

CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|---------------|------------|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|-------|
| CnorvA | TGT | False | False | 23 | -0.612754 | 10 | 14 | False | 1 | 1 | 0.000 |
| BcaliA | TGT | False | False | 21 | -0.605209 | 10 | 14 | False | 1 | 1 | 0.000 |
| BcustA | TGC | False | False | 2 | 1.079208 | 2 | 1 | False | 1 | 1 | 0.073 |
| CjadiA | TGT | False | False | 24 | -0.540045 | 10 | 14 | False | 1 | 1 | 0.055 |
| SjapoA | TGT | False | False | 9 | 0.527215 | 5 | 3 | False | 1 | 1 | 0.009 |
| CchauA | TGC | False | False | 1 | 3.368932 | 1 | 0 | False | 1 | 1 | 0.469 |
| CsuavA | TGT | False | False | 24 | -0.502908 | 10 | 13 | False | 1 | 1 | 0.000 |
| CzempA | TGT | False | False | 14 | -0.562674 | 6 | 6 | False | 1 | 1 | 0.000 |
| PkluyA | TGT | False | False | 24 | -0.781593 | 9 | 15 | False | 1 | 1 | 0.000 |
| OanguA | TGT | False | False | 17 | -0.838275 | 9 | 10 | False | 1 | 1 | 0.000 |
| PcanaA | TGT | False | False | 21 | -0.605209 | 10 | 14 | False | 1 | 1 | 0.000 |
| CvartA | TGT | False | False | 25 | -0.524604 | 10 | 14 | False | 1 | 1 | 0.000 |