

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>CsantA</b>	<b>GTT</b>	False	False	21	-1.715291	8	12	False	1	1	0.000
<b>LstarA</b>	<b>GTT</b>	False	False	62	-1.682041	9	36	False	1	2	0.000
<b>HuvarA</b>	<b>GTA</b>	False	False	53	-1.606431	6	24	False	1	1	0.055
	<b>GTT</b>	False	False	54	-1.454513	6	22	False	1	1	0.055
<b>BcustA</b>	<b>GTA</b>	False	False	103	-1.696386	12	48	False	1	1	0.000
<b>CjadiA</b>	<b>GTA</b>	False	False	84	-1.769787	12	51	False	1	1	0.000
<b>SjapoA</b>	<b>GTA</b>	False	False	19	-1.332187	4	7	False	1	1	0.000
<b>SpombA</b>	<b>GTA</b>	False	False	34	-1.386344	6	16	False	2	2	0.000
<b>CzempA</b>	<b>GTA</b>	False	False	40	-1.489738	7	20	False	2	2	0.000
<b>SoctoA</b>	<b>GTT</b>	False	False	49	-1.480889	6	24	False	2	2	0.000
<b>PkluyA</b>	<b>GTA</b>	False	False	126	-1.729811	13	65	False	1	1	0.000
<b>ScryoA</b>	<b>GTA</b>	False	False	41	-1.508653	7	20	False	2	2	0.000