

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	GGT	False	False	119	-1.678241	12	107	False	1	1	0
ScereD	GGA	False	False	29	-1.711582	5	28	False	1	1	0
SarboA	GGA	False	False	34	-1.683421	6	32	False	1	1	0
LanspA	GGA	False	False	36	-1.672496	6	33	False	1	1	0
ScastA	GGA	False	False	32	-1.660730	6	30	False	1	1	0
WmucoA	GGA	False	False	88	-1.774121	12	83	False	1	1	0
CsantA	GGT	False	False	107	-1.593045	13	93	False	2	2	0
ScereC	GGA	False	False	28	-1.645467	6	26	False	1	1	0
ScereA	GGA	False	False	26	-1.619221	6	24	False	1	1	0
SpombA	GGA	False	False	31	-1.407248	6	26	False	1	1	0
	GGT	False	False	87	-1.397696	7	67	False	2	2	0
SpastA	GGA	False	False	27	-1.600772	6	25	False	1	1	0
BcaliA	GGA	False	False	95	-1.724027	13	88	False	1	1	0
ScereE	GGA	False	False	27	-1.690151	5	26	False	1	1	0
SoctoA	GGA	False	False	40	-1.436208	6	30	False	1	1	0
	GGT	False	False	70	-1.483427	7	59	False	1	1	0
WpijpA	GGA	False	False	75	-1.794468	7	72	False	1	1	0
PcanaA	GGA	False	False	70	-1.752804	11	66	False	1	1	0
ScryoA	GGA	False	False	40	-1.341917	7	29	False	1	1	0
	GGT	False	False	72	-1.522920	6	61	False	1	1	0