CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|--------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|------|
| BcustA | GCT | False | False | 66 | -1.547920 | 11 | 51 | False | 1 | 1 | 0 |
| SjapoA | GCT | False | False | 61 | -1.563616 | 6 | 37 | False | 1 | 1 | 0 |
| SpombA | GCA | False | False | 22 | -1.616249 | 6 | 10 | False | 1 | 1 | 0 |
| CzempA | GCA | False | False | 35 | -1.562177 | 6 | 27 | False | 1 | 1 | 0 |
| SoctoA | GCT | False | False | 68 | -1.583085 | 6 | 37 | False | 1 | 1 | 0 |
| ScryoA | GCT | False | False | 54 | -1.463029 | 6 | 27 | False | 1 | 1 | 0 |