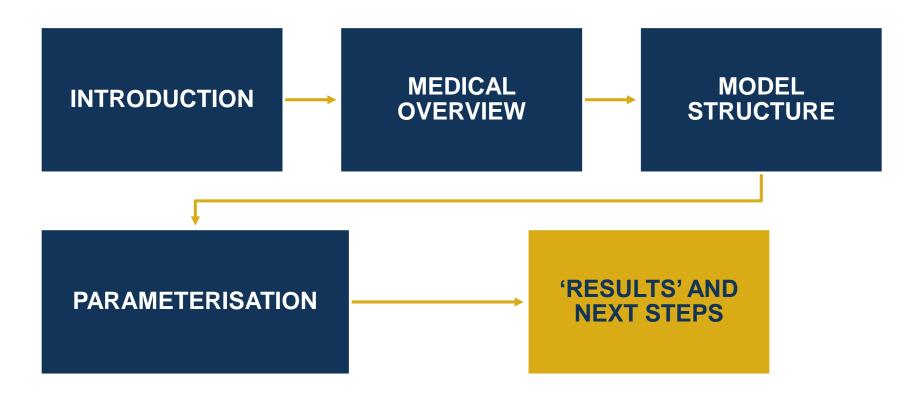


When the drugs don't work...

Matthew Edwards, Nicola Oliver and Ross Hamilton (IFoA Antibiotic Resistance Working Party)

Agenda





Working party background

ABR Event Staple Inn May 2016



Institute

and Faculty

- Develop a simple modelling framework with plausible parameterisation to allow actuaries to develop their own views on likely and stress mortality impacts
- This framework would be developed in a UK context but would be expected to be readily transferable to other countries
- Working party started in January 2017

Working party members

Name	Role	Firm
Matthew Edwards	Chair	Willis Towers Watson
Nicola Oliver	Medical input & Deputy Chair	Medical Intelligence
Sheridan Fitzgibbon	Model structure & parameterisation	Legal & General
Craig Armstrong	Parameterisation (2017)	Aviva
Ross Hamilton	Model development	Lloyds Banking Group
Irene Merk	General	SCOR
Roshane Samarasekera	Model development	GAD
Soumi Sarkar	General	Legal & General
Katherine Fossett	General	Barnett Waddingham



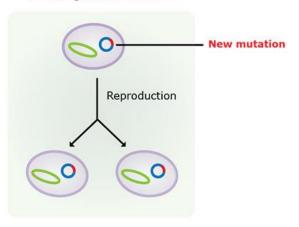


Medical overview

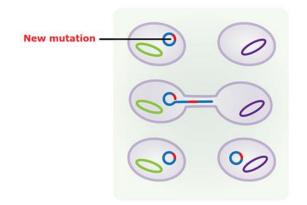
What is antibiotic resistance...

Normal bacteria Resistant bacteria How antibiotic resistance occurs Dead bacteria 2. 3. 1. 4. Bacteria cells in the Antibiotics kill Antibiotic resistant Antibiotic human body. Some are bacteria but resistant bacteria multiply resistance spreads drug resistant strains remain

