

National Centre for Animal Health, Bhutan





# Identification of Echinococcus spp. in definitive and intermediate hosts in Bhutan

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#### Introduction

Earlier pilot studies indicated the presence of *Echinococcous* spp. in community/stray dogs and cystic echinococcosis (CE) in intermediate hosts in Bhutan. Records from Jigme Dorji Wangchuck National Referral Hospital (JDWNRH) also revealed the occurrence of CE in humans between the years 2006-2013 (Pelden 2013). However no epidemiological studies have been conducted to identify *Echinococcus* spp. prevalent in community/stray dogs and in intermediate hosts.



Community/stray dogs in Bhutan

## **Materials & Methods**

Through environmental sampling, 138 dog fecal samples collected from community/stray dogs from Thimphu, Gelegphu and Tsirang areas and also 28 fecal samples collected from the forest area near Mithun breeding farm from the central region were analyzed for the presence of taeniid eggs by the floatation and sieving method (F/Si) and by multiplex PCR (Trachsel *et al.* 2007).

In total, 55 cyst samples were collected from slaughterhouses (Tsirang), yak slaughter places (Thimphu, Paro, Haa) and imported beef carcass from checkpoints (Phuntsholing, Gelegphu, S/jongkha) in 2011-2012. The fertility status' of the cysts were determined by the presence of protoscoleces (PSC) in the hydatid fluid. The cysts without PSC as well as suppurated or calcified lesions were considered as infertile. The sediment containing PSC or germinal layer was collected and subjected to molecular diagnosis.

#### Molecular analysis

Genomic DNA was extracted from each sample (n=58) using the tissue protocol described in the Qiagen mini kit (Qiagen, Hilden, Germany). All samples were subjected to PCR targeting two mitochondrial genes, NADH dehydrogenase subunit 1 (nad1) and cytochrome oxidase subunit 1 (cox1) using the primers and the conditions described by Bart *et al.* (2006).

In order to assess the species level for *Taenia* spp. (fecal samples) and *Echinococcus* spp. (cyst samples), the amplicons of each positive sample were sent for direct sequencing, after purification using the MinElute PCR purification kit (Qiagen, Hilden, Germany) according to the manufacturer's instructions (for further details see Thapa *et al.* 2017).

Table 1. Identification of Echinococcus spp. in fecal samples of canids

Sample origin	Taeniid	Multiplex – PCR findings	Sequence findings
	positive		

## **Results & Discussion**

Table 1 depicts the findings of the microscopic examinations of the fecal samples, Multiplex-PCR and sequencing analysis and Table 2 shows the findings of the cyst samples.

*E. granulosus* s.l. has been identified in the community/stray dogs in Bhutan which could be due to improper slaughter practices of the animals. Similar findings have been reported from other parts of Asia including northern India and Nepal (Reece 2005).

From the cysts, *Echinococcus granulosus* (G1–3) was the prominent species found, the findings are in accordance with earlier studies conducted in Nepal (Joshi *et al.* 1997). In Bhutan, sheep is reared for mainly wool purposes in small numbers. Therefore, samples could not be collected from sheep to confirm a dog sheep cycle. However, the presence of *E. ortleppi* in cattle indicates the existence of a dog cattle cycle in Bhutan. Cyst samples from humans will be analyzed in the near future and further studies are required to elucidate the epidemiology of *Echinococcus* in Bhutan.



A Yak (Bos grunniens) from Bhutan

#### Table 2. *Echinococcus* spp. identified in cysts (n=54) from cattle or yaks

Origin	Infectivity of cysts	E. granulosus	E. ortleppi
		(G1-G3)	(G5)
Imported beef	Fertile	4	1
(India, Nepal)	Sterile	31	0
Cattle slaughtered	Fertile	1	1
in Bhutan	Sterile	6	0
Yaks slaughtered	Fertile	0	0
in Bhutan	Sterile	10	0

Stray or				
community				
dogs (n=138)				

Chronicar

Forest area surrounding Mithun farm (n=28)

20	16 non-Echinococcus	1 T. hydatigena,
	cestodes	9 Spirometra sp.,
		6 not sequenced
	10 <i>E. granulosus</i> sensu lato	Not further
	(8 mixed infection with non-	characterized
	Echinococcus cestodes)	
	2 negative	
14	8 non-Echinococcus	6 T. hydatigena,
	cestodes	1 H. taeniaeformis,
		1 not sequenced
	1 E. granulosus s.l.	Not sequenced
	6 negative	

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