

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CsubhA	TAC	False	False	132	-0.860227	13	122	False	1	1	0
CgallA	TAC	False	False	27	-0.746860	12	25	False	1	1	0
	TAT	False	False	112	-0.834666	13	100	False	1	1	0
HuvarA	TAC	False	False	47	-0.688401	6	39	False	1	1	0
CphanA	TAT	False	False	124	-0.779484	12	109	False	3	3	0
CgigaA	TAT	False	False	31	-0.747067	10	28	False	1	1	0
CcoryA	TAT	False	False	138	-0.890696	13	127	False	1	1	0
CdefoA	TAC	False	False	28	-0.713381	12	25	False	1	1	0
	TAT	False	False	114	-0.830054	13	101	False	2	2	0
PpastA	TAT	False	False	133	-0.925354	12	123	False	1	1	0
YlipoB	TAT	False	False	114	-0.804643	13	102	False	2	2	0
YlipoA	TAT	False	False	114	-0.804643	13	102	False	2	2	0
CalimA	TAC	False	False	25	-0.878218	10	21	False	1	1	0
	TAT	False	False	117	-0.815360	12	106	False	1	1	0
CchauA	TAT	False	False	139	-0.857695	12	125	False	1	1	0
SpombA	TAT	False	False	55	-0.673620	6	41	False	2	2	0
BcustA	TAT	False	False	139	-0.854957	12	122	False	2	2	0
CzempA	TAT	False	False	82	-0.805286	6	64	False	2	2	0
CsubhB	TAC	False	False	132	-0.860227	13	122	False	1	1	0
SoctoA	TAT	False	False	56	-0.706268	7	42	False	2	2	0
PkluyA	TAT	False	False	117	-0.906070	12	106	False	2	2	0
OanguA	TAT	False	False	124	-0.880496	12	112	False	2	2	0
ScryoA	TAC	False	False	27	-0.666667	4	19	False	1	1	0
	TAT	False	False	46	-0.702159	7	36	False	1	1	0