

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	GCT	False	False	75	-1.035614	12	61	False	1	1	0.000
CsuavA	GCT	False	False	71	-1.004174	11	53	False	1	1	0.000
CviswA	GCA	False	False	44	-0.933696	9	39	False	1	1	0.000
BcustA	GCA	False	False	51	-0.916287	10	39	False	1	1	0.000
	GCT	False	False	66	-0.876706	11	51	False	1	1	0.000
CjadiA	GCT	False	False	82	-1.032403	12	62	False	1	1	0.000
CalimA	GCT	False	False	109	-1.074578	12	69	False	1	1	0.000
SjapoA	GCA	False	False	12	-0.673611	3	7	False	1	1	0.000
SpombA	GCT	False	False	68	-0.974901	6	40	False	2	2	0.000
WmucoA	GCA	False	False	47	-0.979260	11	40	False	1	1	0.000
BcaliA	GCT	False	False	77	-0.940989	12	59	False	1	1	0.000
CvartA	GCT	False	False	80	-1.036707	11	64	False	1	1	0.000
SoctoA	GCA	False	False	23	-0.741909	5	12	False	1	1	0.056
	GCT	False	False	68	-1.049157	6	37	False	2	2	0.046
PcanaA	GCA	False	False	46	-0.966744	10	39	False	1	1	0.000
ScryoA	GCT	False	False	54	-1.001244	6	27	False	2	2	0.046