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Mechanistic Modeling of Emergency Events: Assessing the Impact of Hypothetical Releases of Anthrax

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Abstract

A modular system for source-to-dose-to-effect modeling analysis has been developed based on the modeling environment for total risk studies (MENTOR),(1) and applied to study the impacts of hypothetical atmospheric releases of anthrax spores. The system, MENTOR-2E (MENTOR for Emergency Events), provides mechanistically consistent analysis of inhalation exposures for various release scenarios, while allowing consideration of specific susceptible subpopulations (such as the elderly) at the resolution of individual census tracts. The MENTOR-2E application presented here includes atmospheric dispersion modeling, statistically representative samples of individuals along with corresponding activity patterns, and population-based dosimetry modeling that accounts for activity and physiological variability. Two hypothetical release scenarios were simulated: a 100 g release of weaponized *B. anthracis* over a period of (a) one hour and (b) 10 hours, and the impact of these releases on population in the State of New Jersey was studied. Results were compared with those from simplified modeling of population dynamics (location, activities, etc.), and atmospheric dispersion of anthrax spores. The comparisons showed that in the two release scenarios simulated, each major approximation resulted in an overestimation of the number of probable infections by a factor of 5 to 10; these overestimations can have significant public health implications when preparing for and responding effectively to an actual release. This is in addition to uncertainties in dose-response modeling, which result in an additional factor of 5 to 10 variation in estimated casualties. The MENTOR-2E system has been developed in a modular fashion so that improvements in individual modules can be readily made without impacting the other modules, and provides a first step toward the development of models that can be used in supporting real-time decision making.

Keywords

Anthrax spores; dose; emergency events; exposure; MENTOR; modeling; risk

1. INTRODUCTION

During the *B. anthracis* attacks on the U.S. postal system in the fall of 2001, six letters, each containing 1 to 2 g of anthrax spores, caused anthrax infections in 22 individuals, and resulted in five deaths,(2) despite aggressive treatment and care.(3) In addition to the human cost, the economic costs of the *B. anthracis* attacks were significant. U.S. postal facilities contaminated by *B. anthracis* required more than two years of decontamination at a cost of more than \$200 million.(4) Understandably, concerns have been expressed that small

amounts of powdered *B. anthracis* inserted into the air intakes of subways, airports, shopping malls, sports arenas, and other public complexes could be devastating,(5) with long-term consequences potentially posing greater challenges than the short-term impact.(6)

Several studies have attempted to quantify the overall impact of large-scale releases of *B. anthracis*: concluding (a) an aircraft release of 50 kg of *B. anthracis* over an urban area would result in tens to hundreds of thousands of deaths,(7) (b) an airborne release of 100 kg of *B. anthracis* upwind of Washington, DC, would result in 130,000 to 3 million deaths,(8) and (c) 1.49 million people out of 11.5 million people (13.1%) would be infected downwind from a point of release of 1 kg of *B. anthracis*.(9) The economic costs were estimated to reach over \$26 billion per 100,000 people exposed,(10) not including the cost of decontamination. However, unlike nuclear weapons, the consequences of *B. anthracis* attacks can be mitigated.(5)

Though the consequences of a bioterrorism event are often enormous, it is prudent to not overestimate their impact,(11) as the scenario may appear unmanageable to responding agencies, leaving both the public and responders feeling helpless. Proper planning based on more complete interpretation and evaluation of available knowledge and experience can reduce public anxiety as well as increase confidence on behalf of the professionals responding to an adverse event.(12)

Planning and responding to a bioterrorism event, including vaccinations and postexposure prophylaxis, is a challenging task. Vaccines for anthrax exist,(13,14) but in practice several obstacles are present in the form of potential adverse effects,(15,16) lengthy dose regimen requirements,(17) and widely varying perceptions of risk.(18) Furthermore, vaccination of the majority of populations in large urban areas would be impractical due to limited availability of prophylactics, and the costs associated with procuring and handling large supplies.

1.1. Planning for Emergency Response

Several public health issues need to be addressed in planning and responding to emergency events.(19,20) While postattack response to bioterrorism is vital, preattack measures, such as detection, planning, and training, are just as important.(5) Spatial and temporal patterns of detected levels of biological agents can be combined with diagnostic modeling methods and real-time meteorological data to estimate release source strength and location and, consequently, high impact areas. This in turn will allow targeted emergency response planning, including targeted prophylaxis administration in high impact areas, because timely initiation of treatment is a critical factor.(21,22) As a first step toward rapid detection and response to bioterrorism, a national monitoring network in the United States, called BioWatch, has been set up.(23,24) It is being expanded to detect the presence of a number of airborne biological agents, including anthrax spores; however, currently the results are not available in real time. Recent modeling efforts have also focused on evaluating response based on syndromic surveillance and on prioritizing threats.(25-27)

After the anthrax attacks in the United States in 2001, several modeling studies focused on estimating effects of *B. anthracis* releases, with varying levels of model detail and scale. Reshetin and Regens(28) modeled the release of *B. anthracis* inside a 50-story building. Webb and Blaser(29) modeled the transmission of *B. anthracis* through cross-contamination of postal letters, whereas Ho and Duncan(30) modeled the aerosol hazards from a letter containing anthrax spores. Fowler *et al.*(31) performed a probabilistic comparison of vaccination and antibiotic prophylaxis based on probabilities of exposure. Wein *et al.*(9) and Craft *et al.*(32) performed detailed modeling analyses linking *B. anthracis* dispersion over large areas with infection and disease progression mechanism, hospital response logistics,

and corresponding outcomes. More recent efforts include Biowar,(33,34) an agent-based model of bioattacks that links simple dispersion models with models for social networks and behavioral attributes such as health-care-seeking behaviors and pharmaceutical purchases.

One of the limitations of existing *B. anthracis* modeling studies is that they have employed substantial simplifying assumptions. For example, Craft *et al.*(32) assumed uniform population density over a large area, uniform population demographics, Gaussian dispersion with uniform wind speed and direction, constant inhalation rates for all individuals, and personal exposure concentrations being the same as outdoor concentrations. The rationale for such simplifying assumptions is that the equations become mathematically tractable and result in analytical solutions. However, such simplifications limit the model applicability to only a small set of potential scenarios and conditions and minor changes in modeling assumptions may require rewriting underlying model equations.

2. APPROACH

This study presents a prototype modeling system for performing source-to-dose-to-effect analysis of inhalation anthrax infections from airborne weaponized anthrax spore releases. The main principle guiding the development of the present system is that prioritized exposure analysis(35,36) should be conducted in order to minimize misclassification of exposure. Consistent quantification of exposures and doses for different release scenarios, as well as for different emergency response strategies, provides a sound scientific basis for developing and then implementing plans for response strategies. The main objective of this system is to utilize available databases of distributions of demographics, human activity patterns, etc., and to provide sufficient flexibility so that this system can be useful in planning multiple emergency response scenarios for training purposes and for rapid decision making in the aftermath of emergency events.

A modular, mechanistic framework that links available models and databases for characterizing exposures and adverse impacts would improve risk assessment in terms of (a) providing consistency, (b) allowing assessments on multiple scales, and incorporating the important processes from release source to dose received by individuals, (c) optimizing the use of the most up-to-date models and databases for individual processes, and (d) allowing systematic sensitivity and uncertainty analyses to identify the important factors that affect the outcomes the most. The underlying person-oriented exposure modeling approach has been utilized and evaluated in the past for different types of contaminants.(37,38)

2.1. Important Factors Influencing the Impact of *B. anthracis* Releases

There are several factors that influence inhalation exposures and doses of *B. anthracis* to humans from airborne releases.

2.1.1. Release Source(s)—The magnitude, location of source(s), and patterns of release are the primary factors and major unknowns in performing dispersion modeling for emergency events. An attack scenario can include single or multiple sources (differing in location or time), with anthrax spores released quickly (“instantaneous”) or slowly over time (“continuous”). These are required as initial inputs to the exposure and dose modeling analysis.

