



<http://dx.doi.org/10.11646/phytotaxa.167.3.9>

## Confirmation of *Valeriana nivalis* (Valerianaceae) to the Argentinian flora

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### Abstract

We confirm the occurrence of *Valeriana nivalis* to the Argentinean flora, based on results of both morphological and molecular phylogenetic studies. We also include a taxonomic description, distribution, and additional information to discern *V. nivalis* from *V. pycnantha*.

**Key words:** Argentina, Flora, New record, Taxonomy, *Valeriana*

### Introduction

*Valeriana* Linnaeus (1753: 31) includes ca. 250 species distributed all over the world except Australia and New Zealand. Most *Valeriana* species grow in temperate regions of the northern hemisphere, and particularly along the South American Andes, which is an important center of secondary diversification (Kutschker 2008a, 2011). The Argentinean flora includes 48 species of *Valeriana*, mainly distributed along the Andes and central mountains (Xifreda 1999, Kutschker 2008b, 2011).

Diagnostic characters used to discriminate species in most local taxonomic and floristic works on *Valeriana* (eg. Borsini 1942, 1944, 1966, 1999, Cabrera 1993) are considered inadequate (Reese-Krug & Weberling 1991). In addition, some nomenclatural problems have come to light (Reese-Krug & Weberling 1991, 1996). *Valeriana nivalis* Wedd. was recorded for Argentina by Borsini (1942, 1944, 1966, 1999) and Cabrera (1993). However, those records were based on specimens of *Valeriana pycnantha* A. Gray (Reese-Krug & Weberling 1991) and so *V. nivalis* did not occur in later accounts (Xifreda 1999; Kutschker 2008a).

After examining new collections from Jujuy province, Argentina (specimens from BAA, CONC, F, LIL, SI, ULM) and original descriptions, we confirm the existence of *V. nivalis* to the Argentinean flora. This taxonomic placement was confirmed using molecular phylogenetic analysis, based on Bell et al. (2012). Morphological description, distribution and additional material examined of *V. pycnantha* are also included to update information of a sympatric species that could be easily confused with it. In addition, we include *V. altoandina* Cabrera as a new synonym of *V. nivalis* based on a morphological analysis.

### Materials and methods

We examined collections from BAA, CONC, F, LIL, SI, ULM, original descriptions, floras (Borsini 1942, 1944, 1966, 1999, Cabrera 1993) and taxonomic works (Reese-Krug & Weberling 1991).

We amplified and sequenced the chloroplast marker *trnL-F* and the ribosomal ITS region of *V. nivalis* and *V. pycnantha*. DNA extraction, sequencing and alignment were carried out as described in Bell et al. (2012). We used Maximum Likelihood (ML) analysis to infer phylogeny as performed in Bell et al. (2012). Specimens used in molecular analysis are marked with (\*) in the list of examined material of each species. Taxon names and GenBank accession numbers for ITS and *trnL-F* sequences used in the phylogenetic analysis are given in TABLE 1.

## Acknowledgements

To Raúl Pozner, for critical reading and assistance with English translation. To Fernando Biganzoli (BAA), Alicia Marticorena (CONC) and Gerhard Gottsberger (ULM) for the images of the herbarium vouchers. To Adriana Kutschker & Charles Bell for the molecular data supplying. To the curators of the herbaria cited in this work.

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