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[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

- **Authors
Declaration**
- **Instruction
to Authors**
- **Guide for
Authors**
- **Copyright
Statement**
- **Submission**

For Reviewers

- **Guide for
Reviewers**
- **Reviewers
Login**

Subscription

Czech J. Genet. Plant Breed.

**M., Ismaili H., Vrapı H.,
Jaupi A., Smýkal P.:**

**Genetic diversity of
Albanian pea (*Pisum
sativum* L.) landraces
assessed by
morphological traits
and molecular markers**

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In order to investigate the genetic diversity present in the pea germplasm stored in the Albanian genebank, we analyzed 28 local pea genotypes of Albanian origins for 23 quantitative morphological traits, as well as 14 retrotransposon-based insertion polymorphism (RBIP) molecular markers. The study of morphological characters carried out during three growing seasons (2010, 2011 and 2012) had the objective of characterization of traits useful in breeding programs. RBIP marker analysis

revealed the genetic similarity in range from 0.06 to 0.45. ANOVA, principal component analysis (PCA) and cluster analysis was used to visualize the association among different traits. Most of the quantitative morphological traits showed significant differences. PCA and cluster analysis (Ward's method) carried out for morphological traits divided the local pea genotypes into three clusters. Finally, the study identified the agronomically important traits which will facilitate the maintenance and agronomic evaluation of the collections.

Keywords:

clusters analysis; genetic similarity; landraces; morphological traits; pea, retrotransposon

[[fulltext](#)]

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