

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CdublA	GCT	False	False	88	-0.949108	12	83	False	1	1	0.000
ScereD	GCA	False	False	38	-0.854391	6	30	False	1	1	0.000
SarboA	GCA	False	False	38	-0.872795	6	31	False	1	1	0.000
LstarA	GCT	False	False	71	-1.027773	10	46	False	1	1	0.000
ScastA	GCC	False	False	6	-0.138264	4	3	False	1	1	0.000
SpastA	GCA	False	False	45	-0.885696	5	36	False	1	1	0.000
CalbiB	GCT	False	False	78	-0.947245	11	74	False	1	1	0.000
CsantA	GCA	False	False	37	-0.907861	9	25	False	1	1	0.000
ScereC	GCA	False	False	38	-0.734551	6	28	False	2	2	0.000
SjapoA	GCT	False	False	61	-0.983404	6	37	False	1	1	0.000
ScereA	GCA	False	False	39	-0.735817	6	29	False	2	2	0.000
SpombA	GCT	False	False	68	-0.927242	6	40	False	1	1	0.000
CzempA	GCA	False	False	35	-0.892245	6	27	False	1	1	0.000
ScereE	GCA	False	False	39	-0.872931	6	30	False	1	1	0.000
SoctoA	GCA	False	False	23	-0.660166	5	12	False	1	1	0.000
CalbiD	GCT	False	False	76	-0.943164	10	72	False	1	1	0.000
CalbiC	GCT	False	False	78	-0.947245	11	74	False	1	1	0.046
OanguA	GCA	False	False	49	-0.997674	9	41	False	1	1	0.000
CalbiA	GCT	False	False	75	-0.947270	10	71	False	1	1	0.000
ScryoA	GCA	False	False	32	-1.021628	6	20	False	1	1	0.000