

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CphanA	GGA	False	False	69	-1.796747	13	66	False	1	1	0.000
CgallA	GGA	False	False	68	-1.757380	13	61	False	1	1	0.000
CdefoA	GGA	False	False	56	-1.748926	11	51	False	1	1	0.000
YlipoB	GGA	False	False	63	-1.730444	12	57	False	1	1	0.000
YlipoA	GGA	False	False	62	-1.735204	12	57	False	1	1	0.000
CalimA	GGA	False	False	69	-1.768034	12	61	False	1	1	0.000
SpombA	GGT	False	False	87	-1.733955	7	67	False	1	1	0.051
BcustA	GGA	False	False	108	-1.758225	11	95	False	1	1	0.000
SoctoA	GGA	False	False	40	-1.609408	6	30	False	1	1	0.055
ScryoA	GGA	False	False	40	-1.502623	7	29	False	1	1	0.055