
REVIEW ARTICLE

Systematic review of surveillance systems and methods for early detection of exotic, new and re-emerging diseases in animal populations

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SUMMARY

In this globalized world, the spread of new, exotic and re-emerging diseases has become one of the most important threats to animal production and public health. This systematic review analyses conventional and novel early detection methods applied to surveillance. In all, 125 scientific documents were considered for this study. Exotic ($n = 49$) and re-emerging ($n = 27$) diseases constituted the most frequently represented health threats. In addition, the majority of studies were related to zoonoses ($n = 66$). The approaches found in the review could be divided in surveillance modalities, both active ($n = 23$) and passive ($n = 5$); and tools and methodologies that support surveillance activities ($n = 57$). Combinations of surveillance modalities and tools ($n = 40$) were also found. Risk-based approaches were very common ($n = 60$), especially in the papers describing tools and methodologies ($n = 50$). The main applications, benefits and limitations of each approach were extracted from the papers. This information will be very useful for informing the development of tools to facilitate the design of cost-effective surveillance strategies. Thus, the current literature review provides key information about the advantages, disadvantages, limitations and potential application of methodologies for the early detection of new, exotic and re-emerging diseases.

Key words: Animal health, surveillance, early detection, emerging diseases, exotic diseases, systematic review.

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INTRODUCTION

In this globalized world, the spread of new, exotic and re-emerging diseases has become one of the most important threats to animal production and public health, thus requiring the collaborative effort of many disciplines ('One Health' approach) [1]. Some examples include the appearance of 'new' threats, such as Schmallenberg virus (SBV) or bioterrorism agents, the spread of West Nile disease across naive populations of birds, horses and humans in the USA, previously considered 'exotic' (i.e. limited to the Eastern hemisphere), or the 're-emergence' of the avian influenza pandemic. The occurrence of these threats has highlighted the continuous change in disease spread patterns, perhaps as the result of the changing environment, the fact that the world population has increased, and the intensification of livestock productions and global trade [2–5]. Containing the spread of such diseases in this changing, ever-interconnected world requires new surveillance strategies for the rapid recognition of outbreaks and the diagnosis of causative agents, but also for the cost-effective allocation and implementation of resources for an efficient, timely response [6].

Early detection (or early-warning surveillance) can be defined as the 'surveillance of health indicators and diseases in defined populations in order to increase the likelihood of timely detection of undefined (new) or unexpected (exotic or re-emerging) threats' [7]. Early detection of disease (re-)introduction minimizes the potentially devastating consequences of disease spread [8]. For instance, the emergence of a new coronavirus responsible for the severe acute respiratory syndrome (SARS) pandemic in 2002–2003 resulted in thousands of human cases in Vietnam, Canada, Singapore, and Hong Kong, with more than 800 human deaths worldwide in less than 1 year [9]. This scenario demonstrated that proper infrastructure, protocols and techniques to quickly trace the origin and spread of the outbreak and accelerate its control were lacking [10]. Another example of the lack of effective techniques, in this case for a re-emerging disease, is the outbreak of foot-and-mouth disease (FMD) in the UK in 2001, which affected more than 10 million livestock and brought about economic losses of more than US\$12 billion, mainly due to farmer compensations and trade restrictions [11].

Historically, animal disease surveillance activities have focused on measuring the effects of specific endemic diseases or on detecting the introduction of

given exotic diseases [12]. These conventional disease-specific surveillance systems included the passive detection of cases combined with active sampling in the target population. Therefore, these unifocal strategies may result in delayed detection of new, exotic or re-emerging infectious diseases which, by definition, are unexpected and may be caused by unusual, or previously unknown, pathogens [13, 14]. The resulting poor timeliness, influenced by the time between the onset of an outbreak and its detection [15], is a major concern for highly contagious diseases [10, 14]. Other limitations of conventional surveillance systems include the high costs and poor sensitivity of active surveillance based on random sample surveys, and the frequent need to combine passive surveillance with an active surveillance component to improve the ability to detect hazards early [16, 17]. However, as these conventional methods might be still very efficient, the development of tools and strategies for more effective implementation is needed.

In recent years, new approaches have been developed in response to these emerging needs, which have led to the risk-based selection of hazards and the screening of populations at increased risk [18]. Although these novel epidemiological methodologies have the potential benefit to improve surveillance systems, they have not yet been sufficiently recognized to become a standard component of current internationally recognized surveillance systems. Thus, the RISKSUR project was initiated 'to find efficient and practical solutions by taking advantage of the novel scientific methodologies' and 'to define frameworks and integrated tools that allow the design and implementation of epidemiological and economically optimized animal health surveillance systems'. Consequently, the aim of this paper is to review conventional and novel methods that have already been developed and used to improve the likelihood of early detection of new, exotic and re-emerging diseases taking any legal requirements into account. A special focus of the review was the risk-based nature of the approaches, as well as to extract the main applications, benefits and limitations of each methodology. Wherever appropriate, gaps in the methodological concepts that can be targeted by new developments have been identified, as well as potential synergies between the described methodologies.

METHODS

Literature sources and search strategy

The systematic review targeted scientific documents on surveillance methods, systems and approaches

Table 1. *Terms used for the systematic search of scientific literature on early detection*

Section	Search terms
General theme (animal health surveillance)	surveillance OR monitor* AND animal* OR livestock OR veterinary* OR fish* OR wildlife OR 'food system*' OR herd* OR farm* OR cattle OR cow* OR bovine OR ruminant* OR pig* OR porcine OR swine OR sheep OR goat* OR poultry OR bird* OR avian OR horse OR equine OR equid* OR cat* OR dog* AND disease* OR health OR infection* OR outbreak AND
Exotic and emerging diseases	exotic OR emerg* OR
Early detection	'early warning' OR 'early detection' OR
Novel methodologies	syndromic OR participatory OR sentinel OR scanning

* Wildcards (searches for any word that includes the stem presented).

related to early detection. The literature search was performed on 21 January 2013 using the CAB Abstracts and the Scopus databases, which cover more than 90% of veterinary journals [19]. It was restricted to documents written in English (for reviewing convenience) that have been published over the last 20 years (i.e. 1993–2012) to represent the most up-to-date scientific information. This search included published articles, conference proceedings and reports.

A list of search terms was combined into a Boolean query in order to identify articles containing information on animal health surveillance and methodologies for early detection (Table 1). The search terms were searched for in the titles and abstracts of the documents.

Definitions

For the purpose of this review, we defined the epidemiological terms and concepts in accordance with the report from a workshop held by international experts on surveillance (<http://www.animalhealthsurveillance.org/index.php?n=Main.TerminologyFinal>). This report distinguishes between the patterns of disease occurrence as follows: (i) 'New': a previously undefined (unknown) disease or condition, which may result from the evolution or a change in an existing pathogen or parasite, which causes a change of strain, host range, vector, or an increase in pathogenicity; or it may imply the occurrence of any other previously undefined condition; (ii) 'Exotic': a previously defined (known) disease that crosses political boundaries to occur in a country or region in which it is

not currently recorded as being present; and (iii) 'Re-emerging': a previously defined (known) disease that is currently absent or present at a low level in the population in a defined geographical area that reappears or significantly increases in prevalence. In the 'Re-emerging' category, we also considered those new or exotic vector-borne diseases that reoccur in subsequent vector seasons after their first introduction into a region.

Study selection

The literature selection process is illustrated in Figure 1. A two-step process was followed to apply the primary and secondary exclusion criteria in order to select articles that are relevant for the purpose of this review. The first step consisted in excluding the papers that fulfilled at least one of the primary exclusion criteria when reading the title and abstract (Table 2). Then, the full text of each selected paper was assessed in a second screening round for eligibility by five reviewers, who reviewed different sets of articles independently (i.e. the sets of articles were exclusive per reviewer), but all followed common criteria. These reviewers selected the eligible articles by reading the full text and after applying both the primary and secondary exclusion criteria (Table 2).

A list of 47 variables was defined for the purpose of this review (see Supplementary Table S1). The selected variables allowed to extract the following information from the selected articles: disease or threat considered, species involved, area and time-frame of the study, type and characteristics of the approach used (e.g.

