CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|--------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|-------|
| CzempA | AGA | False | False | 21 | -1.651355 | 6 | 16 | False | 1 | 1 | 0.002 |
| CphanA | AGA | False | False | 51 | -1.810886 | 12 | 46 | False | 1 | 1 | 0.003 |
| CgallA | AGA | False | False | 51 | -1.810886 | 12 | 46 | False | 1 | 1 | 0.003 |
| CdefoA | AGA | False | False | 51 | -1.810886 | 12 | 46 | False | 1 | 1 | 0.003 |
| YlipoB | AGA | False | False | 50 | -1.807976 | 12 | 45 | False | 1 | 1 | 0.003 |
| YlipoA | AGA | False | False | 50 | -1.807976 | 12 | 45 | False | 1 | 1 | 0.003 |
| CalimA | AGA | False | False | 51 | -1.810886 | 12 | 46 | False | 1 | 1 | 0.003 |
| SpombA | AGA | False | False | 27 | -1.805811 | 7 | 23 | False | 1 | 1 | 0.002 |
| BcustA | AGA | False | False | 41 | -1.732694 | 11 | 37 | False | 1 | 1 | 0.003 |
| CglabA | AGA | False | False | 34 | -1.822398 | 7 | 31 | False | 1 | 1 | 0.003 |
| CsubhB | CGC | False | False | 2 | 1.729885 | 2 | 1 | False | 1 | 1 | 0.042 |
| CsubhA | CGC | False | False | 2 | 1.729885 | 2 | 1 | False | 1 | 1 | 0.042 |
| NdelpA | AGA | False | False | 34 | -1.822398 | 7 | 31 | False | 1 | 1 | 0.003 |