

The use of microsatellites markers in molecular characteristics of hybrids between Pulsatilla species

Kateryna Fyałkowska ^{1*}, Anna Tereba¹, Grzegorz Łazarski²

*corresponding author: k.fyalkowska@ibles.waw.pl

² Institute of Biological Sciences, Siedlce University of Natural Sciences and Humanities, Bolesława Prusa 14 Street, PL

Introduction

Hybridization is widespread in plants. The genus Pulsatilla contains more than 30 species of herbaceous perennials. It is characterized by a relatively high hybridization rate. Spontaneous hybridization between P. patens and P. pratensis occurs in the natural habitats, most often in sites together with parental forms. Completed plastid genome provide molecular evidence for the hybrid origin of Pulsatilla ×hackelii [1] P. ×hackelii is morphologically intermediate between the pure P. patens and P. pratensis. The distinction between hybrid individuals and pure Pulsatilla species requires a whole set of morphological features. In our study we present the for P. vulgaris [2]. Some of loci can crossamplified only for one species from P. patens and P. pratensis.



Pulsatilla × hackelii leaf and flower. Fot. G. Łazarski

Methods and results

A hybrid specimen was found in the Świętokrzyskie Mountains, near Bocheniec village (southern Poland), in the vicinity of a forest road in a fresh mixed coniferous forest (Querco roboris-Pinetum). We amplified a set of ten microsatellites in two multiplex PCR reactions. From the ten analyzed loci, we successfully amplified eight. The analyzed microsatellites in population of three Pulsatilla species showed some species specific alleles. In case of the P. xhackelii sample, we obtained PCR products in the case of five loci developed for P. vulgaris, but successfully crossamplified in P. patens and P. pratensis. Additionally, we obtained results for two loci, which crossamplified only for P. patens and only one for P. pratensis (Figure 1). Two of the analyzed loci were polymorphic (PV32 and PV65).

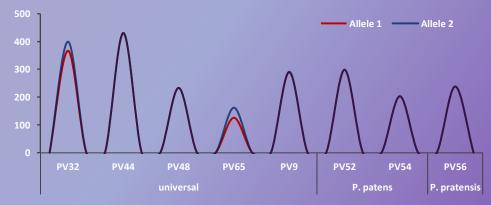


Figure 1. Genotypic characteristics of *Pulsatilla ×hackelii* based on microsatellites markers

Discussion and conclusions

Among the investigated markers, two were monomorphic (PV44 and PV48) and the remaining six were polymorphic. Markers PV32, PV65 and PV9 are highly polymorphic, what making them highly informative in population studies of pasque-flower species as P. patens, P. pratensis and P. vernalis. The obtained results indicated that analyzed microsatellites markers can be used for hybrids identification, although there is a need for further research on a larger number of hybridized individuals.

Pulsatilla pratensis.

References 1. Szczecińska, M.; Łazarski, G.; Bilska, K.; Sawicki, J. The complete plastid genome and nuclear genome markers provide molecular evidence for the hybrid origin of Pulsatilla × hackelii Pohl.. Turkish Journal of Botany 2017 41. doi:10.3906/bot-1610-28 2. Dileo, M.F; Graf, R.; Holderegger, R.; Rico, Y.; Wagner, H.H. Highly polymorphic microsatellite markers in Pulsatilla vulgaris (Ranunculaceae) using next-generation sequencing. Applications in Plant Sciences 2015 3 (7): 1500031.

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