Modelling the spread, detection and control of exotic and endemic plant pests

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National Fruit Fly Council webinar, 6th September 2021

Overview

Background on the AADIS animal disease model The APPDIS pest modelling framework Case study 1: Oriental fruit fly Case study 2: yellow crazy ant



Australian Government

AADIS origins



- Department of Agriculture, Water and the Environment
- Department of Agriculture, Water and the Environment funded a project (2012-2015) to develop a national-scale model of emergency animal disease.
- The outcome was AADIS an agent-based model that can simulate the introduction of virus into any herd in Australia at any time.
- AADIS simulates:
 - -> spread
 - -> detection
 - -> control
 - -> 'proof' of freedom
 - -> resourcing & costing
- www.aadis.org.au





Bradhurst et al (2015). A hybrid modelling approach to simulating foot-and-mouth disease outbreaks in Australian livestock. Frontiers in Environmental Science, 3(17).

AADIS-FMD (foot-and-mouth disease)



How might a model inform policy?

- A simulation model such as AADIS allows very specific questions to be posed:
 - what are the benefits of early detection?
 - what are the consequences of late detection?
 - what resources needed for a big outbreak?
 - how will resource shortfalls impact control?
 - are the controlled areas the right size?
 - what is the impact of illegal movements?
 - will suppressive vaccination help?
- Outbreak & policy experiments are conducted via user dials, i.e., no need for specialist model reformulation or programming.







Australian Government

Department of Agriculture,

Water and the Environment

AADIS collaborators

AUSTRALIAN ANIMAL DISEASE SPREAD MODEL







CSIRO





United States

Department of

Agriculture



Canada





Ministry for Primary Industries Manatū Ahu Matua

Biosecurity

Queensland

Queensland

Government



Doherty Institute

Department of Primary Industries and **Regional Development** GOVERNMENT OF WESTERN AUSTRALIA

AUSVET



VETERINARY & AGRICULTURAL SCIENCES

AADIS: evolution

- contagious livestock diseases (e.g., foot-and-mouth disease)
- insect vector-borne livestock disease (e.g., bluetongue)
- disease in both domestic & wild populations (e.g., African swine fever)
- plant and environmental pests (e.g., tramp ants, fruit flies)
- other countries (Europe (EuFMDiS), USA, Canada, NZ)
- classroom tool (Australia, Europe, Malaysia)
- student projects (honours, Masters, PhD)



Expansion of AADIS to model the spread and control of pest species

- APPDIS (Australian Priority Pest & Disease model) arose through a DAWE/CEBRA project (2018/19).
- Whereas AADIS is (herd) agent-based, APPDIS is a geographic automata where a user-configurable grid overlays a study area. The modelling unit of interest is the cell.
- A cell can have a range of biotic/abiotic attributes.
- A cell may also have a population of the pest under study.

Bradhurst et al (2021). A generalised and scalable framework for modelling incursions, surveillance and control of plant and environmental pests. Environmental Modelling & Software, 139.

Geographic automata

File Control Navigate Layers Config Database Reports

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model status scenario name: bt_demo seeding mode: endemic scenario end mode: fixed		~\					-	
scenario max length: 585 days spread pathways: Jump Diffusion fixed start date: Thu. 01 Sep. 2016 current date: Thu. 01 Sep. 2016								
elapsed real time: 0 secs scenario status: initialised sim day: 0 (0.0 sim years) run number: 1 of 1 convergence (IPs/duration/cost): 0% / 0% / 0%								
detection mode: disabled detection state: silent spread infected: NT controlled:								Byron Bay
standstill: mpacted:								cell ID: 91594
								north boundary lat: -28.658 south boundary lat: -28.748 west boundary long:153.513
ulicoides presence um Culicoides suitable cells: 57515 um Culicoides free cells: 63150								east boundary long: 153.605 cell height: 10.00 km cell width: 8.93 km
um Culicoides quiescent cells: 0 um Culicoides active cells: 21329 um BTV infected cells: 0								cell area: 8,929.53 ha (89.30 km2) nearest weather station ID: 58216 ► region: 5
num undefined transmissions: 0 num Culicoides diffusion transmissions: 0 num Culicoides jump transmissions: 0								Culicoides suitability: raw=44.52 trans=0.74 mean weekly temperature (celsius): 15.89 mean monthly temperature (celsius): 17.48
								annual raintail (mm): 2,141 mean weekly wind speed (km/hr): 26.73 elevation (m): 88
cell 91594 n=0.37 s=1.00 e=0.00 i=0.00 r=0.00 t=15.9 β=0.	000 🖨 🔳 🧯							human population: 0.0000 (0.0000) land use: undefined seed cell with Culicoides ►
р 0.74								infect cell with BTV ► display population curves ► initial population: 12935780 (0.3696)
r e v a								current population: 12935780 (0.3696) max population: 25871559 (0.7392) population state: active
					Grafton			quiescence days: infestation pathway: endemic infestation days: 0
0 110 180 360	540							
sin uay								

Lat, Lon (-28.714, 153.562) - x, y (1060,293)

Modelling pest populations with a geographic automata

- how is the pest distributed? (i.e., which cells are initially populated?)
- how does within-cell abundance change over time?
- when/how might the population spread between cells?
- how effective are surveillance activities at detecting cell populations?
- how effective are treatment programs at controlling/eradicating cell populations?
- how effective are surveillance activities at detecting residual populations?
- how do resource constraints affect surveillance & control effectiveness?

