

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	GGA	False	False	52	-2.718232	9	49	False	1	1	0.000
CmetaC	GGA	False	False	52	-2.718232	9	49	False	1	1	0.000
CmetaA	GGA	False	False	52	-2.718232	9	49	False	1	1	0.000
CmetaF	GGA	False	False	52	-2.718232	9	49	False	1	1	0.000
CmetaD	GGA	False	False	52	-2.718232	9	49	False	1	1	0.000
CmetaE	GGA	False	False	52	-2.718232	9	49	False	1	1	0.000
CbuenA	GGA	False	False	49	-2.795046	10	43	False	1	1	0.000
CviswA	GGT	False	False	113	-2.794794	12	106	False	1	1	0.000
MguilA	GGA	False	False	51	-2.695488	10	45	False	1	1	0.000
CpseuA	GGA	False	False	15	-2.587273	1	12	False	1	1	0.092
CchauA	GGA	False	False	75	-2.796666	11	70	False	1	1	0.000
CalbiB	GGA	False	False	47	-2.754657	11	45	False	1	1	0.000
CdublA	GGA	False	False	41	-2.691450	10	39	False	1	1	0.000
CorthB	GGA	False	False	50	-2.760634	10	47	False	1	1	0.000
CorthC	GGA	False	False	50	-2.749843	10	47	False	1	1	0.000
CalbiE	GGA	False	False	28	-2.661813	6	26	False	1	1	0.000
Ctetra	GGA	False	False	30	-2.681300	8	27	False	1	1	0.000
CalbiC	GGA	False	False	47	-2.754657	11	45	False	1	1	0.000
ClusiA	GGT	False	False	122	-2.742979	12	115	False	1	1	0.000
CalbiA	GGA	False	False	40	-2.714286	10	38	False	1	1	0.000
CalbiD	GGA	False	False	36	-2.693446	10	34	False	1	1	0.000
CcoryA	GGT	False	False	138	-2.824738	12	131	False	1	1	0.000
CbohiA	GGA	False	False	41	-2.763528	9	37	False	1	1	0.000
CparaB	GGT	False	False	143	-2.832827	12	138	False	1	1	0.000
CparaE	GGT	False	False	143	-2.833030	12	138	False	1	1	0.000
CparaD	GGT	False	False	143	-2.832827	12	138	False	1	1	0.000
CparaF	GGT	False	False	143	-2.833030	12	138	False	1	1	0.000
CsantA	GGT	False	False	107	-2.623615	13	93	False	2	2	0.000
CsaraA	GGC	False	False	1	1.954545	1	0	False	1	1	0.310
CtropB	GGG	False	False	18	-2.370705	8	15	False	1	1	0.000
CtropA	GGG	False	False	19	-2.401922	8	16	False	1	1	0.000
MfariA	GGA	False	False	76	-2.787477	12	68	False	1	1	0.000
PkluyA	GGA	False	False	143	-2.762583	13	125	False	1	1	0.000
CalaiA	GGT	False	False	130	-2.834938	11	124	False	1	1	0.000
DhansA	GGA	False	False	92	-2.789860	12	82	False	1	1	0.000
BarxA	GGA	False	False	17	-2.410695	1	15	False	1	1	0.092
CoxycA	GGT	False	False	31	-2.741197	1	28	False	1	1	0.091
BcustA	GGA	False	False	108	-2.805237	11	95	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>CorthA</b>	<b>GGA</b>	False	False	46	-2.819869	10	43	False	1	1	0.000
<b>CsubhA</b>	<b>GGA</b>	False	False	72	-2.838358	12	68	False	1	1	0.000
<b>CsubhB</b>	<b>GGA</b>	False	False	72	-2.838358	12	68	False	1	1	0.000
<b>CtheaA</b>	<b>GGA</b>	False	False	59	-2.766801	11	56	False	1	1	0.000
<b>OanguA</b>	<b>GGT</b>	False	False	94	-2.832493	12	86	False	1	1	0.000
<b>LelonA</b>	<b>GGA</b>	False	False	56	-2.761281	11	53	False	1	1	0.000
<b>CsakeA</b>	<b>GGT</b>	False	False	122	-2.827232	12	115	False	1	1	0.000
<b>CjiufA</b>	<b>GGA</b>	False	False	62	-2.774974	12	60	False	1	1	0.000
<b>CblacA</b>	<b>GGT</b>	False	False	37	-2.766470	1	33	False	1	1	0.097
<b>CmaltA</b>	<b>GGT</b>	False	False	98	-2.831552	13	96	False	1	1	0.000
<b>CparaC</b>	<b>GGT</b>	False	False	143	-2.833030	12	138	False	1	1	0.000