



# Be-FAST – Between Farm Animal Spatial Transmission: An epidemiological model for studying the spread and the economic impact of animal diseases.



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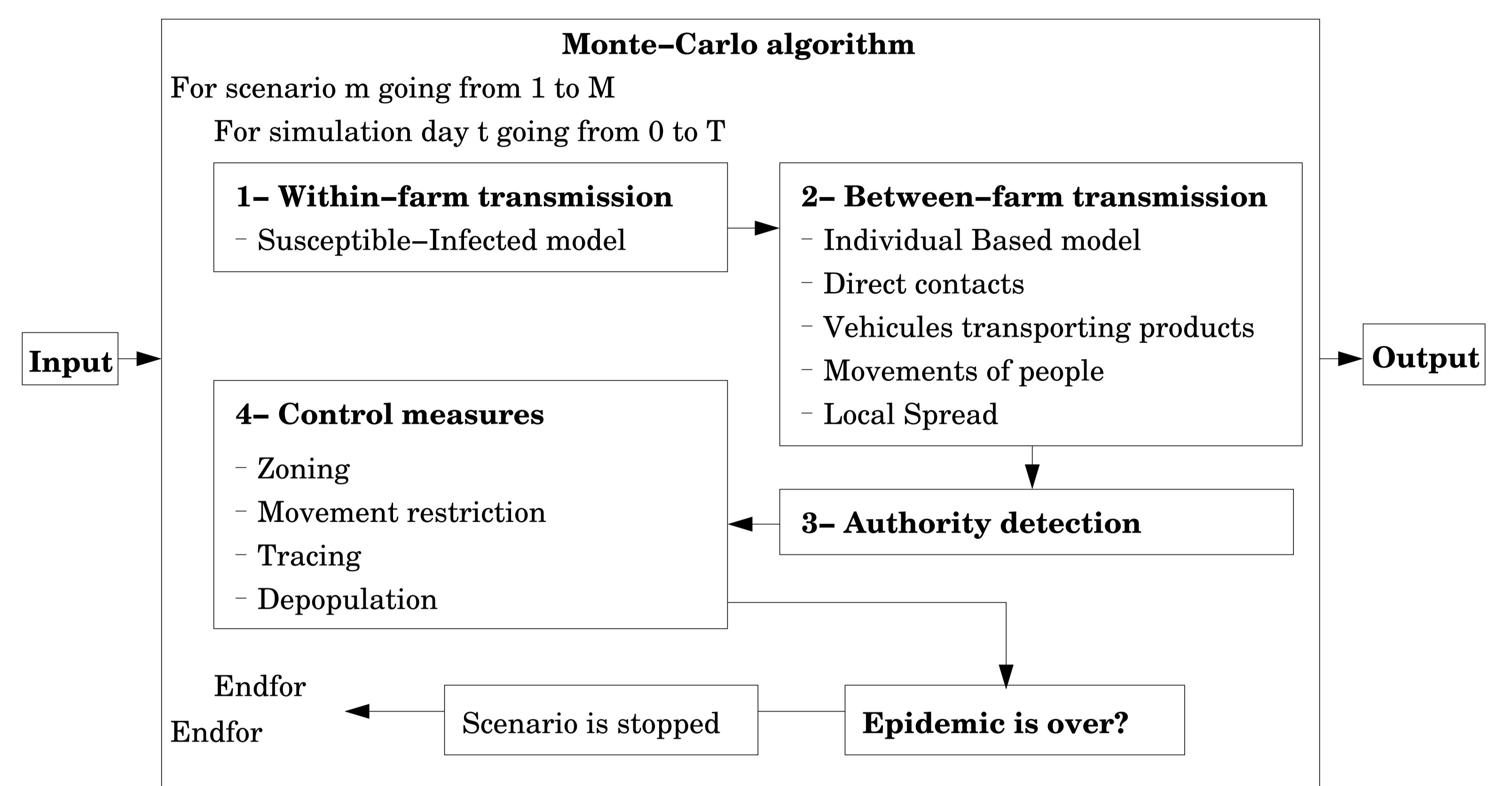
Work founded by: Spanish National Project MTM2011-22658 – European Project CSFV go-DIVA (KBBE-227003) – Proyecto regional integrado para el control progresivo de la fiebre aftosa (GCP/RLA/178/SPA Y GTSF/RLA/172/ITA) – \* E-mail: ivorra@mat.ucm.es



## 1. Main characteristics of Be-FAST

- Simulate both the **between farm** and **within farm** spread of the considered disease in a given region.
- Evaluation of the **economical impact** of a disease outbreak by considering direct and indirect costs.
- Adaptable to **different animal diseases**.
- Possibility to use **real and complex database** for farms and transports for simulating realistic commercial contacts between farms.
- Possibility to use **dynamic coefficients** based, for instance, on the number of infected animals.
- **Large choice of output**, such as: risk maps, R0 values, statistics on the magnitude and duration of epidemic, etc.

