

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	CCA	False	False	37	-0.820117	10	36	False	1	1	0.000
CmetaC	CCA	False	False	36	-0.814850	10	35	False	1	1	0.000
CmetaA	CCA	False	False	37	-0.820117	10	36	False	1	1	0.000
CmetaF	CCA	False	False	37	-0.820117	10	36	False	1	1	0.000
CgallA	CCT	False	False	58	-0.881477	12	57	False	1	1	0.000
CmetaE	CCA	False	False	37	-0.820117	10	36	False	1	1	0.000
CdefoA	CCT	False	False	55	-0.874484	12	54	False	1	1	0.000
SjapoA	CCT	False	False	60	-0.863018	7	54	False	1	1	0.000
SpombA	CCT	False	False	41	-0.644052	5	36	False	2	2	0.000
ClusiA	CCT	False	False	93	-0.885888	12	92	False	1	1	0.000
CmetaD	CCA	False	False	37	-0.820117	10	36	False	1	1	0.000
LstarA	CCT	False	False	70	-0.897464	9	65	False	1	1	0.000
CalaiA	CCA	False	False	24	-0.685104	8	22	False	1	1	0.000
DhansA	CCT	False	False	86	-0.874724	13	85	False	1	1	0.000
BarxA	CCT	False	False	14	-0.704545	1	12	False	1	1	0.098
CphanA	CCT	False	False	49	-0.845255	12	47	False	1	1	0.000
YlipoB	CCA	False	False	60	-0.853632	12	57	False	1	1	0.000
BcustA	CCA	False	False	51	-0.840886	11	47	False	1	1	0.000
	CCT	False	False	68	-0.838415	12	65	False	1	1	0.000
CsalmA	CCT	False	False	79	-0.914306	9	74	False	1	1	0.000
YlipoA	CCA	False	False	60	-0.853632	12	57	False	1	1	0.000
ScryoA	CCG	False	False	2	2.190909	1	0	False	1	1	0.054
CsakeA	CCA	False	False	61	-0.886833	12	60	False	1	1	0.000
HuvarA	CCT	False	False	33	-0.690490	5	29	False	1	1	0.051
CalimA	CCA	False	False	52	-0.836505	11	47	False	1	1	0.000