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BioCat 2.0

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September 2013



Pacific Northwest
NATIONAL LABORATORY

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Pacific Northwest National Laboratory
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Abstract

The U.S. Department of Homeland Security (DHS) National Biosurveillance Integration Center (NBIC) was established in 2008 with a primary mission to “(1) enhance the capability of the Federal Government to (A) rapidly identify, characterize, localize, and track a biological event of national concern by integrating and analyzing data relating to human health, animal, plant, food, and environmental monitoring systems (both national and international); and (B) disseminate alerts and other information to Member Agencies and, in coordination with (and where possible through) Member Agencies, to agencies of State, local, and tribal governments, as appropriate, to enhance the ability of such agencies to respond to a biological event of national concern; and (2) oversee development and operation of the National Biosurveillance Integration System (NBIS).” Inherent in its mission then and the broader NBIS, NBIC is concerned with the identification, understanding, and use of a variety of biosurveillance models and systems.

The goal of this project is to characterize, evaluate, classify, and catalog existing disease forecast and prediction models that could provide operational decision support for recognizing a biological event having a potentially significant impact. Additionally, gaps should be identified and recommendations made on using disease models in an operational environment to support real-time decision making.

Acronyms and Abbreviations

DHS	U.S. Department of Homeland Security
HHS	U.S. Department of Health and Human Services
IT	information technology
LANL	Los Alamos National Laboratory
NBIC	National Biosurveillance Integration Center
NBIS	National Biosurveillance Integration System
SME	subject matter expert
SMW	Semantic MediaWiki
TRL	Technology Readiness Level
USDA	U.S. Department of Agriculture

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1.0 Introduction

The National Biosurveillance Integration Center (NBIC) mission is to enhance the capability of the Federal Government to rapidly *identify, characterize, localize, and track a biological event of national concern* by integrating and analyzing data relating to human health, animal, plant, food, and environmental monitoring systems (both national and international); *and disseminate alerts* and other information to Member Agencies and to agencies of State, local, and tribal governments, to enhance the ability of such agencies to respond to a biological event of national concern.. PNNL has supported the operational biosurveillance modeling needs of NBIC in a project funded by the U.S. Department of Homeland Security (DHS) Science and Technology Directorate (S&T) to identify, characterize, and catalog models that could be used in an operational environment.

A rich and diverse field of infectious disease modeling has emerged in the past 60 years and has advanced our understanding of population- and individual-level disease transmission dynamics, including risk factors, virulence, and spatio-temporal patterns of disease spread. These modeling techniques span domains such as biostatistical, massive agent-based, biophysical, ordinary differential equation, and ecological-niche models. Diverse data sources are being integrated into these models as well, including demographics, remotely sensed measurements and imaging, environmental measurements, and surrogate data such as news alerts and social media. Moreover, nascent research is occurring at the omics-level to aid in forecasting future epidemics; such research includes phylogenetic techniques for predicting pathogen mutations, algorithms for microbial identification in next-generation technologies, meta-genomics, and multi-scale systems biology. Yet emerging infectious diseases continue to impact health and economic security across the globe. There remains a gap in the sensitivity and specificity of these modeling forecasts designed not only to track infectious disease events but also to predict disease occurrence. For an example, one needs to look no further than the 2009 H1N1 influenza pandemic. The latency between identification and characterization of the virus's pathogenicity and transmissibility caused, perhaps, unnecessary mitigation measures such as school and business closures. Moreover, there are strong indicators that dynamics and emergence of vector-borne diseases are in flux because of, among other factors, changes in land use, human behavior, and climate.

The goal of this systematic review was to identify areas for research to characterize the viability of biosurveillance models to provide operationally relevant information to decision makers about disease events. We define a disease event to be a biological event within the context of One Health and all-hazard paradigms. These events are characterized by evidence of condition and risk. These are neither mutually exclusive nor limited to the following examples for evidence of condition: person-to-person transmission (e.g., *Mycobacterium tuberculosis*), zoonoses (e.g., *Francisella tularensis*), foodborne pathogens (e.g., *Salmonella*), vector-borne pathogens (e.g., *equine encephalitis virus*), waterborne pathogens (e.g., *Vibrio cholerae*), airborne pathogens (e.g., influenza), animal-to-animal transmission (e.g., *Aphthae epizooticae*), and plant pathogens (e.g., soybean and wheat rusts). Examples for evidence of risk are accidental or deliberate events affecting air or water quality (e.g., volcanic ash, pesticide runoff), economically motivated adulteration of the food and pharmaceutical supply, and intentional release. In the context of this article, a **biosurveillance model** is broadly defined as an abstract computational, algorithmic, statistical, or mathematical representation that produces informative output related to all-hazards event detection or event risk. The model is formulated with a priori knowledge and may ingest, process, and analyze data. A biosurveillance model may be proactive or anticipatory (e.g., used to detect or forecast an event, respectively), it may assess risk, or it may be descriptive (e.g., used to understand the dynamics or

drivers of an event). The evaluation of biosurveillance models and data sources provides key information to enable the identification of gaps and recommendations for improvements to existing models to optimize the utility of the models in an operational environment. This scientifically driven process ensures that the capabilities and limitations of biosurveillance models and data streams are understood by potential users and stakeholders, and result in a path forward toward enhancing biosurveillance capabilities.

For reference, the statement of work for PNNL's Fiscal Year 2013 (FY13) effort is provided below. The report aligns with the logical flow of our work rather than task by task.

Task 1: Continue identification and characterization of operational biosurveillance models

PNNL will identify, evaluate, and catalog existing syndromic surveillance, indicator systems, and event-detection biosurveillance models published during **FY12–13**. The focus of this task will be to use the attributes and approach developed in the previous work with NBIC to identify, evaluate, and catalog existing event-detection biosurveillance models.

Subtask 1.1: Identify syndromic surveillance, indicator systems, and operational event-detection biosurveillance models. We will first identify event-detection biosurveillance models operationally deployed by government, industry, academia, and foreign and global organizations.

Subtask 1.2: Evaluate models identified in Subtask 2.1 with the model characterization plan developed in the first BioCat effort. We will expand the attributes and develop quantitative metrics that will then be applied to prediction and forecast biosurveillance models.

Subtask 1.3: Catalog models. We will maintain and store the biosurveillance model catalog within a MediaWiki open-source wiki framework that was created under the previous work with NBIC. The catalog is hosted at <http://biocat.pnnl.gov> and is accessible via the web. and access is controlled through user/password authentication.

Subtask 1.4: Collaboration and Sharing. We will travel to meet with the Sponsor, collaborate with analysts to ensure BioCat's operational relevance. Additionally, PNNL may submit and participate in biosurveillance- or biodefense-related conferences to promote the NBIC/NBIS and BioCat for community use.

Task 2: Back-identify, characterize and catalog operational biosurveillance models

PNNL will identify, evaluate, and catalog existing syndromic surveillance, indicator systems, and event-detection biosurveillance models, published during calendar year 2011 and 2012 (CY11–12). The focus of this task will be to use the attributes and approach developed in the previous work with NBIC to identify, evaluate, and catalog existing event-detection biosurveillance models.

Subtask 2.1: Identify syndromic surveillance, indication, and operational event-detection biosurveillance models. We will first identify event-detection biosurveillance models operationally deployed by government, industry, foreign and global organizations during CY11–12.

Subtask 2.2: Evaluate models identified in Subtask 2.1 with the model characterization plan developed in BioCat Phase 1. Leveraging the quantitative analysis and through following the vetted characterization plan, each event-detection biosurveillance model will be evaluated. Additionally, PNNL will evaluate models not yet evaluated discovered during BioCat Phase 1.

Subtask 2.3: Review the discovered models from BioCat phase 1 and identify models with peer-reviewed conference publications and abstracts. Then evaluate those models with the same rubric as in Task 1.

Subtask 2.4: Catalog models. We will maintain and store the biosurveillance model catalog within a MediaWiki open-source wiki framework that was created under the previous work with NBIC. The catalog is hosted at <http://biocat.pnnl.gov> and is accessible via the web, and access is controlled through user/password authentication.

Task 3: Maintain the BioCat for NBIS/NBIC, and Update Software

PNNL will host the BioCat for the DHS Office of Health Affairs NBIS/NBIC. The model catalog continues to provide value to biosurveillance stakeholders across the government and academia. BioCat is located at PNNL on a webserver that was purchased by DHS S&T and is accessible via the Internet. The software framework on which BioCat is built must be updated periodically with security patches and modified configurations.

Subtask 3.1: PNNL will ensure the BioCat remains operational and accessible via the internet.

Subtask 3.2: A variety of projects at PNNL utilize the Knowledge Encapsulation Framework (KEF) software and fund the development of new software and upgrades. Funding under this subtask will ensure that the upgrades developed for other programs are integrated into BioCat.

Subtask 3.3: Coordinate with interagency biosurveillance efforts, and where possible leverage interagency resources and integrate within BioCat. PNNL will also integrate the NBIC's open RSS feed within the BioCat to be viewable by BioCat users.

2.0 Literature Selection and Processing

To provide a comprehensive list of biosurveillance models, the PNNL team took two approaches. In the first approach, PNNL staff contacted subject matter experts (SMEs) in academia and government to discuss existing biosurveillance models. A literature search (the active identification of existing information sources most relevant to the research question or need) was performed in the second approach to comprehensively identify biosurveillance models in the scientific, grey, and limited-distribution literature. One of the main goals of the FY13 funding for BioCat was to identify relevant biosurveillance literature since the end of the last increment of funding in 2011. We identified and pursued several lines of inquiry.

1. Identify people and institutions working in the domain of biosurveillance.
2. Scope agents, pathogens and toxins of interest to NBIC.

3. Conduct targeted literature searching in salient databases from 2011 to present.
4. Vet literature with subject matter experts.
5. Acquire materials for evaluation.
6. Refine list of papers for evaluation based on NBIC prioritization.
7. Import literature citations into Drupal (open source content management platform).

The first task was to update our people and institutions list. This effort involved reviewing the original list from the previous effort and updating contact information and websites of experts and research institutions focused on the topic of biosurveillance. This information is provided in Appendix A.

The second task was to understand the agents, pathogens, and toxins of interest to NBIC. We received nine lists, including several from the Centers for Disease Control and Prevention (CDC) and the U.S. Department of Agriculture (USDA) as provided in Appendix B. Extensive effort was taken to de-duplicate these lists and develop a list of synonyms for each (Appendix C).

Following the de-duplication of the lists of interest, targeted literature searches were conducted. Twelve research databases were queried (Table 2.1). Four additional databases were used in the first round of BioCat literature identification in 2010–2011. By the time targeted searches were performed in 2013, GeoRef, Pollution Abstracts, and Toxicology Abstracts had been cancelled at PNNL due to low use.

Table 2.1. Databases Used to Identify Biosurveillance Modeling Literature

Database	Database Provider
Academic Search Complete	EBSCOHost; Open Source Center
ASFA: Aquatic Sciences and Fisheries Abstracts	ProQuest; PNNL Library
BioOne Abstracts and Indexes	BioOne, PNNL Library
Compendex	Engineering Village, PNNL Library
INSPEC	Engineering Village, PNNL Library
NTIS	Engineering Village, PNNL Library
DTIC (private version)	Defense Technical Information Center
Environment Index	EBSCOHost; Open Source Center
GreenFILE	EBSCOHost; Open Source Center
Google Scholar	Google
Homeland Security Digital Library	Naval Postgraduate School
International Security & Counter Terrorism Reference Center	EBSCOHost; Open Source Center
MEDLINE (through Web of Knowledge portal)	Thomson Reuters, PNNL Library
Military & Government Collection	EBSCOHost; Open Source Center
Oceanic Abstracts	ProQuest; PNNL Library
Scirus	
Scopus	EBSCOHost; Open Source Center
Water Resources Abstracts	ProQuest; PNNL Library
Web of Science (Science edition), Conference Proceedings Citation Index (Science; Social Sciences & Humanities)	Thomson Reuters, PNNL Library

The agents, pathogens and toxins listed in Appendix B were combined with additional keywords (Table 2.2) to refine search results. An example search string: [one disease, toxin, agent, pathogen from Appendix C and all its synonyms (Boolean Operator OR linking the synonyms together)] AND [one or more keywords from Table 2.2]. The dates were limited to 01-01-2011 to the date that the search was executed. Methods of truncation, search limiters, etc. used were dependent upon the database requirements. In some cases it was useful to refine the search results with the NOT Boolean Operator to exclude extraneous citations focused on AIDS, HIV, STD, cancers, etc. The resulting relevant literature was downloaded into EndNote X6 bibliographic management software for portability, ease of use, export capability, and document storage solutions.

Table 2.2. Additional Keywords Used to Refine and Identify Biosurveillance Modeling Literature

Search Terms and Phrases	
Adulteration and food	Early recognition
Adulteration and pharmaceuticals	Epidemic model*
Biolog* or pathogen or toxin or disease model characterization	Event detection and biolog* or pathogen or toxin
Biological risk assessment	Event detection biosurveillance
Biosurveillance	Event-based biosurveillance
"Biosurveillance intelligence"	Foodborne disease* detection or surveillance or outbreak
Bioterror* agents	Forecast + GIS or GPS or remote sensing
Bioterror* and model	Geodemographics
Bioterror* food model	Geoinformatics biosurveillance
CBRN model*	Geosurveillance
Communicable disease detection or model*	Global biosurveillance
Crop disease model*	Global surveillance
Disease and weather forecast	Health intelligence
Disease detection	Health security
Disease epidemic	"Infectious disease" surveillance or model*
Disease forecast	Insect risk + model
Disease forecast* model	Model +predictive, +probabilistic, +projection, +forecast
Disease indicators and/or warnings	Outbreak prediction
Disease management	Outbreak forecast*
Disease map*	Pathogen detection/recognition/surveillance/prediction + model*
Disease modeling	Population dynamic* and outbreak
Disease or bioterror* risk	Predict* or project* or forecast* or "early warn*" or "early detect*" and model*
Disease or outbreak origin	Predictive model
Disease outbreak or emergence or spread or transmission	Predictive model* and disease + infectious + epidemic
Disease patterns	Remote sensing and disease and/or forecast and/or model
Disease regulation	Risk map* and disease model*
Disease risk or forecast or detection	Sentinel surveillance
Disease simulation	Social disruption and disease or pathogen or toxin

Table 2.2. (contd)

Search Terms and Phrases	
Disease source	Social networks, social network analysis and biosurveillance
Disease surveillance and infectious	Spatial analysis disease
Disease transmission	Spatial disease model*
Disease warning	Spatial epidemiology
Drug contamination or surveillance or outbreak	Surveillance or security
Early detection	Syndromic surveillance
	Vector borne (or vector-borne) disease model

After each database in Table 2.1 was searched, a batch export of the citations with abstracts was generated in Microsoft Word[®]. The citations were evaluated by SMEs for potential to contain predictive or forecasting models focused on biosurveillance. Relevant abstracts were flagged with a color: green for Yes, yellow for Maybe. Abstracts that lacked a clear relevance to the research effort were not flagged. Each citation batch was vetted twice, with the first pass conducted by a postdoctoral scientist and the second pass was conducted by a research staff scientist with expertise in epidemiology or microbiology. If the SME disagreed with the initial assessment, the SME judgment was accepted.

Once the batch assessments were complete, the results were reviewed by the data analyst. The data analyst created Group folders (Yes, Maybe, No) in each database's respective EndNote library and moved each citation to the appropriate folder based on the assessment completed in the previous step. Literature in the Yes and Maybe categories were acquired in PDF format. Once these materials were received, the PDF files were stored with the citation in the EndNote database.

The literature citations were prepared for import into Drupal, the web-based content management platform used by BioCat. All of the flagged Yes citations from each database EndNote library were copied into a new library file. Because of the overlap between abstracting/indexing coverage in major scientific databases, duplicate records were identified and removed. The same process was applied to the Maybe citations. This effort resulted in master EndNote libraries consisting of only those citations that met our newly vetted evaluation metrics criteria. Each of these libraries was imported into Drupal separately. As the Yes citations were imported, they were tagged Yes (n=447). As the Maybe citations were imported, they were tagged Maybe (n=1383). The citations that lacked a clear relevance to the research effort were not imported to BioCat (n=26,824).

After the previous steps were completed, a prioritized list of agents was requested from NBIC. From the 281 unique pathogens, toxins, and agents extracted from the nine priority lists (see Appendix B and Appendix C), the client provided PNNL with a smaller subset list of sixteen prioritized agents. These agents are those for which the NBIC has pressing information needs (see Appendix D). Using this smaller prioritized list of agents, the PNNL team queried the EndNote database to identify all of the citations matching these agents and meeting the Yes criteria (n=96). See Section 3.0 for detailed description of the model evaluation process and results.

3.0 Model Evaluations

This work was conducted in part to characterize published models that are capable of predicting disease events in order to determine opportunities for expanded research and to define operational readiness levels. Ultimately, practitioners need to be able to access and then operate these models without significant reprogramming. The model's forecasts or predictions need to have a known precision or error rate. An operationally ready model must also be mature. The PNNL team has expended a significant effort to address this last requirement. In partnership with the Los Alamos National Laboratory (LANL), a set of metrics was developed that enhance the attributes of biosurveillance models that PNNL co-authored during the Phase 1 BioCat project. The LANL team was qualified to contribute to developing the model metrics through their leadership in a Defense Threat Reduction Agency-sponsored effort to evaluate biosurveillance *data* sources. Over the course of three virtual meetings, the PNNL and LANL teams derived evaluation metrics to be used to assess disease models. The result of the discussions led to the PNNL team developing multiple metrics organized into three levels with each tier requiring greater domain knowledge in order to complete the evaluation. The metrics we recommend are enumerated in Appendix E, *Disease Forecast and Prediction Model Evaluation Metrics & Definitions*.

