

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	TTC	False	False	50	2.839100	6	45	False	1	1	0.000
	TTT	False	False	66	2.839575	6	60	False	2	2	0.000
SpombA	TTC	False	False	76	2.484967	7	52	False	4	6	0.000
	TTT	False	False	43	1.838193	6	28	False	1	1	0.000
SpastA	TTC	False	False	53	2.777758	6	47	False	2	2	0.000
	TTT	False	False	59	2.901365	6	55	False	1	1	0.000
AacerA	TTC	False	False	28	2.826345	2	25	False	1	1	0.015
	TTT	False	False	22	2.693144	2	19	False	1	1	0.015
CmetaD	TTC	False	False	76	2.896440	11	69	False	4	4	0.000
	TTT	False	False	76	2.837006	12	72	False	1	1	0.000
CgigaA	TTC	False	False	124	2.827451	12	115	False	2	2	0.000
CsantA	TTC	False	False	30	2.692070	10	23	False	2	3	0.000
	TTT	False	False	109	2.490798	13	88	False	2	3	0.000
CtropB	TTC	False	False	123	2.980370	13	113	False	4	5	0.000
CtropA	TTC	False	False	124	3.002961	13	114	False	4	5	0.000
WpijpA	TTC	False	False	73	2.883687	7	65	False	3	3	0.000
	TTT	False	False	42	2.795227	6	37	False	1	2	0.000
BarxA	TTC	False	False	17	3.050267	1	14	False	1	2	0.116
CoxycA	TTC	False	False	21	3.298268	1	20	False	1	1	0.062
CsubhA	TTC	False	False	147	2.991386	13	131	False	7	10	0.000
CsubhB	TTC	False	False	147	2.991386	13	131	False	7	10	0.000
OanguA	TTC	False	False	66	2.659195	12	55	False	3	3	0.000
	TTT	False	False	107	2.658199	13	83	False	6	8	0.000
SarboA	TTC	False	False	42	3.076526	6	37	False	3	3	0.000
	TTT	False	False	72	2.721918	7	65	False	1	1	0.000
CcastA	TTC	False	False	39	2.407471	6	31	False	1	1	0.000
	TTT	False	False	76	2.804575	7	66	False	4	4	0.000
KtherA	TTC	False	False	58	2.799016	6	52	False	1	1	0.000
	TTT	False	False	58	2.920225	6	52	False	3	3	0.000
CnorvA	TTT	False	False	128	2.810329	13	109	False	6	7	0.000
CviswA	TTC	False	False	63	2.936586	13	55	False	2	3	0.000
	TTT	False	False	93	2.986799	12	86	False	4	5	0.000
MstarA	TTC	False	False	40	2.458344	8	31	False	2	2	0.000
	TTT	False	False	127	2.863578	13	97	False	5	13	0.000
NbaciA	TTC	False	False	50	2.849772	6	45	False	1	1	0.000
	TTT	False	False	68	2.421305	7	55	False	2	2	0.000
CalbiD	TTC	False	False	76	2.939890	11	73	False	2	2	0.000
CbohiA	TTC	False	False	40	3.105030	11	38	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LstarA	TTT	False	False	113	2.862001	12	105	False	3	3	0.000
	TTC	False	False	62	2.865284	10	46	False	4	7	0.000
MfariA	TTT	False	False	92	2.631215	9	59	False	7	14	0.000
	TTC	False	False	67	2.928461	12	60	False	3	3	0.000
MfariB	TTT	False	False	94	2.885643	12	87	False	3	3	0.000
	TTC	False	False	69	3.039896	12	62	False	6	6	0.000
CalaiA	TTT	False	False	93	2.942844	11	85	False	3	3	0.000
	TTC	False	False	34	2.795414	8	31	False	1	1	0.000
LanspB	TTT	False	False	108	2.932531	11	99	False	2	4	0.000
	TTC	False	False	47	2.809082	6	42	False	1	1	0.000
