759 | 10 | 14 | False | 2 | 2 | 0.000 |
| PkluyA | TGT | False | False | 24 | 0.121795 | 9 | 15 | False | 4 | 5 | 0.198 |
| MfariB | TGT | False | False | 14 | -0.595532 | 9 | 13 | False | 1 | 1 | 0.000 |
| PcanaA | TGT | False | False | 21 | -0.633299 | 10 | 14 | False | 1 | 1 | 0.000 |
| BcaliA | TGT | False | False | 21 | -0.633299 | 10 | 14 | False | 1 | 1 | 0.000 |
| LanspB | TGT | False | False | 14 | -0.721027 | 5 | 9 | False | 1 | 1 | 0.000 |

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|--------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|-------|
| CphanA | TGT | False | False | 16 | -0.144881 | 8 | 8 | False | 2 | 2 | 0.096 |
| LanspA | TGT | False | False | 10 | -0.588235 | 5 | 6 | False | 1 | 1 | 0.000 |
| CsojaA | TGT | False | False | 7 | -0.106312 | 5 | 6 | False | 1 | 1 | 0.000 |
| YlipoB | TGT | False | False | 16 | -0.144881 | 8 | 8 | False | 2 | 2 | 0.096 |
| YlipoA | TGT | False | False | 16 | -0.144881 | 8 | 8 | False | 2 | 2 | 0.096 |
| CsalmA | TGT | False | False | 24 | -0.460055 | 10 | 11 | False | 2 | 2 | 0.030 |
| CjadiA | TGT | False | False | 24 | -0.574497 | 10 | 14 | False | 2 | 2 | 0.043 |
| AgossB | TGT | False | False | 13 | -0.316928 | 6 | 9 | False | 2 | 2 | 0.101 |
| BcustA | TGT | False | False | 24 | 0.282439 | 10 | 13 | False | 4 | 5 | 0.001 |
| LkluyC | TGT | False | False | 13 | -0.709918 | 5 | 9 | False | 1 | 1 | 0.000 |
| OanguA | TGT | False | False | 17 | 0.070755 | 9 | 10 | False | 3 | 3 | 0.000 |
| ScryoA | TGT | False | False | 15 | -0.552803 | 6 | 4 | False | 1 | 1 | 0.000 |
| CneerA | TGT | False | False | 14 | -0.573209 | 8 | 13 | False | 1 | 1 | 0.000 |
| LnothA | TGT | False | False | 14 | -0.721027 | 5 | 9 | False | 1 | 1 | 0.000 |
| CsakeA | TGT | False | False | 16 | -0.629479 | 10 | 15 | False | 1 | 1 | 0.000 |
| HuvarA | TGT | False | False | 12 | -0.391410 | 4 | 5 | False | 1 | 1 | 0.001 |
| KtherA | TGT | False | False | 13 | -0.668161 | 5 | 9 | False | 1 | 1 | 0.000 |
| KlactA | TGT | False | False | 12 | -0.712097 | 4 | 8 | False | 1 | 1 | 0.001 |
| CzempA | TGT | False | False | 14 | 0.073120 | 6 | 6 | False | 2 | 3 | 0.190 |
| SpastA | TGT | False | False | 10 | -0.708861 | 6 | 7 | False | 1 | 1 | 0.000 |
| SoctoA | TGT | False | False | 14 | -0.450419 | 6 | 4 | False | 1 | 1 | 0.000 |
| LdasiA | TGT | False | False | 14 | -0.721027 | 5 | 9 | False | 1 | 1 | 0.000 |
| MingeA | TGT | False | False | 18 | -0.329448 | 10 | 9 | False | 2 | 2 | 0.023 |
| LkluyB | TGT | False | False | 2 | 0.726027 | 1 | 1 | False | 1 | 1 | 0.334 |