

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyerA	AAT	False	False	50	-2.672453	6	46	False	1	1	0.000
LdasiA	AAT	False	False	46	-2.672886	6	43	False	1	1	0.000
LkluyF	AAT	False	False	48	-2.679414	6	45	False	1	1	0.000
LkluyE	AAT	False	False	48	-2.679414	6	45	False	1	1	0.000
LkluyD	AAT	False	False	47	-2.672735	6	44	False	1	1	0.000
LkluyC	AAT	False	False	48	-2.679414	6	45	False	1	1	0.000
LkluyA	AAT	False	False	48	-2.679414	6	45	False	1	1	0.000
SpastA	AAC	False	False	8	-1.897619	6	6	False	1	1	0.000
ScereC	AAT	False	False	47	-2.647119	7	43	False	1	1	0.000
ScereA	AAT	False	False	47	-2.647119	7	43	False	1	1	0.000
ScereD	AAT	False	False	47	-2.647119	7	43	False	1	1	0.000
ScereE	AAT	False	False	47	-2.647119	7	43	False	1	1	0.000
LanspB	AAT	False	False	50	-2.672509	6	46	False	1	1	0.000
LanspA	AAT	False	False	47	-2.689732	7	44	False	1	1	0.000
AgossA	AAT	False	False	51	-2.618382	7	44	False	1	1	0.000
AgossB	AAT	False	False	51	-2.618382	7	44	False	1	1	0.000
LnothA	AAT	False	False	48	-2.682260	6	45	False	1	1	0.000
SarboA	AAT	False	False	47	-2.639301	7	43	False	1	1	0.000
HuvarA	AAT	False	False	29	-2.183295	6	17	False	1	1	0.000
ScastA	AAT	False	False	50	-2.574667	7	44	False	1	1	0.000
KtherA	AAT	False	False	47	-2.671478	7	44	False	1	1	0.000
KlactA	AAT	False	False	48	-2.628594	6	43	False	1	1	0.000
CzempA	AAT	False	False	64	-2.600927	7	45	False	1	1	0.001