



ASSESSING THE ECOLOGICAL COVARIATES RELATED TO TICK-BORNE ENCEPHALITIS EMERGENCE IN EUROPE



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BACKGROUND AND AIM METHODS • TBE spatial distribution is showing altitudinal and longitudinal 3. DATA COLLECTION 5. STATISTICAL ANALYSIS **1. LITERATURE SEARCH** shifts, with **new foci** emerging in non-endemic countries^{1,2} • Data of TBE human infections provided by • Single-variable linear regression to (MEDLINE, EMBASE, TESSy (ECDC) assess the relationship between SCISEARCH, BIOSIS, • Bioclimatic, environmental, and ecological TBE incidence and each covariate HCAPLUS, SCOPUS). · To geo-locate all known and potential new risk areas it is raw data • Multi-variable linear regression to fundamental to.. identify the **best parsimonius** model



RESULTS

LITERATURE SEARCH

• We retrieved Information about **covariates** from 62 articles



BEST PARSIMONIOUS MODEL (LMM)

Log(TBE_incidence) ~ Mean winter temp + Mean diurnal temp range + Autumnal cooling rate + % of forested area + EVI + M. glareolus + A. flavicollis + Deer presence + Deer presence² + (1|Reporting Country)



Figure 1 – Main characteristics of the studies included. a) Geographical distribution. b) Type of covariates adopted. c) Number of articles by year of publication and type of analysis.

• We selected for further analysis **31 covariates**, *i.e.* the ones adopted in at least 2 articles:



confidence interval, dots: observed data.). Actual covariate values are shown in the maps.

REMARKS

Environmental drivers (a) Forest cover (b) EVI

Climatic drivers (c) Autumnal cooling rate (d) Mean winter temperature (e) Mean diurnal temperature range

Hosts (f) A. flavicollis (yellow-necked mouse) (g) M. glareolus (bank vole) (h) Deer (D. dama, C.elaphus, C.capreolus) Establishing the precise ecological conditions that favor TBE spread is a challenge, which is reflected in the high heterogeneity of covariates that have been investigated in existing literature. This work will therefore provide essential inputs for the implementation of modern modeling approaches aimed at predicting the risk of spatio-temporal disease spread.

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