Table 2. Frequency and proportion of each exclusion criterion used in the article selection process

	Exclusion criteria	Count	Percentage
Primary	The paper does not focus on animal health	14	9.59%
	The paper focuses more on control measures than on surveillance	9	6.16%
	The disease is endemic in the study area	8	5.48%
	The paper does not focus on surveillance as defined for this project	8	5.48%
	The paper focuses on molecular characterizations of a pathogen	7	4.79%
	The paper is only descriptive (historical trend of the disease, pathogenicity, etc.)	7	4.79%
	The paper is a general review of a particular disease	6	4.11%
	The paper focuses on experimental infections	3	2.05%
	Year of publication is before 1992	3	2.05%
	The paper does not focus on early detection	2	1.37%
	The paper focuses on diagnostic test evaluations	2	1.37%
	The papers focuses on the early detection of antimicrobial resistance	1	0.68%
	The paper focuses on the early detection of new species rather than of new diseases	0	0.00%
	The paper focuses on the evaluation of vaccine efficacy	0	0.00%
The paper is a case report	0	0.00%	
Secondary	The paper is a general review of (a) method(s) for surveillance	29	19.86%
	The paper is a general review of a group of diseases	22	15.07%
	Insufficient information to allow an evaluation of the method described	16	10.96%
	No surveillance design/methods described	9	6.16%

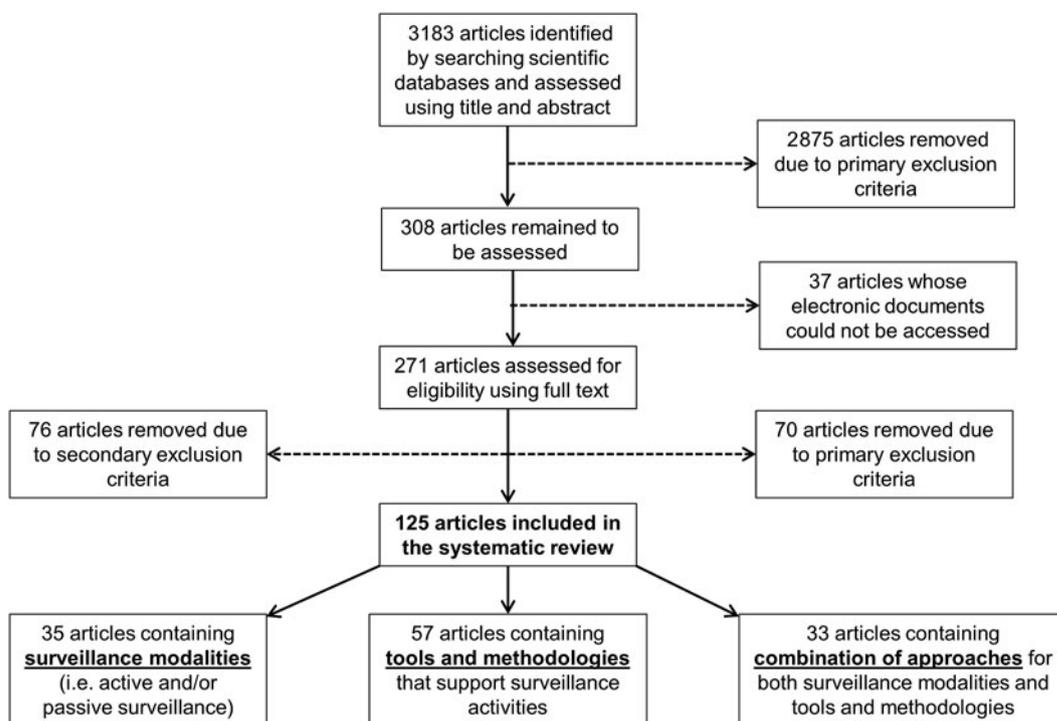


Fig. 1. Flowchart summarizing the article selection process.

risk-based nature, objective, data source and management, sampling strategy, main results, etc.), stage of surveillance that the approach can support, cost-effectiveness, limitations, future prospects, and method(s) used for validation/evaluation.

Certain variables deserved further explanations. First, the variable 'Pattern of disease occurrence' was composed of the categories 'New', 'Exotic', 'Re-emerging' (see Definitions section above), 'Endemic' and 'Not applicable (NA)'. Second, the

variable ‘Disease presence’ complemented the information of the previous one as far as the disease status in the study region was concerned. This variable included the categories ‘Present’, ‘Absent’, ‘Suspected’ (i.e. present, but neither detected nor yet confirmed), ‘Unknown’, and ‘Early detection’. This last category entailed those articles which focused on describing early detection methods that had been applied in multiple diseases or in diseases whose status was not clearly defined, which were linked to the category ‘NA’ of the ‘Pattern of disease occurrence’ variable. Third, the variable ‘Surveillance scope’ was related to the target of surveillance in terms of only one (i.e. ‘Single’) or several threats. In the latter case, two further options were considered: (a) ‘General’, when the surveillance system targeted a wide range of threats without specifying any disease (e.g. livestock diseases); (b) ‘Multi-objective’, when the aforementioned system focused on two or more specific threats (e.g. porcine circovirus-associated disease and porcine reproductive and respiratory syndrome).

Grouping of surveillance approaches

The papers included in this review included different types of surveillance approaches, which were divided in two categories. While some described the data collection, performance and/or results of surveillance activities (i.e. surveillance modalities), there were other papers that described different epidemiological methods for improving such activities. Thus, in order to allow a proper analysis and to create an inventory of surveillance approaches, we grouped all papers into the two main categories ‘Surveillance modalities’ and ‘Tools and methodologies supporting the implementation of surveillance activities’. The first one was divided into the two sub-categories ‘Active’ and ‘Passive’ surveillance, which were further divided into several types of active and passive strategies, respectively. Regarding the second category, the specific tools and methodologies used in each article were specified, as well as the stage of surveillance that they support. For the purpose of this review, we considered the following stages: (i) ‘Risk profiling’: procedures to estimate the probability of occurrence of diseases or threats or to determine the risk for certain population strata; (ii) ‘Sampling design’: procedures focused on the selection of representative members of the population that should be tested (e.g. when and where to sample, number and type of samples); (iii) ‘Data collection and management’: procedures to capture,

processing, and store relevant surveillance information and data; (iv) ‘Data analysis’: procedures to explore the information and to test hypotheses; and (v) ‘Evaluation’: procedures to assess the accomplishment of surveillance criteria to specified requirements or to compare performance of different surveillance systems.

All these categories and sub-categories were not exclusive, so combinations of them were indicated in detail for each article for further analysis.

Data extraction

Relevant information was extracted from the articles and was stored in an Excel (Microsoft Corp., USA) database. To ensure the harmonization of the criteria adopted to extract the information from the articles read by the reviewers and to minimize potential reviewer bias, two additional reviewers validated the Excel databases. These two additional reviewers read all full-text articles, cross-checked the entered variables, and resolved any potential disagreement.

Data analysis

Descriptive analyses of the collected data were performed. Variables were analysed separately and in combination in order to obtain as much information as possible from the articles. A special focus was the identification of the main applications, benefits and constraints of each type of methodology found in the review.

RESULTS

Selection process

The results of the selection process are shown in [Figure 1](#). The search made in the scientific databases returned 3183 papers. By applying the primary criteria, 308 papers were selected for which we were able to access the full text for 271. In the second screening round, 70 articles were excluded based on the primary exclusion criteria, which had not been possible based on the information provided in the title and abstract ([Table 2](#)). In addition, 76 papers were excluded based on the secondary exclusion criteria, meaning that the provided information was not appropriate for the purpose of this review. The frequency of exclusion in this second screening round is shown in [Table 2](#). The secondary criterion that most commonly led to exclusion was reviews of methods

Table 3. Frequency of the different surveillance modalities found in the review, also identifying the use of risk-based methods and the number of papers addressing multiple threats (for both, expressed as a count and percentage in parentheses)