Given the potential importance of these models to global health security, it is essential to develop a categorization scheme that defines a model's viability for use in an operational setting. none exists, but one possibility is illustrated in the following discussion, based upon the readiness level" (TRL) originally developed by NASA.¹ NASA TRL levels were not designed to apply to modeling and simulation, much less biosurveillance models, so the definitions require modification. An initial modification of these definitions is shown in

Table 3.1. In such a scheme, the models are characterized based on how the model was validated, what type of data was used to validate the model, and the validity of data used to create the model.

Currently, the verification and validation (V&V) of predictive models, regardless of realm of application, is an area that requires better definition and techniques. The results of model V&V can be used in the definition of model operational readiness; however, the readiness level definitions must also be accompanied by data validation, uncertainty quantification, and model fitness-for-use evaluations, many of which are areas of active research. Clarity in the definition of readiness levels and the criteria to achieve upper levels will enable confidence in the ability to consistently assign readiness levels to models and lead to enhanced value to decision makers. Further development of the operational readiness level definitions and assignment criteria for disease prediction models are needed in order to evaluate the usefulness of this approach in an operational environment. Public health analysts and decision makers are in need of evidence-based guidance and the value of operational readiness levels for the models on which they depend cannot be overstated. PNNL recommends rigorous investigation to fully outline the criterion for each of the proposed operational readiness levels defined in

Table 3.1. It is also recommended that the viability of a particular model should be assessed through analytical case studies before its use in an operational setting.

¹ Mankins JC. 1995. *Technology Readiness Levels A White Paper*. Advanced Concepts Office, Office of Space Access and Technology, National Aeronautics and Space Administration.

Table 3.1. Initial Definitions of Operational Readiness Levels for Disease Prediction Models

Level	Definition
1	Research only reported on observed information
2	A constructed model which has yet to be applied to data (or The model theory has been developed based on observed or hypothesized information)
3	The model has been created but has not been validated
4	The model has been verified and validated
5	The model has been demonstrated as useful but for only its original location (pathogen or population) and is still being adjusted
6	The model has been demonstrated as useful in both its original location (pathogen or population) as well as an independent location (pathogen or population)
7...n	Further study is needed to explicitly delineate the criterion for additional Technology Readiness Levels

As revealed in Section 2.0, NBIC identified sixteen prioritized agents of interest (Appendix D). Using this list in conjunction with the papers that met our SME requirements, we identified 96 papers ready for evaluation. Based on time and staffing constraints, Web of Science and Google Scholar were used to identify the number of times these articles were cited since publication. The articles with post-publication citations (n=61) were considered candidates for evaluation. Twenty-four articles were fully evaluated and curated in BioCat using the new evaluation metrics (Appendix E). These articles and the evaluation metrics are discussed in Section 3.0.

Table 3.2. Client Priority Agent List

Agent	No. of Articles	No. Articles with Citations	No. Evaluated
<i>Bacillus anthracis</i>	4	0	0
Botulinum toxin	0	0	0
<i>Burkholderia mallei</i>	0	0	0
<i>Burkholderia pseudomallei</i>	0	0	0
Chlorpyrifos (Dursban)	-	-	-
Citrus Greening; Huanglongbing (HLB) of citrus; <i>Candidatus liberibacter americanus</i>	0	0	0
Endosulfan	-	-	-
Foot and Mouth Disease (FMD)	13	6	2
<i>Francisella tularensis</i>	2	1	1
Influenza (pandemic)	52	35	16
<i>Ralstonia solanacearum</i>	0	0	0
Rinderpest virus	0	0	0
SARS	8	7	0
Variola (major and minor)	3	2	2
Viral hemorrhagic fevers including Ebola and Marburg	6	4	3
<i>Yersinia pestis</i>	8	6	0
Total YES citations	96 Articles	61 with Citations	24 Evaluated

The following are the citations for the 24 articles that were fully evaluated.

Influenza, pandemic

Araz, OM, et al. 2011, "A Simulation Model for Policy Decision Analysis: A Case of Pandemic Influenza on a University Campus." *Journal of Simulation* 5(2):89-100.

Araz, OM, et al. 2012, "Simulation Modeling for Pandemic Decision Making: A Case Study with Bi-Criteria Analysis on School Closures." *Decision Support Systems* 55(2):564-75.

Baker, PRA, et al. 2011, "Epidemiologic Modeling with Flusurge for Pandemic (H1N1) 2009 Outbreak, Queensland, Australia." *Emerging Infectious Diseases* 17(9):1608-14.

Dorigatti, I, et al. 2012, "A New Approach to Characterising Infectious Disease Transmission Dynamics from Sentinel Surveillance: Application to the Italian 2009-2010 A/H1N1 Influenza Pandemic." *Epidemics* 4(1):9-21.

Gupta, SD, et al. 2011, "Modeling of H1N1 Outbreak in Rajasthan: Methods and Approaches." *Indian Journal of Community Medicine* 36(1):36-38.

Inaida, S, et al. 2011, "Geographic Trends and Spread of the Pandemic (H1N1) 2009 in the Metropolitan Areas of Japan Studied from the National Sentinel Data." *Japanese Journal of Infectious Diseases* 64(6):473-81.

Jin, Z, et al. 2011, "Modelling and Analysis of Influenza A (H1N1) on Networks." *BMC Public Health* 11(Suppl 1): 59.

Lee, VJ, et al. 2011, "Comparability of Different Methods for Estimating Influenza Infection Rates over a Single Epidemic Wave." *American Journal of Epidemiology* 174(4):468-78.

Lee, VJ, et al. 2011, "A Clinical Diagnostic Model for Predicting Influenza among Young Adult Military Personnel with Febrile Respiratory Illness in Singapore." *PLoS ONE* 6(3): e17468.

Loganathan, P, et al. 2011, "Towards Forecasting Flu Dynamics Using a Regionalized State Space Model." In *International Symposium on Advanced Control of Industrial Processes, ADCONIP 2011*, pp. 175-80.

Matrajt, L, and IM Longini, Jr. 2012, "Critical Immune and Vaccination Thresholds for Determining Multiple Influenza Epidemic Waves." *Epidemics* 4(1):22-32.

Nishiura, H, G Chowell, and C Castillo-Chavez. 2011, "Did Modeling Overestimate the Transmission Potential of Pandemic (H1N1-2009)? Sample Size Estimation for Post-Epidemic Seroepidemiological Studies." *PloS ONE* 6(3): e17908.

Rios-Soto, KR, B Song, and C Castillo-Chavez. 2011, "Epidemic Spread of Influenza Viruses: The Impact of Transient Populations on Disease Dynamics." *Mathematical Biosciences and Engineering* 8(1):199-222.

Sharomi, O, et al. 2011, "Modelling the Transmission Dynamics and Control of the Novel 2009 Swine Influenza (H1n1) Pandemic." *Bulletin of Mathematical Biology* 73(3): 515-548.

Smieszek, T, et al. 2011, "Reconstructing the 2003/2004 H3n2 Influenza Epidemic in Switzerland with a Spatially Explicit, Individual-Based Model." *BMC Infectious Diseases* 11:115.

Torres, SF, et al. 2012, "High Mortality in Patients with Influenza A pH1N1 2009 Admitted to a Pediatric Intensive Care Unit: A Predictive Model of Mortality." *Pediatric Critical Care Medicine* 13(2): e78-e83.

Francisella tularensis

Rydén, P, et al. 2012, "Outbreaks of Tularemia in a Boreal Forest Region Depends on Mosquito Prevalence." *Journal of Infectious Diseases* 205(2):297-304.

Foot and Mouth Disease

Dion, E, L VanSchalkwyk, and EF Lambin. 2011, "The Landscape Epidemiology of Foot-and-Mouth Disease in South Africa: A Spatially Explicit Multi-Agent Simulation." *Ecological Modelling* 222(13):2059-72.

Sanson, RL, et al. 2011, "Foot and Mouth Disease Model Verification and 'Relative Validation' through a Formal Model Comparison." *Revue scientifique et technique (International Office of Epizootics)* 30(2):527-40.

Variola

Fuller, T, et al. 2011, "Using Remote Sensing to Map the Risk of Human Monkeypox Virus in the Congo Basin." *EcoHealth* 8(1):14-25.

McKinley, TJ, et al. 2013, In Press, "Simulation-Based Bayesian Inference for Epidemic Models." *Computational Statistics and Data Analysis*.

Viral Hemorrhagic Fevers

Gale, P, et al. 2012, "Impact of Climate Change on Risk of Incursion of Crimean-Congo Haemorrhagic Fever Virus in Livestock in Europe through Migratory Birds." *Journal of Applied Microbiology* 112(2):246-57.

Medeiros, LCdC, et al. 2011, "Modeling the Dynamic Transmission of Dengue Fever: Investigating Disease Persistence." *PLoS Neglected Tropical Diseases* 5(1):e942.

Wei, L, et al. 2011, "Using Geographic Information System-Based Ecologic Niche Models to Forecast the Risk of Hantavirus Infection in Shandong Province, China." *American Journal of Tropical Medicine and Hygiene* 84(3):497-503.

After the 24 articles were evaluated, the PNNL SMEs (one epidemiologist and one microbiologist) identified several papers with distinctive or significant methodology or research conclusions. These papers are summarized in the following bullets.

- Inaida et al. (2011) applied kriging to influenza-like illness data to produce smoother spatial estimates of influenza incidence.
- Araz et al. (2011) evaluated the efficacy of social distancing and other mitigation practices for influenza transmission on a university campus, based on the campus response plan. They performed sensitivity analysis of the model parameters to identify the primary drivers of the changes in transmission.
- Dion et al (2011) performed a landscape analysis of cattle-buffalo contact near Kruger National Park in South Africa. Although not directly modeling foot and mouth disease, the identification of areas of greater likelihood for cattle-buffalo contact may serve as a proxy estimate of disease transmission risk. In the sensitivity analysis of the model parameters, they included two-way interaction terms.
- Gale et al. (2011) predicted and forecast the geographic spread of Crimean-Congo hemorrhagic fever, taking into account the impact of bird and tick populations in disease movement.
- Wei et al. (2011) do an excellent job clearly describing the model, the input data (source of input data), as well as type of data being fed into the model (e.g., continuous versus categorical). The high transparency of the model (not so clear in other publications) makes it possible for others to provide an independent verification and validation of the predictions and forecasts made in the publication.

4.0 BioCat 2.0 Website

BioCat has been developed and managed as a web-based system to maximize access by end users. Using a content management system was an important first step, because it allowed the use of existing code development and technology to create a structure for the website. The use of a content management system also enabled easy to use search and organization of information stored in the BioCat database. Login-based user permissions limit access and view/edit privileges.. Since the initial development of the BioCat website, the needs and implementation of the website have evolved, as described in this section.

4.1 Needs and Limitations of the BioCat 1.0 Website

In the first version of BioCat, Semantic MediaWiki (SMW) was the content management system. This system enables protecting the BioCat information from individuals who have not specifically been granted access. However, SMW also brought a number of limitations and challenges. The team wanted to have the ability to give different users permission to edit only certain parts of the information. SMW only allows someone to edit all the information or none of it. The team also wanted to track which users were editing what information and to enable the evaluation of models in an organized workflow (explained later). SMW did not support these functions.

4.2 Updating Software to Meet the Needs of BioCat 2.0

Given the limitations presented by SMW, a new content management system was sought for BioCat 2.0. The development team suggested another web-based content management platform called Drupal. Like SMW, Drupal is open source software, meaning that it is free to download and use. The open source model enables developers to constantly collaborate to update, fix and expand Drupal features and capabilities. As a result Drupal has a greater capacity for growth, variety and the ability to meet a far wider range of uses.

4.2.1 Benefits of the Drupal Web-Based Content Management Platform

Drupal has the benefit of being well supported and widely used. It has been adopted as a favorite among content management systems, especially in the U.S. government, including DHS. The community that has been built around Drupal allows collaboration, support, and development among hundreds of thousands of developers with different skill sets and experience. Drupal allows the development of new website add-on features that work with the original system, and those features can then be shared and used by others. Drupal is powerful enough to make many different features possible on a website, so Drupal is well positioned to support future BioCat development if needed.

One add-on feature that is already implemented in BioCat is called “Biblio.” Biblio is a bibliography tool for the management and display of lists of scholarly publications. It supports a number of formats when importing, exporting, or displaying the publications information, including BibTex, which was the format used for BioCat publications. Biblio also handles formatting directly on the webpages, so we can display it in APA, Chicago, and many other scholarly publication formats. This is handled by Drupal directly, rather than requiring the information to be manually entered in all the desired formats.

Drupal enables greater flexibility and control over user access and editing privileges. Different users can be access different content areas, rather than all or none. Different users access to change only certain parts of content, rather than all or none. Drupal also tracks how much work has been completed. These options allow BioCat to have a workflow in which individuals with different levels of knowledge can review specific content according to their permissions and then pass it on to the next person in a multi-stage review process that more efficiently uses different levels of expertise.

4.2.2 Implementation of Publications Review Workflow for BioCat

The “workflow” concept for BioCat was implemented in Drupal. The publication review process has been divided into multiple defined levels of review, each of which must be performed before an evaluation is considered complete. Once the publications are imported, they are deemed “Level 1.” At Level 1, interns completed the preliminary information. If the publication cannot be reviewed for some reason, for example if it is in a language no one can read, it is marked as “cannot evaluate,” and it will be removed from the workflow. If they are unsure about any information that they have filled in, the document can be flagged as needing a second look, which notifies upper-level reviewers to double check the inputted information. Publications deemed to be neither an event prediction nor an event forecast model are removed from the workflow when the Level 1 review is marked complete. All other completed Level 1 publications progress to Level 2 review.

In Level 2, additional information is entered about the model. As in Level 1, a second look may be requested if there is uncertainty regarding the review. Once the Level 2 review is marked complete, it passes to Level 3. In Level 3, SMEs add additional information to complete the review. Once the publication review is marked as complete the publication is removed from the workflow.

Each of these Levels has a web page listing all the publications that are in that Level, so anyone with access to those workflow levels can readily view the publications in the respective workflows. There are also pages to list publications that are completed, closed, unable to be evaluated, and needing second looks. This enables anyone in the workflow process to quickly identify tasks and access the relevant publications.

A user registration process is used to control access to the BioCat site and to the workflow levels. Users must create an account and be logged in before seeing any content on the website. In addition, accounts also must be approved, which enables to the ability to allow or deny each user's access. In addition to controlling site access, user registration enables tracking who uses and edits the site. Without user registration, there is no way to track this information. Moreover, content interactions and changes can be tracked; it is possible to revert content to an earlier version if necessary and to identify which user(s) is responsible if the site is misused.

Each user account can have its own set of permissions that determine what the user can see or change with great granularity (e.g., a publication date). Certain groups of people can be given similar sets of permissions. In BioCat, all users with an approved account can see any publications. Users with the corresponding permissions can edit citations. Users whose account is marked as Level 1 can only see the Level 1 workflow pages, and can only add/edit Level 1 review information. Users with Level 2 accounts can see and edit Levels 1 and 2, while Level 3 users can see and edit information at all three levels.

4.2.3 Site Administration Features

Site administrators can see and edit all workflow levels and can edit any part of a publication being reviewed. In addition, there are a number of pages with access restricted only to administrators. One of these pages contains user information and enables administrators to see all the accounts for the website. In addition to a list of users, demographics are also provided, including charts displaying e-mail domain type and organization affiliation type. This information is collected as part of the account creation process. Site administrators also have access to pages that list what publications have been marked as closed, completed, or cannot be evaluated.

4.2.4 Filtered Display of Content with Faceted Browsing

For anyone viewing a list of all the publications on the site, one of the most helpful tools that Drupal enables us to implement is “faceted browsing”. Faceted browsing filters a publications list and displays only those publications that meet user-selected criteria.² Normally, the list of publications that any logged-in user can see simply shows every publication on the website. A user looking for a specific publication would need to search for it, which means they would need to know some part of its name. However, users may often be looking for certain *types* of publications, not necessarily a particular one. Or

² Stefaner, M. (2009). *Dynamic Taxonomies and Faceted Search: Theory, Practice and Experience*. New York, NY, Springer .

they may not know exactly what publication they are looking for, but they may know certain things about it, such as what year it was published or what type of publication it is. With Drupal, publication information can be evaluated and the information that might matter to a browsing user can be expressed as content criteria. We can provide these content criteria as options (facets) next to the list. When a user selects their criteria of interest, the publication list automatically updates to show only publications that match those criteria. For example, if someone only wants to view publications that are conference papers, they simply select “Conference paper” under the “Filter by Publication Type” facet. The list will then show only conference papers, eliminating the need to browse through the full publication list looking for only conference papers.

