

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	AAT	False	False	89	-0.363434	13	68	False	2	2	0.000
LmeyerA	AAT	False	False	50	-0.158718	6	46	False	1	1	0.000
CgallA	AAT	False	False	62	-0.111635	13	49	False	2	2	0.000
CdefoA	AAT	False	False	62	-0.191564	13	47	False	2	2	0.000
CsubhB	AAC	False	False	67	-0.107320	11	64	False	1	1	0.000
MstarA	AAT	False	False	79	-0.217876	12	66	False	1	1	0.000
SjapoA	AAT	False	False	70	-0.040005	7	36	False	4	7	0.000
NbaciA	AAT	False	False	51	-0.119626	7	45	False	1	1	0.000
CvartA	AAT	False	False	75	-0.295300	11	57	False	2	2	0.000
AacerA	AAT	False	False	21	-0.126532	2	19	False	1	1	0.013
KtherA	AAC	False	False	8	0.324387	4	7	False	1	1	0.000
MguilA	AAT	False	False	61	-0.033538	11	59	False	1	1	0.000
WmucoA	AAC	False	False	25	-0.143408	8	22	False	1	1	0.000
CsantA	AAC	False	False	29	-0.018253	9	19	False	2	2	0.000
	AAT	False	False	55	0.006549	12	44	False	3	3	0.000
LelonA	AAT	False	False	47	-0.119205	10	41	False	1	1	0.000
CsaraA	AAT	False	False	71	-0.049146	12	65	False	2	2	0.000
BcaliA	AAT	False	False	87	-0.349373	12	66	False	2	2	0.000
MfariA	AAT	False	False	57	-0.019055	12	54	False	1	1	0.000
PkluyA	AAC	False	False	25	-0.022626	9	19	False	1	1	0.000
	AAT	False	False	56	-0.108773	11	46	False	2	2	0.000
MfariB	AAT	False	False	65	-0.141130	11	59	False	1	1	0.000
CalaiA	AAT	False	False	70	-0.031507	10	63	False	2	2	0.000
PcanaA	AAT	False	False	83	-0.368948	12	64	False	1	1	0.000
DhansA	AAT	False	False	54	-0.049485	11	51	False	1	1	0.000
BarxA	AAT	False	False	18	0.303030	1	14	False	1	2	0.038
LanspB	AAT	False	False	50	-0.159649	6	46	False	1	1	0.000
CphanaA	AAC	False	False	20	0.055775	7	17	False	1	1	0.000
	AAT	False	False	64	-0.223637	13	49	False	1	1	0.000
LdasiA	AAC	False	False	9	0.340336	4	8	False	1	1	0.000
YlipoB	AAT	False	False	61	-0.106747	13	50	False	2	2	0.000
YlipoA	AAT	False	False	61	-0.106747	13	50	False	2	2	0.000
CsalmaA	AAT	False	False	75	-0.209884	10	51	False	3	3	0.000
CjadiA	AAC	False	False	18	0.039106	8	16	False	1	1	0.000
	AAT	False	False	70	-0.339828	12	55	False	1	1	0.000
AgossB	AAT	False	False	51	-0.175802	7	44	False	1	1	0.000
CsubhA	AAC	False	False	67	-0.107320	11	64	False	1	1	0.000
BcustA	AAT	False	False	68	-0.063636	11	57	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>SoctoA</b>	<b>AAT</b>	False	False	53	-0.271710	6	33	False	3	3	0.000
<b>OanguA</b>	<b>AAC</b>	False	False	16	0.152542	6	12	False	1	1	0.000
	<b>AAT</b>	False	False	68	-0.162518	12	53	False	2	2	0.000
<b>ScryoA</b>	<b>AAC</b>	False	False	25	-0.097635	7	14	False	2	2	0.000
	<b>AAT</b>	False	False	41	-0.384039	6	27	False	1	1	0.000
<b>AgossaA</b>	<b>AAT</b>	False	False	51	-0.175802	7	44	False	1	1	0.000
<b>CsakeA</b>	<b>AAT</b>	False	False	67	0.025743	12	63	False	1	2	0.000
<b>LnothA</b>	<b>AAC</b>	False	False	7	0.445964	4	6	False	1	1	0.000
<b>CsuavA</b>	<b>AAC</b>	False	False	23	0.042028	9	21	False	1	1	0.000
<b>HuvarA</b>	<b>AAC</b>	False	False	35	-0.403375	6	23	False	2	2	0.000
	<b>AAT</b>	False	False	29	-0.690089	6	17	False	2	2	0.000
<b>CjiufA</b>	<b>AAT</b>	False	False	50	-0.102429	10	47	False	1	1	0.000
<b>CalimA</b>	<b>AAT</b>	False	False	65	-0.123623	12	52	False	2	2	0.000
<b>CblacA</b>	<b>AAT</b>	False	False	17	0.388770	1	15	False	1	2	0.038
<b>CzempA</b>	<b>AAT</b>	False	False	64	-0.170903	7	45	False	3	3	0.016
<b>SpombA</b>	<b>AAT</b>	False	False	58	-0.037416	6	40	False	3	4	0.000
<b>WpijpA</b>	<b>AAC</b>	False	False	37	-0.058292	7	35	False	1	1	0.000
<b>MingeA</b>	<b>AAT</b>	False	False	80	-0.295402	12	64	False	1	1	0.000