

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
HuvarA	GAT	False	False	38	-0.781763	5	28	False	1	1	0.000
ScereD	GAT	False	False	38	-0.848040	5	37	False	1	1	0.000
SarboA	GAT	False	False	37	-0.843693	5	36	False	1	1	0.000
MtetrA	GAC	False	False	48	-0.897062	10	44	False	1	1	0.051
LstarA	GAT	False	False	40	-0.890924	7	36	False	1	1	0.000
ScereA	GAT	False	False	38	-0.848040	5	37	False	1	1	0.000
SpastA	GAT	False	False	35	-0.837349	5	34	False	1	1	0.000
ScastA	GAT	False	False	42	-0.840779	5	39	False	1	1	0.000
AgossA	GAT	False	False	37	-0.853665	5	36	False	1	1	0.055
ScereC	GAT	False	False	38	-0.848040	5	37	False	1	1	0.000
AgossB	GAT	False	False	37	-0.853665	5	36	False	1	1	0.055
NbaciA	GAT	False	False	37	-0.861694	5	36	False	1	1	0.000
CglabA	GAT	False	False	39	-0.852153	5	38	False	1	1	0.000
ScereE	GAT	False	False	38	-0.848040	5	37	False	1	1	0.000
CcastA	GAT	False	False	40	-0.857349	5	37	False	1	1	0.000
NdelpA	GAT	False	False	37	-0.852370	5	36	False	1	1	0.000