

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	TAT	False	False	149	-0.798365	13	130	False	1	1	0.000
CgallA	TAT	False	False	112	-0.791202	13	100	False	1	1	0.000
CdefoA	TAT	False	False	114	-0.798157	13	101	False	1	1	0.000
MstarA	TAT	False	False	145	-0.804510	13	129	False	1	1	0.000
SjapoA	TAC	False	False	5	0.199262	2	4	False	1	1	0.000
CvartA	TAT	False	False	127	-0.802141	12	111	False	1	1	0.000
CzempA	TAT	False	False	82	-0.733248	6	64	False	1	1	0.000
MmagnB	TAC	False	False	115	-0.787376	13	104	False	1	1	0.000
MmagnA	TAC	False	False	119	-0.794255	13	108	False	1	1	0.000
LstarA	TAT	False	False	87	-0.714818	10	74	False	1	1	0.000
WmucoA	TAT	False	False	120	-0.811360	12	104	False	1	1	0.000
MtetrA	TAC	False	False	136	-0.821043	13	115	False	1	1	0.000
CsuavA	TAT	False	False	120	-0.806233	12	104	False	1	1	0.000
PcanaA	TAT	False	False	140	-0.791875	12	122	False	1	1	0.000
WpijpA	TAT	False	False	32	-0.678924	6	27	False	1	1	0.000
BarxA	TAT	False	False	21	-0.537658	1	17	False	1	1	0.047
CphanA	TAT	False	False	124	-0.774365	12	109	False	1	1	0.000
YlipoB	TAT	False	False	114	-0.788165	13	102	False	1	1	0.000
YlipoA	TAT	False	False	114	-0.788165	13	102	False	1	1	0.000
CjadiA	TAT	False	False	119	-0.813940	12	104	False	1	1	0.000
MingeA	TAT	False	False	142	-0.778857	13	124	False	1	1	0.000
CalimA	TAT	False	False	117	-0.793892	12	106	False	1	1	0.000
BcaliA	TAT	False	False	143	-0.804607	13	125	False	1	1	0.000