General category	Specific category	Description	Count	Risk-based	Multiple nature	References
Active	Vector surveillance	Search for the vectors implied in the transmission of diseases	18	2 (11.11)	6 (33.33)	[20–37]
	Sentinel surveillance	Repeated collection of information from the same selected sites or groups of animals to identify changes in the health status of a specified population over time	15	9 (60.00)	1 (6.67)	[20–22, 24, 25, 33, 35, 38–45]
	Serological determination	Search for antibodies against a specific pathogen	13	5 (38.46)	1 (7.69)	[21, 46–57]
	Pathogen determination	Search for a specific pathogen (or its antigens or nucleic acids)	13	2 (15.38)	3 (23.08)	[20, 22, 33, 35, 46, 51, 53, 57–62]
	Participatory surveillance	Use of interviews with key informants for collecting qualitative health data from defined populations on health events, risks, impacts and control opportunities	7	1 (14.29)	5 (71.43)	[63–69]
	Others	Approaches not included in the above-mentioned categories	3	1 (33.33)	2 (66.67)	[70–72]
Passive	Clinical investigation	Monitoring the clinical signs compatible with disease(s)	13	0 (0.00)	0 (0.00)	[20, 22, 32, 33, 47, 49, 52, 55, 59, 73–76]
	Syndromic surveillance	Use of health-related information that may be indicative of a probability of change in the health of a population that merits further research or enables a timely impact assessment and action requirement	10		10 (100.00)	[74, 76–84]
	Mortality investigation	Monitoring an unexpected or increased mortality rate	5		0 (0.00)	[20, 33, 35, 49, 55]
	Parameter monitoring	Screening of biological indicators (e.g. animal temperature, animal activity or food/water intake)	2		1 (50.00)	[85, 86]
	Others	Approaches not included in the above-mentioned categories	3		2 (66.67)	[72, 75, 87]

for surveillance ($n = 29$). While these excluded articles revised the use of and provided good knowledge about surveillance methods, the information presented was too general (i.e. applications not provided, not focused on early detection, lack of examples of new, exotic and/or re-emerging diseases). After completion of the screening rounds, 125 articles were included in the review (Tables 3 and 4).

Descriptive results

Most of the documents included in the review were research articles ($n = 99$), although poster proceedings

($n = 13$), reviews ($n = 5$), reports ($n = 4$) and oral communication proceedings ($n = 4$) were also included. The majority of the documents were published after 2005, particularly in 2011 ($n = 33$) and 2012 ($n = 26$).

Exotic ($n = 49$) and re-emerging ($n = 27$) diseases were the most commonly represented health threats in this review, while new diseases appeared less frequently ($n = 11$). In addition, the majority of studies were related to zoonoses ($n = 66$). Particularly, West Nile fever ($n = 22$), bluetongue ($n = 16$), and the ‘avian influenza’ group ($n = 15$) stood out from the rest of diseases or categories of threats (Table 5). In addition, 17 articles dealt with several rare or diverse

Table 4. Frequency of the different methodologies and tools found in the review, also identifying the surveillance purposes the papers address

Specific category	Description	Count	Risk profiling	Sampling design	Data collection and management	Data analysis	Evaluation
Spatial epidemiology (GIS)	Spatial epidemiological approaches, especially those that are GIS-based	27	13 [26, 27, 29, 30, 48, 53, 91, 95, 98, 100–102, 104]	2 [48, 58]	10 [77, 88–90, 92–94, 96, 97, 99]	4 [24, 49, 101, 103]	0
Basic statistics	Application of statistical tests, formulas, or equations to a set of data	20	0	3 [98, 106, 107]	0	16 [24, 38, 42, 45, 50, 51, 61, 67, 74, 80, 83, 86, 96, 103, 105, 109]	6 [74, 98, 105–108]
Regression modelling	Mathematical models that are used to explore epidemiological and disease data and/or to evaluate distinct health measures	20	5 [32, 49, 101, 111, 115]	8 [49, 103, 109, 110, 112, 114, 116, 117]	0	15 [27, 50, 53, 61, 72, 79, 96, 103, 109–112, 114, 116, 117]	4 [49, 112, 113, 115]
Digital surveillance	Tools that use the Internet and computer technologies for collecting and processing health data (e.g. outbreak reports, surveillance results, etc.)	18	0	0	18 [72, 74, 77, 80, 82–84, 88–90, 92–94, 96, 97, 99, 118, 119]	0	0
Simulation modelling	Models to simulate disease dynamics (and their potential consequences) (e.g. SIR models, Markov models)	10	5 [41, 97, 120, 122, 123]	2 [107, 121]	0	4 [30, 119, 123, 124]	5 [41, 107, 121–123]
Decision modelling	Models to identify values, uncertainties and other parameters from a set of decisions, their rationale and the consequent optimal decision (e.g. scenario tree models, info-gap model, neural networks)	9	3 [124, 127, 128]	0	0	1 [81]	7 [59, 75, 120, 124–127]
Cluster analysis	Study of the spatial and/or temporal pattern of disease occurrence based on different scan statistics (e.g. Bernoulli, normal, Poisson, space–time permutation)	9	6 [32, 77, 87, 88, 98, 129]	1 [88]	0	4 [77, 87, 101, 103]	0

Table 4 (cont.)

Specific category	Description	Count	Risk profiling	Sampling design	Data collection and management	Data analysis	Evaluation
Environmental modelling	Models predicting the climatic and environmental factors that affect the replication and survival of pathogens or their vectors.	8	8 [26, 44, 48, 91, 93, 104, 130, 131]	0	0	0	0
Expert opinion	Professional judgement from a specialist on a specific subject of disease epidemiology or surveillance	5	2 [66, 125]	0	0	0	4 [66, 68, 132, 133]
Risk assessment	Statistical models characterizing the likelihood of exposure and/or release of threats given certain risk pathways	5	5 [98, 100, 134–136]	0	0	0	0
Literature review	Search for, and subsequent variable extraction from, a subset of scientific documents dealing with a specific topic	4	0	0	0	2 [137, 138]	2 [5, 139]
Data mining	Methodology to identify patterns in large datasets	4	1 [141]	0	0	3 [81, 82, 140]	0
Network analysis	Study and characterization of animal movement networks	2	2 [97, 123]	0	0	0	0
Others	Approaches not included in the above-mentioned categories	2	0	1 [143]	1 [142]	0	0

Table 5. Diseases or threats studied in the articles, grouped according to the general categories used in the systematic review

Diseases	Count	Zoonotic	Pattern of disease occurrence (frequency) in the study areas
West Nile fever	22	Yes	Rem (12), Ex (9), NA (1)
Livestock diseases and syndromes	17	Yes/no	NA (12), ND (2), New (1), End (2), Ex (2)
Bluetongue	16	No	Ex (10), Rem (3), New (2), NA (1)
Avian influenza	15	Yes	Ex (8), Rem (5), New (2)
Other new, exotic and/or re-emerging diseases	7	Yes/no	NA (7)
Rift Valley fever	5	Yes	Ex (4), End (2), Rem (1)
Diseases of aquatic animals	4	No	NA (4)
Foot-and-mouth disease	4	No	Ex (2), Rem (1), End (1)
Zoonoses	4	Yes	ND (1), NA (3)
Bovine tuberculosis	3	Yes	Rem (1), NA (2)
Food safety, animal and human health	3	Yes/no	Ex (1), ND (1), NA (1)
Abortions	2	Yes/no	NA (2)
Arboviral diseases	2	Yes	NA (2)
Bovine spongiform encephalopathy	2	Yes	Ex (1), NA (1)
Classical swine fever	2	No	Rem (2)
Lyme disease	2	Yes	Ex (2)
Porcine circovirus-associated disease	2	No	New (2)
Porcine reproductive and respiratory syndrome	2	No	New (2)
Salmonellosis	2	Yes	New (1), Ex (1)
A potential new or emerging disease	1	Yes/no	New (1)
African swine fever	1	No	Ex (1)
Bioterrorism agents	1	Yes	ND (1)
Colony collapse disorder	1	No	NA (1)
Eosinophilic meningitis (angiostromyiasis)	1	Yes	Ex (1)
Epizootic haemorrhagic disease	1	No	Ex (1)
Equine infectious anaemia	1	No	NA (1)
Erysipelosis	1	Yes	ND (1)
Febrile illnesses	1	Yes/no	NA (1)
Japanese encephalitis	1	Yes	Rem (1)
Newcastle disease	1	No	NA (1)
Pigeon paramyxovirus infection	1	No	New (1)
Rabies	1	Yes	Ex (1)
Rinderpest	1	No	Ex (1)
Scrapie	1	No	NA (1)
St Louis encephalitis	1	Yes	Rem (1)
Swine influenza	1	No	New (1)
Usutu virus infection	1	Yes	Ex (1)
Wildlife diseases and syndromes	1	Yes/no	NA (1)

For the 'Pattern of disease occurrence' variable, we considered different categories: endemic (End), new (New), exotic (Ex), re-emerging (Rem), not applicable (NA) and not defined (ND). The 'NA' category was indicated for those articles dealing with early detection methods to be applied in multiple diseases or in diseases whose status was not clearly defined for the study areas. Note that the total count is higher than the number of papers considered in the review, since some papers deal with several diseases. Additionally, the sum of the frequencies might be higher than the count since the same disease might be included in the same study, but for several regions with different patterns of disease occurrence.

diseases affecting livestock, so it was not possible to include them in any specific category (i.e. considered in the category 'Livestock diseases and syndromes').