4.2.5 Current Software for BioCat 2.0 Website

The following are Linux server requirements for BioCat as it has been built in Drupal:

- A minimum of 15 MB of disk space (60 MB preferred) for the installation, more for the database
- Apache, Nginx, or Microsoft IIS (server software)
- MySQL 5.0.15 or higher with PHP data objects (PDO), PostgreSQL 8.3 or higher with PDO, SQLite 3.3.7 or higher (open source database)
- PHP 5.2.5 or higher (scripting and programming language)
- Drush (a command line shell and scripting interface for Drupal)
- PHP modules: mod rewrite, apc, curl, GD library, ImageMagick, MB String, ldap, mysql, mssql, pear, xml serializer.

4.3 Visual Tour of the BioCat 2.0 Website

An unregistered site visitor cannot log in and does not have access to any content on the BioCat website. The only links available to visitors are to register/login, to contact us, or to use the DHS navigation at the bottom of the page.

The screenshot shows the top portion of the BioCat website. At the top right, there are links for [Login](#) and [Register](#). Below these, the DHS logo and the word "BioCat" are displayed. A "Home" button is visible. On the right side, there are links for [Contact Us](#) and [Site Map](#). The main content area features a large circular seal of the U.S. Department of Homeland Security. To the left of the seal, the text reads "National Biosurveillance Integration System" followed by a paragraph: "The NBIS enhances the identification, location and tracking of biological events potentially impacting homeland security by uniquely integrating information and data and leveraging interagency communications and relationships. NBIS supports prevention and mitigation of such events by providing timely notifications and ongoing situational awareness to enhance response of government agencies." Below this text, a message states "You are not authorized to access this page." At the bottom, a navigation menu is displayed with the following categories and links:

TOPICS	GET INVOLVED	HOW DO I?	U.S. DEPARTMENT OF HOMELAND SECURITY	NEWS	ABOUT DHS	SITE LINKS
Border Security	Blue Campaign	For the Public		Blog	The Secretary	Contact Us
Citizenship and Immigration Services	Citizen Corps	For Businesses		Comunicado de Prensa	Budget & Performance	DHS Component Websites
Civil Rights and Civil Liberties	If You See Something Say Something	For Travelers At DHS		Data	Careers	En Español
Cybersecurity	Ready.gov	A-Z Index		Events	Contact Us	Privacy Policy
Disasters	Stop.Think.Connect.			Fact Sheets	Doing Business with DHS	Notices
Economic Security	U.S. Coast Guard Auxiliary			Media Contacts	History	Plug-in
Homeland Security Enterprise				Multimedia	Laws & Regulations	FOIA
Homeland Security Jobs				Press Releases	Mission	Inspector General
Human Trafficking				Publications	Organization	Site Map
Immigration and Enforcement				Social Media		GobiernoUSA.gov
International Engagement				Speeches		USA.gov
Law Enforcement Partnerships				Testimony		The White House
Preventing Terrorism						
Transportation Security						

What an Unregistered Visitor to BioCat Sees

On the registration page, BioCat collects a username and email, as well as some preliminary information about the user. The registration must be approved by someone with administrative privileges on the website, who assigns a role to the new user before the user can log in and use the website.

Home » [User account](#)

User account

[Create new account](#) [Log in](#) [Request new password](#)

Username *

Spaces are allowed; punctuation is not allowed except for periods, hyphens, apostrophes, and underscores.

E-mail address *

A valid e-mail address. All e-mails from the system will be sent to this address. The e-mail address is not made public and will only be used if you wish to receive a new password or wish to receive certain news or notifications by e-mail.

Position/Title

Type of Agency

- None -

CAPTCHA

This question is for testing whether or not you are a human visitor and to prevent automated spam submissions.

Math question *

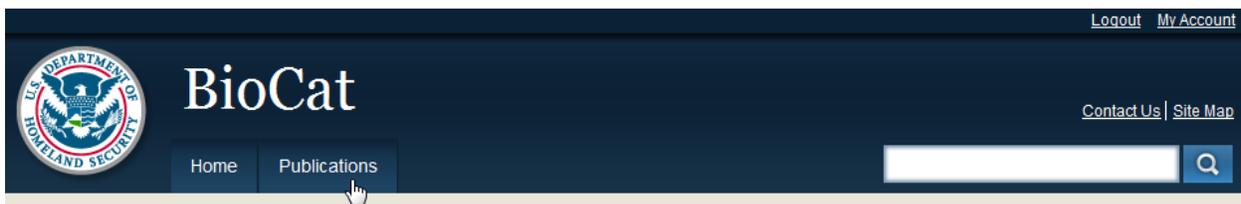
4 + 8 =

Solve this simple math problem and enter the result. E.g. for 1+3, enter 4.

[Create new account](#)

Registration Page for New User Accounts

Once users have an account and can log in, access to the site is based on the user roles that each user has been assigned by the site administrator. In the example here, a general user now has the option at the top of the page to view publications, as well as the ability to search the website.



Publications List and Search Options Available to General Users

The publications page lists a short summary of all the publications on the site. In addition to the titles, the page lists authors, keywords, and the titles of the publication from which each publication was taken. By default, the page lists all the publications alphabetically, 50 entries at a time.

The screenshot shows a web interface for a publications page. At the top left, there is a 'Home' link. Below it is a 'Navigation' menu with links for 'Biblio', 'My bookmarks', and 'Recent content'. The main content area is titled 'Publications' and contains a list of articles. Each article entry includes a title, authors, keywords, and the publication title. On the right side, there are three filtering sections: 'Filter by publication type:', 'Filter by year:', and 'Filter by authors:'. Each section lists various categories with the number of items in parentheses.

[Home](#)

Navigation

- [Biblio](#)
- [My bookmarks](#)
- [Recent content](#)

Publications

["Google flu trends" and emergency department triage data predicted the 2009 pandemic H1N1 waves in Manitoba](#)
Authors: Malik, Mohammad Tufail, Gumel, Abba, Thompson, Laura H., Strome, Trevor, Mahmud, Salaheddin M.
Keywords: Humans, Disease Outbreaks, *Influenza A Virus, H1N1 Subtype, Index Medicus, Influenza, Human / *epidemiology, Population Surveillance, *virology, Linear Models, Manitoba / epidemiology, *Internet, Emergency Service, Hospital / *organization & administration, *Triage
Publication Title: Canadian journal of public health. Revue canadienne de sante publique

["Wait and see" vaccinating behaviour during a pandemic: a game theoretic analysis](#)
Authors: Bhattacharyya, Samit, Bauch, Chris T.
Keywords: Humans, virology, Models, Statistical, *prevention & control, Influenza Vaccines / administration & dosage, Index Medicus, 0 / Influenza Vaccines, TRANSMISSION, Influenza, Human / *epidemiology, *Pandemics, economics, *immunology, Influenza A Virus, H1N1 Subtype / *immunology, Patient Acceptance of Health Care / *statistics & numerical data, Vaccination / *utilization
Publication Title: Vaccine

[\[A study on indicator system for early-warning on hemorrhagic fever with renal syndrome epidemic\]](#)
Authors: Lu, L. P., Wang, L., Ma, F., Yang, B., Zeng, X. J., Pan, L., Wan, K. L., Li, H.
Keywords: Humans, animal, article, epidemic, human, Animals, Disease Outbreaks, infection control, Disease notification, hemorrhagic fever with renal syndrome, early diagnosis
Publication Title: Zhonghua yu fang yi xue za zhi [Chinese journal of preventive medicine]

[A Bayesian analysis of the 2009 decline in tuberculosis morbidity in the United States](#)
Authors: Chen, M. P., Shang, N., Winston, C. A., Becerra, J. E.
Keywords: BAYESIAN analysis, time series analysis, seasonality, regression, tuberculosis trends, ARIMA, intervention models, change point, states
Publication Title: Statistics in Medicine

[A Bayesian approach for estimating bioterror attacks from patient data](#)
Authors: Ray, J., Marzouk, Y. M., Najm, H. N.
Keywords: Humans, Models, Statistical, Index Medicus, Bayes theorem, *epidemiology, *Disease Outbreaks, Bias (Epidemiology), Dose-Response Relationship, Drug, Russia / epidemiology, Bacillus anthracis, Anthrax / diagnosis, *Bioterrorism, Patients / *statistics & numerical data
Publication Title: Statistics in Medicine

Filter by publication type:

- [Journal Article \(1708\)](#)
- [Conference Proceedings \(53\)](#)
- [Book Chapter \(11\)](#)
- [Conference Paper \(7\)](#)
- [Report \(2\)](#)
- [Thesis \(2\)](#)
- [Miscellaneous \(1\)](#)

Filter by year:

- [2011 \(896\)](#)
- [2012 \(813\)](#)
- [2013 \(67\)](#)
- [2010 \(8\)](#)

Filter by authors:

- [Sánchez-Vizcaino, J. M. \(9\)](#)
- [Brownstein, John S. \(7\)](#)
- [Ferguson, Neil M. \(7\)](#)
- [Martínez-López, B. \(7\)](#)
- [Nishiura, Hiroshi \(7\)](#)
- [Patil, A. P. \(7\)](#)
- [Wu, Jianhong \(7\)](#)
- [Castillo-Chavez, Carlos \(6\)](#)
- [Fisman, David N. \(6\)](#)
- [Lee, Bruce Y. \(6\)](#)
- [Longini, Ira M., Jr. \(6\)](#)
- [Moqhadas, Seved M. \(6\)](#)
- [Morse, A. P. \(6\)](#)
- [Viboud, Cecile \(6\)](#)

Publications Page with Default Alphabetical Listing and Filtering Criteria Options

Criteria on the right side of the publications list enable users to narrow their search via faceted browsing. Each of these criteria list some of the most common results for each category. The exact number of list entries in that category are listed in parentheses after it. For example, if someone only wanted to see conference papers, they would select **conference paper** under **Publication Type**:

**Filter by
publication type:**

- [Journal Article \(1708\)](#)
- [Conference Proceedings \(53\)](#)
- [Book Chapter \(11\)](#)
- [Conference Paper \(7\)](#)
- [Report \(2\)](#)
- [Thesis \(2\)](#)
- [Miscellaneous \(1\)](#)

Choosing to Filter the Publications List by **Conference Paper** Publication Type

The page will update according to what criteria the user chooses as a search filter. In this case, the page reloads to show only those seven conference papers, eliminating the need to search through all the entries for this single publication type.

Home » Conference Paper

Navigation

- [Biblio](#)
- [My bookmarks](#)
- [Recent content](#)

Publications

Current search
Search found 7 items

Filter by publication type:
[\(-\) Conference Paper](#)

Filter by year:
[2011 \(4\)](#)
[2012 \(3\)](#)

Filter by authors:
[Almeida, V \(1\)](#)
[Benevenuto, F \(1\)](#)
[Bi, J. \(1\)](#)
[Cao, Z. \(1\)](#)
[Celoso, A \(1\)](#)
[Dong, W. \(1\)](#)
[Duan, W. \(1\)](#)
[Ferraz, F \(1\)](#)
[Ge, Y. \(1\)](#)
[Gomide, J \(1\)](#)
[Heller, K. \(1\)](#)
[Meira, W \(1\)](#)
[Neng, Z \(1\)](#)
[Pentland, A. \(1\)](#)
[Qiu, X. \(1\)](#)
[Shi, M. W. \(1\)](#)
[Teixeira, M. G. \(1\)](#)
[Tian, Z. \(1\)](#)
[Wu, Q. \(1\)](#)
[Xu, L. \(1\)](#)

[Dengue surveillance based on a computational model of spatio-temporal locality of Twitter](#)
Authors: Gomide, J, Celoso, A, Meira, W, Almeida, V, Benevenuto, F, Ferraz, F, Teixeira, M. G.
Publication Title: ACM WebSci'11 (June 14-17, 2011)

[Experimental study on modified model of indoor bio-aerosol dispersion](#)
Authors: Yan, D., Tian, Z., Neng, Z.
Keywords: Atmospheric aerosols, Forecasting, infectious disease, particulate matter, Bacteria, Technology, Experimental studies, Dispersions, Manufacture, Airflow pattern, Bacteria emanating experiment, Bio-aerosol, Continuous point-source dispersion model, Airflow patterns, Bioaerosols, Control methods, Dispersion models, Indoor environment, Modified model, Office environments, Transmission route, Under floor air supply, Atmospheric movements, Experiments
Publication Title: 2011 International Conference on Manufacturing Science and Technology, ICMST 2011

[Modeling and simulation for the spread of H1N1 influenza in school using artificial societies](#)
Authors: Cao, Z., Duan, W., Ge, Y., Qiu, X.
Keywords: Public Health, Computer simulation, Risk management, Information science, Students, Emergency Management, H1N1 Influenza, Modeling and Simulation, Public Health Security, Agent model, Artificial societies, Emergency measures, Environment models, Hebei Province, Role-based, Simulation experiments, Social relationships, School buildings
Publication Title: Pacific Asia Workshop on Intelligence and Security Informatics, PAISI 2011

[Modeling infection with multi-agent dynamics](#)
Authors: Dong, W., Heller, K., Pentland, A.
Keywords: Dynamics, Health risks, Random processes, Epidemic models, human dynamics, living lab, multi-agent modeling, stochastic process, Cellular Phone, Discrete-time, Multi-Agent Model, Potential risks, Small population, Spread of an infection, Cellular telephones, Mobile phones
Publication Title: 5th International Conference on Social Computing, Behavioral-Cultural Modeling and Prediction, SBP 2012

[Modelling to contain pandemic influenza A \(H1N1\) with stochastic membrane systems: A work-in-progress paper](#)
Authors: Xu, L.
Keywords: Computer simulation, Intervention strategy, infectious disease, pandemic influenza, Stochastic systems, bioinformatics, Decision makers, Mitigation strategy, membrane computing, pandemic influenza A (H1N1), pandemic spread modelling, stochastic membrane systems, Ecological systems, Membrane computing, B systems, PEPECT model, Work in progress, Membranes

Publications List Filtered by **Conference Paper** Publication Type

If the user selects one of the publications, BioCat will display a page that shows more details about it. General users will only be able to see the publication information. Here the user has selected the first publication on the previous list. Note that the links at the bottom allow the user to search Google Scholar for this publication, or to add it as a bookmark so that the user can look it up quickly on the **My bookmarks** page (linked in the left navigation) later when revisiting BioCat.

[Home](#) » [Publications](#)

Navigation

- [Biblio](#)
- [My bookmarks](#)
- [Recent content](#)

Dengue surveillance based on a computational model of spatio-temporal locality of Twitter

Thu, 2013-07-25 14:40 -- DrupalSysop

Title	Dengue surveillance based on a computational model of spatio-temporal locality of Twitter
Publication Type	Conference Paper
Year of Publication	2011
Authors	Gomide J , Celoso A , Meira W , Almeida V , Benevenuto F , Ferraz F , Teixeira M.G
Conference Name	ACM WebSci'11 (June 14-17, 2011)
Conference Location	Koblenz, Germany
Abstract	Twitter is a unique social media channel, in the sense that users discuss and talk about the most diverse topics, including their health conditions. In this paper we analyze how Dengue epidemic is reflected on Twitter and to what extent that information can be used for the sake of surveillance. Dengue is a mosquito-borne infectious disease that is a leading cause of illness and death in tropical and subtropical regions, including Brazil. We propose an active surveillance methodology that is based on four dimensions: volume, location, time and public perception. First we explore the public perception dimension by performing sentiment analysis. This analysis enables us to filter out content that is not relevant for the sake of Dengue surveillance. Then, we verify the high correlation between the number of cases reported by social statistics and the number of tweets posted during the same time period (i.e., $R^2 = 0.9578$). A clustering approach was used in order to exploit the spatiotemporal dimension, and the quality of the clusters obtained becomes evident when they are compared to social data (i.e., RandIndex = 0.8914). As an application, we propose a Dengue surveillance system that shows the evolution of the dengue situation reported in tweets, which is implemented in www.observatorio.inweb.org.br/dengue/ .

[Google Scholar](#) [Add Bookmark](#)

Example of Publication Information for One Publication

If a user has been approved as a member of the review workflow, the user will also be able to see more details about the publication, particularly about how the review process has been filled out so far.

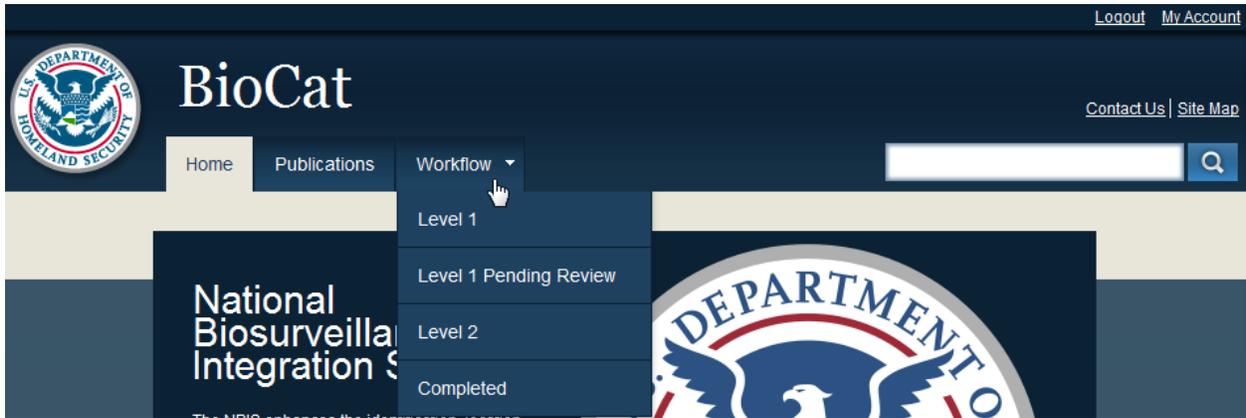
The screenshot displays a workflow information view for a model, organized into three levels. Each level is represented by a dark blue header bar followed by a list of attributes, their descriptions, and their values.