2.1.2. Meteorological Conditions and Topography/Terrain—The meteorological conditions and local topography (rural, urban, etc.) influence the extent of dispersion of anthrax spores and, thus, the overall impact of a release. These include variables such as wind speed, direction, temperature, atmospheric stability, boundary layer thickness, surface

topography, surface roughness, land use, etc.,(39,40) which can together contribute to over a factor of 20 variation in potential casualties.(8)

2.1.3. Population Distributions—The distribution of population in the areas affected by the releases (“downwind locations”) determines the intensity and spread of exposure and potential infections. This is a highly variable and often ignored factor, in the sense that the population distribution changes in time (e.g., commuting to major urban centers or large gatherings at major events). However, it is manageable to an extent, for example, through evacuation of people from downwind locations or enforcing shelter-in-place.

2.1.4. Human Activity Patterns—The location of the individual at any given time (e.g., outdoors versus indoors), and the activity performed (e.g., running versus sleeping or resting), determine the exposure concentrations, the breathing rates, the efficiency of particle uptake, and thus the number of inhaled anthrax spores. These factors are also important in characterizing the potential contact or lack of contact with a contaminant. However, they are often overlooked or poorly characterized in emergency event analyses.(9,28,29,31,32,41)

2.1.5. Physiological Characteristics of Individuals—The base inhalation rates are dependent on the age, gender, and physical characteristics such as body weight and life style patterns. Furthermore, the infection potential of *B. anthracis* is dependent on the age of the individual.(29,32)

Other factors that are important in determining exposures are socioeconomic attributes, such as housing characteristics (age, size, ventilation, etc.), which determine the fraction of outdoor *B. anthracis* that gets entrained indoors.

2.2. Steps and Resources in the Estimation of Exposures, Doses, and Infections

Based on the approach introduced by Georgopoulos *et al.*,(37) several modeling steps (or components, as some of them do not have to be performed in sequence) are needed in assessing the impacts of airborne releases of *B. anthracis*. In general, the following eight steps are needed, as shown in Fig. 1:

1. Estimation of outdoor concentration levels of airborne anthrax spores through one of the following:
 - a. Spatiotemporal analysis of available data. This can involve interpolation of detection data from monitors such as the BioWatch monitoring network(23,24) using statistical techniques such as SpatioTemporal Random Field (STRF)(42) and Bayesian Maximum Entropy (BME).(43)
 - b. Numerical modeling of the atmospheric dispersion of *B. anthracis*. This involves application of atmospheric dispersion models such as California Puff Model (CALPUFF),(44) Hazard Prediction and Assessment Capability (HPAC),(45) or Hybrid Particle and Concentration Transport (HYPACT).(46) These models use dynamic meteorological profiles as inputs (either user-provided or from meteorological data sources) and provide contaminant concentration profiles at different spatial and temporal resolutions. Dispersion modeling at finer scales, for example, within a building or within the vicinity of the release location, can be accomplished through detailed subgrid modeling approaches such as Computational Fluid Dynamics (CFD) based models.(28)
2. Estimation of local *B. anthracis* levels at the scale of interest (such as a census tract) or a conveniently defined grid through one of the following:

- a. Spatiotemporal statistical interpolation of monitor data or outputs of a “coarse-scale” model, using the techniques mentioned in Step 1a.
 - b. Aggregation of the outputs of a “fine-scale” model. This is important when the atmospheric dispersion model provides concentration profiles at a finer scale than the resolution of other model components, such as population distributions, which have a typical resolution of a census tract or a census block.
3. Characterization of attributes of populations (geographic density, age, gender, race, income, etc.) through one of the following:
- a. Selection of a fixed-size sample population (“virtual individuals”) that statistically reproduces essential census demographics.
 - b. Division of the population of interest into an exhaustive set of cohorts based on different relevant population attributes.

The population attributes, such as the distributions of age, gender, employment, and housing, can be developed from available census data (see, e.g., USCB47). Sometimes, relevant databases are available as components of other modeling systems, as in the case of the Air Pollution Exposure Model (APEX),(48) which provides databases for housing as well as for commuting profiles. Depending on the emergency response scenario, relevant adjustments to the distributions of population profiles may be necessary (e.g., changes in the population distribution during special events).

4. Development of activity event (or exposure event) sequences for each member of the sampled population or for each cohort for the exposure period through one of the following:
- a. Existing databases from composites of past studies (for baseline assessment).
 - b. Hypothetical scenario-based or “simulated” activity patterns based on options such as “shelter-in-place” versus different evacuation options.

For baseline assessments, the Consolidated Human Activity Database (CHAD)(49) can be used. It contains over 22,000 person days (diary records) of activity patterns developed from preexisting human activity studies. Each diary record provides a basis for simulating the movement of the “virtual individual” through geographic locations and microenvironments during the simulation period. Each event is defined by geographic location, start time, duration, microenvironment visited, and an activity performed. The attributes of CHAD records include age, gender, employment status, and smoking status of each individual, which can be used for matching the demographic characteristics of each sampled individual.(50) For planning and training purposes, several additional options can be considered for protective action, including evacuation, “shelter-in-place,” or a combination.(51) The corresponding activity profiles can be either synthesized independently or through scenario-specific modifications to existing CHAD diaries.

5. Estimation of personal exposure levels and temporal profiles of *B. anthracis* concentrations in various microenvironments (residences, offices, restaurants, vehicles, etc.) through either one or more of the following methods:
- a. Simple linear, steady-state mass balance.
 - b. Nonlinear, dynamic models.

c. Detailed computational fluid dynamics models.

Several modeling studies of indoor/outdoor relationships of fine particles have been presented in the literature, addressing issues such as contaminant penetration of indoor environment and corresponding particle size dependence.(52-54) The Stochastic Human Exposure and Dose Simulation (SHEDS) application(55) provides distributions of air exchange rates for different types of residential microenvironments, while other models and databases provide distributions for air exchange rates for general nonresidential microenvironments(56) and vehicle microenvironments.(57)

6. Calculation of appropriate inhalation rates for the members of the sample population by combining physiological attributes of study subjects and activities pursued during the individual exposure events. The CHAD diary records also provide information on energy expenditure, which can be used directly to estimate inhalation rates, as discussed in Step 6 of Section 4. Alternatively, probability distributions or tables describing age-specific inhalation rates of humans can also be used.(58-60) The inhalation rates, along with personal exposure levels, provide intake rates of anthrax spores.
7. Calculation of target tissue dose through physiologically-based respiratory deposition modeling by estimating the amount of inhaled *B. anthracis* that is deposited in the lungs. The International Commission on Radiological Protection (ICRP)(61,62) provides deposition fractions of fine particles in different regions of the human lungs. These fractions are age and gender dependent, and are also dependent on the size of the particles.
8. Estimation of probability of infection for each simulated individual based on calculated dose and physiological attributes, and summation of these probabilities to estimate the total number of potential infections for each local area in the simulation domain (e.g., each census tract in the modeling domain). Sparse data are available for describing the dose-response relationships of *B. anthracis* in humans, (29) and several dose-response models have been proposed in the literature, as described in Step 8 in Section 4.

3. MENTOR-2E IMPLEMENTATION

The steps outlined above were implemented in a modular manner as part of the Modeling Environment for Total Risk studies (MENTOR).(1,37) The general approach of MENTOR is to utilize existing models when available and to provide new modules to “fill gaps” in the source-to-dose-to-effect sequence. In that sense, MENTOR is not a “new model”; it can be viewed as a computational toolbox intended to facilitate consistent multiscale risk assessment. In this particular study, several existing models and approaches relevant to exposure estimation have been used. For example, various concepts from the SHEDS approach, which has been applied for studying exposures to particulate matter(55) and pesticides,(63) have been adapted and incorporated into the formulation of different MENTOR modules.

