

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CgallA	GAT	False	False	53	1.114791	9	49	False	2	2	0.000
CbuenA	GAT	False	False	61	1.092596	10	58	False	2	2	0.001
CdefoA	GAT	False	False	54	1.111708	9	50	False	2	2	0.000
PpastA	GAT	False	False	56	1.063345	9	53	False	1	1	0.000
ScastA	GAT	False	False	42	1.095762	5	39	False	1	1	0.000
CpseuA	GAT	False	False	10	1.355778	1	8	False	1	1	0.036
SjapoA	GAT	False	False	38	0.867658	5	27	False	1	1	0.000
LkluyD	GAT	False	False	38	1.109735	5	37	False	1	1	0.000
SpombA	GAT	False	False	37	1.028779	5	27	False	1	2	0.000
LkluyA	GAT	False	False	38	1.109735	5	37	False	1	1	0.000
CorthB	GAT	False	False	54	1.097040	9	53	False	1	1	0.000
CorthC	GAT	False	False	54	1.097040	9	53	False	1	1	0.000
CtetrA	GAT	False	False	51	1.075885	9	49	False	1	1	0.000
ClusiA	GAT	False	False	58	1.041869	9	56	False	1	1	0.000
CoxycA	GAT	False	False	9	1.396360	1	7	False	1	1	0.038
CorthA	GAT	False	False	54	1.097040	9	53	False	1	1	0.000
CgigaA	GAT	False	False	17	1.434594	7	15	False	2	2	0.000
CparaC	GAT	False	False	55	1.095825	9	54	False	1	1	0.000
CparaB	GAT	False	False	55	1.095825	9	54	False	1	1	0.000
CparaE	GAT	False	False	55	1.095825	9	54	False	1	1	0.000
CparaD	GAT	False	False	55	1.095825	9	54	False	1	1	0.000
CparaF	GAT	False	False	55	1.095825	9	54	False	1	1	0.000
MfariA	GAT	False	False	56	1.072525	9	55	False	1	1	0.000
PkluyA	GAT	False	False	53	1.124465	9	52	False	1	1	0.000
LelonA	GAT	False	False	55	1.169875	9	53	False	2	2	0.000
CcastA	GAT	False	False	40	1.101997	5	37	False	1	1	0.000
Cphana	GAC	False	False	3	2.389776	2	2	False	1	1	0.017
	GAT	False	False	53	1.025567	9	50	False	1	1	0.000
YlipoB	GAT	False	False	56	1.197305	9	51	False	3	3	0.000
YlipoA	GAT	False	False	56	1.197305	9	51	False	3	3	0.000
CsalmA	GAT	False	False	62	0.956052	9	50	False	1	1	0.000
CsubhA	GAC	False	False	59	1.074169	9	57	False	1	1	0.055
BcustA	GAC	False	False	2	3.281818	1	1	False	1	1	0.071
	GAT	False	False	56	1.074302	9	52	False	1	1	0.000
CsubhB	GAC	False	False	59	1.074169	9	57	False	1	1	0.055
LkluyC	GAT	False	False	38	1.109735	5	37	False	1	1	0.000
OanguA	GAT	False	False	54	1.086073	9	53	False	1	1	0.000
ScryoA	GAC	False	False	14	0.980110	3	10	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LkluyF	GAT	False	False	27	0.978107	6	20	False	1	1	0.000
	GAT	False	False	38	1.109735	5	37	False	1	1	0.000
HuvarA	GAC	False	False	4	2.055156	3	3	False	1	1	0.000
	GAT	False	False	38	0.985097	5	28	False	3	3	0.000
LkluyE	GAT	False	False	38	1.109735	5	37	False	1	1	0.000
CalimA	GAC	False	False	5	1.935167	3	4	False	1	1	0.001
	GAT	False	False	51	1.017834	9	48	False	1	1	0.000
SoctoA	GAC	False	False	8	1.070133	4	5	False	1	1	0.000
	GAT	False	False	32	0.985123	4	25	False	1	1	0.000