Cphana	TTT	False	False	69	2.860220	7	63	False	2	2	0.000
	TTC	False	False	96	2.543904	13	80	False	2	3	0.000
LanspA	TTT	False	False	77	2.752595	13	59	False	6	7	0.000
	TTC	False	False	56	2.827867	6	52	False	1	1	0.000
CsojaA	TTT	False	False	59	2.973219	6	52	False	3	3	0.000
	TTC	False	False	43	3.021908	8	41	False	1	1	0.000
CglabA	TTT	False	False	55	2.871820	7	51	False	1	1	0.000
	TTC	False	False	51	2.717403	6	44	False	2	2	0.000
ScryoA	TTT	False	False	66	2.672131	6	56	False	3	3	0.000
	TTC	False	False	78	2.412139	6	55	False	4	5	0.000
CneerA	TTT	False	False	38	2.176902	6	23	False	2	3	0.000
	TTC	False	False	72	2.987681	12	70	False	1	1	0.000
CsakeA	TTT	False	False	84	2.866184	12	74	False	3	4	0.000
	TTC	False	False	50	2.883934	11	45	False	2	2	0.000
HuvarA	TTT	False	False	108	3.027947	11	98	False	5	7	0.000
	TTC	False	False	57	2.645205	6	45	False	3	3	0.000
CjiufA	TTT	False	False	44	2.507812	6	32	False	3	4	0.000
	TTC	False	False	76	2.847141	12	69	False	3	3	0.000
KlactA	TTT	False	False	80	2.771556	12	74	False	1	1	0.000
	TTC	False	False	68	2.710843	7	60	False	2	2	0.000
CzempA	TTT	False	False	46	2.729769	6	40	False	1	1	0.000
	TTC	False	False	39	2.389645	7	28	False	2	3	0.000
CmetaB	TTT	False	False	81	2.451425	7	59	False	5	5	0.000
	TTC	False	False	76	2.896440	11	69	False	4	4	0.000
CmetaC	TTT	False	False	76	2.837006	12	72	False	1	1	0.000
	TTC	False	False	76	2.896440	11	69	False	4	4	0.000
CmetaA	TTT	False	False	76	2.837006	12	72	False	1	1	0.000
	TTC	False	False	76	2.896440	11	69	False	4	4	0.000
CmetaF	TTT	False	False	76	2.837006	12	72	False	1	1	0.000
	TTC	False	False	76	2.896440	11	69	False	4	4	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CfrijA	TTC	False	False	58	2.984979	11	54	False	2	2	0.000
	TTT	False	False	94	2.883911	11	90	False	1	1	0.000
CgallA	TTC	False	False	109	2.475472	13	90	False	2	3	0.000
	TTT	False	False	70	2.780390	13	50	False	7	8	0.000
CmetaE	TTC	False	False	76	2.896440	11	69	False	4	4	0.000
	TTT	False	False	76	2.837006	12	72	False	1	1	0.000
PpastA	TTC	False	False	46	2.709488	10	38	False	2	2	0.000
	TTT	False	False	125	2.699537	13	101	False	4	8	0.000
SjapoA	TTC	False	False	75	2.569591	6	51	False	5	8	0.000
	TTT	False	False	58	2.026853	6	32	False	4	7	0.000
CalbiE	TTC	False	False	56	2.943534	6	53	False	2	2	0.000
CtetraA	TTC	False	False	85	3.025484	13	79	False	3	4	0.000
	TTT	False	False	68	2.943005	11	63	False	2	2	0.000
CalbiC	TTC	False	False	85	2.886406	12	81	False	2	2	0.000
	TTT	False	False	64	2.955007	11	60	False	1	1	0.000
CalbiB	TTC	False	False	85	2.886406	12	81	False	2	2	0.000
	TTT	False	False	64	2.955007	11	60	False	1	1	0.000
CalbiA	TTC	False	False	78	2.937928	11	75	False	2	2	0.000
MmagnB	TTC	False	False	157	2.630293	13	116	False	5	13	0.000
	TTT	False	False	15	3.054533	7	13	False	1	1	0.000
