

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>CphanA</b>	TTC	False	False	96	-1.734456	13	80	False	2	2	0
<b>Cgalla</b>	TTC	False	False	109	-1.775785	13	90	False	1	1	0
	TTT	False	False	70	-1.737206	13	50	False	1	1	0
<b>CdefoA</b>	TTC	False	False	106	-1.788117	13	88	False	1	1	0
	TTT	False	False	69	-1.743462	12	51	False	1	1	0
<b>YlipoB</b>	TTC	False	False	109	-1.788561	13	91	False	1	1	0
	TTT	False	False	68	-1.734556	12	49	False	1	1	0
<b>YlipoA</b>	TTC	False	False	109	-1.788561	13	91	False	1	1	0
	TTT	False	False	68	-1.734556	12	49	False	1	1	0
<b>CcastA</b>	TTT	False	False	76	-1.813623	7	66	False	1	1	0
<b>CalimA</b>	TTC	False	False	117	-1.765269	13	98	False	2	2	0
<b>SpombA</b>	TTT	False	False	43	-1.519475	6	28	False	1	1	0
<b>BcustA</b>	TTT	False	False	102	-1.679440	13	76	False	2	2	0
<b>CzempA</b>	TTT	False	False	81	-1.705701	7	59	False	1	1	0
<b>SoctoA</b>	TTT	False	False	50	-1.598154	6	33	False	1	1	0
<b>PkluyA</b>	TTT	False	False	94	-1.692838	13	74	False	2	2	0
<b>OanguA</b>	TTT	False	False	107	-1.786848	13	83	False	1	1	0
<b>ScryoA</b>	TTC	False	False	78	-1.700073	6	55	False	1	1	0