- Level 1**
 - Disease Specific**
The causative agent of the event being modelled is disease specific.
Yes
 - Agent/Pathogen/Toxin/Pest**
The agent(s)/pathogen(s)/toxin(s)/pest(s) of the event being modelled.
Influenza
 - Event Prediction**
The model predicts the future occurrence of an event/outbreak.
Yes
 - Event Forecast**
The model makes estimations about the future progress or behavior of an occurring event/outbreak.
No
 - Verified or Validated**
First pass identification of whether or not the model has been verified and/or validated.
Yes
- Level 2**
 - Corresponding Author**
The article's corresponding author as identified in the full-text document.
Vernon Lee
 - Model Creation Year**
The year the model was created.
2010-01-01 00:00:00
 - Population (Species) Affected Identified**
The specific affected population/species identified and modelled in the article.
Yes
 - Type of Validation and Verification**
The method used to verify and/or validate the model's maturity.
Specificity and Sensitivity (verification)
 - In Use in an Operational Environment**
The model is in use in an operational environment.
No
- Level 3**
 - Accessibility**
The degree to which and mechanism by where the model is made available, e.g., plug-and-play, downloadable, etc.
 - Run Time, Complexity, Capabilities Required**
Characteristics required for executing the model effectively.
 - Input Geography**
Geographic area attributes used as model input.
Local
 - Output Geography**

Workflow Information View Accessible to Authorized Review Editors

This workflow information is only accessible to those users who have been selected as editors. In addition, they are only able to see and perform review of publications in levels at or below their appointed role.

Here is shown the workflow view options in the BioCat menu for a person who is part of Level 2 review. (Note that the **Pending Review** options may appear on this menu when publications in those levels have been flagged as needing a second look.)



Workflow Menu Visible to Authorized Review Editors (Example)

Each of the workflow view pages on the **Workflow** menu list publications that are currently in the corresponding level of workflow. For example, the Level 2 editors can see below all the publications that are ready for them to review on the Level 2 page. They can click on the title to see the previously shown detailed view of the publication, or select **Review** to begin their review.

Home » [Workflow](#)

Navigation

- [Add content](#)
- [Biblio](#)
- [My bookmarks](#)
- [Recent content](#)

Review: Level 2

Title

Title	Updated date	
A mathematical model to distinguish sociological and biological susceptibility factors in disease transmission in the context of H1N1/09 influenza	08/01/13	Review
A new methodology for estimating contagious biological agent casualties as a function of time	08/01/13	Review
An individual-based simulation of pneumonic plague transmission following an outbreak and the significance of intervention compliance	08/01/13	Review
Analysing the spatial patterns of livestock anthrax in Kazakhstan in relation to environmental factors: a comparison of local (G(i)*) and morphology cluster statistics	07/26/13	Review
Attempted early detection of influenza A (H1N1) pandemic with surveillance data of influenza-like illness and unexplained pneumonia	07/31/13	Review
Determinants of the incidence of hand, foot and mouth disease in China using geographically weighted regression models	07/26/13	Review
Development of a resource modelling tool to support decision makers in pandemic influenza preparedness: The AsiaFluCap Simulator	08/02/13	Review
Excess mortality associated with influenza A and B virus in Hong Kong, 1998-2009	08/02/13	Review
Foot and mouth disease revisited: re-analysis using Bayesian spatial susceptible-infectious-removed models	07/26/13	Review
H3N2v and Other Influenza Epidemic Risk Based on Age-Specific Estimates of Sero-Protection and Contact Network Interactions	08/01/13	Review
Impact of weekday social contact patterns on the modeling of influenza transmission, and determination of the influenza latent period	08/02/13	Review
Likely effectiveness of pharmaceutical and non-pharmaceutical interventions for mitigating influenza virus transmission in Mongolia	07/29/13	Review
Modeling Disease Spreading on Complex Networks	08/02/13	Review
Multivariate Bayesian modeling of known and unknown causes of events—An application to biosurveillance	07/26/13	Review
Predicting AH1N1 2009 influenza epidemic in Southeast Europe	08/01/13	Review

Workflow View Page for a Level 2 Review Editor

Upon selecting the **Review** option for a specific publication, users can add/change information up to their level of permission. For example, a Level 2 user can edit Level 1 data and enter Level 2 data but cannot see or edit level 3 data.

Edit Biblio A mathematical model to distinguish sociological and biological susceptibility factors in disease transmission in the context of H1N1/09 influenza

View Edit Revisions

Title *
A mathematical model to distinguish sociological and biological susceptibility factors in disease transmission in the context of H1N1/09 influenza

Publication Type *
Journal Article

Article Attachment
 Simon2011.pdf (349.48 KB) Remove

Review Required *
 Yes
 No
 Maybe
 Cannot Evaluate

Level 1

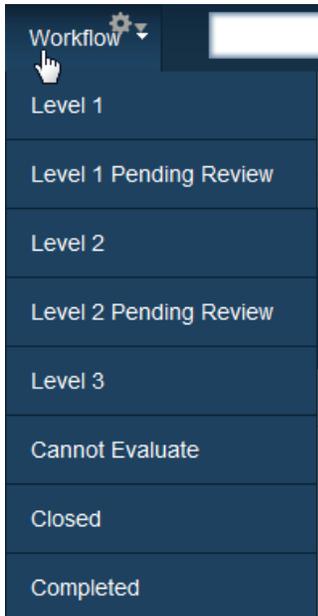
Level 2

Save

Example Form for Editing One Publication by a Level 2 Editor

The user can then save current information, flag it for a second-look review, or mark it as ready for the next level of review. Depending on the criteria previously stated, the workflow tool on the website will organize it according to its level, or take it out of the workflow process if it has been completed or closed.

Users that have been selected as BioCat administrators can see any level of the workflow and access the workflow views through the Workflow menu as shown here. They also have another option on the general menu, **User Admin**.



Workflow Menu with a Full Set of Review Options



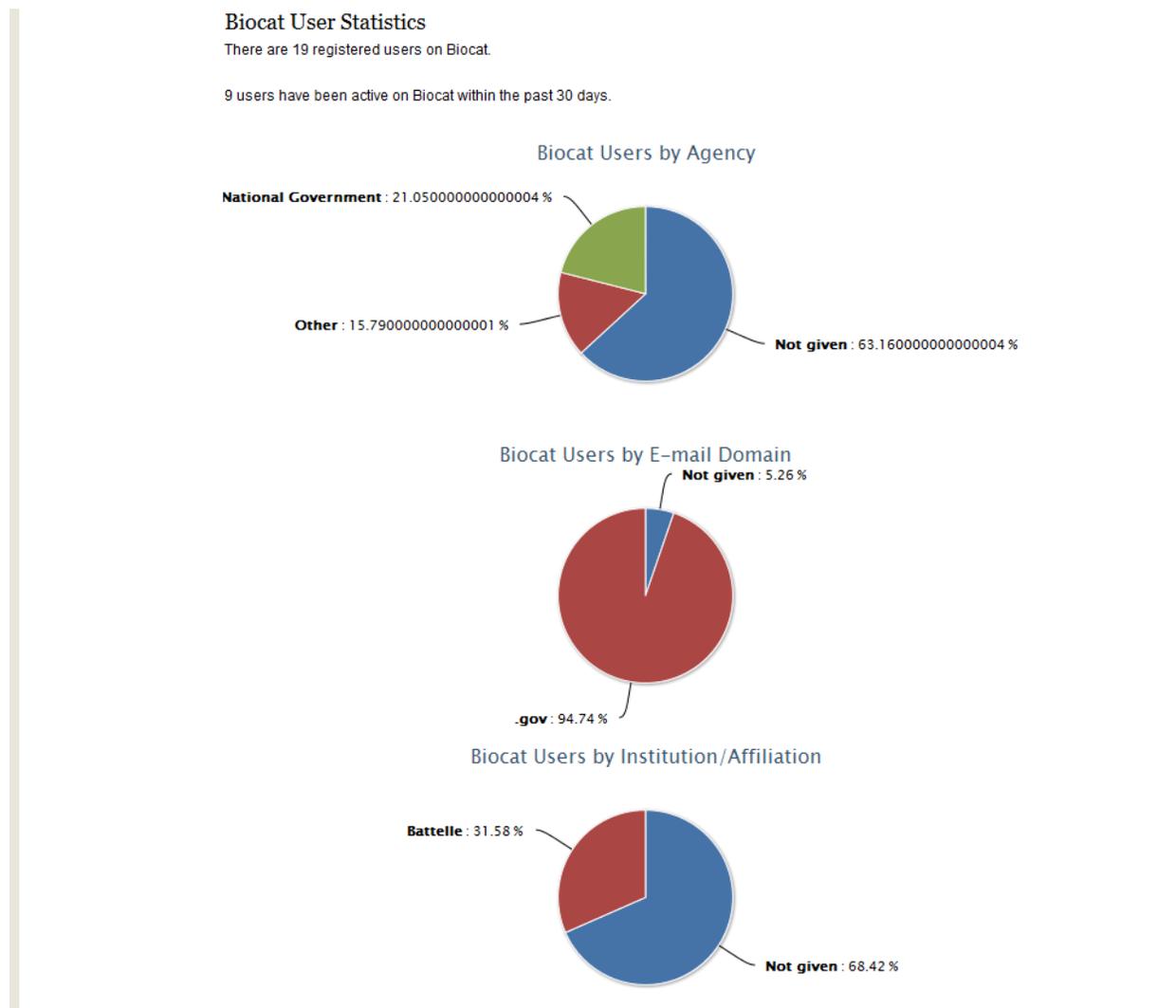
User Admin Option on the BioCat Menu

The user administration view page allows administrators to see a list of the users that have been registered on the site.



User Administration View Page (Example)

In addition, the user administration view page also lists statistics and charts about the users, automatically gathered by the site when users are registered.



User Statistics Charts for All the Users of the BioCat Website

5.0 Future Work

5.1 Integration of BioCat 1.0 into BioCat 2.0

Future funding efforts will enable the PNNL team to merge data collected in the first phase of research with the current work. This is a multi-step process that could be accomplished in either of two ways. The first path would consist of importing the literature citations from the first year effort into the new Drupal-based BioCat website. This can be completed relatively quickly with a small investment of funds and staffing. Literature citations would be migrated from EndNote X4 to EndNote X6 and then exported for ingestion into BioCat. The result would be that the new BioCat website would contain all the literature citations of the Phase 1 implementation, including those that may not be of confirmed interest.

The second path would be more in line with this year's efforts. The proposed first step would be to conduct a vetting process of the literature using the newly established assessment metrics, importing only those citations that survive the vetting and then evaluating a subset based on NBIC's priority list (Appendix D).

5.2 Enabling BioCat Accessibility on Mobile Devices

Another development direction that will likely prove useful for future use is mobile-compatible web design. Currently, the site is only made to display properly when viewed with a larger screen, such as on a desktop computer. When viewed at the smaller resolutions necessary for a tablet or phone, certain elements of the page do not display correctly, and the size of the webpage requires those with small devices to scroll left and right in addition to scrolling up and down. With additional development, the site would be adapted to automatically adjust to the width of the viewing device and to display the information in the most appropriate format according to the device being used. This would be a very helpful feature from the perspective of both users and administrators.

5.3 Additional Drupal Development Features

In addition to enabling mobile access, Drupal also supports additional features for development that could be useful to the BioCat user base. Examples of these features include *multilingual capabilities* and *automatic content ingestion*. Multilingual capabilities allow for the integration of more than one language into the website. This integration may be done through the navigation of the site, within the content of the site, or both. Automatic content ingestion enables Drupal to visit selected websites and social media sources, then extract content from those sources into BioCat. This could be a good way to discover new articles, refresh general site content from DHS or other agencies, or even monitor social media or other sources for mentions of topics and keywords of interest.

As time goes on, the Drupal community at large will continue to release new versions of the content manager. Ongoing maintenance of BioCat is critical to both management of day-to-day functionality and site security. The migration of BioCat into Drupal 8 (the next, developing version of Drupal) should be explored as that technology matures. This strategy will give BioCat access to significantly re-engineered features and new capabilities and it will keep the site current with the rest of the Drupal development and implementation community.

Finally, as users interact with BioCat, it is likely that additional "views" of the data will be requested to improve user work processes and to address user needs. Examples of existing views can be found on the various Workflow pages or the publications page. Examples of new views might include additional visualizations of publication information, reports, raw data for integration with other applications, or content workflow updates.

Appendix A

Biosurveillance Modelers and Stakeholders

The following table provides contact information and websites of the experts and research institutions that were identified as focused on the topic of biosurveillance.

Biosurveillance Modelers and Stakeholders

Name	Agency	Division	Email	Phone	Address	Position	Website(s)
Anderson, Sir Roy M	Imperial College London	School of Public Health	roy.anderson@imperial.ac.uk	44 (0)20 7594 3399	G22, Norfolk Place, St. Mary's Campus	Chair in Infectious Disease Epidemiology	
Anyamba, Assaf	NASA	Goddard Earth Sciences Technology Center at NASA's Goddard Space Flight Center	Assaf.Anyamba@nasa.gov	(301) 614-6601	NASA Goddard Building 33, Room G308 NASA GSFC, Code 618 Greenbelt, MD 20771 USA	Associate Research Scientist	http://gimms.gsfc.nasa.gov/
Arthur, Ray	HHS/CDC	Global Disease Detection Operations Center	rca8@cdc.gov; ray.arthur@cdc.hhs.gov	(404) 639-3855	Atlanta, Georgia	Director	
Barrett, Christopher L.	Virginia Tech	Virginia Bioinformatics Institute	cbarrett@vbi.vt.edu	(540) 231-1745	Research Building XV Virginia Tech Blacksburg, VA 24061 USA	Director of Network Dynamics and Simulation Science Laboratory	http://ndssl.vbi.vt.edu/people/cbarrett.html
Bartrand, Timothy A.	Tetra Tech Clancy Environmental Consultants, Drexel University	Environmental Information Services Group	tab32@drexel.edu; tbartrand@clancyenv.com	(610) 668-9227	109 Llanfair Rd Bala Cynwyd, PA 19004 USA	Environmental Engineer, Visiting research assistant professor	
Beardsley, Kristine	FBI	Bioterrorism Prevention Program in the Weapons of Mass Destruction Directorate	kristine.beardsley@ic.fbi.gov			Supervisory Special Agent	
Beckham, Tammy	Texas A&M	Texas Veterinary Medical Diagnostic Laboratory	trbeckham@ag.tamu.edu; tbeckham@tvmdl.tamu.edu	979-845-2855	1500 Research Parkway, Ste. 130 College Station, TX 77845	Director of Department of Homeland Security's Center of Excellence for Foreign Animal and Zoonotic Diseases (FAZD Center)	

Biosurveillance Modelers and Stakeholders

Name	Agency	Division	Email	Phone	Address	Position	Website(s)
Blackburn, Jason	University of Florida	Emerging Pathogens Institute	jkblackburn@ufl.edu	352-392-0494	Emerging Pathogens Institute University of Florida, Gainesville, FL		
Boghossian, Aida	USDA	DOD/NCMI	aboghossian@ncmi.detrick.army.mil	301-619-8539	Ft Detrick, Frederick, MD	USDA/APHIS rep	
Boots, Mike	University of Exeter, UK	Centre for Ecology and Conservation, College of Life and Environmental Sciences	m.boots@exeter.ac.uk	44 (0) 13 2625 5735	Daphne du Maurier Centre for Ecology and Conservation College of Life and Environmental Sciences University of Exeter, Cornwall Campus TR10 9 EZ UK	Professor of Disease Biology	http://people.exeter.ac.uk/mb437/
Bright, Patti	Department of the Interior	Contaminant Biology Program	pbright@usgs.gov	703-648-4238		Acting USGS Contaminant Biology Program Coordinator	
Buckeridge, David L.	University Clinic - Royal Victoria Hospital, Canada	Department of Epidemiology, Biostatistics and Occupational Health	david.buckeridge@mcgill.ca	514-934-1934 Ext. 32991	Pine 1140 1140 avenue des Pins Ouest Montreal H3A 1A3 Quebec Canada	Associate Professor	http://surveillance.mcgill.ca/index.html
Butel, Mike	DoD	Global Health Surveillance	Michael.Butel@ha.osd.mil		U.S. Department of Defense Arlington, VA	NIWG Principal, Assistant Secretary of Defense (Health Affairs)	

Biosurveillance Modelers and Stakeholders

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Codeco, Claudia T.	Oswaldo Cruz Foundation, Rio de Janeiro, Brazil	The Scientific Computing Program	codeco@malaria.occ.fiocruz.br; codeco@fiocruz.br; codeco@icict.fiocruz.br	55-21-598-4399	Programa de Computação Científica/ Fundação Oswaldo Cruz Av. Brasil, 4365, Manguinhos Rio de Janeiro, RJ, Brasil, 21045-900	Associate Researcher	
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Biosurveillance Modelers and Stakeholders

