PublisherInfo				
PublisherName		BioMed Central		
PublisherLocation		London		
PublisherImprintName	:	BioMed Central		

Downsizing the Paramecium genome

ArticleInfo		
ArticleID	:	3826
ArticleDOI	:	10.1186/gb-spotlight-20001109-02
ArticleCitationID	:	spotlight-20001109-02
ArticleSequenceNumber	:	263
ArticleCategory	:	Research news
ArticleFirstPage	:	1
ArticleLastPage	:	2
ArticleHistory	:	RegistrationDate: 2000–11–09OnlineDate: 2000–11–09
ArticleCopyright	:	BioMed Central Ltd2000
ArticleGrants	:	
ArticleContext	:	130591111

The elimination of specific DNA elements is a common feature in the formation of a transcriptionally active macronucleus during the sexual reproduction of ciliated protozoa. The micronuclear DNA of *Paramecium tetraurelia* contains some 50,000 internal eliminated sequences (IES), each of which is flanked by TA dinucleotide repeats and inverted sequences. In the November Molecular and Cellular Biology Ku *et al.* describe an in vivo method to analyse IES excision (*Mol Cell Biol* 2000, **20**:8390-8396). They injected DNA containing a 28 base pair IES and a drug-resistance gene into mated ciliate cells. They selected for transformants and assayed for excision by simple PCR amplification. Analysis of various deletions showed that flanking sequences are essential for excision, whereas mutations within the IES had no effect. This study reveals similarities between *Paramecium* and DNA excision events in Tetrahymena.

References

1. Genome downsizing during ciliate development: nuclear division of labor through chromosome restructuring.

- 2. Molecular and Cellular Biology, [http://intl-mcb.asm.org]
- 3. Programmed DNA deletions in Tetrahymena: mechanisms and implications.