

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>MmagnB</b>	<b>TTC</b>	False	False	157	-2.674471	13	116	False	1	1	0
<b>CphanA</b>	<b>TTT</b>	False	False	77	-2.677840	13	59	False	1	1	0
<b>CgallA</b>	<b>TTT</b>	False	False	70	-2.548333	13	50	False	2	2	0
<b>MmagnA</b>	<b>TTC</b>	False	False	157	-2.680418	13	117	False	1	1	0
<b>CdefoA</b>	<b>TTT</b>	False	False	69	-2.658624	12	51	False	1	1	0
<b>MingeA</b>	<b>TTT</b>	False	False	125	-2.674421	13	95	False	1	1	0
<b>MstarA</b>	<b>TTT</b>	False	False	127	-2.682369	13	97	False	1	1	0
<b>YlipoB</b>	<b>TTT</b>	False	False	68	-2.626609	12	49	False	1	1	0
<b>WmucoA</b>	<b>TTT</b>	False	False	104	-2.772298	13	86	False	1	1	0
<b>CalimA</b>	<b>TTC</b>	False	False	117	-2.752567	13	98	False	1	1	0
<b>MtetrA</b>	<b>TTC</b>	False	False	163	-2.690657	13	122	False	1	1	0
<b>SjapoA</b>	<b>TTC</b>	False	False	75	-2.536789	6	51	False	1	1	0
<b>SpombA</b>	<b>TTC</b>	False	False	76	-2.573783	7	52	False	1	1	0
<b>YlipoA</b>	<b>TTT</b>	False	False	68	-2.626609	12	49	False	1	1	0
<b>CzempA</b>	<b>TTC</b>	False	False	39	-2.557851	7	28	False	1	1	0
<b>SoctoA</b>	<b>TTC</b>	False	False	68	-2.553484	6	47	False	1	1	0
	<b>TTT</b>	False	False	50	-2.539319	6	33	False	1	1	0
<b>ScryoA</b>	<b>TTC</b>	False	False	78	-2.563632	6	55	False	1	1	0