The MENTOR-2E system has been coded in Matlab(64) (www.mathworks.com), while various relevant programs such as the CALPUFF model(44) have been linked in a pipeline manner. The model code and the underlying data files are available upon request. The computational time for the CALPUFF simulation is dependent on the extent of the modeling domain, the grid resolution, and the duration of the simulated period. For the case of 250 m resolution, and an area covering the entire State of New Jersey, the CALPUFF simulation required about 2 CPU hours on a 3 GHz Pentium Processor. The computational time for

calculation of exposures, doses, and effects of *B. anthracis* release is dependent mainly on the number of virtual individuals simulated per census tract, the number of census tracts, and the time period considered in the simulation. This study used 500 virtual individuals per census tract, and the exposure and dose calculations required about 4 minutes of CPU time per census tract for population sampling, exposures, and dose calculations. However, the calculations involving activity patterns, exposures, and doses are run in a distributed manner on a computer cluster, as these calculations for each census tract are independent of those for other census tracts.

4. CASE STUDY APPLICATION

The location and timing of the hypothetical scenarios used in this study to demonstrate the MENTOR-2E system were assumed to represent potential variations of the anthrax attacks of 2001 through the postal system in the State of New Jersey. The main difference is in the amount and release characteristics. *B. anthracis* was assumed to be released in the air and the release was assumed to occur starting at 08:00 hours on September 18, 2001 in the vicinity of the Hamilton Post Office (the site and the day of the mailing of letters containing anthrax spores in 2001), as shown in Fig. 2. A hypothetical release of 100 g (1 trillion spherical spores per gram) of weaponized anthrax spores was assumed, similar to the release characteristics assumed by the modeling study of Craft *et al.*(32) Two types of releases were considered in the simulation: a “quick” release, where the spores were assumed to be released over a period of one hour (release scenario A), and a “continuous” release, where the spores were assumed to be released over a period of 10 hours (release scenario B). The study focused on the impact of these releases on the general population of the State of New Jersey, and exposures during the day of September 18, 2001 were simulated. Only the census tracts in New Jersey (totaling 1944 census tracts) were considered here, instead of the entire region of potential impact, to illustrate the application of the system to targeted administrative areas by responding agencies. The inclusion of other census tracts in other states is straightforward.

The specific MENTOR-2E application for this case study used the following eight steps (Fig. 1) for assessing exposures, dose, and potential infections.

Step 1

The ambient *B. anthracis* concentrations were calculated using the CALPUFF model,(44) which is a generalized nonsteady-state air quality model that simulates the transport, transformation, and dispersion processes of “puffs” of material from emission sources, and provides hourly average estimates of concentrations. CALPUFF has been adopted by the USEPA in its Guideline on Air Quality Models(65) as a preferred model for assessing long-range transport of pollutants, and on a case-by-case basis for certain near-field applications involving complex meteorological conditions. Meteorological data were retrieved from the National Oceanic and Atmospheric Administration (NOAA)(66) and converted into CALPUFF input format. In this study, it was assumed that the hourly averages provided by the CALPUFF model at a resolution of 250 m are adequate for characterizing *B. anthracis* exposures and doses, based on the general guidance on CALPUFF modeling.(67) In this study, a 1,000 × 1,000 grid at 250 m resolution was used.

In general, the selection of a particular dispersion model and modeling options depends heavily on the type of release and the response options, including the scale of attack, detection time, complexity of the geography, and population densities in the downwind regions. Some emergency events need to be modeled at high temporal and spatial resolutions. Examples include large releases of chemicals such as chlorine, where high-resolution concentration profiles are important in order to capture peak concentrations.

However, in the case of *B. anthracis* releases, the main metric of concern is the total uptake of anthrax spores, so hourly averages were considered adequate. In general, a tradeoff can be made by considering model resolution, model complexity, and setup and simulation time.

In order to perform comparative evaluation, a simplified dispersion equation was also used(32) to characterize ambient concentrations of anthrax spores. A constant wind speed of 1.5 m/s was assumed blowing toward the northeast, at 45 degrees, corresponding to the general direction and speed of the wind during September 18, 2001, as part of this simplified application.

Step 2

Concentration estimates from CALPUFF at regularly spaced grid points at 250 m × 250 m resolution were aggregated at the level of a census tract; the average of concentrations at all grid points within a census tract was assumed to represent the concentration at the geometrical centroid of the census tract. The spatial averaging approach can sometimes result in artificial “discontinuities” in the census tract level concentrations, as is the case of Fig. 3, where two census tracts close to the plume show a zero concentration, whereas the surrounding census tracts show nonzero concentrations; however, this approach follows mass balance. The spatial averaging approach was used here because detailed allocation of concentrations can be computationally very demanding. It should be noted that for finer-scale spatial resolution (e.g., exposures studies focusing on census block level resolution or on regularly spaced grids), the CALPUFF model outputs can be aggregated or interpolated depending on the resolution. In case of the simplified dispersion modeling, the concentrations were estimated directly at the geometric centroid of each census tract.

Step 3

The attributes of the population under study were retrieved from the 2000 U.S. Census Survey.(47) Due to the variability of the urban population, in order to statistically reproduce essential demographic distributions of age, gender, housing type, and employment status, a rather large statistical sample of 500 “virtual individuals” was sampled for each of the 1944 census tracts under study.(37,55) In this study, the effect of commuting on population distributions was not considered, and the residential distributions were assumed to represent the population distributions throughout.

Step 4

A 24-hour activity diary for each “virtual individual” was selected from the CHAD diaries using the approach described earlier. In this study, the 113 microenvironments in the CHAD diaries were grouped into four categories: home, other indoor, outdoor, and vehicle.

Step 5

The outdoor concentration levels of *B. anthracis*, aggregated at the census-tract level, were used as inputs to the MENTOR modules for estimating microenvironmental concentrations. The estimation of *B. anthracis* levels in the various microenvironments in this study was based on the simple mass balance equation:

$$\frac{dC_{in}}{dt} = f_p \cdot A \cdot C_{out} + \frac{S}{V} - (A + F_d) \cdot C_{in}, \quad (1)$$

where C_{in} is the indoor *B. anthracis* concentration (number/volume), f_p the penetration factor (dimensionless fraction, indicating the amount of *B. anthracis* that can penetrate indoors), A the air exchange rate between outdoor and indoor (1/time), C_{out} the outdoor *B.*

anthracis concentration (number/volume), S the indoor *B. anthracis* source rate (mass/time), V the indoor volume (volume), and F_d the decay rate of *B. anthracis* indoors (via deposition, absorption to walls, etc.; 1/time). This equation was further simplified on the basis of the following assumptions: (1) steady-state approximation (which can provide a reasonable approximation for time durations of hours), (2) $S = 0$ (i.e., no indoor sources), thus resulting in $C_{in} = A f_p C_{out} / (A + F_d)$. Furthermore, the anthrax spores were assumed to behave similar to fine particulate matter in the estimation of penetration and deposition. The resuspension of anthrax spores was assumed to be negligible at the modeling time scale in this study. However, it will be a critical factor in the decontamination process.

For each microenvironment corresponding to the activity event, the parameters in the mass balance equation were generated through random samples. The residential parameters include day- and nighttime air exchange rates, house volumes, and penetration and deposition factors; these were sampled once for each individual from the corresponding distributions (see Table 2 from Burke *et al.*55), and were held constant throughout the simulation. Parameters for other microenvironments, such as the air exchange rates in the vehicles, stores, etc., were sampled separately throughout the simulation from two distributions: one for the general nonresidential microenvironments(56) and one for vehicles References 37 and 68. In the simplified application, all virtual individuals were assumed to be outdoors, and anthrax spore concentration at the census tract centroid is assumed to be the exposure concentration.