MmagnA	TTC	False	False	157	2.654024	13	117	False	5	13	0.000
	TTT	False	False	15	2.816382	7	12	False	1	1	0.000
CparaA	TTC	False	False	42	2.857515	10	38	False	2	2	0.000
	TTT	False	False	78	2.716700	11	72	False	1	1	0.000
CcoryA	TTC	False	False	68	2.923077	12	62	False	3	3	0.000
	TTT	False	False	76	2.892991	12	73	False	1	1	0.000
CparaC	TTC	False	False	56	2.883993	11	51	False	3	3	0.000
	TTT	False	False	97	2.792027	12	91	False	1	1	0.000
CparaB	TTC	False	False	56	2.883993	11	51	False	3	3	0.000
	TTT	False	False	97	2.792027	12	91	False	1	1	0.000
CparaE	TTC	False	False	56	2.883993	11	51	False	3	3	0.000
	TTT	False	False	97	2.792027	12	91	False	1	1	0.000
CparaD	TTC	False	False	56	2.883993	11	51	False	3	3	0.000
	TTT	False	False	97	2.792027	12	91	False	1	1	0.000
CparaF	TTC	False	False	56	2.883993	11	51	False	3	3	0.000
	TTT	False	False	97	2.792027	12	91	False	1	1	0.000
MtetraA	TTC	False	False	163	2.660560	13	122	False	7	14	0.000
	TTT	False	False	10	2.892970	6	8	False	1	1	0.000
CsaraA	TTC	False	False	50	2.930763	11	44	False	3	3	0.000
	TTT	False	False	108	2.847267	11	97	False	3	4	0.000
CsuavA	TTC	False	False	78	2.688985	11	66	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
PkluyA	TTT	False	False	90	2.735202	13	72	False	6	7	0.000
	TTC	False	False	80	2.739579	12	66	False	4	4	0.000
	TTT	False	False	94	2.460009	13	74	False	4	4	0.000
PcanaA	TTT	False	False	124	2.790634	13	102	False	5	8	0.000
YlipoB	TTC	False	False	109	2.514112	13	91	False	2	3	0.000
	TTT	False	False	68	2.830438	12	49	False	7	8	0.000
YlipoA	TTC	False	False	109	2.514112	13	91	False	2	3	0.000
	TTT	False	False	68	2.830438	12	49	False	7	8	0.000
CjadiA	TTC	False	False	82	2.773684	12	70	False	4	4	0.000
	TTT	False	False	88	2.611598	13	68	False	6	6	0.000
AgossB	TTC	False	False	37	3.001788	7	33	False	2	2	0.000
	TTT	False	False	75	2.773744	7	66	False	2	2	0.000
NdelpA	TTC	False	False	63	2.655047	6	51	False	2	3	0.000
	TTT	False	False	54	2.639718	7	48	False	1	1	0.000
LelonA	TTC	False	False	106	2.855460	13	94	False	4	5	0.000
	TTT	False	False	49	2.925495	11	48	False	1	1	0.000
LkluyF	TTC	False	False	34	2.890003	6	31	False	1	1	0.000
	TTT	False	False	80	2.811691	7	72	False	2	2	0.000
CalimA	TTC	False	False	117	2.621890	13	98	False	2	4	0.000
	TTT	False	False	57	2.569458	12	42	False	4	5	0.000
LkluyD	TTT	False	False	78	2.889394	7	70	False	2	3	0.000
BcaliA	TTT	False	False	115	2.807984	13	98	False	5	6	0.000
SoctoA	TTC	False	False	68	2.336605	6	47	False	4	5	0.000
	TTT	False	False	50	2.292828	6	33	False	3	3	0.000
LkluyB	TTT	False	False	23	3.030854	1	22	False	1	1	0.067
CbuenA	TTC	False	False	72	3.005760	13	68	False	2	2	0.000
	TTT	False	False	83	2.794961	11	75	False	2	2	0.000
CdefoA	TTC	False	False	106	2.568770	13	88	False	2	4	0.000