Regarding the geographical location of the studies, most related to North America ($n = 35$) and Europe ($n = 57$, Table 6). In fact, the countries that most frequently appeared (Fig. 2) were the USA ($n = 20$),

Canada ($n = 15$), Germany ($n = 10$), France ($n = 9$), Italy ($n = 9$), the UK ($n = 8$), and Spain ($n = 7$).

The review covered a wide range of domestic and wild target host species (Fig. 3). The commonest species was cattle ($n = 41$). Interestingly, a large proportion of studies published from 2001 onwards (i.e. 28.00%) focused on methods involving wildlife,

Table 6. Frequency of papers according to target diseases in different world areas

Continent	Count	Most relevant disease(s) or category(ies) of threat
Europe	57	Bluetongue ($n = 15$), West Nile fever ($n = 11$), avian influenza ($n = 9$), livestock diseases and syndromes ($n = 5$)
North America	35	West Nile fever ($n = 9$), livestock diseases and syndromes ($n = 6$)
Africa	10	Rift Valley fever ($n = 4$), livestock diseases and syndromes ($n = 5$)
Asia	10	Avian influenza ($n = 2$), livestock diseases and syndromes ($n = 3$)
Oceania	6	Foot-and-mouth disease ($n = 2$)
South America	3	Avian influenza ($n = 2$)
World	2	New, exotic and/or re-emerging diseases ($n = 1$), aquatic diseases ($n = 1$)
Central America	1	West Nile fever ($n = 1$)
(Not applicable)	1	
(No data)	4	

The category 'World' was assigned to those papers whose study areas were located in all the continents. Note that the figures in Figure 2 do not have to coincide with those in this table since this Table indicates the frequency of papers per world region, while Figure 2 indicates the frequency of the study countries (i.e. the study described in an article might have been performed in three countries in Europe).

especially in relation to exotic or re-emerging diseases ($n = 29$).

Description of the approaches used in the articles

We explored the approaches used in the articles by collecting variables about surveillance modalities or tools, data source (i.e. primary and secondary data), risk-based nature and surveillance scope. Twenty-three papers focused on active surveillance strategies, five on passive surveillance strategies, and 57 papers included tools and methodologies that support surveillance strategies. The remaining articles contained more than one of the above-cited categories ($n = 40$, e.g. active surveillance to collect data, followed by the implementation of an epidemiological model).

We identified papers whose data source was primary (i.e. data obtained from their own study; $n = 55$), secondary (i.e. data collected from other studies or sources; $n = 52$), or both ($n = 18$). Most papers were based on real data ($n = 61$), although 40 were based on simulations, and 24 on both.

The 'risk-based' concept already appeared in the eldest articles included in the review (i.e. 1995) (Fig. 4). In total, 60 papers included risk-based approaches, especially those including tools and methodologies (i.e. 50 of 60). When evaluating the risk-based categories (see Supplementary material, section S1), risk-based sampling ($n = 21$) and risk-based requirement ($n = 36$) seemed to be the most frequently used ones. Both types aimed to increase the chance of early detection of hazards by either selecting the target population to be sampled or identifying the critical points of surveillance programmes to intensify and improve their

performance. Risk-based sampling was used mainly when considering sentinel surveillance for exotic diseases. Conversely, risk-based requirement was applied to multiple disease types (i.e. new, exotic and re-emerging) and approaches. In this review, risk-based analyses ($n = 7$) were not highly represented, and risk-based prioritization ($n = 6$) was a minor category that was applied to methodologies dealing with absent diseases.

Regarding the 'surveillance scope', almost half ($n = 61$) of the papers covered multiple diseases, based on either general ($n = 41$) or multi-objective ($n = 20$) approaches. Most of the papers based on general methods were related to syndromic surveillance, which has a generalist nature *per se*. A multi-objective scope was typical of the papers describing methods, active surveillance strategies, and a combination of both. These papers dealt mainly with (i) methods that focused on specific diseases in certain host species ($n = 10$); (ii) methods that focused on a single disease, but can be applicable for others ($n = 5$); and (iii) vector surveillance ($n = 5$).

Specific approaches found in the papers: surveillance modalities

Each active or passive approach found in the review was assigned to a specific category in accordance with the methodologies applied and described in the paper (Table 3).

Active surveillance approaches

The two most often cited active surveillance activities found in the reviewed papers were vector surveillance

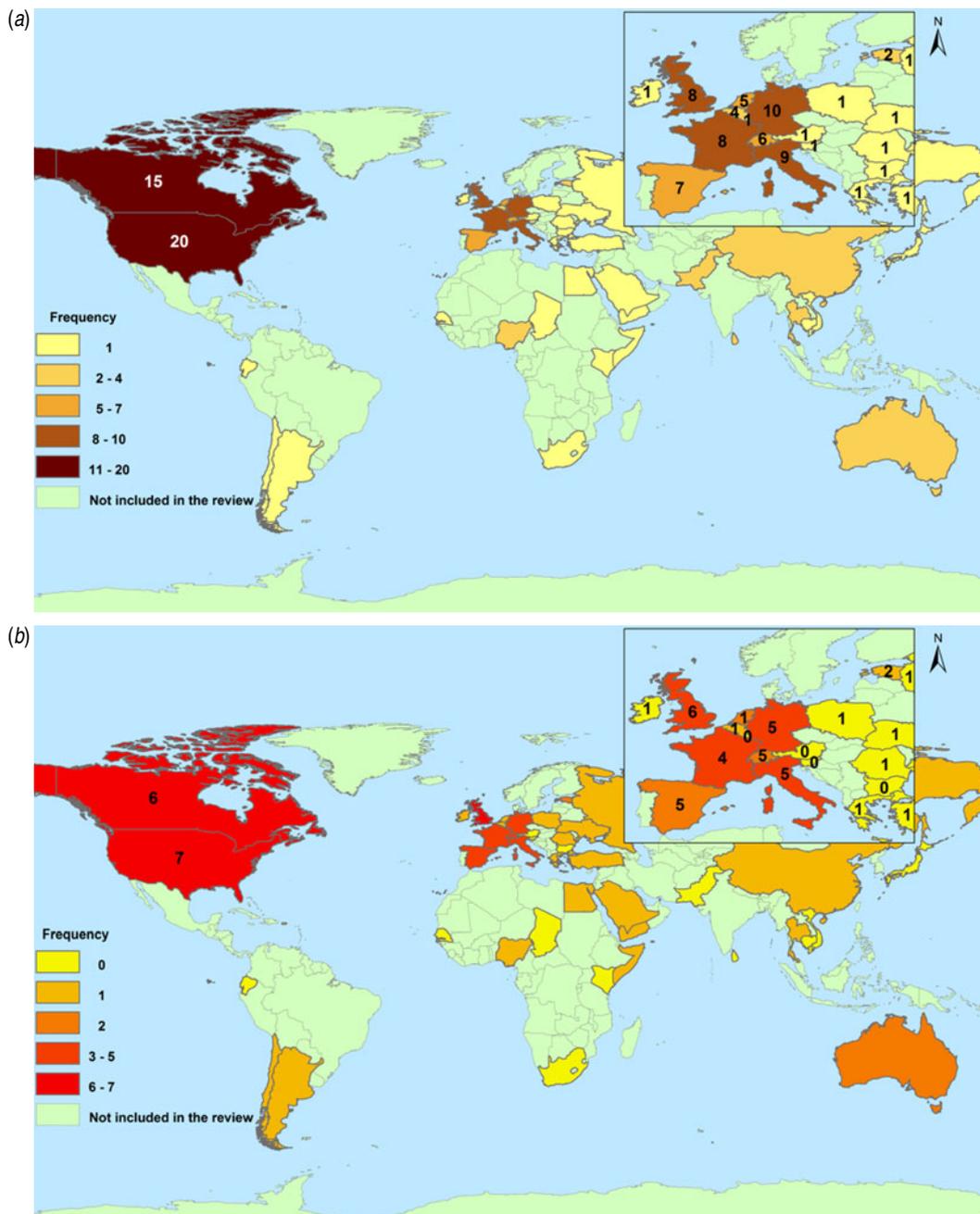


Fig. 2. Frequency of the study countries in the review. (a) Considering all the papers, and (b) considering only the risk-based papers.