Step 6

In this study, exposure to *B. anthracis* was assumed to occur solely through the inhalation of contaminated air. Thus, one of the main factors in *B. anthracis* exposure is a person's inhalation rate. For each activity event of a virtual individual, inhalation rates were calculated using a combination of age- and gender-dependent ideal body mass, basal metabolic rate, and activity-specific energy expenditure and METs (Metabolic Equivalent of Tasks; described in detail in References 37 and 68). In the simplified application, the breathing rates were assumed to be constant across all individuals.

Step 7

The population-based lung dosimetry model employed by Georgopoulos *et al.*,(37) based on the HUMTRN model,(69) was used to calculate the delivered doses for individuals of both genders and of different ages. The calculated inhalation rates were combined with the corresponding microenvironmental concentrations to estimate the inhaled dose delivered to the lung for each virtual individual. Lung deposition of anthrax spores was calculated for three regions of the lungs: nasal-pharyngeal (NP), tracheobronchial (TB), and pulmonary (P), using empirical values of deposition fractions from the International Commission on Radiological Protection (ICRP) databases.(61,62) Total uptake of *B. anthracis* for each virtual individual was estimated for the day of September 18, 2001, from the sum of event-based doses inhaled by the individual during the exposure event sequence. In the simplified application, all inhaled anthrax spores were assumed to be initially deposited in the lung.

Step 8

Calculation of the probability of infection for each sampled individual in a census tract through the application of a *B. anthracis* dose-response model, and scaling that to the total population of the census tract. Several empirical dose-response models have been proposed for *B. anthracis*,(9,29,32,41) and the range of potential infections spans an order of magnitude, solely based on the choice of the dose-response relationship.(41) Six different dose-response models were used in this study to estimate the probability $P(s, a)$ that an individual of age a would be infected from a dose of s spores. Here, the dose is assumed to

represent the number of spores “initially deposited” into the lung, whereas the dose-response models are assumed to account for the subsequent, time-dependent ciliary clearance.

1. $P(s, a) = \min\left(1, \frac{s}{c_1 - c_2 a_1}\right)$, based on Craft *et al.*, (32) where $c_1 = 38,000$ and $c_2 = 450$, $a_1 = \min(a, A_{\text{cut}})$, and A_{cut} is the cut-off age of 80 years, beyond which the dose response is assumed to plateau with age.
2. $P(s, a) = \Phi(\alpha + \beta \cdot \log(s) + \gamma \cdot a + \delta \cdot a^2)$, an age-dependent probit model based on Wein *et al.* (9) where Φ is the *cdf* of the normal distribution, and parameters $\alpha = -9.733$; $\beta = 1.025$; $\gamma = -0.016/\text{year}$; and $\delta = 0.0006/\text{year}$. (2)
3. $P(s, a) = \Phi(\alpha + \beta \cdot \log(s))$, an age-independent probit model from Wilkening, (41) where $\alpha = -2.6361$, and $\beta = 0.291$, corresponding to an ID_{50} of 8,600 spores, and a probit slope of 0.67.
4. Another age-independent probit model from Wilkening, (41) of the form $P(s, a) = \Phi(\alpha + \beta \log(s))$, with $\alpha = 5.6263$, and $\beta = 0.621$, corresponding to an ID_{50} of 8,600 spores, and a probit slope of 1.43.
5. $P(s, a) = \exp\left(\frac{-\lambda s}{\lambda + \theta}\right)$, an exponential model that accounts for probability of spore destruction and spore germination in the lungs, (41,70) where $\theta = 0.109/\text{day}$, and $\lambda = 8.8 \times 10^{-8}$.
6. $P(s, a) = \frac{\beta \cdot (\exp(s/\alpha) - 1)}{1 + \beta \cdot (\exp(s/\alpha) - 1)}$, an age-dependent logit model from Webb and Blaser, (29) where α and β are derived from age-dependent values for ID_{50} and ID_{10} , classified into four age groups.

In principle, the uncertainties in the dose-response relationship modeling directly translate into the corresponding uncertainties in the number of probable infections. However, the age-dependence of dose-response and activity patterns warrant explicit characterization of these uncertainties.

The total number of potential infections in each census tract was obtained by summing the probabilities of infections across all the virtual individuals in that census tract, and scaling that value to the population of the census tract. The summation is possible because infections of different individuals represent independent events.

$$X_j = \left(\sum_{i=1}^{N_{\text{sample},j}} P(s_{ij}, a_{ij}) \right) \frac{N_j}{N_{\text{sample},j}}, \quad (2)$$

where X_j is the number of potential infections in census tract j , $N_{\text{sample},j}$ the number of sampled virtual individuals in census tract j , s_{ij} the dose of inhaled *B. anthracis* for sampled virtual individual i in census tract j , a_{ij} the age of sampled virtual individual i in census tract j , and N_j the total population of census tract j .

The impact of *B. anthracis* releases on sensitive population subgroups, such as the elderly, can also be studied easily using MENTOR-2E. Since the virtual individuals are described via several attributes that include age, gender, body weight, employment status, etc., the potential infections within subgroups of interest can be obtained by selecting the virtual individuals that belong to that subgroup. In general, for a subgroup M , the corresponding potential infections can be calculated through

$$X_{j,M} = \left(\sum_{i \in M} P(s_{ij}, a_{ij}) \right) \frac{N_j}{N_{\text{sample},j}}. \quad (3)$$

5. RESULTS

Fig. 3 shows the average outdoor *B. anthracis* concentrations during 08:00–09:00 hours for both release scenarios. Fig. 4 shows the average outdoor concentrations during 14:00–15:00 hours; this corresponds to six and a half hours after the start of the *B. anthracis* release, and five and half hours after the end of the one-hour release. As the maps show, the pattern of *B. anthracis* concentrations varies significantly depending on the pattern of the release. Though the initial concentrations are high in the quick-release case (scenario A), high-airborne concentrations of *B. anthracis* remain in the slow-release case (scenario B).

Median and 95th percentile values for individual biological doses per census tract for the general population are shown in Fig. 5 (for release scenario A) and in Fig. 6 (for release scenario B). The MENTOR-2E system also allows focusing on susceptible subpopulations; the 95th percentile doses for the elderly (65 years and older) are shown for both release scenarios in Fig. 7.

5.1. Effect of Simplifying Approximations Used

In order to illustrate the differences in estimates arising out of differences in model assumptions, the MENTOR-2E results were compared with those obtained using two approximations: (a) the constant inhalation rate of $5 \times 10^{-4} \text{ m}^3/\text{s}$ (1.944 m^3/h) used by Craft *et al.*,(32) and (b) a reference constant inhalation rate of 0.78 m^3/h from the USEPA exposure factors handbook (EFH).(71) The corresponding average breathing rate for the study population in the MENTOR-2E simulation was about 0.71 m^3/h , which is approximately equal to the inhalation rate from the EFH. In employing the Craft *et al.* and EFH approximations, the outdoor concentration values were assumed to represent the personal exposure concentrations, and the corresponding doses and probabilities of infection were estimated.

The estimated percentiles of biological doses of *B. anthracis* (i.e., number of deposited anthrax spores) for the individuals in the entire study population are shown in Fig. 8 (release scenario A) and Fig. 9 (scenario B). The corresponding percentiles of probabilities of infection shown in Fig. 10 (release scenario A) and Fig. 11 (release scenario B). The total numbers of probable infections for the entire study region were also estimated, as shown in Table I. In both release scenarios, the infection estimates derived using the constant inhalation approaches are about five to ten times higher than those derived through MENTOR-2E. The dependence of the potential infection estimates on the choice of the dose response is also strong, as shown in Table II. The estimates span an order of magnitude, solely based on the choice of the dose-response model. This, coupled with the order of magnitude change in estimates based on inhalation rates and microenvironmental calculations, results in a range of two orders of magnitude for the estimates. Further approximations, such as constant wind speed, are likely to overestimate the impacts in some areas, while underestimating the impacts in others. In fact, as shown in Table II, the estimates using a constant wind speed and simplified dispersion equation employed by Craft *et al.*(32) are substantially higher. In fact, the corresponding doses are high enough that the differences due to various dose-response formulations become negligible. As noted earlier, such overestimations and underestimations can have significant adverse impact on response to emergency events.

