PLANT

The nucleotide sequence and genome structure of the 
geminivirus miscanthus streak virus

M. Chatani, Y. Matsumoto, H. Mizuta, M. Ikegami, M. I. Boulton and
J. W. Davies

Nucleotide sequence and gene organization of the 3'- 
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The sequence between nucleotides 161 and 512 of cowpea 
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Complete nucleotide sequence and genetic organization of 
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Non-replicating deletion mutants of brome mosaic virus 
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Reciprocal phenotype alterations between two satellite 
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Coat protein gene sequences of two cucumber mosaic virus 
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The 5'-terminal sequence of potato leafroll virus RNA: 
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Comparison of viral nucleic acid intermediates at early and 
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S. N. Covey and D. S. Turner

ANIMAL

Transcriptional activity across the Epstein–Barr virus 
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E. A. Kirchner, G. W. Bornkamm and A. Polack
Isolation and sequencing of the Epstein-Barr virus BNLF-1 gene (LMP1) from a Chinese nasopharyngeal carcinoma

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*In vivo* detection of metabolic changes in a mouse model of scrapie using nuclear magnetic resonance spectroscopy

A reovirus in the brown planthopper, *Nilaparvata lugens*

A single amino acid change in the E2 glycoprotein of Venezuelan equine encephalitis virus affects replication and dissemination in *Aedes aegypti* mosquitoes

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Efficacy of individual measles virus structural proteins in the protection of rats from measles encephalitis

Defective synthesis of envelope proteins by temperature-sensitive mutants representing complementation groups B and D of respiratory syncytial virus

Role of the *gag* and *pol* genes of human immunodeficiency virus in the morphogenesis and maturation of retrovirus-like particles expressed by recombinant vaccinia virus: an ultrastructural study

Neutralizing activity of anti-peptide antibodies against the principal neutralization domain of human immunodeficiency virus type 1

Construction and functional characterization of mutants of the bovine leukaemia virus trans-activator protein p34\textsuperscript{\textit{tax}}

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W. Goldmann, N. Hunter, G. Benson, J. D. Foster and J. Hope

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M. Sakurai, A. Taneda, H. Nagoya and K. Sekikawa

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Sequencing data on the long control region of human papillomavirus type 16

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Evolution of herpesvirus thymidine kinases from cellular deoxycytidine kinase

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