# A REGION-WIDE SURVEY IN AOSTA VALLEY FOR THE PRESENCE OF TICKS AND TICK-BORNE DISEASES



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## **1. BACKGROUND AND RATIONALE OF THE STUDY**

- Ticks are important vectors of many viruses, bacteria and protozoa that can cause serious infections in humans and animals.

- The epidemiology of vector-borne diseases (VBD) is rapidly changing [1] and VBD are becoming a global public health/veterinary issue that needs active surveillance [2].

- Mountain areas are preferential sites to study tick ecology as climatic conditions are exasperated and subject to more extreme changes [2]

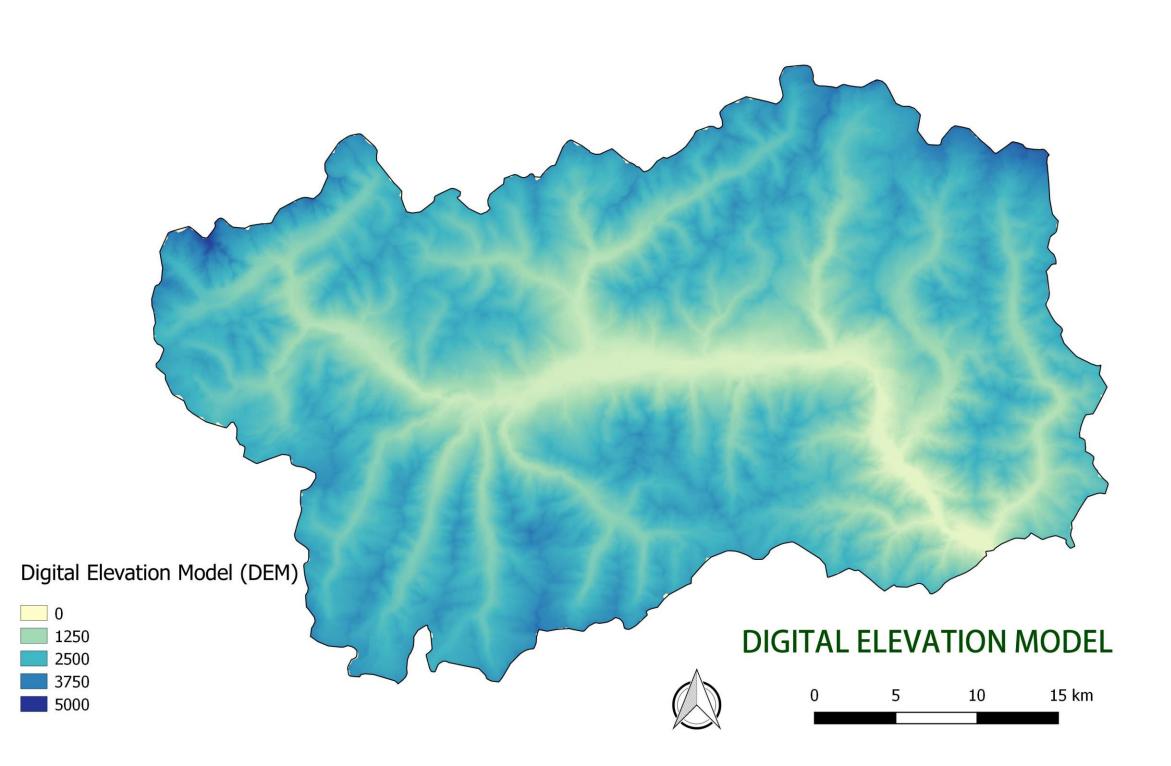
## 2. METHODS

## 2a. Identification of sampling sites

GIS-based analysis using Q-GIS software was performed to identify suitable sampling sites on the basis of Altitude, Solar Radiance and Land Cover.

## GOALS

- survey distribution and seasonal abundance of ticks in Aosta Valley



## 3. RESULTS

**3a.** 535 ticks were collected and identified as Ixodes ricinus (n=533) and Ixodes hexagonus (n=2)

(Northwestern Italy)

- analyze the collected ticks for the presence of Babesia spp., Theileria spp., Anaplasma spp. and Ehrlichia spp.

#### 2b. Tick sampling

- Environmental dragging was performed monthly from May 2016 to April 2017 (12 consecutive months) in 34 locations.

