

Cluster 15 - uncorrected

Seq118617	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	C	G	A	C	A	200
Seq68009	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq108154	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq238089	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq168357	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq202010	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq406671	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq356254	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	C	200
Seq295851	AGT	A	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq243825	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq43837	AGT	C	G	A	A	T	C	T	C	T	G	A	T	C	A	C	T	C	G	A	G	A	C	200
Seq27829	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq33413	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq204866	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq258270	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq386905	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq407120	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200

Cluster 15 - corrected

Seq118617	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	N	G	A	C	A	200
Seq68009	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq108154	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq238089	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq168357	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq202010	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq406671	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq356254	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	N	200
Seq295851	AGT	N	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq243825	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq43837	AGT	C	G	A	A	T	C	T	C	T	N	A	T	C	A	C	T	C	G	A	G	A	C	200
Seq27829	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq33413	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq204866	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq258270	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq386905	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200
Seq407120	AGT	C	G	A	A	T	C	T	C	T	A	T	C	A	C	T	C	G	A	G	A	C	A	200