Initial pest distribution

APPDIS provides three methods to distribute a pest:

- 1) <u>point incursion</u> one or more cells can be explicitly seeded with a pest propagule, e.g., post-border arrival of an exotic fruit fly at a port.
- <u>established population</u> externally generated population densities/counts per cell, perhaps informed by trapping data, census data, or habitat suitability studies.
- 3) <u>built-in species distribution model</u> (SDM) mechanistic estimation based on configurable ranges of environmental criteria such as temperature, vegetation, elevation, rainfall, land use, etc.

Distribution of feral pigs via an external habitat model

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Distribution of *Culicoides brevitarsis* via the built-in SDM

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Within-cell abundance

- APPDIS provides two ways of estimating within-cell pest abundance over time:
- 1) mathematically e.g., temperaturedependent logistic growth with carrying capacity based on cell habitat suitability
- 2) externally generated 'time slices' that specify each cell's population count at a configurable interval





Natural dispersal between-cells

- As the pest population in a cell increases so too does the dispersal pressure to other cells.
- Natural short-range dispersal (e.g., tramp ant budding, fruit fly unaided flight) is modelled via a stochastic spatial diffusion kernel. Spread depends on the source cell pest density and destination cell suitability
- Optional dependencies include land use, temperature, elevation, gradient, wind speed & rainfall.

Assisted dispersal between cells

- Sporadic longer-range dispersal (e.g., hitchhiking, rafting, nuptial flight, specific market/agricultural activities) is modelled via one or more stochastic jump processes.
- Spread depends on the source cell pest density & destination cell suitability.
- Optional dependencies include human population density, land use, watercourses, temperature, elevation, gradient, wind direction/speed, rainfall.

Jump-diffusion spread of *Culicoides brevitarsis*



Detection and control

- Once the spread mechanisms have been calibrated to the satisfaction of domain experts the next step is configuring detection & control mechanisms.
- APPDIS simulates:
 - \rightarrow general surveillance
 - \rightarrow early detection surveillance (via a permanent trapping grid)
 - \rightarrow delimiting surveillance in response to detections
 - \rightarrow treatment programs in response to detections
 - \rightarrow post-treatment surveillance (to assess absence/presence)
- Control actions are user-configurable with respect to mode, extent, periodicity, cost & effectiveness.
- Response imperfections are included e.g., false positive reports, resource constraints, hard-to-detect residual populations, escapes from managed areas.

Case Study 1 Oriental fruit fly

- National grid with 5x5 km² cells.
- Point introduction an oriental fruit fly propagule (arbitrarily n=25) at the Port of Cairns.
- Within-cell abundance:
 - \rightarrow temperature dependent logistic growth
 - → carrying capacity determined by suitability layer (land use and NDVI)
- Spread: natural dispersal (diffusion) + human-mediated hitchhiking (jumps)
- Surveillance and control:
 - \rightarrow general surveillance
 - → early detection surveillance (trapping grid)
 - → delimiting surveillance (50km radius)
 - → treatment (10km radius)
 - \rightarrow post-treatment surveillance

Detection and control of Oriental fruit fly

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Risk map of Oriental fruit fly spread after introduction in Cairns



Case Study 2: yellow crazy ant

- Regional grid with 10 ha cells
- Established population across 1540 ha
- Within-cell abundance:
 - \rightarrow temperature independent logistic growth
 - ightarrow very simple suitability layer
- Spread: budding (diffusion) + hitchhiking, agricultural & rafting jumps
- Surveillance and control:
 - \rightarrow general surveillance
 - \rightarrow delimiting surveillance
 - \rightarrow treatment
 - \rightarrow post-treatment surveillance

Control of yellow crazy ant



Lat, Lon (-17.253, 146.227) - x, y (1338,633)

Post-treatment surveillance of yellow crazy ant



- The post-treatment surveillance trap spacing was systematically varied between 2 m and 100 m
- The model suggests that an 18 m spacing was the most cost-efficient.

Risk map of yellow crazy ant spread after 30 years



Acknowledgements



Australian Government

Department of Agriculture, Water and the Environment



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