

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	TAT	False	False	149	-0.818078	13	130	False	2	2	0
HuvarA	TAC	False	False	47	-0.709120	6	39	False	1	1	0
CfrijA	TAT	False	False	145	-0.871277	13	138	False	1	1	0
CgallA	TAC	False	False	27	-0.743814	12	25	False	1	1	0
	TAT	False	False	112	-0.800463	13	100	False	2	2	0
CdefoA	TAC	False	False	28	-0.710026	12	25	False	1	1	0
	TAT	False	False	114	-0.808513	13	101	False	2	2	0
PpastA	TAT	False	False	133	-0.876534	12	123	False	2	2	0
MstarA	TAT	False	False	145	-0.798543	13	129	False	3	3	0
SjapoA	TAT	False	False	62	-0.860969	6	45	False	2	2	0
NbaciA	TAT	False	False	65	-0.831835	7	59	False	1	1	0
CvartA	TAT	False	False	127	-0.831884	12	111	False	2	2	0
MtetrA	TAC	False	False	136	-0.849084	13	115	False	3	3	0
ClusiA	TAT	False	False	130	-0.859047	12	125	False	1	1	0
MmagnB	TAC	False	False	115	-0.778231	13	104	False	2	2	0
MmagnA	TAC	False	False	119	-0.786048	13	108	False	2	2	0
LstarA	TAC	False	False	14	-0.715343	3	12	False	1	1	0
	TAT	False	False	87	-0.710452	10	74	False	2	2	0
WmucoA	TAT	False	False	120	-0.830783	12	104	False	2	2	0
CsantA	TAC	False	False	42	-0.671176	11	34	False	2	2	0
	TAT	False	False	120	-0.824123	13	100	False	2	3	0
ScereC	TAT	False	False	64	-0.845092	7	59	False	1	1	0
ScereA	TAT	False	False	66	-0.849527	7	61	False	1	1	0
CtropB	TAT	False	False	69	-0.824277	12	64	False	1	1	0
CsuavA	TAT	False	False	120	-0.845065	12	104	False	2	2	0
ScereD	TAT	False	False	65	-0.848060	7	60	False	1	1	0
CtropA	TAT	False	False	68	-0.829190	12	63	False	1	1	0
MfariA	TAC	False	False	26	-0.750624	12	25	False	1	1	0
PkluyA	TAT	False	False	117	-0.888117	12	106	False	1	1	0
CcastA	TAT	False	False	67	-0.864099	7	62	False	1	1	0
PcanaA	TAT	False	False	140	-0.815347	12	122	False	2	2	0
WpijpA	TAT	False	False	32	-0.724365	6	27	False	1	1	0
CphanA	TAC	False	False	18	-0.761669	9	17	False	1	1	0
	TAT	False	False	124	-0.771141	12	109	False	2	2	0
SpastA	TAT	False	False	69	-0.861018	7	64	False	1	1	0
ScereE	TAT	False	False	67	-0.841020	7	61	False	1	1	0
YlipoB	TAC	False	False	25	-0.792339	11	23	False	1	1	0
	TAT	False	False	114	-0.787128	13	102	False	2	2	0

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
YlipoA	TAC	False	False	25	-0.792339	11	23	False	1	1	0
	TAT	False	False	114	-0.787128	13	102	False	2	2	0
CsalmA	TAT	False	False	133	-0.878241	12	112	False	3	3	0
CjadiA	TAT	False	False	119	-0.850816	12	104	False	2	2	0
BcustA	TAT	False	False	139	-0.828054	12	122	False	2	2	0
CglabA	TAT	False	False	71	-0.838496	7	63	False	1	1	0
SoctoA	TAT	False	False	56	-0.643067	7	42	False	4	4	0
OanguA	TAT	False	False	124	-0.836997	12	112	False	2	3	0
NdelpA	TAT	False	False	74	-0.847350	6	66	False	1	1	0
ScryoA	TAT	False	False	46	-0.624607	7	36	False	3	3	0
CneerA	TAT	False	False	144	-0.876598	13	137	False	1	1	0
SarboA	TAT	False	False	69	-0.861211	7	64	False	1	1	0
MingeA	TAT	False	False	142	-0.766279	13	124	False	4	4	0
ScastA	TAC	False	False	10	-0.591699	5	9	False	1	1	0
CalimA	TAC	False	False	25	-0.835309	10	21	False	1	1	0
	TAT	False	False	117	-0.792784	12	106	False	2	2	0
BcaliA	TAT	False	False	143	-0.831669	13	125	False	2	2	0
Spomba	TAT	False	False	55	-0.617178	6	41	False	3	3	0