

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CsakeA	CGT	False	False	14	0.265275	8	13	False	1	1	0.000
MfariB	AGA	False	False	38	0.092281	11	37	False	1	1	0.001
CgigaA	CGA	False	False	2	1.422727	1	1	False	1	1	0.090
CbuenA	AGA	False	False	32	0.143870	10	30	False	1	1	0.001
CcoryA	CGT	False	False	48	0.126110	11	44	False	1	1	0.000
CbohiA	CGT	False	False	42	0.113660	10	39	False	1	1	0.000
MguilA	CGT	False	False	16	0.219205	7	15	False	1	1	0.000
CchauA	CGT	False	False	40	0.152378	11	36	False	1	1	0.000
CblacA	CGT	False	False	5	0.578182	1	4	False	1	1	0.053
CsubhB	CGT	False	False	11	0.277311	7	9	False	1	1	0.000
MfariA	CGT	False	False	16	0.218110	8	15	False	1	1	0.000
CsubhA	CGT	False	False	11	0.277311	7	9	False	1	1	0.000
CtetrA	AGA	False	False	28	0.139818	10	26	False	1	1	0.058
DhansA	CGT	False	False	16	0.219205	7	15	False	1	1	0.000