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Cooke, William H.	Mississippi State University	GeoResources Institute	whc5@geosci.msst ate.edu	601-325-9479	GeoResources Institute Box 9652 Mississippi State, MS 39762	Associate professor, research scientist, interim department head	
Corcoran, Patrick	US Postal Service	Dangerous Mail Investigations	PJCorcoran@uspis. gov			Inspector in Charge for Operational Support	
Craft, David L.	Massachusetts General Hospital and Harvard Medical School	Department of Radiation Oncology and Operations Research Center	dcraft@alum.mit.e du; dcraft@partners.or g		Massachusetts General Hospital Northeast Proton Center - NPTC 55 Fruit St Boston, MA 02114	Assistant Professor	
Craigmile, Peter F.	Ohio State University, University of Glasgow, Scotland	Department of Statistics	pfc@stat.osu.edu; peter.craigmile@gl asgow.ac.uk		School of Mathematics and Statistics University Gardens University of Glasgow Glasgow, G12 8QQ United Kingdom	Associate Professor, Reader	http://www.stats.gla.ac.uk/~pcraigmile/
Culotta, Aron	Northeastern Illinois University	Department of Computer Science	A- Culotta@neiu.edu	(773) 442-4720 (department)	5500 N. St. Louis Avenue Chicago, IL 60625	Assistant Professor	http://cs.neiu.edu/~culotta
Daszak, Peter	EcoHealth Alliance; Columbia University	Center for Infection and Immunity	daszak@ecohealtha lliance.org	212-380-4473; 815-365-8337	460 West 34th St, 17th floor, New York, NY 10001- 2320	President and Disease Ecologist, adjunct professor	
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Biosurveillance Modelers and Stakeholders

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Eagle, Nathan	MIT; txteagle Inc., Northeastern University, The Santa Fe Institute, Harvard University	MIT Media Lab, Department of Epidemiology at Harvard; Computer Science at Northeastern	nathan@mit.edu; eagle@hsph.harvard.edu	857-776-3279; (505) 204-6637	MIT Design Laboratory 77 Massachusetts Ave Bldg E15-383 Cambridge MA 02139 The Santa Fe Institute 1399 Hyde Park Rd Santa Fe, NM 87501	CEO of txteagle, Adjunct Assistant Professor at Harvard, visiting assistant professor at MIT Media lab, research assistant professor at Northeastern, Omidyar Fellow at Santa Fe Institute	
Eisen, Rebecca J.	CDC	Division of Vector Borne Infectious Diseases, National Center for Zoonotic, Vector Borne, and Enteric Diseases, Ft. Collins, Co.	dyn2@cdc.gov; rebecca.eisen@cdc.hhs.gov	(970) 266-3523	3150 Rampart Rd Fort Collins CO 80521	Research biologist	
Elder, Bret D.	LSU	Department of Biological Sciences	elder@lsu.edu	225-578-6733	206 Life sciences Annex Baton Rouge, LA 70803	Assistant professor	
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Biosurveillance Modelers and Stakeholders

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Fang, Haiping	Chinese Academy of Sciences, Shanghai	Shanghai Institute of Applied Physics	fanghaiping@sinap.ac.cn	86-21-39194785	Shanghai Institute of Applied Physics Chinese Academy of Sciences P.O. Box 800-204 Shanghai 201800, China	Professor in Physics	
Ferguson, Neil M	Imperial College London	Medical Research Council Centre for Outbreak Analysis and Modeling, School of Public Health	neil.ferguson@imperial.ac.uk	44 (0) 20 7594 3296	Medical School, St. Mary's Campus	Director, Centre for Outbreak Analysis and Modeling, Professor	
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Biosurveillance Modelers and Stakeholders

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Biosurveillance Modelers and Stakeholders

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Biosurveillance Modelers and Stakeholders

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Biosurveillance Modelers and Stakeholders

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Johansson, Anders	Swedish Defense Research Agency, Umea University	Department of Clinical Microbiology, Bacteriology and Infectious Diseases	anders.johansson@climi.umu.se; anders.johansson@infdis.umu.se	46 90 785 17 32	Department of Hospital Infection Control SE-901 85 Umea Sweden	Senior lecturer	
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Keim, Paul	Northern Arizona University	Pathogen Genomics Division, Biology, The Translational Genomics Research Institute	pkeim@tgen.org	928-523-1078		Professor of Biology, Director	
Kim, Deok Ryun	International Vaccine Institute,	Translational Research Division	drkim@ivi.int		South Korea Institute of Hygiene and Epidemiology, Vietnam	Senior Statistical Analyst	

Biosurveillance Modelers and Stakeholders

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Liao, Chung-Min	National Taiwan University	Department of Bioenvironmental Systems Engineering	cmliao@ntu.edu.tw	886-2-2363-4512	National Taiwan University Taipei, Taiwan 10617 ROC		
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Biosurveillance Modelers and Stakeholders

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Longini, Ira	University of Florida	College of Public Health	ilongini@ufl.edu	404-275-5156	Emerging Pathogens Institute University of Florida Gainesville, FL Fort Collins, CO	Professor	http://csquid.org/people/ira-m-longini-jr-phd/
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Biosurveillance Modelers and Stakeholders

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Biosurveillance Modelers and Stakeholders

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Ortega, Neli Regina Siqueira	University of Sao Paulo	Medical Informatics, Faculty of Medicine	neli@dim.fm.usp.br	(11) 30617682	University of Sao Paulo, Faculty of Medicine, University of Birmingham, Department of Pathology Rua Teodoro Sampaio 115 - Discipline of Informatics - LIM01 Cerqueira Cesar 05405-000 - Sao Paulo, SP - Brazil	Tenured Professor	http://lattes.cnpq.br/600175225658473 5
Pascual, Mercedes	University of Michigan	Dept of Ecology and Evolutionary Biology	pascual@umich.edu	734-615-9808	2039/2041 Kraus Natural Science Bldg, 830 North University, Ann Arbor, MI 48109-	Professor	

Biosurveillance Modelers and Stakeholders

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					1048		
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Biosurveillance Modelers and Stakeholders

Name	Agency	Division	Email	Phone	Address	Position	Website(s)
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Biosurveillance Modelers and Stakeholders

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Biosurveillance Modelers and Stakeholders

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Biosurveillance Modelers and Stakeholders

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Weaver, Curtis	HHS/CDC	Office of Surveillance, Epidemiology and Laboratory Services, Public Health Surveillance	atr8@cdc.gov		CDC COTPER/BCU Atlanta, GA 30333 USA	Acting Director of Biosurveillance Coordination Activity	

Biosurveillance Modelers and Stakeholders

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Zhou, Yicang	Xi'an Jiaotong University, China	Department of Applied Mathematics	zhouyc@mail.xjtu.edu.cn		Department of Mathematics, Xi'an Jiaotong University, Xi'an, 710049, P. R. China		

Appendix B

Client-Identified Source Lists for Agents, Toxins, and Pathogens of Interest¹

List Name	List Location	NBIC Target Domains
HHS and USDA Select Agents and Toxins	http://www.selectagents.gov/resources/List_of_Select_Agents_and_Toxins_2012-12-4.pdf	Human, Animal, Plant, Food
CDC - The Top 5 Causes of Waterborne Outbreaks in Private Groundwater Wells	http://www.cdc.gov/healthywater/drinking/private/wells/diseases.html	Human, Environment, Food
EPA - Regulated Microorganisms (pathogens - excludes indicator organisms)	http://water.epa.gov/drink/contaminants/index.cfm#Microorganisms	Human, Environment, Food
CDC - Bioterrorism Agents/Diseases	http://www.bt.cdc.gov/agent/agentlist-category.asp	Human, Animal, Environment, Food
CDC - Healthcare-Associated Infections (HAIs)	http://www.cdc.gov/HAI/organisms/organisms.html	Human, Food
CDC - List of Nationally Notifiable Conditions	http://wwwn.cdc.gov/nndss/document/2012_Case%20Definitions.pdf	Human, Animal, Environment, Food
USDA - National Animal Health Reportable Disease List	http://www.aphis.usda.gov/animal_health/nahrs/downloads/2011_nahrs_dz_list.pdf	Human, Animal, Environment, Food
USDA - Pathogens that Threaten U.S. Agriculture	http://www.ars.usda.gov/research/docs.htm?docid=14271	Food, Plant
Pests and Pathogens Threatening North American Forests, Trees	http://www.dontmovefirewood.org/	Food, Plant

¹ These lists were received from the client via email on December 13, 2012.

Appendix C

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1	Abrin	Abrin; Abrin; abrin-a; Proteopedia 1abr; Abrin poisoning; rosary pea; jequirity pea; <i>Abrus precatorius</i>
5	<i>Acinetobacter</i>	<i>Acinetobacter</i> ; <i>Acinetobacter baumannii</i>
1, 7	African horse sickness virus	African horse sickness virus; African horse sickness; AHSV; AHS
1, 7	African swine fever virus	African swine fever virus; African swine fever
1	Akabane virus	Akabane virus; Akabane disease; Tinaroo virus; Sabo virus; Yaba-7
9	Alder Dieback	Alder Dieback; <i>Phytophthora alni</i> ; alder die back
8, 9	Ambrosia beetle	Ambrosia beetle; <i>Xyleborus glabratus</i> ; laurel wilt
6, 7	Anaplasmosis (<i>Anaplasma marginale</i> , <i>A. centrale</i>)	Anaplasmosis; <i>Anaplasma marginale</i> ; <i>Anaplasma centrale</i> ; gall sickness
9	Asian Gypsy Moth	Asian Gypsy Moth; <i>Lymantria dispar asiatica</i> ; <i>Lymantria dispar japonica</i>
9	Asian Longhorned Beetle	Asian Longhorned Beetle; <i>Anoplophora glabripennis</i> ; Asian Longhorned beetle
7	Aujeszky's Disease (Pseudorabies)	Aujeszky's Disease; Pseudorabies; PRV
7	Avian chlamydiosis (psittacosis and ornithosis, <i>Chlamydia psittaci</i>)	Avian chlamydiosis; psittacosis; ornithosis; <i>Chlamydia psittaci</i> ; avian chlamydiosis
7	Avian infectious bronchitis	Avian infectious bronchitis; IB
7	Avian infectious laryngotracheitis	Avian infectious laryngotracheitis; ILT
1	Avian influenza virus	Avian influenza virus; highly pathogenic avian influenza virus; HPAI virus; HPAI A(H5N1); bird flu
6, 7	Babesiosis	Babesiosis; <i>Babesia bovis</i> ; <i>Babesia bigemina</i> ; <i>Babesia microti</i>
1, 4, 6, 7	<i>Bacillus anthracis</i>	<i>Bacillus anthracis</i> ; anthrax; woolsorter's disease; ragpicker's disease
1	<i>Bacillus anthracis</i> Pasteur strain	<i>Bacillus anthracis</i> Pasteur strain

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
9	Balsam Woolly Adelgid	Balsam Woolly Adelgid; <i>Adelges piceae</i>
9	Banded Elm Bark Beetle	Banded Elm Bark Beetle; <i>Scolytus schevyrewi</i>
9	Beech Bark Disease	Beech Bark Disease; <i>Nectria coccinea</i> var. <i>faginata</i> Lohman; beech scale
1, 7	Bluetongue virus (exotic)	Bluetongue virus exotic; bluetongue virus; BTV; catarrhal fever
1, 4, 6	Botulinum neurotoxin producing species of Clostridium*	Botulinum neurotoxin producing species of Clostridium; <i>Clostridium botulinum</i> ; <i>Clostridium butyricum</i> ; <i>Clostridium baratii</i> ; <i>Clostridium argentinense</i> ; botulism
1, 4	Botulinum neurotoxins	Botulinum neurotoxins; BoNT; botulinum toxin; botulinum toxin type A; botulinum toxin type B; botulinum toxin type C1; botulinum toxin type D; botulinum toxin type E; botulinum toxin type F; botulinum toxin type G; BTX-A; BTX-C1; BTX-D; BTX-E; sausage poison; fatty poison
7	Bovine Genital Campylobacteriosis (<i>Campylobacter fetus venerealis</i>)	Bovine Genital Campylobacteriosis; <i>Campylobacter fetus venerealis</i> ; bovine venereal campylobacteriosis; BVC
1, 7	Bovine spongiform encephalopathy agent	Bovine spongiform encephalopathy agent; bovine spongiform encephalopathy prion; CJD prion; Bovine spongiform encephalopathy; BSE; variant Creutzfeldt–Jakob disease; CJD; v-CJD; new variant Creutzfeld–Jakob disease; nv-CJD; mad cow disease
7	Bovine Tuberculosis (<i>Mycobacterium bovis</i>)	Bovine Tuberculosis; <i>Mycobacterium bovis</i> ; Bovine TB
7	Bovine viral diarrhea (BVD)	Bovine viral diarrhea; BVD; pestivirus genus; BVDV; BVDV-1; BVDV-2; hog cholera virus
9	Bromeliad Weevil	Bromeliad Weevil; <i>Metamasius callizona</i>
9	Brown Longhorned Spruce Beetle	Brown Longhorned Spruce Beetle; <i>Tetropium fuscum</i>
1, 7	<i>Brucella abortus</i>	<i>Brucella abortus</i> ; bovine brucellosis; brucellosis; Bang's disease; Crimean fever; Gibraltar fever; Malta fever; Maltese fever; Mediterranean fever; rock fever; undulant fever; contagious abortion; infectious abortion

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1, 7	<i>Brucella suis</i>	<i>Brucella suis</i> ; brucellosis; Porcine brucellosis; Enzootic Abortion; Contagious Abortion, Undulant Fever
1, 4, 6, 7	Brucellosis (<i>Brucella</i> species)	Brucellosis (<i>Brucella</i> species)
1, 7	Brucellosis, Ovine and Caprine	Brucellosis, Ovine; Brucellosis, Caprine; <i>Brucella melitensis</i> ; <i>Brucella ovis</i> ; ovine epididymitis; Malta Fever
5	<i>Burkholderia cepacia</i>	<i>Burkholderia cepacia</i>
1, 4, 7	<i>Burkholderia mallei</i> *	<i>Burkholderia mallei</i> ; <i>Pseudomonas mallei</i> ; glanders
1	<i>Burkholderia pseudomallei</i> *	<i>Burkholderia pseudomallei</i> ; <i>Pseudomonas pseudomallei</i> ; melioidosis; Whitmore's disease
9	Butternut Canker	Butternut Canker; <i>Ophiognomonia clavignenti-juglandacearum</i>
9	Cactus Moth	Cactus Moth; <i>Cactoblastis cactorum</i> ; Prickly pear moth
1	Camel pox virus	Camel pox virus; CMPV; Camelpox
2	<i>Campylobacter jejuni</i>	<i>Campylobacter jejuni</i> ; campylobacteriosis; campy; campylobacter enteritis; gastroenteriti; <i>Campylobacter fetus</i> subsp. Jejuni
8	<i>Candidatus Liberibacter asiaticus</i> , <i>L. africanus</i> , and <i>L. americanus</i>	<i>Candidatus Liberibacter asiaticus</i> ; <i>L. africanus</i> ; <i>L. americanus</i> ; Citrus greening disease; huanglongbing; yellow dragon disease
7	Caprine arthritis and encephalitis (CAE)	Caprine arthritis and encephalitis; CAE; CAEV; Caprine arthritis and encephalitis virus
1	Cercopithecine herpesvirus 1 (Herpes B virus)	Cercopithecine herpesvirus 1; Herpes B virus; Herpesvirus simiae; Macacine herpesvirus 1; Cercopithecine Herpesvirus 1; Herpes Simian B virus; Herpes B virus; herpes B; monkey B virus; herpesvirus simiae; herpesvirus B; CHV-1; CeHV-1; SHBV
6	Chancroid	Chancroid; <i>Haemophilus ducreyi</i>
9	Chestnut Blight	Chestnut Blight; <i>Cryphonectria parasitica</i>
9	Chestnut Gall Wasp	Chestnut Gall Wasp; <i>Dryocosmus kuriphilus</i>
6	<i>Chlamydia trachomatis</i> infection	<i>Chlamydia trachomatis</i> infection; <i>Chlamydia trachomatis</i> ; chlamydia
6	Cholera	Cholera; <i>Vibrio cholera</i>