($n = 18$) [20–37] and sentinel surveillance ($n = 15$) [20–22, 24, 25, 33, 35, 38–45].

Vector surveillance. This approach focused on three main groups of vectors of interest: *Culicoides* spp. ($n = 8$), mosquitoes ($n = 8$, especially *Culex* spp.) and ticks ($n = 2$). *Culicoides* midges are relatively frequently represented due to widespread surveillance efforts focusing on bluetongue.

Mosquito monitoring has been a surveillance component for several diseases, such as Rift Valley fever, West Nile fever, Usutu virus, or St Louis encephalitis, all of which have been important new ($n = 2$), exotic ($n = 9$) or re-emerging ($n = 4$) diseases in recent years. The main benefit of these approaches was their characterization of the distribution of vectors, not only for the target disease, but also for other similar diseases. Hence

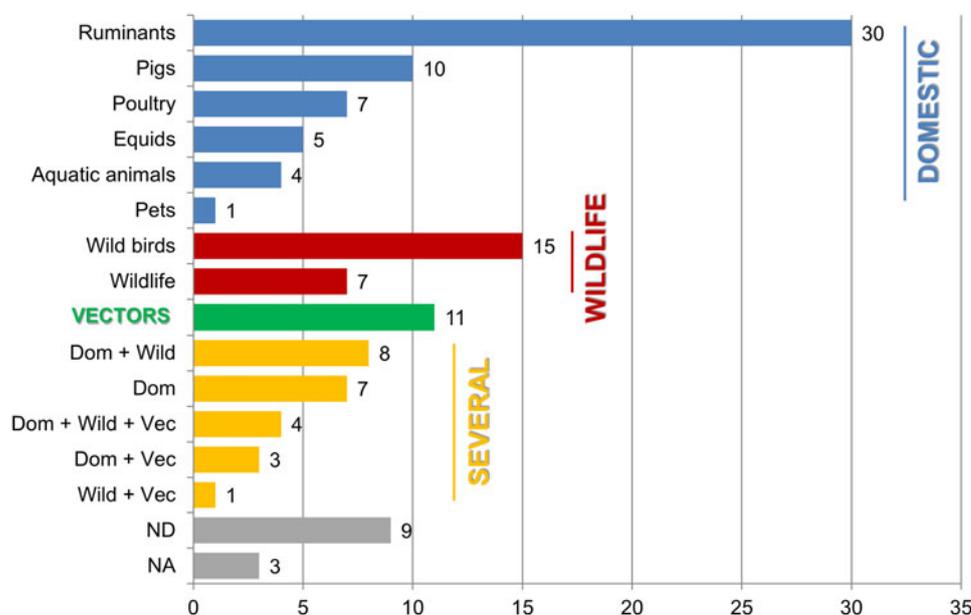


Fig. 3. Distribution of the number of papers according to target host species. The main host species include domestic animals (Dom, in blue), wildlife (Wild, in red), vectors (Vec, in green), and a combination of several of those (in yellow). The papers in which the target host was not specifically defined (ND), or those in which the species was not applicable (NA), are also indicated (in grey).

their potential spread can be predicted. Another issue, when deciding to place collection traps in areas at high risk of disease occurrence, was that vector density may have been overestimated. Therefore, most studies recommended increasing the sampling area. This strategy, when combined with sentinel surveillance ($n = 7$), seemed sufficiently effective for early warning and for setting up control measures.

Sentinel surveillance. The incorporation of sentinels into surveillance systems raised the probability of detection of the first incursion of a particular disease to the earliest possible time since they were in place to provide early information on the species ($n = 5$), time periods ($n = 3$) and/or areas ($n = 8$) at higher risk. Therefore, it could be a more effective system than conventional active surveillance activities. However, difficulties can arise to efficiently implement sentinel surveillance, even when objectives have been appropriately designed.

Pathogen and serological determinations. These categories were cited in many studies in an attempt to detect antigens or nucleic acids of new, exotic or re-emerging pathogens ($n = 13$) [21, 46–57] or antibodies against them ($n = 13$) [20, 22, 33, 35, 46, 51, 53, 57–62]. Different applications of these studies were found: (i) as a permanent component of surveillance systems, as occasional complementary activity ($n = 10$);

(ii) as a data source for the parameterization of statistical models ($n = 4$); or (iii) as a preliminary screening for new, exotic or re-emerging pathogens (namely West Nile, avian influenza, emerging salmonellosis, pigeon paramyxovirus, simian African viruses or eosinophilic meningitis) in order to obtain a preliminary overview of the sanitary situation before applying specific measures.

Participatory surveillance. A novel technique within the sub-category active surveillance (i.e. earliest publication included in the review in 2005) was the use of participatory surveillance ($n = 7$) [63–69]. The application of this method can result in enhanced collaboration and communication among different sectors and institutions to better understand the causes of success or failure (i.e. strengths or weaknesses) of current surveillance programmes to help policy reforms, or to obtain a quick overview of the epidemiological situation in a given area. Most were conducted in developing countries ($n = 4$). The main limitation was the non-response rate, mainly due to reluctance to answer or lack of trust in such approaches. This lack of participation might impede further analyses being undertaken in most situations. In addition, this approach relied on the reporting by observers (e.g. animal owners, veterinarians) to be efficient when, in fact, outbreak reporting may be

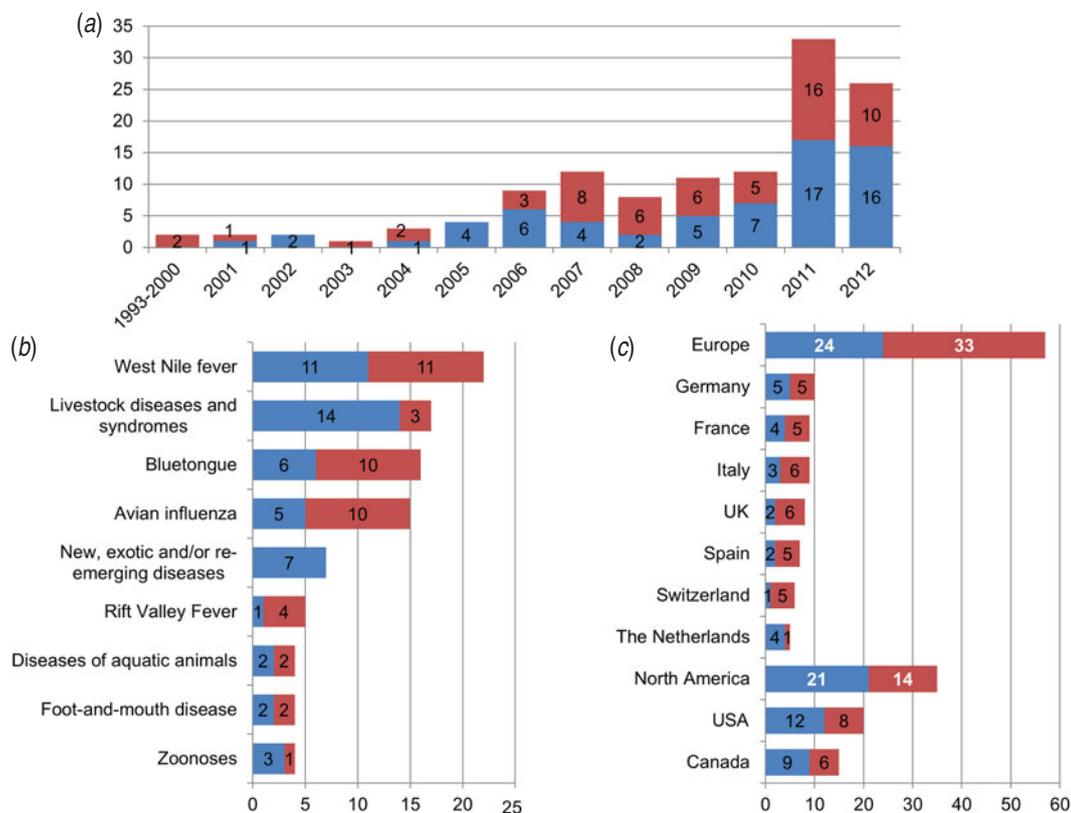


Fig. 4. Distribution of the papers according to a risk-based (in red) or non-risk-based (in blue) approach. (a) Considering the year of publication. (b) Considering the most relevant threats. (c) Considering the most representative countries of Europe and North America.

often disincentivized because of the potential adverse economic impact on livestock holdings or on the sector. Other issues could be the poor representativeness of interviewees or inadequate questionnaire design, which could require a second round to be solved.