Vertical gene transmission

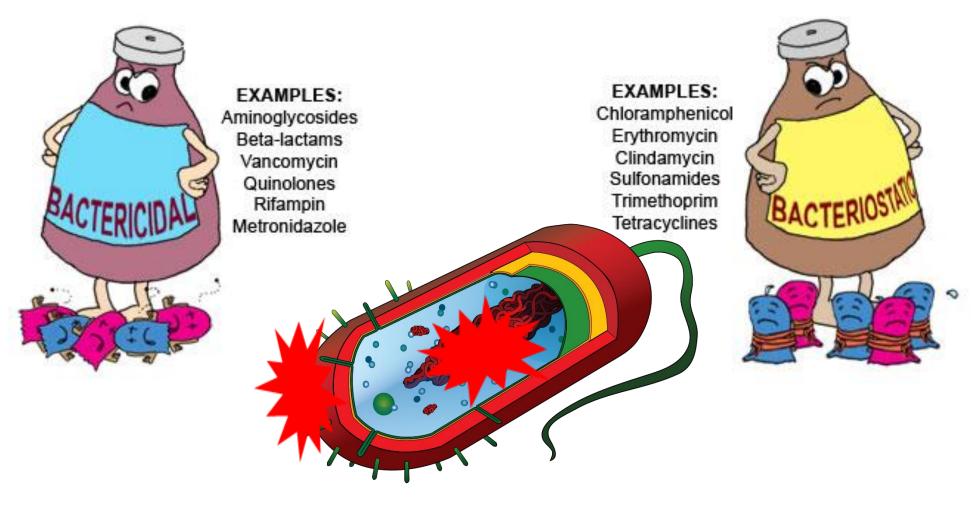


Horizontal gene transmission



"The thoughtless person playing with penicillin treatment is morally responsible for the death of the man who succumbs to infection with the penicillin-resistant organism." Sir Alexander Fleming, 1928



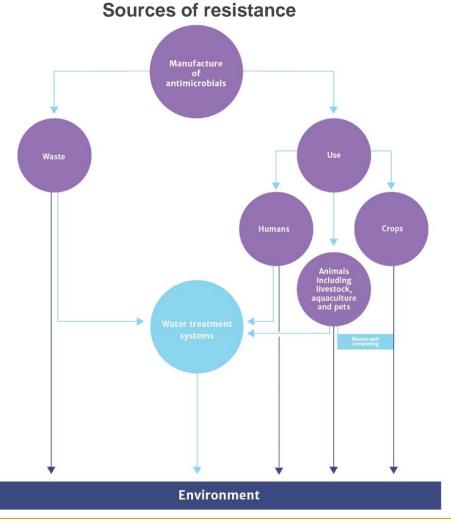


How do antibiotics work? (the science!)

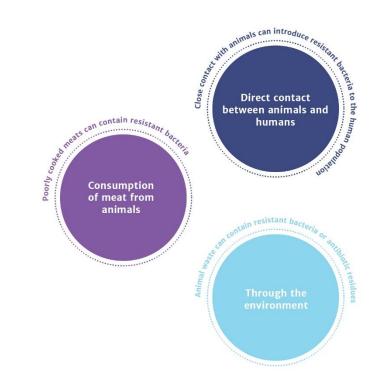


What are the sources of resistance?

Correct of resistance



How animals can pass on resistant bacteria

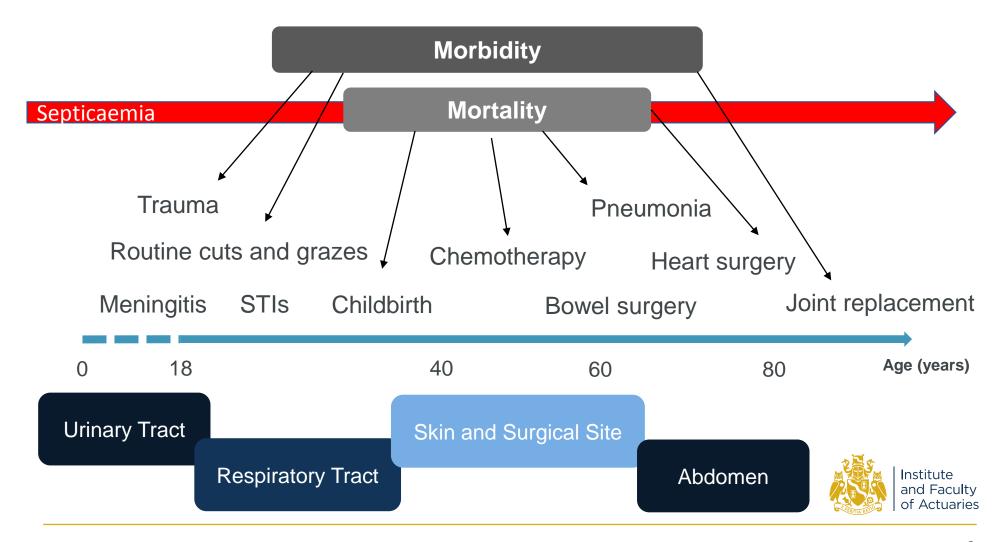


Infographics sourced from "Review on Antimicrobial Resistance" 2014



8

How does ABR affect people and our work?





