

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	CAT	False	False	61	0.925274	9	55	False	1	1	0.000
CparaB	CAT	False	False	45	1.020224	9	44	False	1	1	0.000
CfrijA	CAT	False	False	58	1.018332	9	55	False	1	2	0.000
CgallA	CAT	False	False	42	1.127392	8	38	False	2	2	0.000
CbuenA	CAT	False	False	54	0.961914	9	52	False	1	1	0.000
CviswA	CAT	False	False	54	0.867673	9	50	False	1	1	0.000
CdefoA	CAT	False	False	40	1.057338	8	38	False	1	1	0.000
MstarA	CAT	False	False	56	1.065684	8	52	False	2	2	0.000
SjapoA	CAT	False	False	39	1.056483	6	33	False	3	3	0.000
CchauA	CAC	False	False	11	1.284685	4	10	False	1	1	0.000
	CAT	False	False	49	1.113212	9	47	False	2	2	0.000
SpombA	CAT	False	False	31	0.869051	5	27	False	1	1	0.000
CvartA	CAT	False	False	53	0.954753	9	48	False	1	1	0.000
CorthB	CAT	False	False	42	1.007900	9	41	False	1	1	0.000
CorthC	CAT	False	False	43	1.007685	9	42	False	1	1	0.000
CorthA	CAT	False	False	42	1.007900	9	41	False	1	1	0.000
ClusiA	CAT	False	False	47	1.078471	10	46	False	1	1	0.000
AacerA	CAT	False	False	18	1.178627	2	17	False	1	1	0.224
CgigaA	CAC	False	False	32	0.939021	8	30	False	1	1	0.000
ScereC	CAT	False	False	43	0.958109	5	40	False	1	1	0.000
CparaA	CAT	False	False	36	1.028634	8	35	False	1	1	0.000
CcoryA	CAT	False	False	55	1.022684	9	54	False	1	1	0.000
CparaC	CAT	False	False	45	1.020224	9	44	False	1	1	0.000
LstarA	CAT	False	False	44	1.014370	7	38	False	2	2	0.000
CparaE	CAT	False	False	45	1.020224	9	44	False	1	1	0.000
CparaD	CAT	False	False	45	1.020224	9	44	False	1	1	0.000
WmucoA	CAT	False	False	53	0.967109	9	47	False	2	2	0.000
CparaF	CAT	False	False	45	1.020224	9	44	False	1	1	0.000
CsantA	CAC	False	False	16	1.153703	7	15	False	1	1	0.000
	CAT	False	False	42	0.832610	8	36	False	1	1	0.000
Mtetra	CAC	False	False	54	0.704125	8	45	False	1	1	0.000
CsaraA	CAT	False	False	56	0.960965	9	54	False	1	1	0.000
CtropB	CAC	False	False	26	0.937355	8	24	False	1	1	0.000
	CAT	False	False	34	1.031671	8	33	False	1	1	0.000
CsuavA	CAT	False	False	57	0.951693	9	52	False	1	1	0.000
ScereD	CAT	False	False	42	1.030776	5	40	False	1	1	0.000
CtropA	CAC	False	False	25	0.934386	8	23	False	1	1	0.000
	CAT	False	False	35	1.030551	8	34	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
PkluyA	CAT	False	False	45	1.079088	9	44	False	1	1	0.000
CalaiA	CAT	False	False	55	1.134945	8	53	False	2	2	0.001
PcanaA	CAT	False	False	59	1.028536	9	54	False	2	2	0.000
WpijpA	CAC	False	False	25	1.156654	5	22	False	2	2	0.000
CphanaA	CAT	False	False	40	1.136573	8	36	False	2	2	0.000
LanspA	CAT	False	False	43	1.072780	5	41	False	1	1	0.000
CsojaA	CAT	False	False	42	1.013788	6	39	False	2	2	0.000
ScereE	CAT	False	False	43	1.028450	5	41	False	1	1	0.000
YlipoB	CAC	False	False	17	0.965640	7	15	False	1	1	0.000
	CAT	False	False	42	1.041730	8	40	False	1	1	0.000
YlipoA	CAC	False	False	17	0.965640	7	15	False	1	1	0.000
	CAT	False	False	42	1.041730	8	40	False	1	1	0.000
CsalmaA	CAT	False	False	51	1.111429	9	48	False	2	2	0.000
CjadiA	CAT	False	False	57	0.960198	9	52	False	1	1	0.000
AgossB	CAT	False	False	45	1.122972	5	42	False	2	2	0.004
CsubhA	CAC	False	False	52	0.970826	10	50	False	1	1	0.000
BcustA	CAT	False	False	52	1.045890	9	49	False	1	1	0.000
CsubhB	CAC	False	False	52	0.970826	10	50	False	1	1	0.000
OanguA	CAT	False	False	43	1.109306	9	42	False	1	1	0.000
LelonA	CAT	False	False	45	0.981087	8	43	False	1	1	0.000
CneerA	CAT	False	False	57	0.932126	9	55	False	1	1	0.000
AgossA	CAT	False	False	45	1.122972	5	42	False	2	2	0.004
SarboA	CAT	False	False	42	1.039771	5	40	False	1	1	0.000
MingeA	CAT	False	False	52	0.997796	8	48	False	1	1	0.000
ScereA	CAT	False	False	43	1.028450	5	41	False	1	1	0.000
CcastA	CAT	False	False	42	1.052055	5	41	False	1	1	0.000
CalimA	CAT	False	False	45	1.049591	8	43	False	1	1	0.000
BcaliA	CAT	False	False	60	0.969137	9	53	False	2	2	0.000
CmaltA	CAT	False	False	51	0.916512	9	48	False	1	1	0.000
SpastA	CAT	False	False	42	1.039799	5	40	False	1	1	0.000