CoReTracker report

| | | Clad. Val | Reassigned | Tot. count | Telford | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob |
|--------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|-------|
| CsantA | AAA | False | False | 34 | -0.842313 | 12 | 26 | False | 1 | 1 | 0.116 |
| HuvarA | AAA | False | False | 20 | -0.761058 | 6 | 12 | False | 1 | 1 | 0.181 |
| CsalmA | AAA | False | False | 39 | -0.825029 | 11 | 33 | False | 1 | 1 | 0.208 |
| CzempA | AAA | False | False | 19 | -0.600514 | 7 | 14 | False | 1 | 1 | 0.209 |