

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	AAT	False	False	48	0.079936	10	46	False	1	1	0
CnorvA	AAT	False	False	89	0.099951	13	68	False	5	6	0
CmetaA	AAT	False	False	48	0.079936	10	46	False	1	1	0
CmetaF	AAT	False	False	48	0.079936	10	46	False	1	1	0
CmetaD	AAT	False	False	48	0.079936	10	46	False	1	1	0
CmetaC	AAT	False	False	48	0.079936	10	46	False	1	1	0
CdefoA	AAC	False	False	24	-0.002461	9	20	False	1	1	0
	AAT	False	False	62	0.058348	13	47	False	3	3	0
PpastA	AAC	False	False	8	0.265700	6	6	False	1	1	0
MstarA	AAT	False	False	79	0.002852	12	66	False	1	1	0
SjapoA	AAT	False	False	70	-0.090012	7	36	False	2	4	0
SpombA	AAT	False	False	58	0.006593	6	40	False	2	2	0
CvartA	AAT	False	False	75	0.078434	11	57	False	3	5	0
CorthB	AAT	False	False	52	0.071173	10	50	False	1	1	0
CorthC	AAT	False	False	51	0.072751	10	49	False	1	1	0
CorthA	AAT	False	False	52	0.071173	10	50	False	1	1	0
ClusiA	AAT	False	False	62	0.101504	11	59	False	1	1	0
MmagnB	AAC	False	False	74	0.014560	12	62	False	1	1	0
CgallA	AAC	False	False	22	0.004946	9	18	False	1	1	0
	AAT	False	False	62	0.041230	13	49	False	2	2	0
MmagnA	AAC	False	False	74	0.033095	12	63	False	1	1	0
CparaA	AAT	False	False	41	0.088426	9	39	False	1	1	0
CparaC	AAT	False	False	51	0.069447	10	49	False	1	1	0
CmetaE	AAT	False	False	48	0.079936	10	46	False	1	1	0
CparaE	AAT	False	False	51	0.069447	10	49	False	1	1	0
CparaD	AAT	False	False	51	0.069447	10	49	False	1	1	0
WmucoA	AAT	False	False	60	0.160594	10	48	False	3	4	0
CparaF	AAT	False	False	51	0.069447	10	49	False	1	1	0
CsantA	AAC	False	False	29	-0.125626	9	19	False	1	1	0
	AAT	False	False	55	-0.024013	12	44	False	1	1	0
MtetraA	AAC	False	False	81	-0.011799	12	68	False	1	1	0
CsuavA	AAC	False	False	23	0.050693	9	21	False	1	1	0
	AAT	False	False	66	0.089415	10	51	False	2	4	0
PcanaA	AAT	False	False	83	0.073896	12	64	False	3	4	0
DhansA	AAT	False	False	54	0.094022	11	51	False	1	1	0
CphanA	AAT	False	False	64	0.065376	13	49	False	3	4	0
YlipoB	AAC	False	False	21	0.142302	8	16	False	1	1	0
	AAT	False	False	61	0.021138	13	50	False	2	2	0

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
YlipoA	AAC	False	False	21	0.142302	8	16	False	1	1	0
	AAT	False	False	61	0.021138	13	50	False	2	2	0
CsalmaA	AAT	False	False	75	0.015337	10	51	False	2	3	0
CjadiA	AAT	False	False	70	0.129511	12	55	False	3	5	0
BcustA	AAT	False	False	68	-0.015789	11	57	False	1	1	0
CglabA	AAT	False	False	49	0.066265	7	45	False	1	1	0
CtheaA	AAT	False	False	52	0.056476	10	50	False	1	1	0
NdelpA	AAT	False	False	51	0.035708	7	45	False	1	1	0
ScryoA	AAT	False	False	41	-0.049146	6	27	False	1	1	0
CneerA	AAT	False	False	71	0.027538	12	66	False	1	1	0
HuvarA	AAC	False	False	35	-0.244148	6	23	False	1	1	0
CparaB	AAT	False	False	51	0.069447	10	49	False	1	1	0
MingeA	AAT	False	False	80	-0.004045	12	64	False	2	2	0
ScastA	AAT	False	False	50	0.066857	7	44	False	1	1	0
CalimA	AAC	False	False	19	0.414477	9	15	False	1	2	0
	AAT	False	False	65	-0.086470	12	52	False	1	1	0
KlactA	AAT	False	False	48	0.052548	6	43	False	1	1	0
LstarA	AAT	False	False	57	0.116664	8	44	False	2	2	0
BcaliA	AAT	False	False	87	0.071546	12	66	False	4	5	0
SoctoA	AAT	False	False	53	-0.143167	6	33	False	1	1	0