Discovery, research, and development of new antibiotics: the WHO priority list of antibiotic-resistant bacteria and tuberculosis

Evelina Tacconelli, Elena Carrara", Alessia Savoldi", Stephan Harbarth, Marc Mendelson, Dominique L. Monnet, Céline Pulcini,
Gunnar Kahlmeter, Jan Kluytmans, Yehuda Carmeli, Marc Ouellette, Kevin Outterson, Jean Patel, Marco Cavaleri, Edward M. Cox, Chris R. Houchens,
M. Lindsay Grayson, Paul Hansen, Nalini Singh, Ursula Theuretzbacher, Nicola Magrini, and the WHO Pathogens Priority List Working Group†

Criteria

Mortality

Health-care burden

Community burden

Prevalence of resistance

10-year trend of resistance

Transmissibility

Preventability in the community

Preventability in health-care setting

Treatability

Pipeline

"The major objective of the global priority pathogens list (global PPL) is to guide the prioritization of incentives and funding, help align R&D priorities with public health needs and support global coordination in the fight against antibiotic-resistant bacteria"



Priority 1: CRITICAL#

Acinetobacter baumannii, carbapenem-resistant

Pseudomonas aeruginosa, carbapenem-resistant

Enterobacteriaceae*, carbapenem-resistant, 3rd generation cephalosporin-resistant

Priority 2: HIGH

Enterococcus faecium, vancomycin-resistant

Staphylococcus aureus, methicillin-resistant, vancomycin intermediate and resistant

Helicobacter pylori, clarithromycin-resistant

Campylobacter, fluoroquinolone-resistant

Salmonella spp., fluoroquinolone-resistant

Neisseria gonorrhoeae, 3rd generation cephalosporin-resistant, fluoroquinolone-resistant

Priority 3: MEDIUM

Streptococcus pneumoniae, penicillin-non-susceptible

Haemophilus influenzae, ampicillin-resistant

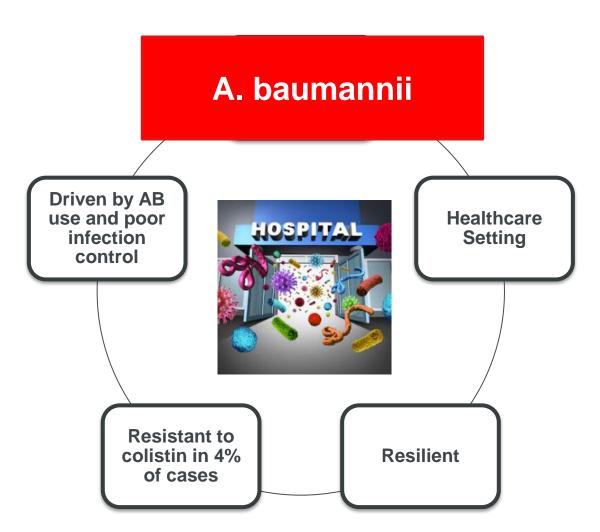
Shigella spp., fluoroquinolone-resistant