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
8	Citrus leprosis virus	Citrus leprosis virus; Nail-head rust; nail-head spot; scaly bark
9	Citrus Longhorned Beetle	Citrus Longhorned Beetle; <i>Anoplophora chinensis</i>
1, 7	Classical swine fever virus	Classical swine fever virus; classical swine fever; hog cholera; CSFV; pig plague
5	<i>Clostridium difficile</i>	<i>Clostridium difficile</i> ; CDF; C diff
1, 4	<i>Clostridium perfringens</i> epsilon toxin	<i>Clostridium perfringens</i> ,epsilon toxin; <i>Clostridium perfringens</i> enteritis; Clostridium enterotoxin poisoning
5	<i>Clostridium sordellii</i>	<i>Clostridium sordellii</i>
1, 6	Coccidioides posadasii/Coccidioides immitis	Coccidioides posadasii; Coccidioides immitis; coccidioidomycosis; Valley Fever; San Joaquin Valley Fever; desert bumps; desert rheumatism; Posadas' disease
9	Common Pine Shoot Beetle	Common Pine Shoot Beetle; <i>Tomicus piniperda</i>
1	Conotoxins (Short, paralytic alpha conotoxins containing the following amino acid sequence X1CCX2PACGX3X4X5X6CX7)	Conotoxins; short paralytic alpha conotoxins; amino acid sequence X1CCX2PACGX3X4X5X6CX7; α -conotoxin; δ -conotoxin; κ -conotoxin; μ -conotoxin; ω -conotoxin; alpha conotoxin; delta conotoxin; kappa conotoxin; mu conotoxin; omega conotoxin
7	Contagious agalactia (<i>Mycoplasma agalactiae</i> , <i>M. Capricolum capricolum</i> , <i>M. putrefaciens</i> , <i>M. mycoides mycoides</i> , <i>M. mycoides mycoides</i> LC)	Contagious agalactia; <i>Mycoplasma agalactiae</i> ; <i>Mycoplasma Capricolum capricolum</i> ; <i>Mycoplasma putrefaciens</i> ; <i>Mycoplasma mycoides mycoides</i> ; <i>Mycoplasma mycoides mycoides</i> LC
7	Contagious bovine pleuropneumonia (<i>Mycoplasma mycoides mycoides</i>)	Contagious bovine pleuropneumonia; <i>Mycoplasma mycoides mycoides</i> ; CBPP
7	Contagious caprine pleuropneumonia (<i>Mycoplasma capricolum capripneumoniae</i>)	Contagious caprine pleuropneumonia; <i>Mycoplasma capricolum capripneumoniae</i> ; Contagious caprine pleuropneumonia; CCPP; Mycoplasma strain F38
7	Contagious equine metritis (<i>Taylorella equigenitalis</i>)	Contagious equine metritis; <i>Taylorella equigenitalis</i> ; CEM
8	Cotton leaf curl virus	Cotton leaf curl virus; cotton leaf curl; Cotton leaf curl geminivirus; CLCuV
1, 4, 6, 7	<i>Coxiella burnetii</i>	<i>Coxiella burnetii</i> ; Q fever; Query fever

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
3	coxsackieviruses A	coxsackieviruses A; Hand, foot, and mouth disease; HFMD; Cosackie A virus
3	coxsackieviruses B	coxsackieviruses B; Coxsackie B virus
7	Crayfish plague (<i>Aphanomyces astaci</i>)	Crayfish plague; <i>Aphanomyces astaci</i>
1, 7	Crimean-Congo hemorrhagic fever virus	Crimean-Congo hemorrhagic fever virus; Crimean-Congo hemorrhagic fever; CCHF
8	<i>Cronartium flaccidum</i>	<i>Cronartium flaccidum</i> ; Scots pine blister rust; Cronartium rust; blister rust; pine-stem rust; resin canker; resin top disease; two-needle pine blister rust
3, 6	Cryptosporidiosis	Cryptosporidiosis; <i>Cryptosporidium enteritis</i> ; crypto; <i>Cryptosporidium parvum</i>
9	Cycad Aulacaspis Scale	Cycad Aulacaspis Scale; <i>Aulacaspis yasumatsui</i>
6	Cyclosporiasis	Cyclosporiasis; <i>Cyclospora cayetanensis</i> ; traveller's diarrhea; cyclospora
7	Cysticercosis (<i>Cysticercus cellulosae</i> , metacestode stage of <i>Taenia solium</i>)	Cysticercosis; <i>Cysticercus cellulosae</i> ; metacestode stage of <i>Taenia solium</i> ; pork tapeworm; cysticercosis; neurocysticercosis; ophthalmic cysticercosis; subcutaneous cysticercosis
6	Dengue virus infections	Dengue virus infections; dengue virus; dengue fever; dengue hemorrhagic fever; dengue infection
1	Diacetoxyscirpenol	Diacetoxyscirpenol; 4,15-diacetoxyscirpenol
6	Diphtheria	Diphtheria; <i>Corynebacterium diphtheriae</i>
9	Dogwood Anthracnose Disease	Dogwood Anthracnose ; <i>Discula destructiva</i>
7	Dourine (<i>Trypanosoma equiperadum</i>)	Dourine; <i>Trypanosoma equiperadum</i>
7	Duck viral hepatitis	Duck viral hepatitis; duck hepatitis
9	Dutch Elm Disease	Dutch Elm Disease; <i>Ophiostoma ulmi</i> ; <i>Ophiostoma himal-ulmi</i> ; <i>Ophiostoma novo-ulmi</i>
2	<i>E. coli</i> 0157:H7	<i>E. coli</i> 0157:H7; <i>Escherichia coli</i> ; <i>E. coli</i> ; Shiga toxin-producing <i>E. coli</i> ; STEC; enterohemorrhagic <i>E. coli</i> ; EHEC; <i>E. coli</i> gastroenteritis
1	Eastern Equine Encephalitis virus	Eastern Equine Encephalitis virus; EEEV; EEE

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1	Ebola virus	Ebola virus; Ebola; Ebola hemorrhagic fever; EHF; Ebola virus; EBOV; Zaire ebolavirus; ZEBOV; Sudan ebolavirus; SEBOV; Reston ebolavirus; REBOV; Cote d'Ivoire ebolavirus; CIEBOV; Bundibugyo ebolavirus
7	Echinococcosis / hydatidosis	Echinococcosis; hydatidosis; <i>Echinococcus granulosus</i> ; <i>Echinococcus multilocularis</i> ; <i>Echinococcus oligarthrus</i> ; <i>Echinococcus vogeli</i> ; <i>Echinococcus shiquicus</i> ; hydatid disease; cystic echinococcosis
3	echoviruses	echoviruses; Enteric Cytopathic Human Orphan virus
1	Ehrlichia ruminantium (Heartwater)	<i>Ehrlichia ruminantium</i> ; Heartwater; <i>Cowdria ruminatum</i> ; cowdriosis
6	Ehrlichiosis	Ehrlichiosis; Human monocytic ehrlichiosis; HME; Human granulocytic ehrlichiosis; HGE; Human granulocytic anaplasmosis; HGA
9	Emerald Ash Borer	Emerald Ash Borer; <i>Agrilus planipennis</i>
5	<i>Enterobacteriaceae</i> (carbapenem-resistant)	<i>Enterobacteriaceae</i> ; carbapenem-resistant; gram negative bacteria
3	enteroviruses	enteroviruses
7	Enzootic abortion of ewes (Ovine psittacosis, <i>Chlamydia psittaci</i>)	Enzootic abortion of ewes; Ovine psittacosis; <i>Chlamydia psittaci</i> ; Enzootic abortion of ewes
7	Enzootic bovine leukosis (BLV)	Enzootic bovine leukosis; BLV
7	Epizootic hematopoietic necrosis	Epizootic hematopoietic necrosis; EHN; EHNV
7	Epizootic hemorrhagic disease (EHD)	Epizootic hemorrhagic disease; EHD
7	Epizootic ulcerative syndrome	Epizootic ulcerative syndrome; EUS
7	Equine herpesvirus myeloencephalopathy (EHV1 - EHM)	Equine herpesvirus myeloencephalopathy; EHV1; EHV1-EHM
7	Equine infectious anemia	Equine infectious anemia; EIA; EIAV
7	Equine influenza (Virus Type A)	Equine influenza (Virus Type A); horse flu
7	Equine piroplasmosis (Babesiosis, <i>Babesia [Piroplasma] equi</i> , <i>B. caballi</i>)	Equine piroplasmosis (Babesiosis, <i>Babesia [Piroplasma] equi</i> , <i>Babesia caballi</i>); Equine piroplasmosis; EP

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
7	Equine rhinopneumonitis (EHV 1 and 4)	Equine rhinopneumonitis; Equine viral rhinopneumonitis; EHV1; EHV4; Equine abortion virus
7	Equine viral arteritis (EVA)	Equine viral arteritis; EVA
9	Erythrina Gall Wasp	Erythrina Gall Wasp; <i>Quadrastichus erythrinae</i>
9	Eurasian Nun Moth	Eurasian Nun Moth; <i>Lymantria monacha</i>
9	European Gypsy Moth	European Gypsy Moth; <i>Lymantria dispar</i> ; EGM
9	European Larch Canker	European Larch Canker; <i>Lachnellula (Dasyscypha) willkommii</i> (Hartig)
9	European Oak Bark Beetle	European Oak Bark Beetle; <i>Scolytus intricatus</i>
9	European Spruce Bark Beetle	European Spruce Bark Beetle; <i>Ips typographus</i> L.
1, 7	Foot and mouth disease virus	Foot-and-mouth disease virus; <i>Aphthae epizooticae</i> ; FMDV; FMD; FMD serotype
7	Fowl cholera (<i>Pasteurella multocida</i>)	Fowl cholera; <i>Pasteurella multocida</i> ; Pasteurellosis
7	Fowl typhoid (<i>Salmonella gallinarum</i>)	Fowl typhoid; <i>Salmonella gallinarum</i>
1, 4, 6, 7	<i>Francisella tularensis</i> *	<i>Francisella tularensis</i> ; tularemia; Pahvant Valley plague; rabbit fever; deer fly fever; Ohara's fever
9	Pine Pitch Canker	Pine Pitch Canker; <i>Fusarium circinatum</i>
9	<i>Fusarium</i> fungus, unnamed	<i>Fusarium</i> fungus; <i>Fusarium</i> spp.
8	<i>Geosmithia</i> sp.	<i>Geosmithia</i> sp.; Thousand cankers disease; <i>Geosmithia morbida</i>
2, 3, 6	<i>Giardia</i>	<i>Giardia</i> ; <i>Giardia intestinalis</i> ; <i>Giardia lamblia</i> ; <i>Giardia duodenali</i> ; Giardiasis
1, 7	Goat pox virus	Goat pox virus; goatpox; goat pox; <i>Variola caprina</i>
9	Golden Haired Pine Bark Beetle	Golden Haired Pine Bark Beetle; <i>Hylurgus ligniperda</i> ; Golden-haired bark beetle
9	Goldspotted oak borer	Goldspotted oak borer; <i>Agrilus auroguttatus</i> ; GSOB
6	Gonorrhea	Gonorrhea; the clap; <i>Neisseria gonorrhoeae</i>
7	Gyrodactylosis	Gyrodactylosis; <i>Gyrodactylus salaris</i>
6	<i>Haemophilus influenzae</i> , invasive disease	<i>Haemophilus influenzae</i> ; invasive disease; <i>Haemophilus influenzae</i> type B; Hib; Pfeiffer's bacillus; <i>Bacillus influenzae</i>

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
6	Hansen's disease (leprosy)	Hansen's disease; leprosy
4, 6	Hantavirus pulmonary syndrome	<i>Hantavirus</i> pulmonary syndrome; <i>Hantavirus</i> ; HPS
8	<i>Harpophora maydis</i>	<i>Harpophora maydis</i> ; late wilt of maize
9	Harrisia cactus mealybug	Harrisia cactus mealybug; <i>Hypogeococcus pungens</i>
7	Heartwater (<i>Cowdria ruminantium</i>)	Heartwater; <i>Cowdria ruminantium</i> ; <i>Cowdria ruminatum</i> ; <i>Ehrlichia ruminantium</i> ; Cowdriosis; nintas; ehrlichiosis
9	Hemlock Woolly Adelgid	Hemlock Woolly Adelgid; <i>Adelges tsugae</i>
6	Hemolytic uremic syndrome, post-diarrheal	Hemolytic uremic syndrome, post-diarrheal; Hemolytic uremic syndrome; HUS
7	Hemorrhagic septicemia (<i>Pasteurella multocida</i> , serotypes B/Asian or E/African)	Hemorrhagic septicemia; <i>Pasteurella multocida</i> serotype B/Asian; <i>Pasteurella multocida</i> serotype E/African; Shipping Fever; <i>Pasteurella multocida</i> ; HS
1	Hendra virus	Hendra virus; Equine Morbilivirus; <i>Henipavirus</i> ; HeV
2, 6	Hepatitis A	Hepatitis A; HAV; hep A
5, 6	Hepatitis B (acute, chronic or perinatal)	Hepatitis B; HBV; hep B
5, 6	Hepatitis C	Hepatitis C; HCV; hep C
8	<i>Heterodera</i> spp. and <i>Meloidogyne</i> spp.	<i>Heterodera</i> spp.; <i>Meloidogyne</i> spp.; cyst and root-knot nematodes
7	Highly pathogenic avian influenza	Highly pathogenic avian influenza; HPAI
5, 6	Human Immunodeficiency Virus (HIV)	Human Immunodeficiency Virus; HIV; AIDS
7	Infection with abalone herpes-like virus	Infection with abalone herpes-like virus; AbHV; abalone viral ganglioneuritis; AVG
7	Infection with <i>Bonamia exitiosa</i>	<i>Bonamia exitiosa</i>
7	Infection with <i>Bonamia ostreae</i>	<i>Bonamia ostreae</i> ; bonamiosis
7	Infection with <i>Marteilia refringens</i>	<i>Marteilia refringens</i> ; Aber disease; marteiliosis
7	Infection with <i>Perkinsus marinus</i>	<i>Perkinsus marinus</i> ; Dermo; Perkinsosis
7	Infection with <i>Perkinsus olseni</i>	<i>Perkinsus olseni</i> ; <i>Perkinsus atlanticus</i>
7	Infection with <i>Xenohalictis californiensis</i>	<i>Xenohalictis californiensis</i> ; withering syndrome; abalone rickettsiosis

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
7	Infectious bovine rhinotracheitis/infectious pustular vulvovaginitis (IBR/IPV)	Infectious bovine rhinotracheitis/infectious pustular vulvovaginitis; IBR/IPV; Bovine herpesvirus 1; bovine rhinotracheitis; infectious pustular vulvovaginitis
7	Infectious bursal disease (Gumboro disease)	Infectious bursal disease; Gumboro disease; IMB; IBDV
7	Infectious hematopoietic necrosis	Infectious hematopoietic necrosis; IHN; Infectious hematopoietic necrosis virus; IHNV
7	Infectious hypodermal and haematopoietic necrosis	Infectious hypodermal and haematopoietic necrosis; IHHN; Infectious hypodermal and haematopoietic necrosis virus; IHHNV
7	Infectious myonecrosis	Infectious myonecrosis; IMN; IMNV; Infectious myonecrosis virus
7	Infectious salmon anemia	Infectious salmon anemia; ISA; ISAV; European genotype (genotype I); North American genotype (genotype II)
5	Influenza	Influenza; flu; H1N1; H3N2; Influenza A; Influenza B; Influenza C
6	Influenza-associated mortality, pediatric	Influenza-associated mortality, pediatric
1, 7	Japanese encephalitis virus	Japanese encephalitis virus; JEV; Japanese encephalitis; JE
5	<i>Klebsiella</i>	<i>Klebsiella</i> ; <i>Klebsiella pneumoniae</i>
7	Koi herpesvirus disease	Koi herpesvirus disease; Koi herpes virus; KHV; cyprinid herpesvirus-3; CyHV-3
9	Larch Casebearer	Larch Casebearer; <i>Coleophora laricella</i>
9	Larger Pine Shoot Beetle	Larger Pine Shoot Beetle; common pine shoot beetle; <i>Tomicus piniperda</i>
1	Lassa fever virus	Lassa fever virus; Lassa fever; Lassa viral hemorrhagic fever
8, 9	Laurel wilt	Laurel wilt; <i>Raffaelea lauricola</i> ; Laurel wilt of avocado; Laurel wilt of redbay
6	Lead, exposure screening test result	Lead, exposure screening test result
3, 6	Legionellosis / <i>Legionella</i>	Legionellosis; Legionella; <i>Legionella pneumophila</i> ; Legionnaire's disease; Pontiac fever

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
7	Leptospirosis	Leptospirosis; <i>Leptospira</i> ; Weil disease; Icterohemorrhagic fever; Swineherd's disease; Rice-field fever; Cane-cutter fever; Swamp fever; Mud fever; Hemorrhagic jaundice; Stuttgart disease; Canicola fever
6	Listeriosis	Listeriosis; <i>Listeria monocytogenes</i>
9	Lobate Lac Scale	Lobate Lac Scale; <i>Paratachardina pseudolobata</i>
7	Low pathogenic avian influenza (H5 or H7 subtypes)	Low pathogenic avian influenza; Avian influenza H5; Avian influenza H7; LPAI
1	Lujo virus	Lujo virus; Lujo hemorrhagic fever; Lujo fever
1, 7	Lumpy skin disease virus	Lumpy skin disease virus; Lumpy skin disease; LSDV
6	Lyme disease	Lyme disease; <i>Borrelia burgdorferi</i>
7	Maedi-visna / ovine progressive pneumonia	Maedi-visna/ovine progressive pneumonia; ovine progressive pneumonia; progressive pneumonia; Maedi; Zwoegersiekte; La bouhite; Graaff-Reinet disease; Marsh's progressive pneumonia
8	<i>Magnaporthe grisea</i>	<i>Magnaporthe grisea</i> ; Wheat blast; rice blast; rice rotten neck; rice seedling blight; blast of rice; oval leaf spot of graminea; pitting disease; ryegrass blast; Johnson spot; blast disease, blight disease
6	Malaria	Malaria; <i>Plasmodium falciparum</i> ; <i>Plasmodium vivax</i> ; <i>Plasmodium malariae</i> ; <i>Plasmodium ovale</i> ; <i>Plasmodium knowlesi</i>
1, 7	Malignant catarrhal fever virus (Alcelaphine herpesvirus type 1)	Malignant catarrhal fever virus (Alcelaphine herpesvirus type 1); Malignant catarrhal fever virus; Alcelaphine herpes virus 1; AIHV-1; Ovine herpes virus 2; OHV-2; malignant catarrhal fever; bovine malignant catarrhal fever; BMCF
1	Marburg virus	Marburg virus; Lake Victoria Marburg virus; Marburg hemorrhagic fever; MHF
7	Marek's Disease	Marek's Disease; Gallid herpesvirus 2; GaHV-2; Marek's disease virus; MDV
6	Measles	Measles; measles virus
9	Mediterranean Pine Engraver Beetle	Mediterranean Pine Engraver Beetle; <i>Orthotomicus erosus</i>

