CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|--------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|------|
| CnorvA | AAT | False | False | 89 | -2.469817 | 13 | 68 | False | 3 | 3 | 0 |
| HuvarA | AAC | False | False | 35 | -2.422428 | 6 | 23 | False | 2 | 2 | 0 |
| | AAT | False | False | 29 | -2.116341 | 6 | 17 | False | 1 | 1 | 0 |
| CsalmA | AAT | False | False | 75 | -2.624964 | 10 | 51 | False | 1 | 1 | 0 |
| WmucoA | AAC | False | False | 25 | -2.635698 | 8 | 22 | False | 1 | 1 | 0 |
| | AAT | False | False | 60 | -2.518797 | 10 | 48 | False | 2 | 2 | 0 |
| CjadiA | AAT | False | False | 70 | -2.525551 | 12 | 55 | False | 3 | 3 | 0 |
| SjapoA | AAT | False | False | 70 | -2.450747 | 7 | 36 | False | 2 | 2 | 0 |
| CsaraA | AAT | False | False | 71 | -2.842823 | 12 | 65 | False | 1 | 1 | 0 |
| SpombA | AAC | False | False | 8 | -1.392421 | 4 | 1 | False | 1 | 1 | 0 |
| | AAT | False | False | 58 | -2.554970 | 6 | 40 | False | 1 | 1 | 0 |
| CsuavA | AAT | False | False | 66 | -2.458492 | 10 | 51 | False | 3 | 3 | 0 |
| BcaliA | AAT | False | False | 87 | -2.475877 | 12 | 66 | False | 3 | 3 | 0 |
| WpijpA | AAC | False | False | 37 | -2.818932 | 7 | 35 | False | 1 | 1 | 0 |
| PcanaA | AAT | False | False | 83 | -2.501134 | 12 | 64 | False | 3 | 3 | 0 |
| CvartA | AAT | False | False | 75 | -2.549947 | 11 | 57 | False | 3 | 3 | 0 |