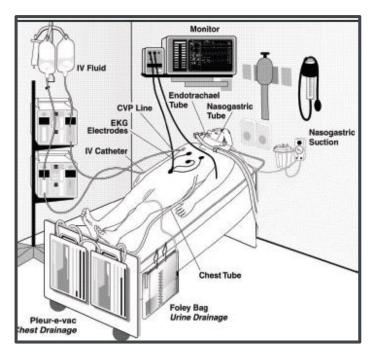
A. baumannii

Pseudomonas

Enterobacteriaceae







Pneumonia

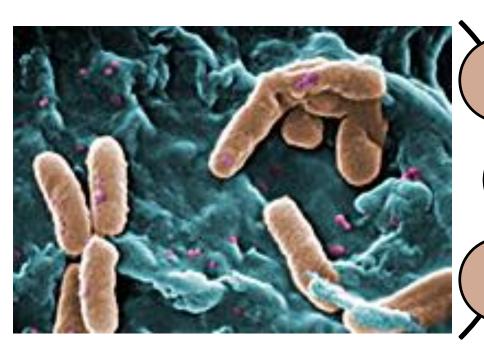
Bloodstream Infection

Wound Infection

Urinary Tract



Pseudomonas



Found widely in the environment

Common cause of mild and serious infections

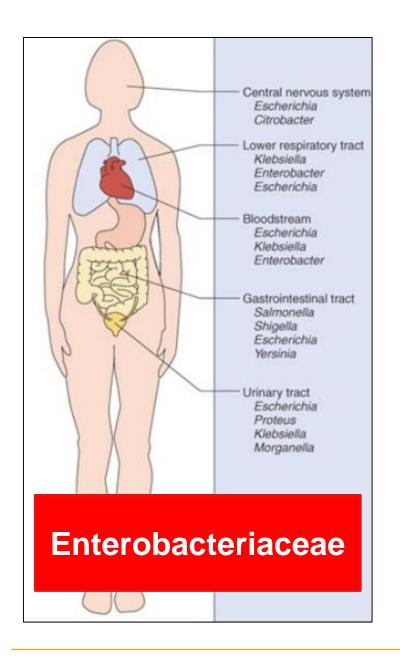
Risk profile similar to A. Baumannii

Pneumonia

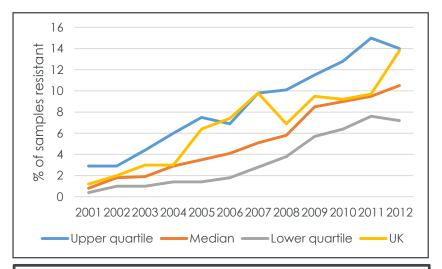
Wound Infection

Bloodstream Infection





These bacteria are associated with higher frequency of inappropriate antimicrobial therapy, poorer clinical response, and longer length of hospital stay

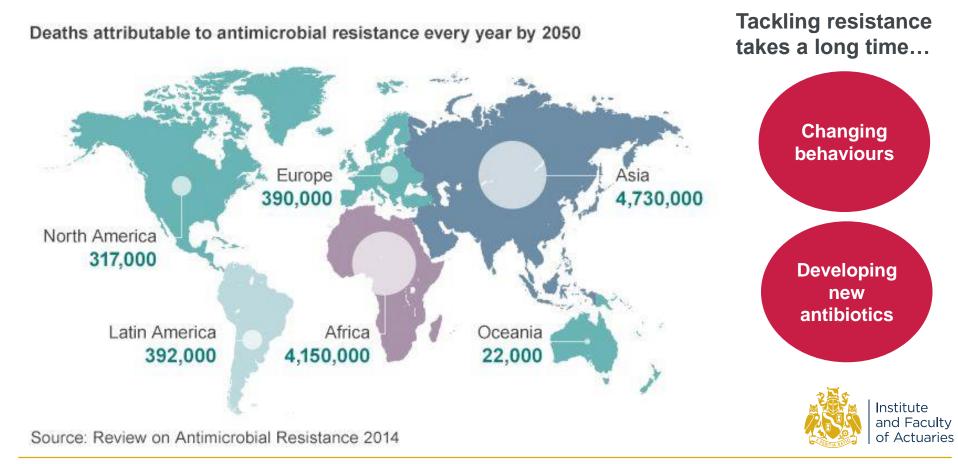


Third-generation cephalosporin resistance rates in *E. coli* across Europe, showing the UK, 1999 to 2012 (Department of Health, 2015)



...and why it is important?