## 2. Structure of the model

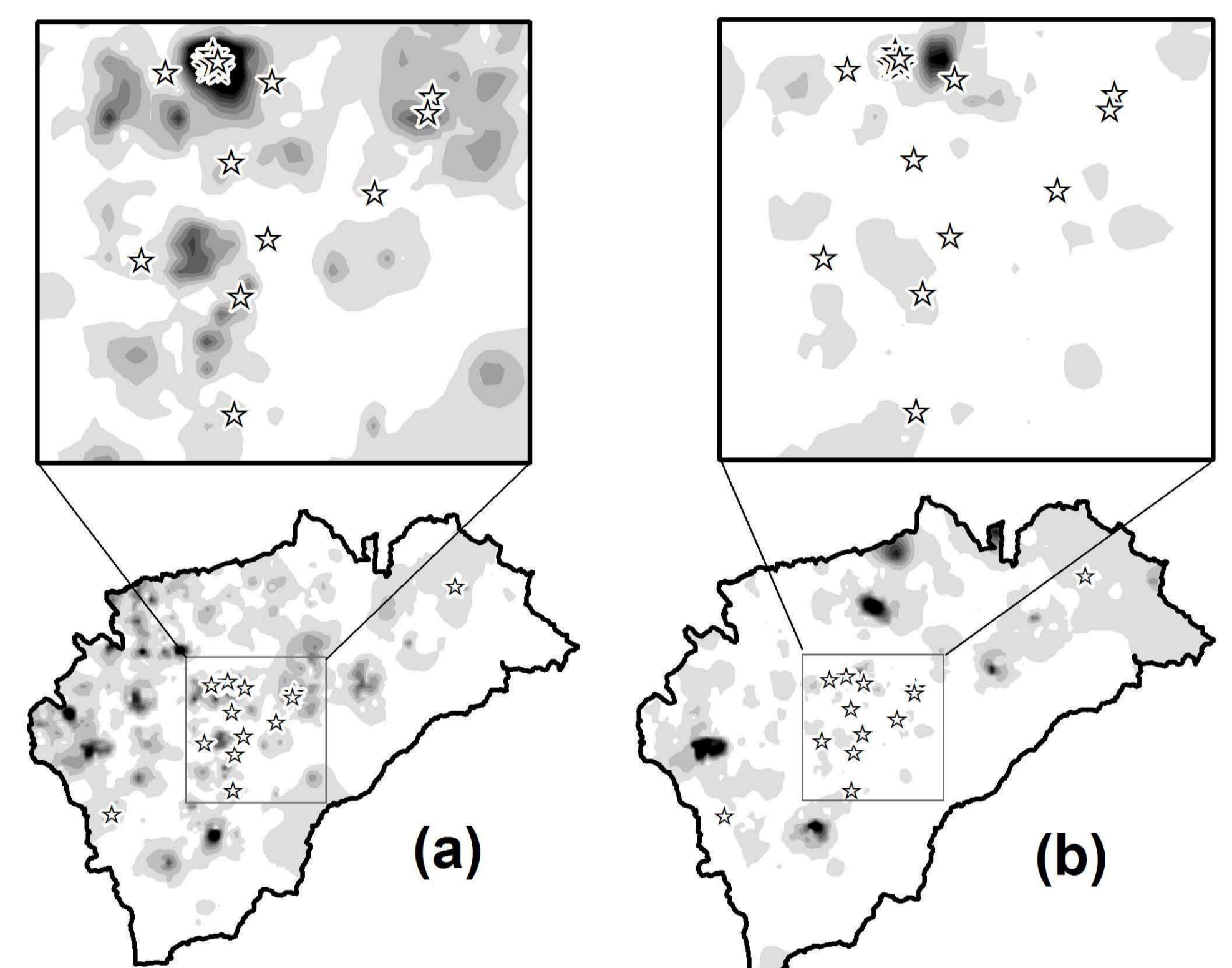


## 3. Model validation and sensitivity analysis: Classical Swine Fever in the Spanish province of Segovia

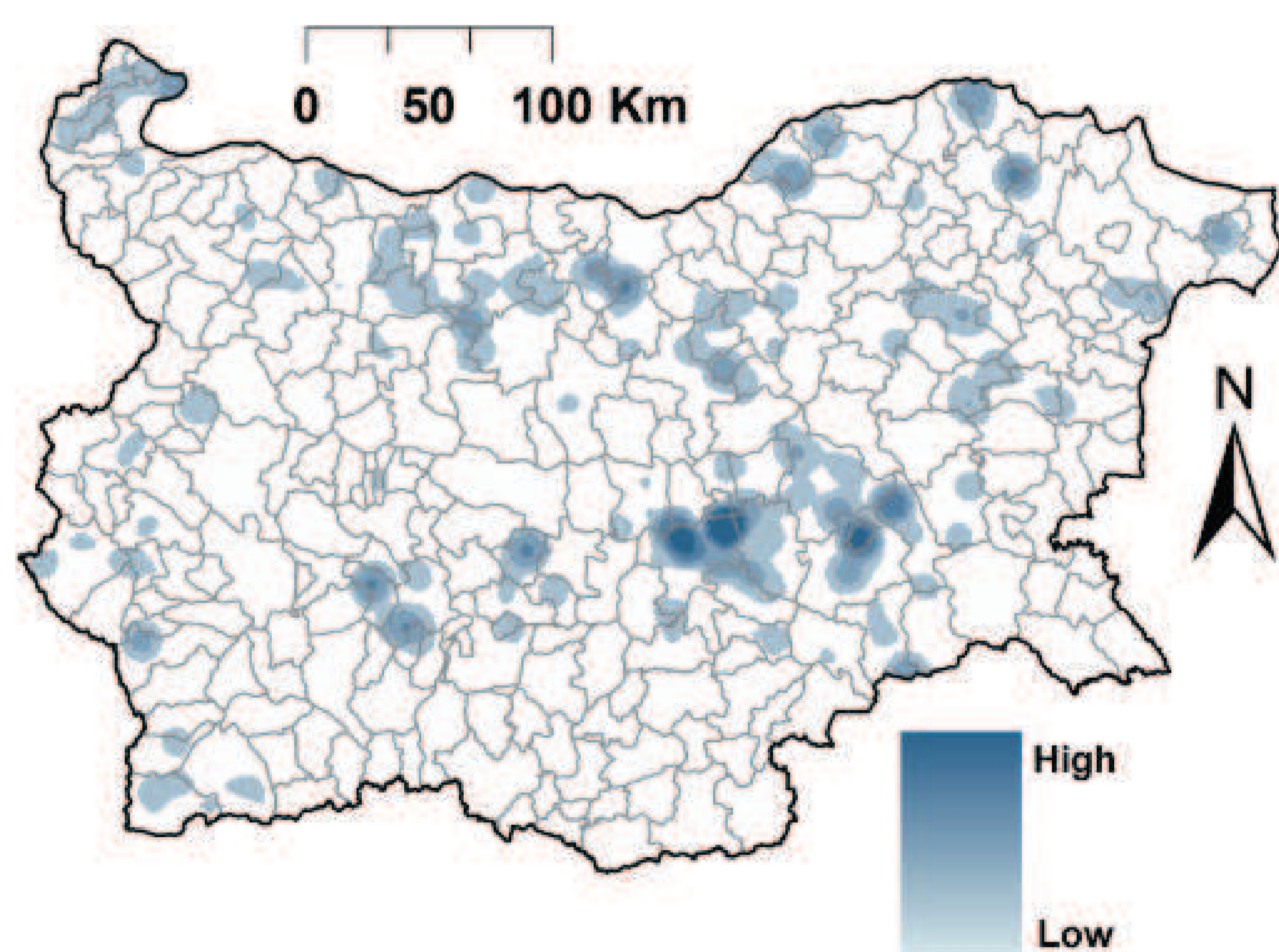
Some results obtained when considering models **Be-FAST** and **InterSpread +** and **Real** outbreaks data occurring in 1997-98:

- **TABLE:** Computational time in hours; % of infection due to each CSFV **Route**: local spread (*LS*), Integrator vehicles (*INT*), Sanitary Defense Association persons (*SDA*) and transport of animals (*TA*); % of detection due to each control **Measure**: observation of clinical signs (*CS*), zoning (*ZO*) and tracing (*TR*)
- **FIGURE:** Risk map of CSFV introduction given by **a)** Be-FAST and **b)** InterSpread +. ★ denotes the location of real infected farms.

