

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LkluyF	ATG	False	False	45	-2.981209	6	39	False	1	1	0.000
LnothA	ATG	False	False	42	-2.939186	6	38	False	1	1	0.000
LmeyeA	ATG	False	False	43	-2.960465	6	38	False	1	1	0.000
LanspB	ATG	False	False	43	-2.960465	6	38	False	1	1	0.000
LanspA	ATG	False	False	43	-2.976770	6	37	False	1	1	0.000
CcoryA	ATG	False	False	66	-2.968914	12	57	False	1	1	0.000
HuvarA	ATG	False	False	54	-2.861764	6	27	False	1	1	0.001
LdasiA	ATG	False	False	42	-2.939186	6	38	False	1	1	0.000
LkluyE	ATG	False	False	45	-2.981209	6	39	False	1	1	0.000
CsantA	ATG	False	False	58	-2.940160	11	37	False	2	2	0.194
KtherA	ATG	False	False	43	-2.957114	6	38	False	1	1	0.000
SjapoA	ATG	False	False	35	-2.700353	5	18	False	1	1	0.000
LkluyD	ATG	False	False	44	-2.973362	6	38	False	1	1	0.000
SpombA	ATG	False	False	40	-3.021584	6	23	False	1	1	0.000
LkluyA	ATG	False	False	45	-2.981209	6	39	False	1	1	0.000
MfariA	ATG	False	False	64	-2.983210	10	56	False	1	1	0.000
SoctoA	ATG	False	False	38	-2.990020	6	24	False	1	1	0.000
LkluyC	ATG	False	False	45	-2.981209	6	39	False	1	1	0.000
ScryoA	ATG	False	False	40	-2.942559	6	24	False	1	1	0.000