

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
<b>CneerA</b>	<b>ATT</b>	False	False	97	-2.818391	13	86	False	1	1	0.000
<b>MingeA</b>	<b>ATA</b>	False	False	147	-2.840428	12	109	False	1	1	0.199
<b>MmagnB</b>	<b>ATT</b>	False	False	41	-2.749692	10	30	False	1	1	0.000
<b>CfrijA</b>	<b>ATC</b>	False	False	23	-2.551188	7	20	False	1	1	0.000
<b>MmagnA</b>	<b>ATT</b>	False	False	44	-2.764076	10	33	False	1	1	0.000
<b>LstarA</b>	<b>ATT</b>	False	False	97	-2.814136	9	73	False	1	2	0.051
<b>MstarA</b>	<b>ATA</b>	False	False	162	-2.897938	11	121	False	1	1	0.199
<b>CsalmaA</b>	<b>ATC</b>	False	False	10	-2.067669	7	6	False	1	1	0.000
<b>CsantA</b>	<b>ATA</b>	False	False	223	-2.745084	13	145	False	2	2	0.199
<b>MtetrA</b>	<b>ATT</b>	False	False	33	-2.730110	8	26	False	1	1	0.000
<b>SjapoA</b>	<b>ATT</b>	False	False	104	-2.753102	7	63	False	1	1	0.000
<b>CzempA</b>	<b>ATA</b>	False	False	91	-2.738519	7	57	False	1	1	0.235
<b>MfariA</b>	<b>ATA</b>	False	False	138	-2.846643	13	120	False	1	1	0.199