

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
HuvarA	GAT	False	False	38	-0.963375	5	28	False	1	1	0
MmagnB	GAT	False	False	21	-0.786991	8	20	False	1	1	0
MmagnA	GAT	False	False	22	-0.799906	8	21	False	1	1	0
LstarA	GAT	False	False	40	-0.930101	7	36	False	1	1	0
CsalmA	GAT	False	False	62	-0.932361	9	50	False	2	2	0
ClusiA	GAT	False	False	58	-0.874766	9	56	False	1	1	0
MtetrA	GAC	False	False	48	-0.917187	10	44	False	1	1	0
SjapoA	GAT	False	False	38	-0.981346	5	27	False	1	1	0
CcastA	GAT	False	False	40	-0.838826	5	37	False	1	1	0