

Identification of tamaraw (*Bubalus mindorensis*) from natural habitat-derived fecal samples by PCR-RFLP analysis of cytochrome b gene

Abstract

Fecal DNA analysis is a useful tool for the investigation of endangered species. Tamaraw (*Bubalus mindorensis*) is endemic to the Philippine island of Mindoro but knowledge of its genetic and ecological information is limited. In this study, we developed a species identification method for tamaraw by fecal DNA analysis. Eighteen feces presumed to be from tamaraw were collected in Mount Iglit-Baco National Park and species-known feces from domestic buffaloes and cattle were obtained from a farm. Additionally, one species-unknown fecal sample was obtained in Mount Aruyan Preserve, where the sighting of tamaraw has not been reported in recent years. Based on DNA sequence data previously reported, the genus *Bubalus*- and tamaraw-specific primers for PCR of cytochrome b gene were newly designed. The *Bubalus*-specific primer yielded a 976 bp fragment of cytochrome b for all fecal samples from tamaraw and domestic buffaloes, but not for cattle, whereas the tamaraw-specific primer yielded a 582 bp fragment for all tamaraw fecal samples and for one of the four domestic buffalo samples. PCR-RFLP (restriction fragment length polymorphism) analysis of the 976 bp PCR fragment with *AvrII* or *BsaXI* provided distinct differences between tamaraw and domestic buffalo. PCR-RFLP analysis also showed that the species-unknown sample obtained in Mount Aruyan Preserve, originates from tamaraw. © 2010 The Authors. Journal compilation © 2010 Japanese Society of Animal Science.