

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	AGT	False	False	73	-1.093416	12	56	False	1	1	0.000
MmagnB	AGC	False	False	22	-0.672489	10	15	False	1	1	0.000
MmagnA	AGC	False	False	20	-0.545455	10	14	False	1	1	0.000
HuvarA	TCT	False	False	26	-0.950923	6	16	False	1	1	0.000
MstarA	AGT	False	False	74	-1.093881	12	56	False	1	1	0.000
CsalmA	AGT	False	False	79	-1.161178	11	54	False	1	1	0.000
MtetrA	AGC	False	False	39	-0.939919	12	31	False	1	1	0.000
KlactA	AGC	False	False	1	3.889908	1	0	False	1	1	0.039
SpombA	TCT	False	False	29	-1.146658	6	17	False	1	1	0.000
SjapoA	AGC	False	False	4	0.819512	3	2	False	1	1	0.000
OanguA	TCT	False	False	46	-0.984211	11	42	False	1	1	0.000
PcanaA	TCT	False	False	33	-0.984520	8	27	False	1	1	0.000
ScryoA	TCA	False	False	37	-1.006452	6	24	False	1	1	0.000