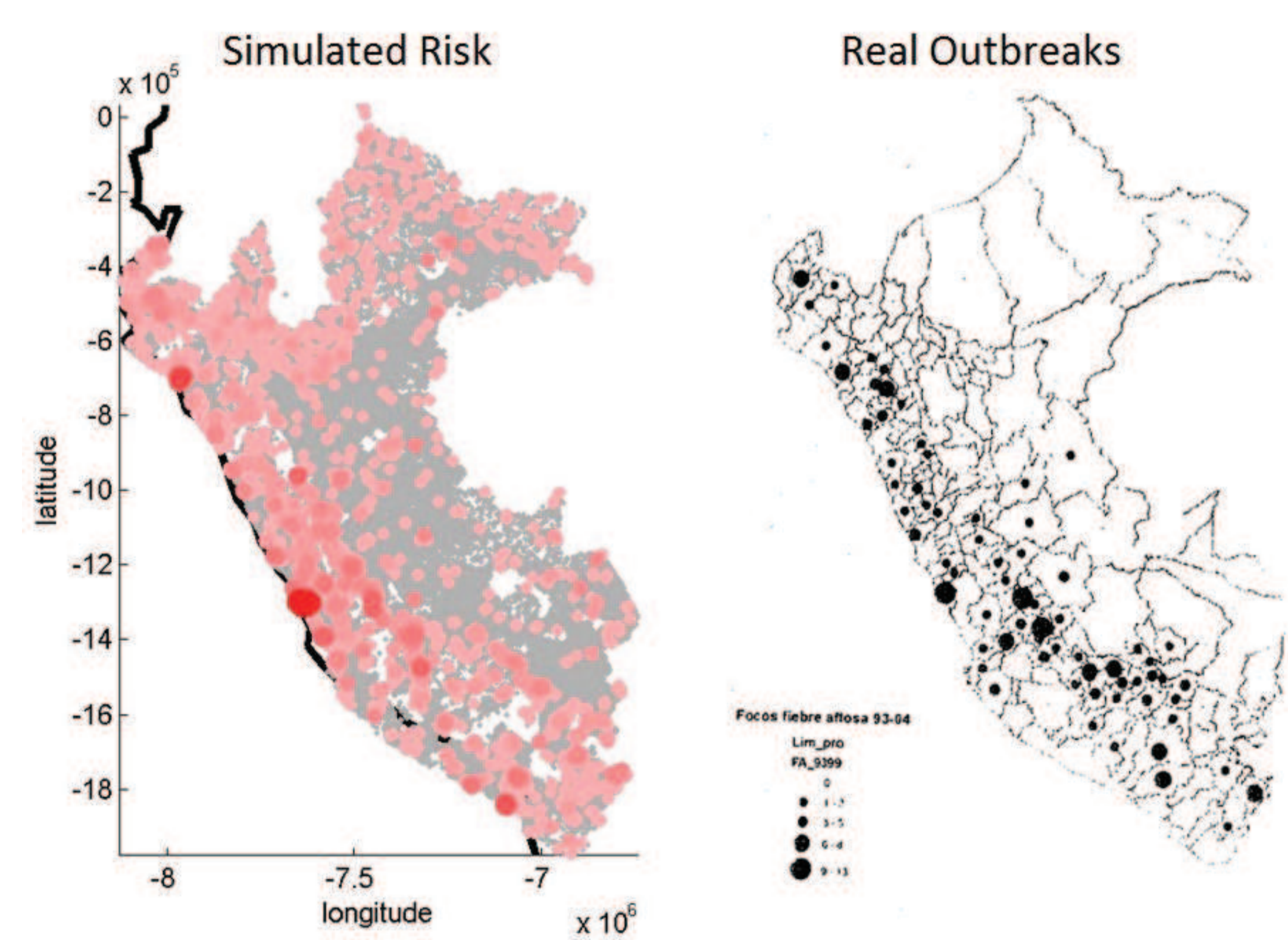
Model	Comp. Time (h)	Route				Measure		
		LS	INT	SDA	TA	CS	ZO	TR
<b>Be-FAST</b>	4	54	26	14	6	47	30	23
<b>Inter S+</b>	3	51	13	10	26	38	50	12
<b>REAL</b>	-	52	24	20	4	55	27	18



## 4. Application: Classical Swine Fever in Bulgaria



## 5. Application: Foot and Mouth Disease in Peru



## 6. Some References

- [1] B. Martínez-López, B. Ivorra, E. Fernández Carrión, T. Alexandrov, A.M. Ramos and J.M. Sánchez-Vizcaíno. *Evaluation of the risk of classical swine fever (CSF) spread from backyard pigs to other domestic pigs by using the spatial stochastic disease spread model Be-FAST: The example of Bulgaria.* **Veterinary Microbiology**. Science Direct Online. 2013
- [2] B. Ivorra, B. Martínez-López, J.M. Sánchez-Vizcaíno and A.M. Ramos. *Mathematical formulation and validation of the Be-FAST model for Classical Swine Fever Virus spread between and within farms.* **Annals of Operations Research**. Online First. 2013
- [3] B. Martínez-López, B. Ivorra, D. Ngom, A.M. Ramos and J.M. Sánchez-Vizcaíno. *A novel spatial and stochastic model to evaluate the within and between farm transmission of classical swine fever virus: II Validation of the model.* **Veterinary Microbiology**, **155**: 21-32. Elsevier. 2012.
- [4] B. Martínez-López, B. Ivorra, A.M. Ramos and J. M. Sánchez-Vizcaíno. *A novel spatial and stochastic model to evaluate the within and between farm transmission of classical swine fever virus: I. General concepts and description of the model.* **Veterinary Microbiology**, **147**: 300-309. Elsevier. 2011.