Other active methods. Other active approaches included in the review that could not be included in any major sub-category were the monitoring of animal shipments and revenue collection [70], the identification of bird species that could be involved in West Nile virus transmission [71], and periodic active clinical inspections in farms [72].

Passive surveillance approaches

As passive surveillance modalities, we found clinical investigations ($n = 13$) [20, 22, 32, 33, 47, 49, 52, 55, 59, 73–76], syndromic surveillance ($n = 10$) [74, 76–84], mortality investigations ($n = 5$) [20, 33, 35, 49, 55], parameter monitoring ($n = 2$) [85, 86] and other passive activities ($n = 3$) [72, 75, 87].

Syndromic surveillance. This review revealed that syndromic surveillance was a novel method (i.e. first articles in 2006) characterized by its general surveillance scope ($n = 10$). Although pure syndromic methods were mainly described in the articles, they were also linked to digital surveillance frameworks ($n = 6$). In general, these articles proposed a method for the early detection of changes in population health (e.g. clinical cases, abnormal mortality rates, post-mortem findings) which were defined as syndromes. In fact, shortening the detection time was highlighted in the majority of the studies as being the main advantage (i.e. even several weeks prior to the laboratory testing). Other benefits were their potential cost-effectiveness, the integration and optimization of data from different sources, or the simplicity of having an easy system covering multiple threats. However, in the reviewed articles these methods always appeared as retrospective approaches since they were designed based on historical data. The retrospective nature of these approaches suggests that future validation would be necessary to determine their full applicability and benefit.

Moreover, criteria harmonization was an important requirement when defining syndromes since data can be obtained from several sources. It was also indicated that the continuous update of syndrome definition would be necessary in order to include new diseases with distinct pathological profiles. These approaches should also include an effective mechanism to prevent ‘false alarms’ (i.e. increase in the number of syndromes identified, which were later not found to be associated with any true hazard). Several authors highlighted that it was not possible for this method to replace conventional surveillance methods, but that it was useful for early warning, supporting planning and policy development.

Other passive methods. Other passive surveillance approaches (i.e. clinical and mortality investigations) identified in the current review consisted mainly of conventional methods used as a source of data input for epidemiological tools and methodologies ($n = 4$) or as a component of a surveillance system for exotic and re-emerging diseases ($n = 8$). Some other minor passive strategies were parameter monitoring (i.e. movement activity, milk yield, feeding, water intake [85], and temperature [86]), conduction of awareness-raising meetings [72], a post-farm-gate passive surveillance scheme for pigs at sale yards and abattoirs [75], and a scanning surveillance system based on submissions where a diagnosis could not be reached (DNR) [87].

Specific approaches found in the papers: tools and methodologies that support surveillance

Each tool and methodology found in the review was assigned to a specific category in accordance with the methodologies applied and described in the paper (Table 4). In Table 4, the stage of surveillance that each tool found in the review supports was also indicated. In this section we described the main applications, benefits, and limitations per tool as directly extracted and compiled from the papers.

Spatial epidemiology and GIS-based approaches

This was the most represented category in the methodological papers ($n = 27$) [24, 26, 27, 29, 30, 48, 49, 53, 58, 77, 88–104]. The main spatial approach found was risk mapping ($n = 12$), which consists in obtaining risk maps for the probability of environmental suitability for a specified vector or disease by spatial overlaying

the relevant risk factors using statistical or other algorithms. This methodology was applied particularly to multi-factorial and/or vector-borne diseases because they allowed the inclusion of multiple factors and resulted in easily interpretable outputs, with direct applications to the cost-effective allocation of surveillance and control resources. However, major constraints included lack of adequate quality data, the risk of missing important information in the spatial model, both of which result in potentially biased model outputs. Thus almost all the studies included a validation step in their approaches ($n = 9$ of the 12).

GIS approaches were also frequently forming part of digital surveillance frameworks ($n = 10$) in order to support the visualization of the surveillance results or to implement certain spatial transformations of the data. In addition, other applications of spatial methodologies to early detection were: (i) supporting active surveillance activities, such as designing sampling sites ($n = 2$) or processing and presenting surveillance results ($n = 8$); and (ii) supplementing epidemiological tools and methodologies ($n = 10$), such as regression models, cluster analysis, risk assessments or simulation modelling, mainly as a tool to represent and interpret their results.

Basic statistics

This category [24, 38, 42, 45, 50, 51, 61, 67, 74, 80, 83, 86, 96, 98, 103, 105–109] included different statistical tests (e.g. Mann–Whitney, χ^2 , Student’s t , Fisher’s exact probability), formulas (e.g. sensitivity, specificity, positive predictive value, incidence rate), or equations (e.g. sample size calculator, cumulative sum control chart, modified two-stage method). These were applied for the analysis of the results from surveillance activities ($n = 11$) or the output of epidemiological tools and methodologies ($n = 5$). These statistics also offered the possibility to enhance the improvement of surveillance activities, by calculating sample sizes according to the required situation ($n = 3$), or evaluating surveillance options by comparing their attributes or their sampling schemes ($n = 6$).

Regression modelling

This category considered mathematical models that were used to explore epidemiological and disease data related to surveillance and/or to evaluate distinct health measures ($n = 20$) [27, 32, 49, 50, 53, 61, 72, 79, 96, 101, 103, 109–117]. We found a wide range of models, such as linear, logistic, binomial, Poisson, time series, or CART. Some of them presented an original use of

these methodologies in the early detection context. For instance, O'Sullivan *et al.* [117] used regression models to illustrate the potential identification of a novel swine disease (i.e. porcine circovirus-associated disease), and employed test requests for porcine reproductive and respiratory syndrome.

In general, this was the category that offered the widest range of applications for surveillance. Most of the regression methodologies described in the papers were applied to data analysis ($n = 15$), either to assess the role of a set of variables to disease epidemiology or to analyse trends and patterns of disease occurrence. Moreover, these methods were applied to explore risk factors or to model disease occurrence ($n = 5$). Another potential consisted in its application for syndromic surveillance ($n = 7$), used for examining the association of certain variables with the defined syndromes or for obtaining the alarms. As for evaluation of surveillance, regression models were used for comparing the performance of newly developed early detection strategies with conventional surveillance activities ($n = 4$). Typically, the utility of the outputs generated by regression models was affected by restrictive or inaccurate model assumptions and/or data acquisition and management problems. However, the review showed the potential utility of these methods for the development of enhanced, risk-based surveillance systems in numerous articles dating back to past decades.

Digital surveillance

Although there is no precise definition of the term, it broadly includes the use of the internet and computer technologies for collecting and processing health information, including outbreak reports and surveillance data [14]. The articles included in this category ($n = 18$) [72, 74, 77, 80, 82–84, 88–90, 92–94, 96, 97, 99, 118, 119] comprised several tools for data collection, management and processing, which normally involved several diseases ($n = 12$). Digital surveillance approaches were normally linked with governments, and with health and research institutions, and were beneficial for inter-institutional relationships and decision-making. Another advantage was their real-time nature, which allowed the constant update of the situation and earlier detection, and intervention should animal health threats occur. As additional benefits, these platforms were considered useful, simple, flexible and timely. Their main limitations were that they were still under development (i.e. pilot phases),

which implies the need for future adjustments and calibrations, and a standardized way to submit safely and easily information and to provide good quality data was lacking.

Simulation modelling

Simulation models ($n = 10$) [30, 41, 97, 107, 119–124] aimed to mimic disease transmission in a population in time, and sometimes in space, terms in specific circumstances. The outputs from the simulations could be used for estimating the risk of disease spread ($n = 5$). In addition, simulation models allowed to evaluate surveillance activities ($n = 5$), by obtaining surveillance performance attributes, as well as comparing the effectiveness of different surveillance strategies for early detection. In case the epidemics were established, they also proved useful to evaluate the effectiveness of different mitigation measures. Moreover, the outputs from the model could be used as inputs for other epidemiological tools ($n = 4$). A novel inclusion in these models was the estimation of the cost or the benefit of surveillance and management activities. For this kind of tools, sensitivity analyses were mostly performed in order to evaluate the approaches and try to overcome the potential degree of subjectivity when assigning model parameter values.

