

Agricultural Journals

Czech Journal of

GENETICS AND PLANT BREEDING

home page about us contact

us

Table of Contents

IN PRESS

CJGPB 2014

CJGPB 2013

CJGPB 2012

CJGPB 2011

CJGPB 2010

CJGPB 2009

CJGPB 2008

CJGPB 2007

CJGPB 2006

CJGPB 2005

CJGPB 2004

CJGPB 2003

CJGPB 2002

CJGPB

Home

Editorial Board

For Authors

- AuthorsDeclaration
- Instruction to Authors
- Guide for Authors
- CopyrightStatement
- Submission

For Reviewers

- Guide for Reviewers
- ReviewersLogin

Subscription

Czech J. Genet. Plant Breed.

Complete genome sequence of a *Brome Mosaic Virus* isolate from the Czech Republic

Czech J. Genet. Plant Breed., 46 (2010): 178-182

An isolate of *Brome mosaic virus* (BMV) was originally isolated from Agropyron repens and maintained in Hordeum vulgare. The full-length genome of this isolate (BMV-CZ) was sequenced. Phylogenetic analysis revealed that BMV-CZ shared a minimum of 95.6% sequence identity, localized in the 5' -UTR of RNA-1 with the other BMV isolates from the database, and a maximum of divergence of 30.8% with Broad bean mottle virus localized in the 5' -UTR of RNA-3. This is the first sequence report of full-length BMV from the Czech Republic.

Keywords:

© 2011 Czech Academy of Agricultural Sciences

Diviv, sequence analysis, variability