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
4	Melioidosis	Melioidosis; <i>Burkholderia pseudomallei</i> ; <i>Pseudomonas pseudomallei</i>
1	Menangle virus	Menangle virus
6	Meningococcal disease (<i>Neisseria meningitides</i>)	Meningococcal disease; <i>Neisseria meningitides</i> ; bacterial meningitis; meningococemia
5	Methicillin-resistant <i>Staphylococcus aureus</i>	Methicillin-resistant <i>Staphylococcus aureus</i> ; <i>Staphylococcus aureus</i> ; MRSA
1	Monkeypox virus	Monkeypox virus; monkeypox
6	Mumps	Mumps; mumps virus; epidemic parotitis
5	<i>Mycobacterium abscessus</i>	<i>Mycobacterium abscessus</i>
1	<i>Mycoplasma capricolum</i>	<i>Mycoplasma capricolum</i> ; <i>Mycoplasma capricolum</i> subspecies <i>capripneumoniae</i> ; contagious caprine pleuropneumonia, mastitis and severe arthritis
1	<i>Mycoplasma mycoides</i>	<i>Mycoplasma mycoides</i>
1	<i>Mycoplasma mycoides</i> subspecies <i>mycoides</i> small colony (<i>Mmm</i> SC) (contagious bovine pleuropneumonia)	<i>Mycoplasma mycoides</i> subspecies <i>mycoides</i> small colony (<i>Mmm</i> SC) (contagious bovine pleuropneumonia); contagious bovine pleuropneumonia
7	Mycoplasmosis (<i>M. gallisepticum</i>)	Mycoplasmosis; <i>Mycoplasma gallisepticum</i> ; mycoplasmosis
7	Mycoplasmosis (<i>M. synoviae</i>)	Mycoplasmosis; <i>Mycoplasma synoviae</i> ; mycoplasmosis
7	Nairobi sheep disease	Nairobi sheep disease; Nairobi sheep disease virus; NSDV
7	Necrotizing hepatopancreatitis	Necrotizing hepatopancreatitis; Necrotising hepatopancreatitis; NHP; Texas pond mortality syndrome; TPMS; Texas necrotizing hepatopancreatitis; TNHP; Peru necrotizing hepatopancreatitis; PNHP
1, 7	Newcastle disease virus	Newcastle disease virus; Virulent Newcastle disease virus 1; Newcastle disease virus; NDV; Newcastle disease; velogenic Newcastle disease; mesogenic Newcastle disease

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1, 4, 7	Nipah virus	Nipah virus; Nipah virus encephalitis
5	Norovirus	Norovirus; stomach flu; winter vomiting bug; Norwalk-like virus
6	Novel Influenza A virus infection	Novel Influenza A virus infection; influenza
9	Oak Dieback	Oak Dieback; <i>Phytophthora quercina</i>
9	Ohi'a Rust	Ohi'a Rust; <i>Puccinia psidii</i> Winter; <i>Puccinia</i> rust
7	Paratuberculosis	Paratuberculosis; <i>Mycobacterium avium</i> paratuberculosis; MAP; Johne's disease
1, 8	<i>Peronosclerospora philippinensis</i> (<i>Peronosclerospora sacchari</i>)	<i>Peronosclerospora philippinensis</i> ; <i>Peronosclerospora sacchari</i> ; Phillipine downy mildew; Philippine downy mildew of maize; PDM; Brown Stripe downy mildew of corn
6	Pertussis	Pertussis; <i>Bordetella pertussis</i> ; <i>Bordetella parapertussis</i> ; Whooping cough
1, 7	Peste des petits ruminants	Peste des petits ruminants; Peste des petits ruminants virus; PPR; ovine rinderpest; PPRV
1, 8	<i>Phoma glycinicola</i> (formerly <i>Pyrenochaeta glycines</i>)	<i>Phoma glycinicola</i> ; Red leaf blotch of soybean; <i>Pyrenochaeta</i> leaf spot; <i>Dactuliophora</i> leaf spot; <i>Pyrenochaeta</i> leaf blotch; RLB; <i>Pyrenochaeta glycines</i> ; <i>Dactuliophora glycines</i> ; <i>Dactuliochaeta glycines</i> of soybean
8, 9	<i>Phytophthora kernoviae</i>	<i>Phytophthora kernoviae</i> ; Ramorum dieback; sudden oak death
9	<i>Phytophthora</i> Root Rot	<i>Phytophthora</i> Root Rot; <i>Phytophthora cinnamomi</i>
9	Pine Flat Bug	Pine Flat Bug; <i>Aradus cinnamomeus</i> Panzer
8	plum pox virus (PPV)	plum pox virus; PPV; plum pox; Sharka; Sharka virus
3, 6	Poliovirus infection	Poliovirus infection; polio; poliomyelitis; paralytic poliomyelitis; poliovirus; PV
9	Polyphagous shot hole borer	Polyphagous shot hole borer; <i>Euwallacea</i> sp.; <i>Fusarium</i> fungus

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
7	Porcine reproductive and respiratory syndrome (PRRSV)	Porcine reproductive and respiratory syndrome (PRRSV); PRRS virus; PRRSV; porcine arterivirus; Blue ear pig disease; mystery swine disease
9	Port-Orford-Cedar Root Disease	Port-Orford-Cedar Root Disease; <i>Phytophthora lateralis</i>
4, 6	Psittacosis	Psittacosis; <i>Chlamydia psittaci</i> ; parrot fever
8	<i>Puccinia graminis</i> f. sp. <i>tritici</i>	<i>Puccinia graminis</i> f. sp. <i>tritici</i> ; Stem rust of wheat (Ug99); black rust
7	Pullorum disease (<i>Salmonella pullorum</i>)	Pullorum disease; Bacillary White Diarrhoea; Pullorum disease; <i>Salmonella pullorum</i> ; Salmonella
6, 7	Rabies	Rabies; Rabies virus; Lyssavirus; Ephemerovirus; Vesiculovirus; Australian bat virus; Mokola virus; Duvenhage virus; European bat virus 1; European bat virus 2; Lagos bat virus
1, 8	<i>Ralstonia solanacearum</i> race 3, biovar 2	<i>Ralstonia solanacearum</i> race 3, biovar 2; <i>Ralstonia</i> bacterial wilt of potato; <i>Ralstonia</i> bacterial wilt of tomato; <i>Ralstonia</i> bacterial wilt of geranium; Brown rot (potato); southern bacterial wilt (tomato); Moko disease (banana); Granville wilt (tobacco); Southern Wilt of Geranium
1, 8	<i>Rathayibacter toxicus</i>	<i>Rathayibacter toxicus</i> ; <i>Clavibacter toxicus</i> ; <i>Rathayibacter</i> poisoning; Annual ryegrass toxicity; ARGT; Yellow slime disease; Rathay's disease; annual ryegrass toxicosis; annual ryegrass staggers; parasitized annual ryegrass; ryegrass toxicity; toxic annual ryegrass; tunicamycin poisoning; wimmera ryegrass toxicity; Black Springs syndrome; flood plain staggers; blown grass/beard grass poisoning; corynetoxin poisoning; corynetoxicosis; Stewarts range syndrome; tunicaminyluracil toxicosis; veldtgrass staggers; bacterial galls; seed gall nematode
1	Reconstructed replication competent forms of the 1918 pandemic influenza virus containing any portion of the coding regions of all eight gene segments (Reconstructed1918 Influenza virus)	Reconstructed replication competent forms of the 1918 pandemic influenza virus containing any portion of the coding regions of all eight gene segments (Reconstructed1918 Influenza virus); 1918-like HINI; Spanish Flu; 1918 influenza; reconstructed 1918 influenza virus

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
7	Red sea bream iridoviral disease (RSIVD)	Red sea bream iridoviral disease; RSIVD; Red sea beam iridovirus
9	Red-Haired Pine Bark Beetle	Red-Haired Pine Bark Beetle; <i>Hylurgus ligniperda Fabricius</i>
1, 4	Ricin	Ricin; Castor bean; <i>Ricinus communis</i>
1, 6	Rickettsiosis, spotted fever	Rickettsiosis, spotted fever; <i>Rickettsia rickettsii</i> ; <i>Rickettsia australis</i> ; <i>Rickettsia</i> ; <i>Rickettsia honei</i> ; <i>Rickettsia japonica</i> ; <i>Rickettsia felis</i> ; Rickettsial spotted fever; Rocky mountain spotted fever; RMSF; Australian tick typhus; Tobia fever; Sao Paulo fever; North Asian tick typhus; Flinders Island spotted fever; Japanese spotted fever
1, 7	Rift Valley fever virus	Rift Valley fever virus; Rift Valley fever; RVF
1, 7	Rinderpest virus	Rinderpest virus; Rinderpest; cattle plague; steppe murrain; RPV
6	Rubella	Rubella; Rubella virus; German measles; congenital rubella syndrome
2	<i>Salmonella</i> serotype Typhimurium	<i>Salmonella</i> serotype Typhimurium; <i>Salmonella</i> Typhimurium; Salmonellosis
6	Salmonellosis	Salmonellosis; <i>Salmonella</i> spp; <i>Salmonella enterica</i>
7	Salmonellosis (<i>Salmonella abortus ovis</i>)	Salmonellosis (<i>Salmonella abortus ovis</i>); <i>Salmonella abortus</i>
1, 6	SARS-associated coronavirus	SARS-associated coronavirus (SARS-CoV); SARS-associated coronavirus; SARS virus; SARS-CoV; SARS; Severe Acute Respiratory Syndrome
1	Saxitoxin	Saxitoxin; saxitoxin; STX; neosaxitoxin; neoSTX; gonyautoxins; GTX; decarbamoylsaxitoxin; dcSTX; paralytic shellfish poison; PSP
1, 8	<i>Sclerophthora rayssiae</i> var <i>zeae</i>	<i>Sclerophthora rayssiae</i> var <i>zeae</i> ; Brown stripe downy mildew; brown stripe downy mildew of maize; brown stripe downy mildew of corn
7	Scrapie	Scrapie; Scrapie prion
7	Screw worm, New World	Screw worm, New World; <i>Cochliomyia hominivorax</i> ; New World screw worm fly; screw worm; myiasis
7	Screw worm, Old world	Screw worm, Old world; <i>Chrysomia bezziana</i> ; screw worm; myiasis

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1, 7	Sheep pox virus	Sheep pox virus; Sheeppox virus; variola ovina; sheeppox; sheep pox
1	Shigatoxin	Shigatoxin; Stx; <i>Shiga</i> -like ribosome inactivating proteins; SLT-1; SLT-2; Stx-1; Stx-2
1, 2, 4, 6	Shigellosis	Shigellosis; <i>Shigella</i> spp.; <i>Shigella flexneri</i> ; <i>Shigella dysenteriae</i> ; <i>Shigella boydii</i> ; <i>Shigella sonnei</i> ; bacillary dysentery; Marlow syndrome
9	Sirex Woodwasp	Sirex Woodwasp; <i>Sirex noctilio</i>
9	Soapberry Borer	Soapberry Borer; <i>Agrilus prionurus</i>
7	Spring viremia of carp	Spring viremia of carp; spring viremia of carp virus; SVCV; Infectious Dropsy of Carp; Infectious Ascites; Hydrops; Red Contagious Disease; Rubella; Hemorrhagic Septicemia
9	Spruce Aphid	Spruce Aphid; <i>Elatobium abietinum</i>
1, 4	Staphylococcal enterotoxins	Staphylococcal enterotoxins; <i>Staphylococcus aureus</i> enterotoxin; Staphylococcal enterotoxins; Staphylococcal enterotoxin B; SEB; Staphylococcal enterotoxins A,B,C,D,E subtypes; staphyloenterotoxigenesis; staphyloenterotoxemia; staphylococcal food poisoning
5, 6	<i>Staphylococcus aureus</i> infection	<i>Staphylococcus aureus</i> infection; staph; <i>staph aureus</i> ; <i>Staphylococcus aureus</i>
6	Streptococcal toxic shock syndrome	Streptococcal toxic shock syndrome (STSS); streptococcal toxic shock syndrome; STSS; toxic shock-like syndrome; TSLS
6	Streptococcus pneumonia, invasive disease	<i>Streptococcus pneumonia</i> , invasive disease (IPD); <i>Streptococcus pneumoniae</i>
9	Sudden Oak Death Syndrome	Sudden Oak Death Syndrome; <i>Phytophthora ramorum</i>
1, 7	Swine vesicular disease	Swine vesicular disease; Swine vesicular disease virus; SVD
1, 8	<i>Synchytrium endobioticum</i>	<i>Synchytrium endobioticum</i> ; Potato wart disease; black scab
6	Syphilis	Syphilis; <i>Treponema pallidum</i> subspecies <i>pallidum</i> ; chancre

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1	T-2 toxin	T-2 toxin; alimentary toxic aleukia; fusariotoxin T2; insariotoxin; mycotoxin T-2; NSC 138780; T-2 mycotoxin
7	Taura syndrome	Taura syndrome; Taura syndrome virus; TSV
6	Tetanus	Tetanus; <i>Clostridium tetani</i> ; lockjaw
1	Tetrodotoxin	Tetrodotoxin; Tetrodotoxin; tetrodox; TTX; anhydrotetrodotoxin; 4-epitetrodotoxin; tetrodonic acid; pufferfish poisoning
7	Theileriasis (<i>Theileria annulata</i> , <i>T. parva</i>)	Theileriasis; <i>Theileria annulata</i> ; <i>Theileria parva</i> ; theileriasis; theileria
9	Thousand Canker Disease	Thousand Canker Disease; <i>Geosmithia morbida</i>
6	Toxic shock syndrome (non-Strep)	Toxic shock syndrome (non-Strep); toxic shock syndrome; TSS
7	Transmissible gastroenteritis	Transmissible gastroenteritis; Transmissible gastroenteritis virus; TGEV; transmissible gastroenteritis; TGE
6, 7	Trichinellosis	Trichinellosis; <i>Trichinella spiralis</i>
7	<i>Tritrichomonas foetus</i>	<i>Tritrichomonas foetus</i>
7	Trypanosomiasis	Trypanosomiasis; <i>Trypanosoma congolense</i> , <i>Trypanosoma vivax</i> ; <i>Trypanosomabrucei brucei</i> ; <i>Trypanosoma evansi</i> ; Surra
5, 6	Tuberculosis	Tuberculosis; TB; <i>Mycobacterium tuberculosis</i> ; <i>Mycobacterium avium</i> ; <i>Mycobacterium bovis</i> ; consumption
7	Turkey rhinotracheitis	Turkey rhinotracheitis; avian metapneumovirus; aMPV
6	Typhoid Fever	Typhoid Fever; <i>Salmonella typhi</i> ; gastric fever; abdominal typhus; infantile remittant fever; slow fever; nervous fever; pythogenic fever
1, 4	Typhus fever (<i>Rickettsia prowazekii</i>)	Typhus fever; <i>Rickettsia prowazekii</i> ; typhus; epidemic typhus; camp fever; jail fever; hospital fever; ship fever; famine fever; putrid fever; petechial fever; epidemic louse-borne typhus; louse-borne typhus
5	Vancomycin-intermediate <i>Staphylococcus aureus</i>	Vancomycin-intermediate <i>Staphylococcus aureus</i> ; <i>Staphylococcus aureus</i> ; VISA
5	Vancomycin-resistant <i>Enterococci</i>	Vancomycin-resistant <i>Enterococci</i> ; <i>Enterococci</i> spp; VRE

