

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MmagnB	GTA	False	False	47	-2.027856	11	31	False	1	1	0
	GTT	False	False	49	-2.025117	9	27	False	1	1	0
MmagnA	GTA	False	False	45	-2.015138	11	30	False	1	1	0
	GTT	False	False	49	-2.023414	9	27	False	1	1	0
MingeA	GTT	False	False	58	-1.883713	12	40	False	1	2	0
MstarA	GTA	False	False	66	-2.031064	10	42	False	1	1	0
	GTT	False	False	50	-1.960728	12	32	False	1	1	0
CsalmA	GTT	False	False	93	-1.947613	11	47	False	1	2	0
CsantA	GTA	False	False	91	-2.038488	12	50	False	1	1	0
MtetrA	GTC	False	False	28	-2.067146	11	11	False	1	1	0
	GTG	False	False	62	-2.025464	11	33	False	1	1	0
SpombA	GTT	False	False	42	-2.043587	6	23	False	1	1	0
SoctoA	GTA	False	False	32	-1.995169	7	14	False	1	1	0
ScryoA	GTA	False	False	41	-1.890318	7	20	False	1	1	0