 Biotic and abiotic variables were recorded at each sampling site: Air and ground humidity (%) Temperature (°C) Vegetation coverage (canopy vs. bush, ammount) Slope and aspect, elevation and slope exposure

#### **2c. Species identification**

Morphological identification of ticks was carried out by means of optical microscopy fallowing keys of Manilla and Iori [4].

#### 2d. Molecular detection of tick-borne pathogens

-Ticks were pooled by sampling transect, species and development stage and total genomic DNA was extracted using TRIZOL (Sigma Aldrich).

- Species-specific PCR protocols were used on pooled samples to assess Minimum Infection Rate (MIR) of: Babesia/Theileria spp. [5] Anaplasma/Ehrlichia spp. [6]





I. hexagonus: 2 adults

I. ricinus: 70 adults, 380 nimphs, 83 larvae

SEX: F- 25 M-19

At least one tick found in 29.41% of transects [n=10/34] (Cl95% 16.83 – 46.17%)

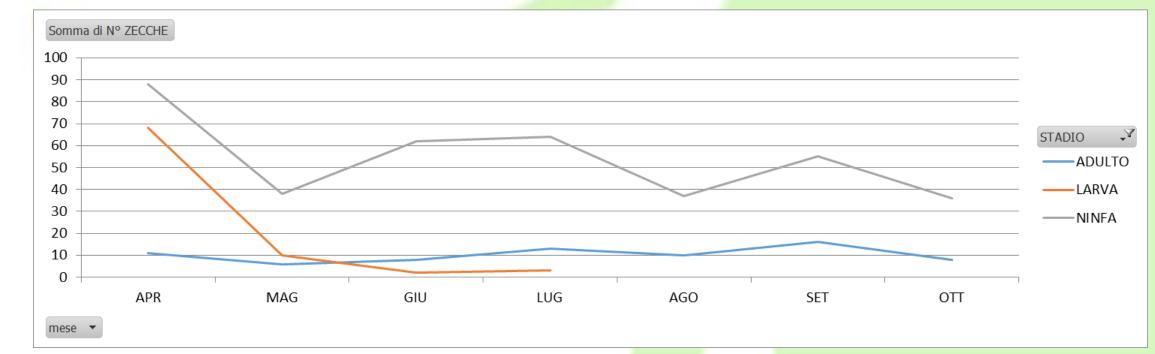
Positive correlationLandCover forest p<0.05</th>Elevation p<0.01</td>Negative correlationSolar radiance p<0.05</td>

Babesia/Theileria sppP= 15.12; Cl95% 9.05% -24.16%)Anaplasma/Ehrlichia sppP= 19.77%; Cl95% 12.72% - 29.40%

MIR A. phagocytophilum **P=0.80**% MIR B. capreoli **P=3.63**% MIR Cd. Neohrlichia mikurensis **P=4.44**% MIR B. microti-like **P=0.40**%

- Sequencing of all PCR positive pools was performed to confirm molecular identification.

### 3b. Seasonal population dynamic



## 4. CONCLUSIONS

- Ticks and VBD wide-spread across the study area.
- Vector-borne pathogens tipically related to small mammals hosts are the most prevalent pathogens in larvae and nimphs.

- Environmental covariates are significant predictors of vector abundance

- GIS-based models can be succesfully used to predict vector and pathogen presence and abundance.



#### [1] Heyman et al. A clear and present danger: tick-borne diseases in Europe. Expert Rev. Anti-Infect. Ther., 8: 33–50, 2010. [2] Mwamuye et al. Novel Rickettsia and emergent tick-borne pathogens: a molecular survey of ticks and tickborne pathogens in Shimba Hills National Reserve, Kenya. Ticks Tick-borne Dis., 8(2): 208–218, 2017.[4] Manilla and Iori . Chiave illustrate delle zecche d'Italia. Parassitologia, 34:83-95, 1992. [5] Zanet et al. Piroplasmosis in Wildlife: Babesia and Theileria Affecting Free-Ranging Ungulates and Carnivores in the Italian Alps. Par. Vectors, 7 (1): 70, 2014. [6] Munderloh et al. Isolation of the equine granulocytic ehrlichiosis agent, *Ehrlichia equi*, in tick cell culture. J. Clin. Microbiol. 34:

