

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>PpastA</b>	<b>GCA</b>	False	False	78	-1.567408	10	64	False	1	1	0
<b>CgigaA</b>	<b>GCT</b>	False	False	64	-1.540519	10	54	False	1	1	0
<b>SarboA</b>	<b>GCT</b>	False	False	50	-1.531968	6	40	False	1	1	0
<b>HuvarA</b>	<b>GCT</b>	False	False	59	-1.352922	6	39	False	1	1	0
<b>ScereA</b>	<b>GCT</b>	False	False	52	-1.491821	6	41	False	1	1	0
<b>SpastA</b>	<b>GCT</b>	False	False	47	-1.492872	6	35	False	1	1	0
<b>ScereC</b>	<b>GCT</b>	False	False	53	-1.469877	6	41	False	1	1	0
<b>KlactA</b>	<b>GCT</b>	False	False	54	-1.547016	6	42	False	1	1	0
<b>CsaraA</b>	<b>GCA</b>	False	False	40	-1.602907	10	37	False	1	1	0
<b>BcustA</b>	<b>GCT</b>	False	False	66	-1.372349	11	51	False	1	1	0
<b>ScereD</b>	<b>GCT</b>	False	False	51	-1.483871	6	40	False	1	1	0
<b>ScereE</b>	<b>GCT</b>	False	False	51	-1.483399	6	40	False	1	1	0
<b>CcastA</b>	<b>GCT</b>	False	False	57	-1.501840	6	46	False	1	1	0