6. DISCUSSION AND CONCLUSIONS

This study presents a demonstration application of the integrated MENTOR-2E system for assessing the impact of airborne *B. anthracis* releases, with a main focus on mechanistically consistent linkage of modules in the source-to-dose-to-effect sequence. Future work can focus on characterization of commuting patterns, population gathering at different events, and custom response scenarios, such as “shelter-in-place” versus evacuation.

The difference in dose estimates between the release scenarios, shown in Fig. 5 (release scenario A) and in Fig. 6 (release scenario B), is solely due to the difference in the concentrations profiles, which in this case are influenced by the *B. anthracis* release patterns. However, variations in meteorological conditions can result in significantly different concentration profiles. Similarly, different forms of release, for example, releases from multiple locations, either simultaneously, or in a staggered manner, influence the concentration profiles. Therefore, for emergency response modeling, improved estimates of concentration profiles are very important. It must be noted that the estimates of infections are “probabilistic” expected values, and do not take into account the time for incubation between the exposure and the actual infection; therefore, the results do not reflect the estimates of possible number of potential hospital admissions within days of the *B. anthracis* release. Furthermore, the exposures modeled here included only those occurring during the day of September 18, 2001, and subsequent exposures were assumed to be negligible as the *B. anthracis* plume expands and becomes significantly diluted.

The simplifying assumptions have a strong impact on estimates of probable infections, as shown in Table II. In both release scenarios considered, the same concentration profiles were used for the three exposure and dose calculation approaches (Table II, Rows 2–7). Furthermore, even higher estimates of infections are possible if simplifying assumptions are used for calculating the concentration profiles (e.g., Table II, Row 8, which shows results for the case where simplified dispersion modeling and simplified population characteristics are assumed, both used by Craft *et al.*(32)).

Wilkening(41) discusses the uncertainty associated with the selection of the dose-response model for inhalation anthrax, and concludes that a factor of 10 uncertainty exists based on the choice of the dose-response model. However, as shown in Fig. 10 and Fig. 11, the impact of simplifying assumptions with respect to microenvironments and activity patterns spans about a factor of 7–10. Therefore, all components of the source-to-dose-to-effect modeling, including dispersion, exposures, and dose response, need to be improved in order to use such models in planning, intervention design, and training.

Integrated modeling applications such as the one presented here need to be considered in the context of multiple sources and types of uncertainties, including natural uncertainty/variability (atmospheric turbulence, population demographics and variability, etc.), model/structural uncertainties (selection of the population units, alternative dose-response models, etc.), input/parameter uncertainties (estimates of source strength, wind speed, direction, etc.), and evaluation data uncertainties (data used in parameter estimation for the underlying models). This study focused on the population variability model uncertainty in exposure calculation approaches, and model uncertainty in dose-response relationships. However, there is a need for further systematic uncertainty analyses focusing on identifying major uncertainties in emergency response planning, and characterizing the contributions of these. Such characterization will further improve the confidence in simulation models used for planning, training, and decision support.

Ongoing work focuses on improving the performance of MENTOR-2E by performing some of the modeling steps beforehand and using those outputs toward achieving a real-time

impact assessment and real-time evaluation of alternative response strategies. The approach as presented here can be easily parallelized (e.g., representative activity pattern generation, sampling of microenvironmental factors, housing characteristics, physiological parameters, etc., can be performed beforehand, and the “concentration profiles” can then be used to calculate the exposures). Furthermore, since each administrative unit is modeled independently (e.g., each census tract level), they can be run in parallel on a distributed cluster of computers.

A comprehensive planning scheme for detecting and responding to a bioterrorism event should consist of effective and efficient monitoring, ability to characterize the release sources based on detected values, and ability to accurately estimate the potential impact, not only in terms of the overall magnitude, but also in the spatial distribution. It should provide means to assess alternative response strategies, e.g., flexibility to define custom activity patterns, modification of population distributions and activity patterns (e.g., during population evacuation from affected areas), as it can be used for training of emergency responders. Furthermore, such a system should be tuned for near real-time application for maximum benefit.

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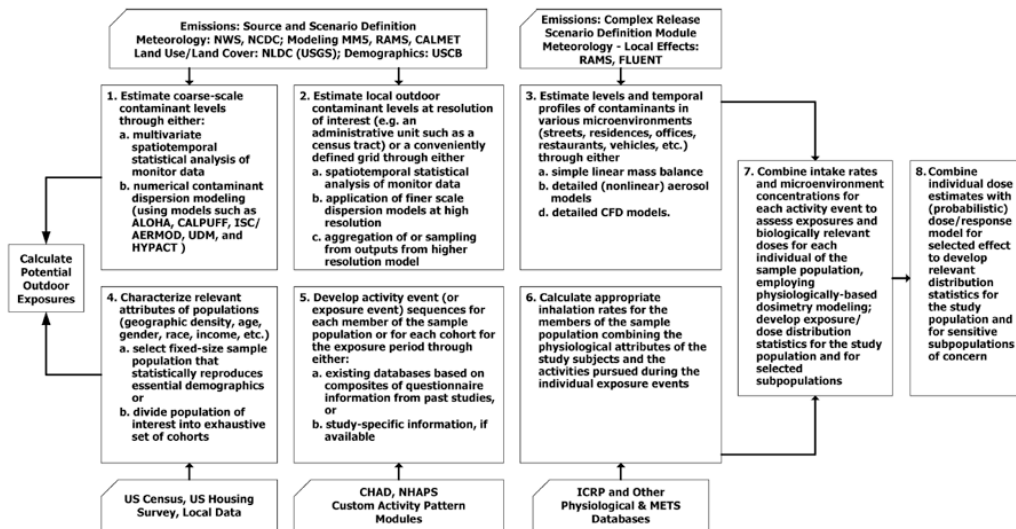


Fig. 1. A generalized eight-step flowchart describing the processes involved in assessing risk using a source-to-dose-to-effect framework, adapted from Georgopoulos *et al.*(37) this is also referred to as person-oriented population-based exposure modeling (POM/PBEM). This flowchart reflects the structure of the MENTOR approach and provides a general “template” for comparing the application of risk assessment systems. A subset of the models appearing in this flowchart has been used in this study.

