PublisherInfo				
PublisherName		BioMed Central		
PublisherLocation		London		
PublisherImprintName		BioMed Central		

Discovering metabolic loci

ArticleInfo		
ArticleID	:	4684
ArticleDOI		10.1186/gb-spotlight-20030123-01
ArticleCitationID		spotlight-20030123-01
ArticleSequenceNumber	\Box	36
ArticleCategory	\Box	Research news
ArticleFirstPage	\Box	1
ArticleLastPage	\Box	2
ArticleHistory	:	RegistrationDate : 2003–1–23 OnlineDate : 2003–1–23
ArticleCopyright	\vdots	BioMed Central Ltd2003
ArticleGrants	:	
ArticleContext		130594411

Jonathan B Weitzman

Email: jonathanweitzman@hotmail.com

Bacterial genomes often contain clusters of genes involved in secondary metabolite biosynthesis that are not expressed until induction by specific chemical or physical stimuli. In an Advanced Online Publication in Nature Biotechnology Zazopoulus *et al.* describe a high-throughput genome-scanning method to detect such clusters in bacterial genomes (*Nature Biotechnology*, 21 January 2003, doi;10.1038/nbt784). Genome sequence tags (GSTs) were screened against a database of microbial gene clusters of metabolic loci in order to rapidly identify novel metabolic loci. Zazopoulus *et al.* used the genome-scanning approach to screen actinomycete strains for genes involved in the biosynthesis of enedyine, a class of anti-tumor antibiotics. They found a conserved cluster of five genes. The putative 'warhead gene cassette' includes a polyketide synthase gene that may be involved in the formation of the reactive chromophore ring structures found in all enediynes.

References

- 1. The calicheamicin gene cluster and its iterative type I enediyne PKS.
- 2. *Nature Biotechnology*, [http://www.nature.com/naturebiotechnology]

This PDF file was created after publication.