## CoReTracker report

|        |     | Clad.<br>Val | Reassigned | Tot.<br>count | Telford   | Tot.<br>Genes | Use.<br>count | Trans.<br>Val | Rea.<br>Genes | Rea.<br>count | Prob |
|--------|-----|--------------|------------|---------------|-----------|---------------|---------------|---------------|---------------|---------------|------|
| HuvarA | CCT | False        | False      | 33            | -0.699424 | 5             | 29            | False         | 1             | 1             | 0    |
| OanguA | CCA | False        | False      | 48            | -0.846120 | 12            | 47            | False         | 1             | 1             | 0    |
| SjapoA | CCA | False        | False      | 18            | -0.648747 | 5             | 14            | False         | 1             | 1             | 0    |
| SpombA | CCA | False        | False      | 41            | -0.795719 | 7             | 36            | False         | 1             | 1             | 0    |
| NbaciA | CCA | False        | False      | 23            | -0.708160 | 5             | 21            | False         | 1             | 1             | 0    |
|        | CCT | False        | False      | 50            | -0.868602 | 7             | 49            | False         | 1             | 1             | 0    |
| CcastA | CCT | False        | False      | 36            | -0.826387 | 6             | 35            | False         | 1             | 1             | 0    |