Decision modelling

This category mostly included scenario tree models ($n = 6$) [59, 75, 120, 125–127], although it also comprised other decision methods such as neural networks [81, 128] or the info-gap decision theory [124]. The scenario tree and the info-gap models reviewed were used for the evaluation of surveillance systems ($n = 7$), and also to demonstrate how the likelihood of early disease detection can be improved by combining the most appropriate passive and/or active strategies in surveillance programmes for exotic and re-emerging threats. Data quality and model assumptions greatly influenced the usefulness of outputs, therefore sensitivity analyses and validation procedures were often performed ($n = 6$). Other applications of decision models consisted in their use in syndrome classification in syndromic systems ($n = 1$), or the definition of risk categories and factors ($n = 3$).

Cluster analysis

Cluster analysis [32, 77, 87, 88, 98, 101, 103, 129] was used for early warning in order to detect clustering of epidemiological events (e.g. disease cases, deaths,

outbreak signals) or estimates (e.g. relative risk) and then lead to a risk characterization ($n = 6$). In terms of data analysis ($n = 4$), this tool proved to be useful for validating the main GIS-based tool used, or if combined with syndromic and scanning surveillance data ($n = 3$). It could even serve for enhancing prospective real-time surveillance strategies in highlighted areas ($n = 1$). However, in general, detected clusters might not represent real outbreaks, so further epidemiological research is required to determine the cause of any spatial, temporal or space-time clustering. For this reason, all the papers validated their models using field data.

Environmental modelling

This category could be considered as a special type of statistical model. These models aimed to predict the climatic and environmental factors that affect diseases directly (e.g. effect on the incubation period or resistance of the pathogen in the environment, $n = 2$) [104, 130], indirectly (e.g. effect on the vector population dynamics or movements; $n = 5$) [26, 44, 48, 91, 131] or both ($n = 1$) [93]. The environmental approaches reviewed here included normalized difference vegetation index (NDVI) anomalies ($n = 3$), air or water temperature variations ($n = 3$), degree-day models ($n = 2$) and a wind-borne spread model. Output from all these tools could be considered to determine areas and/or periods at higher risk. However, it should be noted that the modelled variables constitute single components in the multi-factorial system of disease epidemiology, so results should be applied with caution.

Expert opinion

Expert opinion methods ($n = 5$) [66, 68, 125, 132, 133] were conducted in order to evaluate surveillance systems in place ($n = 4$), to identify risk factors and population strata at higher risk ($n = 1$), or to obtain risk estimates for disease entry and exposure ($n = 1$). Considering that this is a purely subjective method, caution is needed when using the outputs derived from these approaches. However, they could serve as a starting point for future studies.

Risk assessment

Risk assessment was a purely risk-based method ($n = 5$) [98, 100, 134–136], which generated risk estimates to be used to design and assess risk-based surveillance systems, although they focused on the release and/or

exposure of pathogens, and not on surveillance itself. Risk assessments can also provide information about the evaluation of variations in risk, based on a standardized model framework that can be adjusted and updated. This helps improve the effectiveness of surveillance if the disease epidemiology, and therefore, the risk changes over time. Similarly to other methods, the quantity and quality of the data used to parameterize models determined the usefulness of the outputs.

Other minor epidemiological methods

Four literature reviews were included since they provided good knowledge and examples of the use of methodologies for early detection. Specifically, two studies identified patterns of global disease occurrence [137, 138], while the other two reviewed the use of animals as sentinels for bioterrorism [5] or for human disease [139].

Some other methodologies identified in the current review, but only used rarely, were: data mining techniques, applied to syndromic frameworks [81, 82, 140] or investigate disease risk classification as a proxy for compromised biosecurity in cattle [141]; network analysis, which contributed to the assessment of potential spread patterns through animal movements [97, 122]; a framework to integrate veterinary health reporting with public health systems [142]; or studies to select the best sample for early detecting classical swine fever [143]. Although these papers can provide relevant information for use in surveillance design or for modelling purposes, they proved too heterogeneous to allow a proper analysis of the approach as a whole in the remit of this systematic review.

DISCUSSION

The present paper aimed to review approaches and methodologies that have been developed, described and implemented over the last 20 years for the early detection of new, exotic, and re-emerging diseases. Unlike previous reviews, which have usually focused on general or specific aspects of one surveillance type (i.e. syndromic surveillance or participatory surveillance), this systematic review of the scientific literature is the first that aimed to cover as many approaches and methodologies targeting early detection as possible. Several methods were highlighted in the review as the most commonly cited in the papers. It is worth to mention that those are not necessarily

the ones most commonly used for early detection. Special attention was paid to the review of methodologies in order to document the benefits and current limitations of each method.

Traditionally, passive components have formed the core of the surveillance approaches used for the detection of new, exotic and re-emerging threats [16]. These 'traditional' surveillance activities are likely to remain as an essential component of early warning surveillance in the future, but may need to be supplemented with new approaches. The evaluation of conventional surveillance systems in terms of their accuracy (sensitivity or specificity), precision (repeatability), timeliness, multiple utility, value and cost-effectiveness [17] has highlighted the need for improvement, especially after the 2001 FMD outbreak in the UK. In recent years, many new surveillance approaches have been developed.

One of these developments is the use of risk-based surveillance methods, which are likely to be more efficient than conventional methods [18]. When applied together with traditional design approaches, these risk-based methods allow the incorporation of prior information about disease occurrence to surveillance design and, thus, assure an appropriate and cost-effective data collection [18]. The 'risk-based' concept is not new as such surveillance systems have already been implemented as 'targeted surveillance' since the 2000s. Some examples could be the surveillance of bovine spongiform encephalopathy [18] or the surveillance of vector-borne diseases such as bluetongue, including sentinel surveillance in areas at high risk [38, 144]. However, the incorporation of risk-based approaches into existing surveillance programmes is not very frequent and their use over countries has not been properly adapted. As an illustration, this review has identified 60 papers that deal with newly proposed risk-based methods. Conversely, we found that only 10 papers actually reported risk-based methods that have already been implemented into countries' surveillance systems.

The current review shows that risk-based methods have been particularly applied for vector-borne diseases. Risk mapping, environmental models and vector spread simulations are risk-based approaches that have been used to identify the areas and time periods in which surveillance is more likely to detect these threats early and successfully. Other diseases in which risk-based methods have been applied are avian influenza and FMD. For avian influenza, the risk-based approaches analysed focused mainly on early

detection of disease entry by wild birds, and took into consideration environmental factors such as presence of wetland areas. Regarding FMD, the articles analysed mainly described risk-based models for identifying optimal procedures for the early detection of this disease if there was an outbreak and the virus spread.

In summary, risk-based methods have been developed mainly for diseases for which (i) biotic factors exist (such as vectors or wild birds) whose presence depends directly on abiotic environmental factors, and which are essential for the presence and spread of the disease; or (ii) different scenarios pose varied levels of risk for the introduction and spread of the disease. In addition, the application of risk-based approaches was related to the nature of the surveillance. While most of the studies that focus on a specific disease are risk-based (68.75%), it seems that those approaches dealing with multiple diseases (i.e. general or multi-objective) are mainly non risk-based (73.77%). Thus, the potential development of risk-based methods for simultaneous application on multiple diseases is an interesting field for further research. However, identification of suitable risk factors to predict the occurrence of multiple diseases is a considerable challenge, but may explain why risk-based approaches for effective multi-objective surveillance systems have not yet been widely implemented.

The review has shown that a significant amount of work has been done in relation to the development of risk-based approaches for identifying populations at high risk for certain infections, especially by risk mapping. Although risk maps are relatively easy to interpret and provide extremely useful information to allocate preventive measures, there is still work to be done to integrate the results of these analyses into the design of risk-based sampling strategies for early detection.

The same limitation applies to the other approaches identified in this review. Methods such as regression or simulation models, cluster analyses, and risk assessments can be used to inform risk-based surveillance and to provide useful information regarding disease patterns, relevant risk factors and estimates, or to the efficiency of surveillance activities. However, their use in operational surveillance systems is not common [145]. This is mainly due to the complexity of the underlying epidemiological concepts, algorithms, and/or software used, and to decision makers not confiding in the outputs obtained and/or to their inability or unwillingness to work constructively

with the uncertainty estimates associated with these model outputs. Therefore, outputs generated by these quantitative methods need to be communicated more effectively to decision-makers and better evidence for the value of risk-based surveillance strategies is required.