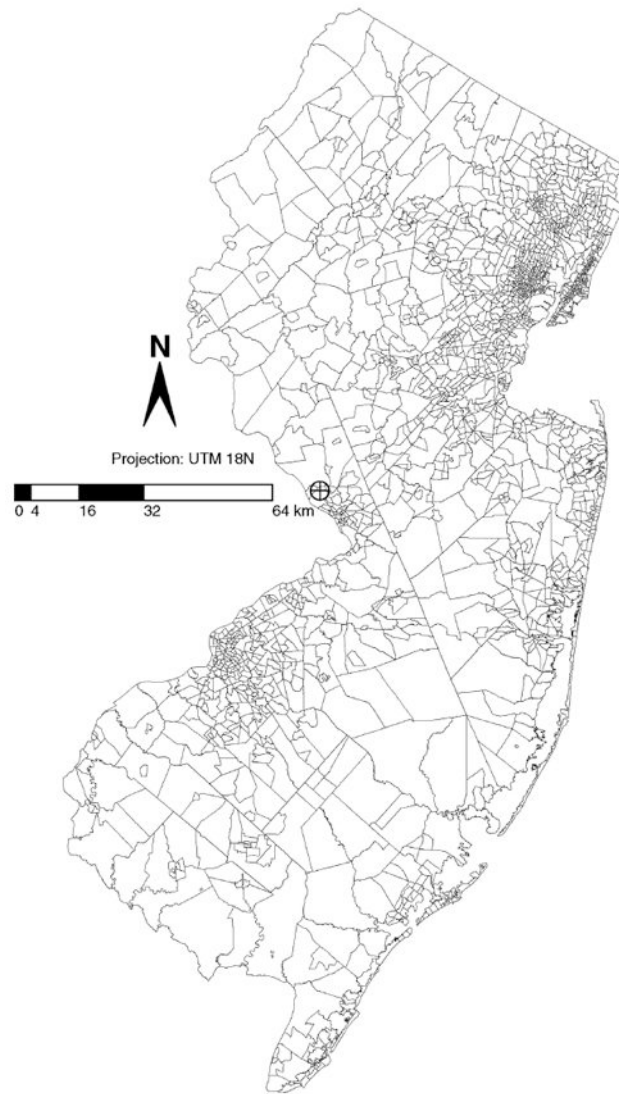


Fig. 2. The location of the hypothetical release of anthrax spores and the distribution of the census tracts within the State of New Jersey. The hypothetical release location corresponds to the Hamilton Post Office, the place from which envelopes containing *B. anthracis* were mailed on September 18, 2001.

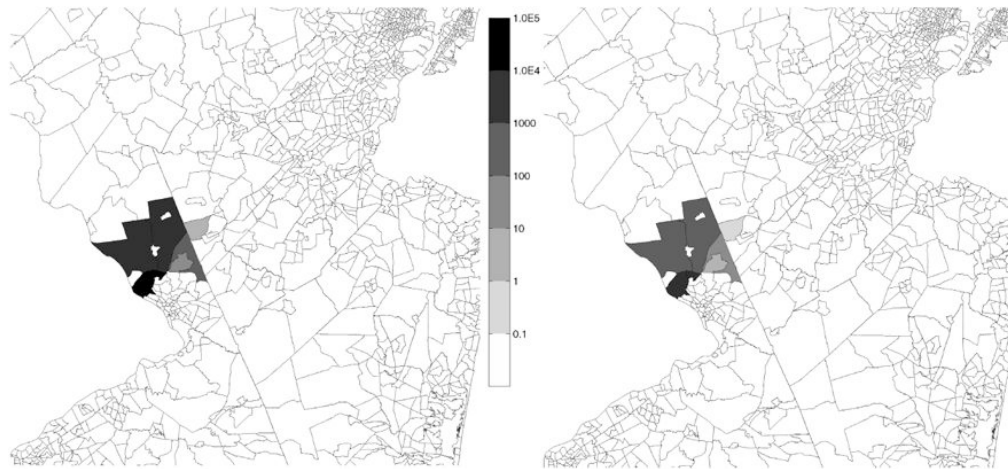


Fig. 3. Outdoor number concentrations of anthrax spores ($1/\text{m}^3$) during 08:00–09:00 calculated by CALPUFF for a release of 100 g of *B. anthracis* over a period of one hour (left) (release scenario A) and 10 hours (right) (release scenario B), for the day of September 18, 2001, with the release starting at 08:00 hours (values less than 1 indicate the probability of finding one spore per m^3).

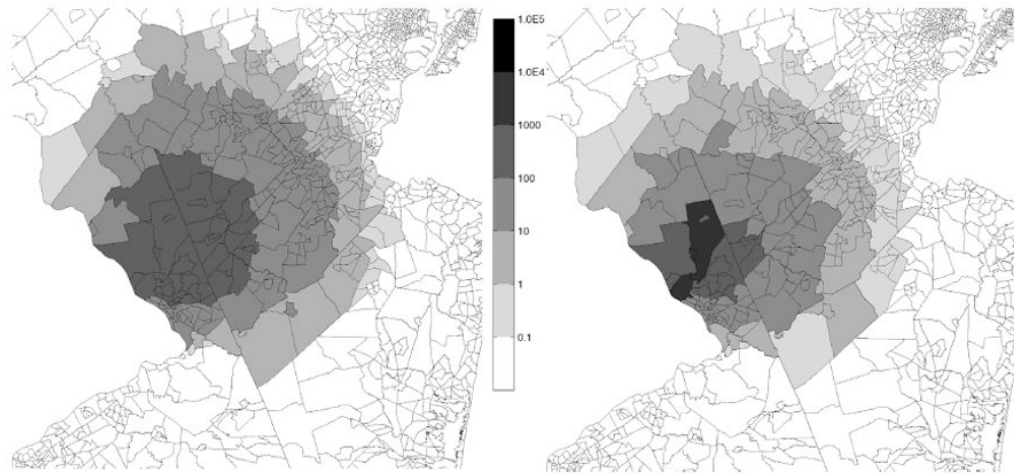


Fig. 4. Outdoor number concentrations of anthrax spores ($1/\text{m}^3$) during 14:00–15:00 calculated by CALPUFF for release scenario A (left) and release scenario B (right). Values less than 1 indicate the probability of finding one spore per m^3 . In the case of scenario A (left), the release has already been completed and no spores were released during the previous five and a half hours.

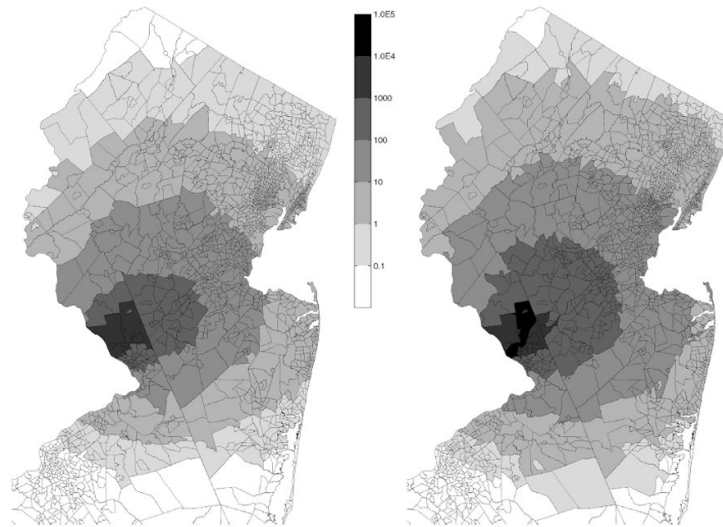


Fig. 5. Estimated individual biological doses (spores/day) of *B. anthracis* in each census tract due to a release scenario A: median (left) and 95th percentiles (right) of doses for the simulated population in each census tract. Values less than 1 indicate the probability of inhaling one spore.

