

# CoReTracker report

|   |     | Clad. Val | Reassigned | Tot. count | Telford   | Tot. Genes | Use. count | Trans. Val | Rea. Genes | Rea. count | Prob  |
|---|-----|-----------|------------|------------|-----------|------------|------------|------------|------------|------------|-------|
| H | ATG | False     | False      | 43         | -0.109043 | 1          | 34         | False      | 1          | 1          | 0.213 |
| D | ATG | False     | False      | 33         | -0.205674 | 1          | 31         | False      | 1          | 1          | 0.132 |
| F | ATG | False     | False      | 40         | -0.189504 | 1          | 37         | False      | 1          | 1          | 0.183 |