CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	TAT	False	False	131	-2.984727	11	123	False	1	1	0.00
CmetaC	TAT	False	False	131	-2.984727	11	123	False	1	1	0.00
CmetaA	TAT	False	False	131	-2.984727	11	123	False	1	1	0.00
CmetaF	TAT	False	False	131	-2.984727	11	123	False	1	1	0.00
CmetaD	TAT	False	False	131	-2.984727	11	123	False	1	1	0.00
CmetaE	TAT	False	False	131	-2.984727	11	123	False	1	1	0.00
CpseuA	TAT	False	False	13	-2.473202	1	11	False	1	1	0.04
SjapoA	TAT	False	False	62	-2.876761	6	45	False	1	1	0.00
CchauA	TAC	False	False	19	-2.526261	8	18	False	1	1	0.00
CorthB	TAT	False	False	137	-2.982465	11	130	False	1	1	0.00
CorthC	TAT	False	False	138	-2.987771	11	130	False	1	1	0.00
CorthA	TAT	False	False	137	-2.987330	11	129	False	1	1	0.00
CcoryA	TAT	False	False	138	-2.995186	13	127	False	1	1	0.00
CbohiA	TAT	False	False	145	-2.986251	13	138	False	1	1	0.00
CparaB	TAT	False	False	138	-2.988701	11	130	False	1	1	0.00
CparaE	TAT	False	False	138	-2.988701	11	130	False	1	1	0.00
CparaD	TAT	False	False	138	-2.988701	11	130	False	1	1	0.00
CparaF	TAT	False	False	138	-2.988701	11	130	False	1	1	0.00
CoxycA	TAT	False	False	13	-2.548660	1	9	False	1	1	0.04
CtheaA	TAT	False	False	138	-2.982734	11	131	False	1	1	0.00
LelonA	TAT	False	False	122	-2.976714	12	115	False	1	1	0.00
HuvarA	TAT	False	False	16	-2.458977	6	12	False	1	1	0.00
CjiufA	TAT	False	False	126	-2.962374	11	121	False	1	1	0.00
CparaC	TAT	False	False	138	-2.988701	11	130	False	1	1	0.00