

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	AAA	False	False	40	-1.747970	11	36	False	1	1	0.116
PpastA	AAA	False	False	44	-1.764096	12	41	False	1	1	0.208
MstarA	AAA	False	False	42	-1.760408	11	38	False	1	1	0.208
SpombA	AAA	False	False	16	-1.867799	6	9	False	1	1	0.116
MmagnB	AAA	False	False	30	-1.672563	11	27	False	1	1	0.116
MmagnA	AAA	False	False	30	-1.672563	11	27	False	1	1	0.116
LstarA	AAA	False	False	28	-1.538398	8	25	False	1	1	0.116
CsantA	AAA	False	False	34	-1.784166	12	26	False	1	1	0.116
MtetraA	AAG	False	False	16	-1.481825	8	14	False	1	1	0.000
MfariA	AAA	False	False	45	-1.799027	11	42	False	1	1	0.208
PkluyA	AAA	False	False	45	-1.803521	13	39	False	1	1	0.208
MfariB	AAA	False	False	43	-1.806269	12	41	False	1	1	0.208
CalaiA	AAA	False	False	41	-1.750708	9	38	False	1	1	0.116
PcanaA	AAA	False	False	43	-1.864977	11	40	False	1	1	0.116
CsalmA	AAA	False	False	39	-1.747655	11	33	False	1	1	0.209
BcustA	AAA	False	False	47	-1.853429	12	40	False	1	1	0.208
OanguA	AAA	False	False	45	-1.664301	13	39	False	2	2	0.116
ScryoA	AAA	False	False	20	-1.228155	6	10	False	2	3	0.153
HuvarA	AAA	False	False	20	-1.445673	6	12	False	2	2	0.344
ScastA	AAA	False	False	20	-1.635005	7	18	False	1	1	0.210
CzempA	AAA	False	False	19	-1.567095	7	14	False	1	1	0.210
SoctoA	AAA	False	False	21	-1.264011	6	10	False	2	3	0.153