

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CsubhA	TCA	False	False	43	-0.060997	12	39	False	1	1	0.000
HuvarA	TCA	False	False	50	-0.671251	6	31	False	1	1	0.000
MmagnB	AGC	False	False	22	-0.123726	10	15	False	1	1	0.000
	TCA	False	False	52	-0.476876	11	34	False	1	1	0.000
CphanA	AGT	False	False	45	-0.087589	12	38	False	1	1	0.000
	TCA	False	False	85	-0.387101	12	63	False	1	1	0.000
CgallA	AGC	False	False	2	2.731132	2	1	False	1	1	0.035
	TCA	False	False	81	-0.347627	12	63	False	1	1	0.000
CalimA	AGT	False	False	45	-0.105764	12	38	False	1	1	0.000
	TCA	False	False	82	-0.334325	12	63	False	1	1	0.000
CdefoA	AGC	False	False	1	5.318182	1	0	False	1	1	0.134
	TCA	False	False	80	-0.299921	12	62	False	1	1	0.000
MingeA	AGT	False	False	73	-0.350008	12	56	False	1	1	0.000
	TCT	False	False	36	-0.312179	9	26	False	1	1	0.000
MstarA	AGT	False	False	74	-0.354829	12	56	False	1	1	0.000
	TCA	False	False	69	-0.404209	11	51	False	1	1	0.000
CsalmA	AGT	False	False	79	-0.445995	11	54	False	1	1	0.000
MmagnA	AGC	False	False	20	-0.034091	10	14	False	1	1	0.000
	TCA	False	False	52	-0.476876	11	34	False	1	1	0.000
YlipoB	AGC	False	False	1	5.318182	1	0	False	1	1	0.134
	TCA	False	False	75	-0.336892	12	57	False	1	1	0.000
CsantA	AGT	False	False	78	-0.353810	11	54	False	2	2	0.000
	TCA	False	False	31	-0.225993	8	23	False	1	1	0.000
MtetrA	AGC	False	False	39	-0.254477	12	31	False	1	1	0.000
	TCT	False	False	25	0.042996	9	21	False	1	1	0.000
SpombA	AGC	False	False	3	0.977778	3	0	False	1	1	0.016
YlipoA	AGC	False	False	1	5.318182	1	0	False	1	1	0.134
	TCA	False	False	75	-0.336892	12	57	False	1	1	0.000
CzempA	AGT	False	False	38	-0.547511	6	21	False	1	1	0.000
	TCT	False	False	34	-0.497483	6	20	False	1	1	0.000
CsubhB	TCA	False	False	43	-0.060997	12	39	False	1	1	0.000
SoctoA	AGT	False	False	32	-0.582862	6	15	False	1	1	0.000
CalaiA	TCT	False	False	40	-0.116392	8	37	False	1	1	0.000
ScryoA	AGT	False	False	31	-0.487618	6	14	False	1	1	0.000