

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
HuvarA	AGA	False	False	26	-0.744563	6	23	False	1	1	0.063
MmagnB	AGA	False	False	33	-0.807029	11	31	False	1	1	0.003
CphanA	AGA	False	False	51	-0.822839	12	46	False	1	1	0.003
CgallA	AGA	False	False	51	-0.822839	12	46	False	1	1	0.003
MmagnA	AGA	False	False	33	-0.807029	11	31	False	1	1	0.003
CdefoA	AGA	False	False	51	-0.822839	12	46	False	1	1	0.003
LstarA	CGT	False	False	1	3.291262	1	0	False	1	1	0.332
MstarA	AGA	False	False	34	-0.765569	10	32	False	1	1	0.003
YlipoB	AGA	False	False	50	-0.822930	12	45	False	1	1	0.003
YlipoA	AGA	False	False	50	-0.822930	12	45	False	1	1	0.003
CalimA	AGA	False	False	51	-0.822839	12	46	False	1	1	0.003
CsantA	AGG	False	False	10	-0.483279	5	8	False	1	1	0.001
MtetrA	AGA	False	False	32	-0.648478	11	29	False	2	2	0.003
SjapoA	AGA	False	False	26	-0.816943	7	21	False	1	1	0.002
SpombA	AGG	False	False	1	3.485437	1	0	False	1	1	0.246
	CGA	False	False	4	0.092910	3	2	False	1	1	0.000
CzempA	AGA	False	False	21	-0.725310	6	16	False	1	1	0.002
SoctoA	AGA	False	False	28	-0.813803	7	23	False	1	1	0.002
MingeA	AGA	False	False	37	-0.828726	10	35	False	1	1	0.003
ScryoA	CGA	False	False	5	-0.197652	2	4	False	1	1	0.001