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
5	Vancomycin-resistant <i>Staphylococcus aureus</i>	Vancomycin-resistant <i>Staphylococcus aureus</i> ; <i>Staphylococcus aureus</i> ; VRSA
6	Varicella	Varicella; <i>Varicella zoster</i> ; Chicken pox virus; varicella zoster virus; zoster virus; human herpes virus type 3; chickenpox
1, 4, 6	Variola major virus (Smallpox virus)	Variola major virus (Smallpox virus); smallpox; <i>Variola major</i> ; Variola virus; VAR; Variola vera
1	Variola minor virus (Alastrim)	Variola minor virus (Alastrim); <i>Variola alastrim</i> ; <i>Variola minor</i> ; alastrim; white pox; kaffir pox; Cuban itch; West Indian pox; milk pox; pseudovariola
1	Venezuelan Equine Encephalitis virus	Venezuelan Equine Encephalitis virus; Venezuelan equine encephalitis; VEE; Venezuelan equine encephalomyelitis virus
1, 7	Vesicular stomatitis virus (exotic): Indiana subtypes VSV-IN2, VSV-IN3	Vesicular stomatitis virus (exotic): Indiana subtypes VSV-IN2, VSV-IN3; Vesicular stomatitis; Vesicular stomatitis virus; VSV; Indiana subtype VSV-IN2; Indiana subtype VSV-IN3; vesicular stomatitis Indiana virus; VSIV
6	Vibriosis	Vibriosis; <i>Vibrio vulnificus</i> ; <i>Vibrio parahaemolyticus</i>
9	Viburnum leaf beetle	Viburnum leaf beetle; <i>Pyrrhalta viburni</i>
7	Viral hemorrhagic septicemia	Viral hemorrhagic septicemia (VHS); Viral hemorrhagic septicemia virus; VHSV; Viral hemorrhagic septicemia; VHS
9	Walnut Twig Beetle	Walnut Twig Beetle; <i>Pityophthorus juglandis</i> ; Thousand Cankers disease
9	White Pine Blister Rust	White Pine Blister Rust; <i>Cronartium ribicola</i>
7	White spot disease	White spot disease; <i>Ichthyophthirius multifiliis</i> ; Ich; Ick
7	White tail disease	White tail disease; <i>Macrobrachium rosenbergii</i> Nodavirus; MrNV; extra small virus; XSV
1, 8	<i>Xanthomonas oryzae</i> pv. <i>oryzae</i> and <i>X. oryzae</i> pv. <i>oryzicola</i>	<i>Xanthomonas oryzae</i> pv. <i>oryzae</i> and <i>X. oryzae</i> pv. <i>Oryzicola</i> ; <i>Xanthomonas oryzae</i> ; Rice blight disease; Bacterial leaf blight; BLB; Kresek disease; Bacterial leaf streak; BLS

De-duplicated List of Agents, Toxins, and Pathogens of Interest

List(s) Source	Disease or Agent Name	Synonyms, Include Disease/Agent Specified (; separated)
1, 8	<i>Xylella fastidiosa</i> (citrus variegated chlorosis strain)	<i>Xylella fastidiosa</i> (citrus variegated chlorosis strain); Pierce's disease; California vine disease; Anaheim disease (grapevine); leaf scorch (almond); dwarf (lucerne); phony disease (peach); leaf scald (plum); leaf scorch (elm, oak, plane, mulberry, maple); variegated chlorosis (citrus)
6	Yellow Fever	Yellow Fever; Yellow fever virus; Yellow Jack; Bronze John
7	Yellowhead disease	Yellowhead disease; Yellowhead virus
1, 4, 6	<i>Yersinia pestis</i>	<i>Yersinia pestis</i> ; Plague bacteria; Plague; bubonic plague; pneumonic plague; Black Death

Appendix D

Prioritized List of Agents for BioCat¹

The recommendation for a prioritized list of biological agents is shown in the table below. The list is based on the Tier 1 agents and toxins identified from the “HHS and USDA Select Agents and Toxins List.”² Tier 1 agents are thought to be at particularly high risk for deliberate misuse.³ Five additional agents are also included: pandemic influenza, endosulfan, chlorpyrifos, *Ralstonia solanacearum*, and citrus greening disease. Influenza is included due to its known potential impacts on human health as well as infrastructure. Endosulfan and chlorpyrifos are highly toxic chemical compounds that have a history of use as pesticides; food and/or water contamination are potential concerns. Two plant diseases were added as well to ensure coverage for this important NBIS domain.

The agents listed below include the human, animal, plant, food, and environment (HAPFE) categories that NBIC is charged with monitoring. Because of significant differences in the means of exposure, the types of models required can vary greatly between agents, and it is acknowledged that there may be gaps in the available model coverage. This list is not meant to be comprehensive but to provide a range of high-priority agents and toxins suitable for analysis using BioCat models.

Table D.1. Prioritized Agents of Interest for BioCat Analysis

Agent	NBIS Domain Affected
<i>Bacillus anthracis</i>	Human, Animal, Environment
Botulinum toxin	Food
<i>Burkholderia mallei</i>	Human, Animal
<i>Burkholderia pseudomallei</i>	Human, Animal
Chlorpyrifos (Dursban)	Environment, Food
Citrus Greening, Huanglongbing (HLB) of citrus, <i>Candidatus Liberibacter americanus</i>	Plants
Endosulfan	Environment, Food
Foot and Mouth Disease (FMD)	Animal
<i>Francisella tularensis</i>	Human, Animal
Influenza (pandemic)	Human, Animal
<i>Ralstonia solanacearum</i>	Plants
Rinderpest virus	Animal
SARS	Human
Variola (major and minor)	Human
Viral hemorrhagic fevers including Ebola and Marburg	Human, Animal
<i>Yersinia pestis</i>	Human, Animal

¹ List received from the client via email on April 10, 2013.

² <http://www.selectagents.gov/select%20agents%20and%20toxins%20list.html>

³ <http://www.asm.org/index.php/public-policy-2/97-policy/documents/8417-sa-10-9-12>

Appendix E

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Tier (1-3)	Term	Definition	Source	Staffing	Notes
1	Event-type I	First-pass identification of the disease, agent and vector of interest in the article.		Tier 1 data collection will be conducted by a research associate.	
1	Disease Specific	The causative agent of the event being modeled is disease specific.			
1	Agent/Pathogen/Toxin/Pest	The agent(s)/pathogen(s)/toxin(s)/pest(s) of the event being modeled.	Client provided lists		
1	Vector	An invertebrate that transmits a pathogen from reservoir to host.			
1	Operational Aspects I	First pass to identify characteristics of the model being evaluated.			
1	Event Prediction	The model predicts an event/outbreak is expected to occur in the future (left - before an event).			
1	Event Forecast	The model makes estimations about an event/outbreak occurring at some specified future date (the index case exists, what is the projected - estimated spread/trajectory/R0/infectivity/transmissibility).			
1	Technological Maturity	The level of maturity of a particular model in comparison to others.			
1	Verified or Validated	First-pass identification of whether or not the model has been verified and/or validated (Does the article mention any type of validation or verification?).			
1	Model Identification I	First-pass identification of the Model discussed in the article.			

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Tier (1-3)	Term	Definition	Source	Staffing	Notes
1	Model Name	The name of the model (e.g., EpiCast, FLuTe, IT-CI, ...).			
2	Model Identification II	Second pass and more detailed information extraction of the Model discussed in the article.		Tier 2 data collection will be conducted by a post-doc, advanced undergrad/Master's level intern or S&E II/III.	
2	Owner	The "owner" of the model used in the article. This can be a person, university, corporation or organization.			
2	Website	The website where additional information about the model can be found.			
2	Corresponding Author	The article's corresponding author as identified in the full-text document.			
2	Corresponding Author Phone Number	The corresponding author's phone number as identified in the full-text document.			
2	Corresponding Author Email Address	The corresponding author's email address as identified in the full-text document.			
2	Model Creation Year	The year the model was created.			
2	Revision	Is this a modification of the original model (same owner or different).			
2	Revision Date	The date the model was revised/modified.			
2	This Model Rocks	This is a subjective opinion of the PNNL SME on whether this model is "cream of the crop" or "pit of the pot."			
2	Event-type II	Second-pass identification of the affected population detailed in the article.			
2	Population (Species) Affected Identified	The specific affected population/species identified and modeled in the article.			
2	Technological Maturity	The level of maturity of a particular model in comparison to others.	Metrics for Evaluation draft report (2010)		

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Tier (1-3)	Term	Definition	Source	Staffing	Notes
2	Type of Validation and Verification	The method used to verify and/or validate the model's maturity.			
2	In Use in an Operational Environment	The model is or is not in use in an operational environment.			
3	Operational Aspects III	Third pass to identify characteristics of the model being evaluated.		Tier 3 data collection will be conducted by an advanced post-doc/S&E III/IV	
3	Spatial	Models which forecast the geographic spread of a disease after it occurs based on the relationship between the outbreak and primary geospatial factors.	PLoS ONE Manuscript		
3	Ecological Niche*	Computational models/algorithms to predict the distribution of species in geographic space on the basis of a mathematical representation of their known distribution in environmental space.	Wikipedia		* Also called environmental niche modeling, species distribution modeling, predictive habitat distribution modeling, and climate envelope modeling.
3	Event-Based	Models which attempt to identify outbreaks either through sentinel groups or through the collection of real-time diagnostic, clinical, or syndromic data.	PLoS ONE Manuscript		
3	Dynamical	Models which examine how a specific disease moves through a population.	PLoS ONE Manuscript		
3	Statistical Models	Models which are a formalization of relationships between variables in the form of mathematical equations.	Wikipedia		

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Tier (1-3)	Term	Definition	Source	Staffing	Notes
3	Agent Based Model	Computational models for simulating the actions and interactions of autonomous agents (both individual or collective entities such as organizations or groups) with a view to assessing their effects on the system as a whole.	Wikipedia		
3	Accessibility	The degree to which and mechanism by which the model is made available, e.g., plug-and-play, downloadable, etc.	Corley et al. (2012) ¹ and modified as needed		
3	Run Time, Complexity, Capabilities Required	Characteristics required for executing the model effectively.			
3	Geographic and Population Coverage	Geographic area and population size of the organism(s) being modeled.			
3	Input Geography	Geographic area attributes used as model input.			
3	Output Geography	Geographic area attributes generated as model output.			
3	Local	A geographic area bounded by city or county limits.	Based on LANL's work and modified as needed		
3	Provincial/State	A geographic area within the political boundaries of one government, such as a province or state.	Based on LANL's work and modified as needed		
3	Regional	A geographic area defined by similar features and location, such as the Columbia Basin or Latin America.	Based on LANL's work and modified as needed		
3	National	A geographic area composed of a country.	Based on LANL's work and modified as needed		

¹ Courtney D. Corley, Mary J. Lancaster, et al. Assessing the Continuum of Event-Based Biosurveillance Through an Operational Lens. Biosecurity and Bioterrorism: Biodefense Strategy, Practice, and Science. March 2012. doi:10.1089/bsp.2011.0096. PNNL-SA-77400

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Tier (1-3)	Term	Definition	Source	Staffing	Notes
3	Global	A geographic area defined as the entire world/international in scope.	Based on LANL's work and modified as needed		
3	Vector Population Size	The size of the vector population.			
3	Modeled Species Population Size	The population size of the species impacted.			
3	Data Input	Data input is a general characterization of the type of model used to produce results.	PLoS ONE and Corley et al. (2012)		
3	Epidemiological Data from the Same Location	Models which take into account epidemiological characteristics of the disease from the same location.	PLoS ONE manuscript		
3	Epidemiological Data from Difference Location	Models which take into account epidemiological characteristics of the disease from different same locations.	PLoS ONE manuscript		
3	Governmental and Non-Governmental Organizations	Models which use data acquired from governmental or non-governmental organizations.	PLoS ONE manuscript		
3	Literature	Models which use any form data excluding epidemiological, weather, or population data.	PLoS ONE manuscript and modified as needed		
3	Satellite/Remote Sensing	Models which use data acquired via satellite or remote sensing technologies.	PLoS ONE manuscript		
3	Simulated	Models which are simulated using software or statistical techniques.	PLoS ONE manuscript and modified as needed		
3	Laboratory Diagnostic	Models developed using laboratory diagnostic data.	PLoS ONE manuscript		
3	Expert Opinion	Models based on expert opinion - does not specify from whom or what type of data was used in the modeling.	PLoS ONE manuscript		
3	Output	Output is the information/data/analysis produced after running a model.	Corley et al. (2012) and modified as needed		

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Tier (1-3)	Term	Definition	Source	Staffing	Notes
3	Output Content	Model output may be quantitative or qualitative and may include probabilities, forecasts, digests, counts, warnings, or lists of outlying observations.	Corley et al. (2012) and modified as needed		
3	Sustainability	A measure of how long model output may be considered valid and confident.	Corley et al. (2012) and modified as needed		
3	Cost	Cost represents the overarching financial investment (in USD) of purchasing the model, purchasing equipment to run the model, staffing expertise, and any other expense required to develop or sustain the model.	Corley et al. (2012) and modified as needed		
3	Research, development, testing, and evaluation	The cost (in USD) to perform research, and to construct, prepare, develop, test, and evaluate the model.	Corley et al. (2012) and modified as needed		
3	Operations	The cost to maintain and operate the model. This includes any projected storage, data acquisition, or staffing expense to maintain model performance.	Corley et al. (2012) and modified as needed		
3	Government funded	A tribal, local, regional, national, or international governmental agency funded the development and/or operations of the model.	Corley et al. (2012) and modified as needed		

Appendix F

BioCat Evaluation & Workflow Data Fields

Triage Level	Metric	Submetric	Required	Type of Field	Allowed Value	Notes	
1	Event-type (I)	Disease Specific	Y	Yes or No (Radio / Drop Down)			
		Agent/Pathogen/Toxin/Pest	N	Check Boxes		If Disease Specific is Y, then pop-up Agent selection	
		Vector	N	String Field (Manual); separated			
	Operational Aspect (I)	Event Prediction	Y	Yes or No (Radio / Drop Down)			
		Event Forecast	Y	Yes or No (Radio / Drop Down)			
		Verified or Validated	Y	Yes or No (Radio / Drop Down)			
	Technological Maturity (I)	Model Name	N	String Field (Manual)			
	Model Identification (I)	Second Look Required	N	Yes or No (Radio / Drop Down)		If Y, trigger workflow AND hide this field from public/user view.	
	2	Model Identification (II)	Owner	N	String Field (Manual)		Firstname Lastname

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Triage Level	Metric	Submetric	Required	Type of Field	Allowed Value	Notes
		Website	N	String Field (Manual)		
		Corresponding Author	Y	String Field (Manual)		
		Corresponding Author Phone Number	N	Phone Number		Phone number with area code - US or international
		Corresponding Author Email Address	N	Email Address		Valid email address
		Model Creation Year	N	Date		Only Year is Required
		Revision	N	Yes or No (Radio / Drop Down)		
		Revision Date	N	Date		Only Year is Required
		This Model Rocks	N	Yes or No (Radio / Drop Down)		Hide this field from public/user view.
	Event-type (II)	Population (Species) Affected Identified	Y	Yes or No (Radio / Drop Down)		If Population Y, prompt to fill in text box for Population Affected (see row below).
		<i>Population (Species) Affected</i>	N	String Field (Manual); separated		If Population Y (above), prompt to fill in text box for names(s) of Population Affected.
	Technological Maturity (II)	Type of Validation and Verification	N	Drop Down	Statistical Verification	Need to be able to choose more than one.
					Sensitivity Analysis (verification)	
					Specificity and Sensitivity (verification)	

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Triage Level	Metric	Submetric	Required	Type of Field	Allowed Value	Notes
					Verification using Training Data	
					Validation using Temporally Independent Data	
					Validation using Spatially and Temporally Independent Data	
					Second Look Required	Trigger workflow AND hide this field from public/user view.
		In Use in an Operational Environment	N	Yes or No (Radio / Drop Down)		
3	Operational aspects (III)	Type of Model	Y	Check Boxes	Spatial	Not mutually exclusive. Need to be able to choose more than one.
					Ecological Niche	
					Event-Based	
					Dynamical	
					Statistical Models	
					Agent Based Model	
					Other	If selected, provide string field (required).
		SME Needed (Y/N)	N	Yes or No (Radio / Drop Down)		

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Triage Level	Metric	Submetric	Required	Type of Field	Allowed Value	Notes
		Accessibility	N	String Field		Pop-up suggestion: (plug-and-play, downloadable, etc.)
		Run Time, Complexity, Capabilities Required	N	String Field		
	Geographic and Population Coverage	Input Geography	N	Drop Down	Local	Need to be able to choose more than one.
					Provincial/State	
					Regional	
					National	
		Output Geography	N	Drop Down	Local	Need to be able to choose more than one.
					Provincial/State	
					Regional	
					National	
	Vector Population Size	N	Drop Down	10s; 100s; 1000s; 10,000s; 100,000s, 1,000,000s		
	Modeled Species Population Size	N	Drop Down	10s; 100s; 1000s; 10,000s; 100,000s, 1,000,000s		
	Data Input (III)	Input Content	Y	Check Boxes	Epidemiological Data from the Same Location	
					Epidemiological Data from Difference Location	

Disease Forecast and Prediction Model Evaluation Metrics & Definitions

Triage Level	Metric	Submetric	Required	Type of Field	Allowed Value	Notes
					Governmental and Non-Governmental Organizations	
					Literature	
					Satellite/Remote Sensing	
					Simulated	
					Laboratory Diagnostic	
					Expert Opinion	
	Output (III)	Output Content	Y	String Field (Manual) ';' separated		Pop-up suggestion: (e.g., structured warning output, probabilities, etc.)
		Sustainability	N	String Field (Manual) ';' separated		Pop-up example: For how long are the forecasts/predictions confident?
	Cost (III)	Research, development, testing, and evaluation	N	Drop Down	\$10s; \$100s; \$1000s; \$10,000s; \$100,000s, \$1,000,000s	
		Operations	N	Drop Down	\$10s; \$100s; \$1000s; \$10,000s; \$100,000s, \$1,000,000s	
		Government funded	Y	Drop Down or Check boxes	Y, N, Not Indicated	



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