

MV (Massive) *Pelophylax bergeri*

Date assessed	2021-10-18
Year published	2021
Eicat category	MV (Massive)
Justification for EICAT assessment	Impact through hybridisation between <i>P. bergeri</i> and <i>P. lessonae</i> is massive, hybrids have completely replaced the native <i>P. lessonae</i> (Dubey et al. 2014; Dufresnes et al. 2007, Dufresnes and Dubey 2020). The deep introgression has collapsed the genetic structure of the native <i>Pelophylax</i> populations (Dubey et al. 2014).
Confidence rating	High
Mechanism(s) of maximum impact	Hybridisation
Countries of most severe impact	France; Switzerland
Description of impact	Hybridisation - genetic introgression between <i>Pelophylax bergeri</i> and the native species <i>P. lessonae</i> and <i>P. esculentus</i> has led to the replacement of most pure native populations.
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Reviewers	EICAT authority
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