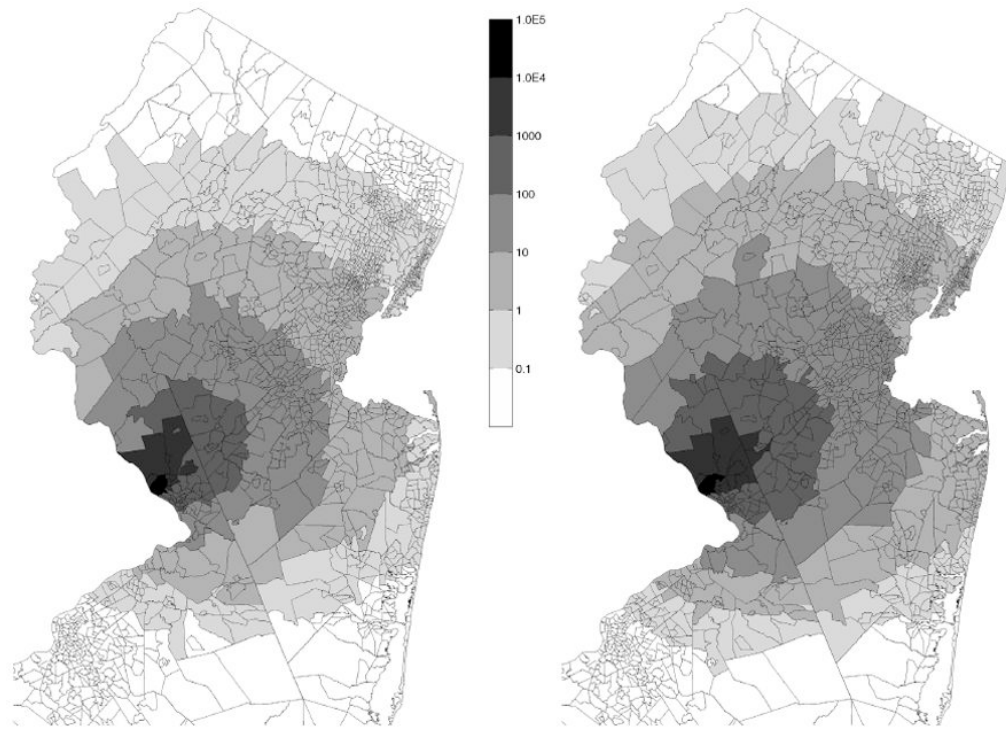


Fig. 6. Estimated individual biological doses (spores/day) of *B. anthracis* in each census tract due to a release scenario B: median (left) and 95th percentiles (right) of doses for the simulated population in each census tract.

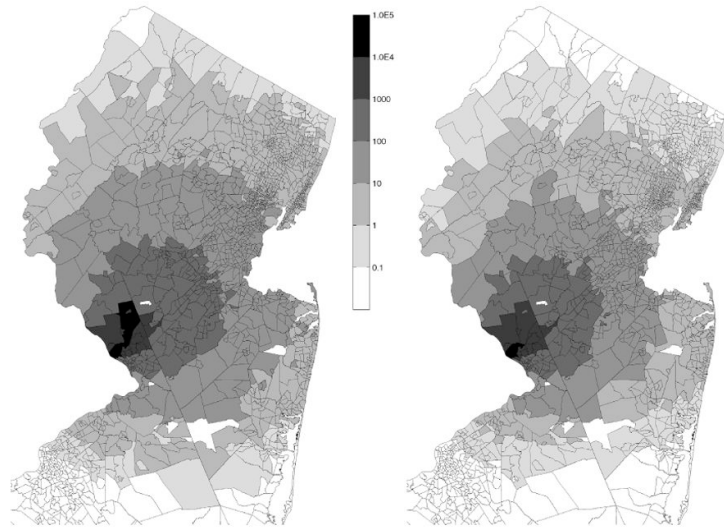


Fig. 7. Estimated 95th percentile biological doses of *B. anthracis* for elderly individuals (over 65) in each census tract due to release scenarios A (left) and B (right).

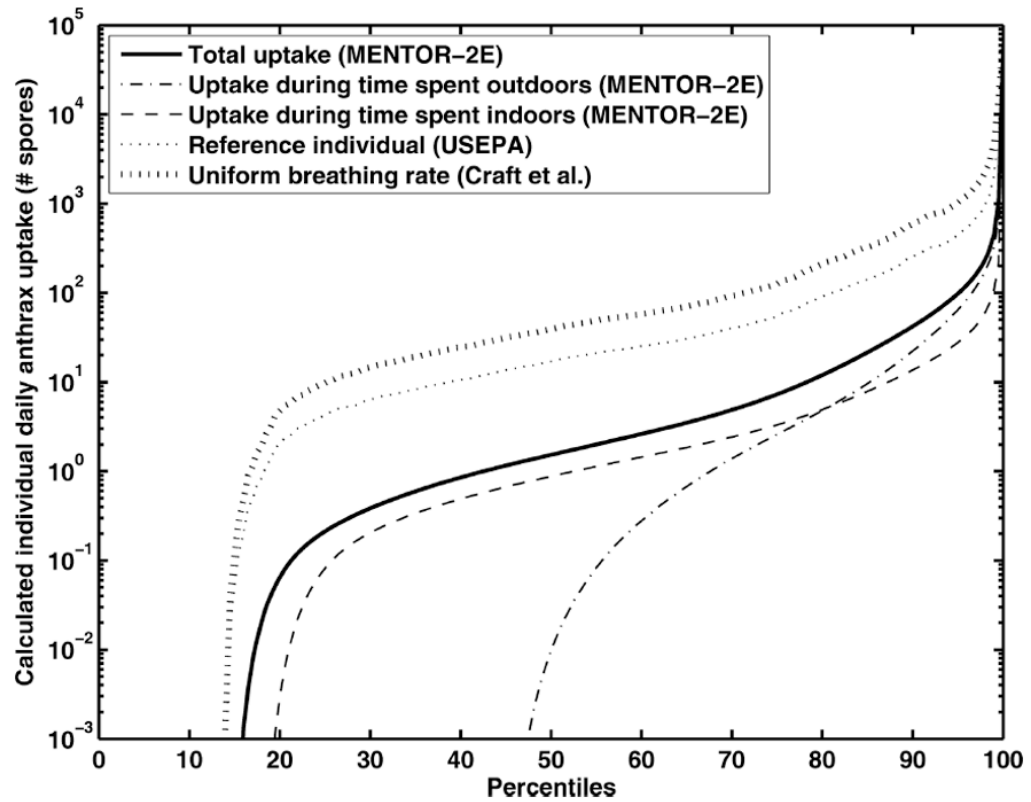


Fig. 8. Comparison of individual biological doses of *B. anthracis* estimated via different approaches. Percentiles of individual doses are shown for the entire study population for release scenario A (100 grams *B. anthracis* released over one hour).

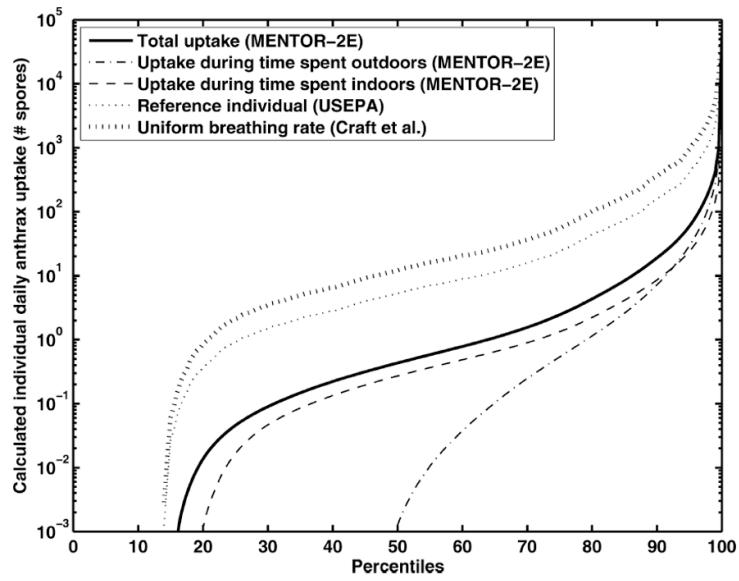


Fig. 9. Comparison of individual biological doses of *B. anthracis* estimated via different approaches. Percentiles of individual doses are shown for the entire study population for release scenario B (100 grams *B. anthracis* released over 10 hours).