Other methodological challenges found in the review include limited sample size, failure to account for clustering in space or time or for imperfect test characteristics, lack of risk estimates, bias in the selection of risk factors, low public awareness of the emerging threat, inclusion of out-of-date surveillance results, multiple data sources and poor quality of data. These limitations can be overcome by appropriately selecting the method to use according to the threat of interest. In addition, those studies including a validation of the particular methods investigated which have used real disease data have demonstrated the benefits of their inclusion in surveillance programmes. The adoption of risk-based methods and other new approaches can be facilitated by providing evidence for their cost-effectiveness. This is essential since it is typically being argued that risk-based designs involve additional costs given their increased complexity if compared to random sampling approaches.

Another recent development in surveillance methods with a clear application to early detection systems is the use of syndromic data. It has been applied for the early detection of seasonal increases in incidence of known human hazards, such as influenza or heat-related mortality (e.g. [146–148]), or for assessing the impact of environmental disasters on the health of populations (e.g. [149, 150]). Although some debate exists, syndromic surveillance could be considered as an ‘enhanced passive’ method, since the provision of data is observed-initiated (i.e. passive), but there is an active involvement of the investigators. The potential role of syndromic surveillance as a part of early detection in animal health requires further research. Syndromic data collection can be useful, but the role it plays in the detection of different health events needs to be further clarified or demonstrated, and systems that are currently being developed need further validation as more data become available. Syndromic data analysis following the alert for a new, exotic or re-emerging disease, or in the face of an increased risk of incursion of a currently absent disease, may also prove useful to speed up the investigation of potential outbreaks in order to limit disease spread. The combination of these methods with rapid systems for data collection or good inter-

institutional communication may be very advantageous. For instance, the incorporation of syndromic approaches into digital surveillance systems, which are still in their very early development stages, could result in a real-time health monitoring system. Nevertheless, it is important to recognize that these systems will still rely on the availability of effective laboratory testing to diagnose the underlying infectious causes of any emerging trends and alarms [151].

Developments have also been made in the active surveillance activities applied for early detection. Although these activities are practically and economically not always the most efficient, the review showed their great potential for early detection and interest in their improvement. Sentinel surveillance, as shown in this review, can be a very useful surveillance system component when dealing with new, exotic or re-emerging diseases as it increases the probability and timeliness of detecting recently infected or newly diseased animals. It is also useful for vector monitoring, an important feature if we consider that most of the emerging diseases in recent years were vector-borne [152, 153]. Thus, the combination of sentinel and vector surveillance could be particularly relevant for monitoring vector-borne diseases with easily distinguishable clinical symptoms and rapid onset. In fact vector surveillance as a monitoring activity has proven very useful for disease early detection and control in recent years. For instance, it is believed that the outbreaks of bluetongue in Europe could have been detected and controlled faster if more information about the vector distribution and competence had been available [154, 155]. As a result of the control efforts for this infection in Europe, *Culicoides* spp. distribution is now well-documented, and this information will also benefit the design of surveillance programmes for other *Culicoides*-borne diseases, such as the recent Schmallenberg disease [156].

Most of the methods developed for surveillance have been traditionally linked to domestic livestock. In recent years, early detection applied to surveillance in wild populations has become an emerging research area [157]. In this review, 34 articles dealing with surveillance methods for wildlife infectious diseases were identified (especially from 2001). However, the meeting organised in 2009 by the European Wildlife Disease Association (EWDA) highlighted that only 14 of the 25 represented European countries performed some kind of surveillance activity in wild animal populations [158], probably due to certain limitations in the implementation of surveillance

activities in wild species [154]. Most of the surveillance strategies that focused on wildlife compiled from this review investigated morbidity and mortality events, identified new pathogens, and monitored the status of known diseases within wild animal populations. These activities may provide insight into infectious disease pathologies and agents, as well as new host ranges [157]. All the conclusions reached from these activities should be quickly shared worldwide in order to prevent the transboundary spread of new, exotic and re-emerging pathogens. Fortunately, the 2011 EWDA meeting set the basis to create an international network for wildlife health surveillance and to foster cooperation between public and animal health professionals [158]. Since the interactions of the interface connecting humans, livestock and wildlife have intensified in the last 20–30 years (mainly as a result of anthropogenic factors), they have become an important driver of disease emergence [159]. Therefore, further studies need to be conducted to establish data collection and diagnostic methods for new, exotic and re-emerging pathogens adapted to this interface.

The expansion of the European Union and the increase in population over the past 20 years have led to important socioeconomic and health consequences for Member States (MS). This development includes the promotion of the ‘open trade’ concept, which means removing regulatory barriers for the purpose of encouraging trade between countries and enhancing competitiveness both within and outside the European Union. Consequently, the risk of introducing new, exotic or re-emerging diseases into MS increases with the number of movements between countries, and the trade of live animals is one of the main introduction routes for infectious diseases. For this reason, MS agreed to follow strict legal and health requirements with a view to minimizing high-risk contacts. The implementation of surveillance programmes for infectious diseases also needs to provide protection against the entrance of new, exotic, and re-emerging diseases through trade and via other entry routes (e.g. contacts with wildlife). However, some of the lessons learnt from recent health crises include the need to further improve the detection of new pathogens, such as SBV [160], and to minimize the time until implementation of control measures for exotic or re-emerging diseases. This depends on local characteristics that can result in delayed detection, such as the outbreaks of classical swine fever in The

Netherlands and Belgium in 1996–1997 [161, 162] or the incursion of FMD in the UK in 2001 [11].

This systematic review of the literature provides a thorough summary of methodologies that can be considered in the design of surveillance programmes aimed at such that their ability to detect new, exotic, and re-emerging diseases early is improved. The number of research publications related to surveillance topics and methods is continuing to increase. In addition, the concepts and terms related to surveillance have been continuously evolving over time. However, we have captured the main tools that could be used for surveillance purposes but we are conscious that new methods may arise in addition to the ones included in this review. Those included could also be combined resulting in different ways of applying such tools. Our objective was to summarize a set of epidemiological tools that have offered useful results for the early detection of diseases without aiming to be prescriptive or limiting the potential expansion of research towards new methodologies. A group of publications that are not included in this systematic review is related to those articles dealing with outbreak preparedness. One could argue that some of the epidemiological tools mentioned in this review have been used for preparedness of exotic diseases. Some examples are the spatio-temporal cluster analysis and the observed-to-expected ratios for H5N1 avian influenza outbreaks in the absence of population data [163], ‘Be-FAST’, a novel simulation model for infectious disease spread both within and between farms, which also considers the specific farm-to-farm contact network [164, 165], and/or the combination of a cluster analysis with a social network analysis to identify farms at high risk for the introduction and spread of infectious diseases [166, 167]. Articles dealing with preparedness would focus on obtaining epidemiological information or on testing hypotheses about disease dynamics and target population structures prior to, during, or after an outbreak of a new, exotic or re-emerging threat rather than identifying populations, locations or timeframes at a higher risk of infection. However, ‘preparedness’ can be interlinked and complementary with surveillance concepts to minimize their consequences in a timely, efficient and cost-effective way. Finally, unpublished literature was not included in order to minimize the potential bias of over-representing those available as the observers’ information source. Therefore, we have covered a detailed list of methods in early detection that allow a reliable overview and analysis.

The complexity and dynamism of the underlying eco-social system within which these pathogens emerge has emphasized the need for interdisciplinary collaboration to improve early warning surveillance systems, thus comprising epidemiologists, producers, field veterinarians, pathologists, animal health consultants, diagnostic laboratories, ecologists, wildlife experts, economists, social scientists, and national and international institutions [168]. The establishment of these 'One Health/Ecohealth' networks will become a key requirement for the effective protection of public health, trade, and animal health and welfare [169]. The approaches identified in this review that follow these holistic perspectives comprise risk-based methods, monitoring health events combined with specific disease outbreaks, enhancing of participatory surveillance and the use and integration of digital information to allow the rapid, cost-effective identification of threats to also improve timeliness. The next steps to improve surveillance systems in the European Union are: (i) the development of evaluation frameworks to study the applicability of these novel methodologies integrated with, or replacing, currently used methods, while maintaining or improving the sensitivity and specificity of current standards of surveillance systems; and (ii) the definition of parameters that should be considered and recorded in each country and which level of detail this information should have (i.e. minimum requirements) to implement novel approaches and risk-based methods. Thus, the present literature review provides key information about the advantages, disadvantages, limitations and potential application of methodologies for the early detection of new, exotic and re-emerging diseases.

SUPPLEMENTARY MATERIAL

For supplementary material accompanying this paper visit <http://dx.doi.org/10.1017/S095026881400212X>.

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DECLARATION OF INTEREST

None.

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