PublisherInfo				
PublisherName		BioMed Central		
PublisherLocation		London		
PublisherImprintName	$\Box$	BioMed Central		

## ATM splicing defect

ArticleInfo			
ArticleID		4422	
ArticleDOI	$\Box$	10.1186/gb-spotlight-20020314-01	
ArticleCitationID	$\Box$	spotlight-20020314-01	
ArticleSequenceNumber	$\Box$	88	
ArticleCategory	$\begin{bmatrix} \vdots \end{bmatrix}$	Research news	
ArticleFirstPage	:	1	
ArticleLastPage	$\Box$	2	
ArticleHistory	:	RegistrationDate : 2002–3–14 OnlineDate : 2002–3–14	
ArticleCopyright	:	BioMed Central Ltd2002	
ArticleGrants	:		
ArticleContext	$\Box$	130593311	

## Jonathan B Weitzman

Email: jonathanweitzman@hotmail.com

Individuals with mutations in the *ATM* gene develop ataxia-telangiectasia, a neurodegenrative disorder characterized by immunological defects and cancer predisposition. In an Advanced Online Publication from Nature Genetics, Pagani *et al.* describe a new kind of *ATM* mutation that leads to an unusual splicing defect (11 March 2002, DOI:10.1038/ng858). The mutant *ATM* allele contains a four-nucleotide deletion (GTAA) within intron 20. This deletion results in the inclusion of a 65 nucleotide 'cryptic exon' in the *ATM* mRNA. The ATM sequence, termed intron-splicing processing element (ISPE), is complementary to U1 snRNA. Experiments with a hybrid minigene confirmed the importance of the ISPE sequence; and interaction with U1 snRNA affected the efficiency of intron removal. Introduction of the ISPE sequence into a different genomic context, exon 9 of the cystic fibrosis transmembrane regulator (*CFTR*) gene, caused exon skipping and splicing defects.

## References

- 1. The genetic defect in ataxia-telangiectasia.
- 2. Nature Genetics, [http://www.nature.com/ng/]

This PDF file was created after publication.