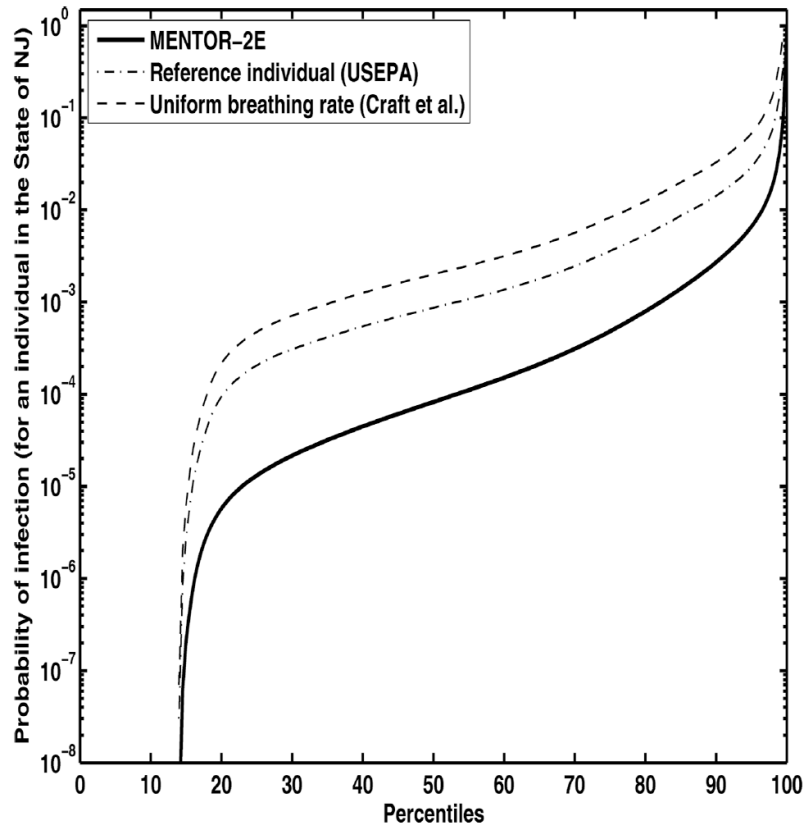


Fig. 10. Comparison of probabilities of anthrax infection estimated via different approaches. Percentiles of probabilities of infection are shown for the entire study population for release scenario A (100 grams *B. anthracis* released over one hour).

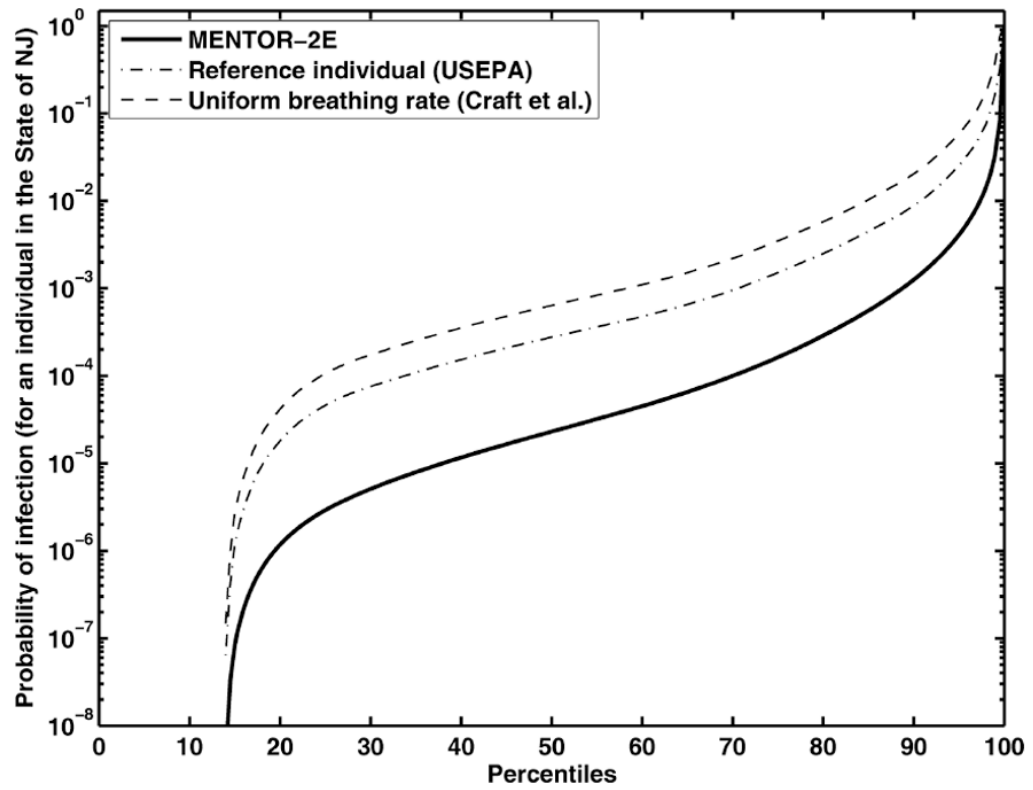


Fig. 11. Comparison of probabilities of anthrax infection estimated via different approaches. Percentiles of probabilities of infection are shown for the entire study population for release scenario B (100 grams *B. anthracis* released over 10 hours).

Table I

Sources and Values of Parameters Used in the MENTOR-EE Application for Simulating Impact of Hypothetic Releases of Anthrax Spores

Parameter	Value and/or Source
Number of anthrax spores released (hypothetical releases over 1 h and 10 h)	1015 [32]
Start time for the simulation (coinciding with mailing time of anthrax letters)	9/18/2001 08:00 h [2]
Location of the release (coordinates calculated using information from Reference 2)	Lat: 40.277; Lon: -74.813 [2]
Variable wind speed (obtained for Mercer County airport for 9/18/2001); the simplified dispersion case used a constant 1.5 m/s blowing toward north-east 0–3 m/s (Reference 66)	0–3 m/s [66]
CALPUFF grid resolution (general guidance from Reference 67)	250 m × 250 m [67]
Number of census tracts modeled (for application to the State of New Jersey)	1944 [47]
Census tract level population distribution (age; gender; occupation)	[47]
Virtual individuals per census tract	500 [37]
Activity patterns/metabolic equivalents (consolidated human activity database)	[49]
Distributions of parameters for residential microenvironments (air exchange rates; housing volumes; penetration and deposition factors; a total of 10 parameters)	Table II in [55]
Distributions of air exchange rates for general nonresidential microenvironments	[56]
Distributions of air exchange rates for vehicle microenvironments	[57]
Distributions for body weights (function of age and gender)	[37]
Empirical factors for deposition of anthrax spores in the lung (“uptake”)	[37,61,62,69]
Parameters for empirical dose-response models (details in Step 8 in Section 4)	[9,29,32,41]

Note: Further details on the exposure parameters are available in Reference 68.

Table II
 A Comparison of the Estimates of Probable Infections Derived Through Different Approaches for Exposure and Dose-Response Modeling

	Dose-Response Model					
	Model 1	Model 2	Model 3	Model 4	Model 5	Model 6
Scenario A; Craft <i>et al.</i> (simplifying assumption regarding breathing rates and location)	166,750	119,910	719,160	161,930	172,230	141,310
Scenario B; Craft <i>et al.</i> (simplifying assumption regarding breathing rates and location)	140,660	109,940	525,920	140,910	145,650	124,870
Scenario A; EPA EFH	85,669	56,020	485,770	75,747	88,549	70,466
Scenario B; EPA EFH	71,954	52,118	355,560	70,046	74,994	61,247
Scenario A; MENTOR-2E	25,013	15,631	187,840	21,059	25,908	20,472
Scenario B; MENTOR-2E	19,747	12,950	129,420	17,416	20,247	16,458
Scenario A/B; Craft <i>et al.</i> (simplifying assumptions regarding breathing rates, location, and plume dispersion)	1,180,932	1,090,121	1,076,816	1,049,390	1,144,563	1,187,637

Note: Total number of probable infections in study area are shown for the two release scenarios (100 g of anthrax released over (A) 1 h and (B) 10 h). The columns correspond to the different dose-response models used, and the rows correspond to different exposure modeling approaches used.