	TTT	False	False	69	2.705946	12	51	False	5	6	0.000
LdasiA	TTC	False	False	61	2.767125	6	56	False	1	1	0.000
	TTT	False	False	55	2.919590	6	49	False	2	2	0.000
CpseuA	TTC	False	False	19	2.854545	1	17	False	1	1	0.068
LkluyE	TTC	False	False	35	2.859002	6	32	False	1	1	0.000
	TTT	False	False	79	2.824627	7	71	False	2	2	0.000
CchauA	TTC	False	False	61	2.990936	12	55	False	3	4	0.000
	TTT	False	False	81	2.920814	12	79	False	1	1	0.000
LkluyC	TTC	False	False	34	2.890003	6	31	False	1	1	0.000
	TTT	False	False	80	2.811691	7	72	False	2	2	0.000
CvartA	TTC	False	False	65	2.749655	10	57	False	2	2	0.000
	TTT	False	False	105	2.708157	13	82	False	6	8	0.000
LkluyA	TTC	False	False	34	2.890003	6	31	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
	TTT	False	False	80	2.811691	7	72	False	2	2	0.000
CdublA	TTC	False	False	67	2.893448	11	63	False	2	2	0.000
CorthB	TTC	False	False	68	2.845575	12	62	False	3	3	0.000
CorthC	TTC	False	False	65	2.832942	12	59	False	3	3	0.000
CorthA	TTC	False	False	70	2.838222	12	64	False	3	3	0.000
	TTC	False	False	56	3.015106	12	53	False	2	2	0.000
ClusiA	TTT	False	False	97	2.956536	12	91	False	4	4	0.000
	TTC	False	False	65	2.979990	11	63	False	2	2	0.000
MguilA	TTT	False	False	90	2.719154	12	79	False	2	2	0.000
	TTC	False	False	55	2.745870	8	47	False	3	3	0.000
WmucoA	TTT	False	False	104	2.791381	13	86	False	4	6	0.000
	TTC	False	False	53	2.809319	6	47	False	2	2	0.000
ScereC	TTT	False	False	60	2.944249	6	55	False	2	2	0.000
	TTC	False	False	56	2.796947	6	50	False	2	2	0.000
ScereA	TTT	False	False	58	2.951554	6	53	False	2	2	0.000
	TTC	False	False	54	2.802886	6	48	False	2	2	0.000
ScereD	TTT	False	False	59	2.952420	6	54	False	2	2	0.000
	TTC	False	False	51	2.803757	6	45	False	2	2	0.000
ScereE	TTT	False	False	61	2.924929	7	56	False	2	2	0.000
	TTC	False	False	76	3.025113	13	70	False	4	4	0.000
DhansA	TTT	False	False	87	2.844615	12	75	False	4	5	0.000
	TTC	False	False	45	2.357737	10	32	False	2	2	0.000
CsalmA	TTT	False	False	110	2.540546	12	81	False	5	9	0.000
	TTC	False	False	59	2.772193	6	54	False	1	1	0.000
LnothA	TTT	False	False	57	2.909504	6	51	False	2	2	0.000
	TTC	False	False	71	2.324285	12	55	False	1	1	0.000
BcustA	TTT	False	False	102	2.569698	13	76	False	9	9	0.000
	TTC	False	False	71	2.908760	11	66	False	3	3	0.000
CtheaA	TTC	False	False	34	2.941767	7	31	False	1	1	0.000
AgossA	TTT	False	False	78	2.809302	7	68	False	3	3	0.000
	TTC	False	False	41	2.551418	9	32	False	2	2	0.000
MingeA	TTT	False	False	125	2.792459	13	95	False	6	12	0.000
	TTC	False	False	42	2.574825	6	35	False	1	1	0.000
ScastA	TTT	False	False	77	2.735606	7	63	False	4	4	0.000
	TTC	False	False	43	2.801682	11	40	False	1	1	0.000
CmaltA	TTT	False	False	115	2.942603	12	102	False	5	6	0.000
KmarxA	TTC	False	False	30	2.941176	1	28	False	1	1	0.067