

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
PpastA	AGT	False	False	58	-1.957709	11	46	False	1	1	0.000
SarboA	TCA	False	False	57	-1.950301	7	50	False	1	1	0.000
CgigaA	TCA	False	False	39	-1.820970	10	34	False	1	1	0.000
CbohiA	TCT	False	False	34	-1.776026	9	32	False	1	1	0.000
LstarA	AGT	False	False	49	-1.867826	8	34	False	1	1	0.000
LkluyE	AGT	False	False	15	-1.694622	5	12	False	1	1	0.000
BcustA	AGT	False	False	47	-1.952983	11	32	False	1	1	0.000
	TCA	False	False	97	-1.953940	12	76	False	1	1	0.000
LkluyF	AGT	False	False	15	-1.694622	5	12	False	1	1	0.000
SjapoA	AGT	False	False	26	-1.830820	6	11	False	1	1	0.000
	TCT	False	False	61	-1.642333	7	38	False	2	2	0.000
LkluyD	AGT	False	False	15	-1.694622	5	12	False	1	1	0.000
LkluyC	AGT	False	False	15	-1.694622	5	12	False	1	1	0.000
LkluyA	AGT	False	False	15	-1.694622	5	12	False	1	1	0.000
CsubhB	AGC	False	False	62	-1.796717	11	55	False	1	1	0.000
MfariA	TCT	False	False	60	-1.900681	12	51	False	1	1	0.000
CsubhA	AGC	False	False	62	-1.796717	11	55	False	1	1	0.000
PkluyA	AGT	False	False	46	-1.940421	11	33	False	1	1	0.000
MingeA	TCA	False	False	73	-1.821908	12	53	False	1	1	0.051
DhansaA	AGC	False	False	1	4.456311	1	0	False	1	1	0.168