"We have reached a critical point and must act now on a global scale to slow down antimicrobial resistance" – Professor Dame Sally Davies, UK Chief Medical Officer





Global increase and geographic convergence in antibiotic consumption between 2000 and 2015

Eili Y. Klein^{a,b,c1}, Thomas P. Van Boeckel^d, Elena M. Martinez^a, Suraj Pant^a, Sumanth Gandra^a, Simon A. Levin^{e,f,g,1}, Herman Goossens^h, and Ramanan Laxminarayan^{a,f,i}

aCenter for Disease Dynamics, Economics & Policy, Washington, DC 20005; Department of Emergency Medicine, Johns Hopkins School of Medicine, Baltimore, MD 21209, "Department of Epidemiology, Johns Hopkins Bloomberg School of Public Health, Baltimore, MD 21205; "Institute of Integrative Biology, ETH Zürich, CH-8006 Zürich, Switzerland; "Department of Ecology and EV

Environmental Institute, Princeton University, Princeton, NJ 08544; 9Beijer Institute Medical Microbiology, Vaccine & Infectious Diseases Institute, University of Antwer of Washington, Seattle, WA 98104

Contributed by Simon A. Levin, February 23, 2018 (sent for review October 3, 201









ດ ≋ ≗ ≡າbinations thwart efforts to curb researchers

Antibiotic-resistant gonorrhoea cases expected to emerge worldwide

Warnings after UK man and two Australians suffer STI untreatable with usual drugs

Sally Wardle | Friday 20 April 2018 18:39 BST | D 10 comments



















Culture-independent discovery of the malacidins as calcium-dependent antibiotics with activity against

multidrug-resistant Gram-posi

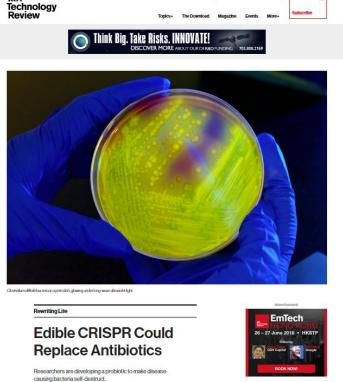
Berglund et al. Microbiome (2017) 5:134 DOI 10.1186/s40168-017-0353-8

Microbiome

Open Access

CrossMark

Bradley M. Hover¹, Seong-Hwan Kim¹, Micah Katz¹, Zachary Melinda A. T MIT and Sean F. F. MIT Technology



RCH

ification of 76 novel B1 metallo-βmases through large-scale screening nomic and metagenomic data

und^{1,2}, Nachiket P. Marathe^{2,3}, Tobias Österlund^{1,2}, Johan Bengtsson-Palme^{2,3}, Stathis Kotsakis^{2,3}, Flach^{2,3}, D G Joakim Larsson^{2,3} and Erik Kristiansson^{1,2*}







Ross Hamilton

Objectives & Research

Define Objectives

- Model ABR impact on:
- Mortality
- Morbidity

Literature Review

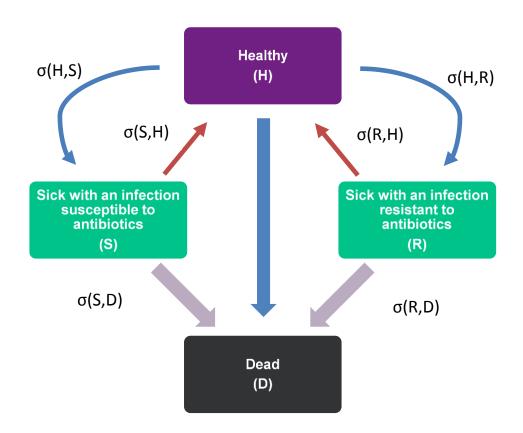
- KPMG / RAND model
- Research papers

Model Structure

- Complex enough to model scenario
- Not overly complex
- Capable of being adapted by users



Chosen model structure



Modelling criteria

- Simplicity
- Availability of data
- Appropriate outputs

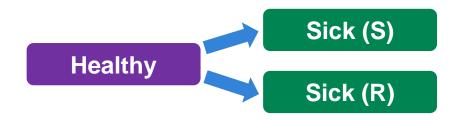
Basic structure decided on:

- Multi-state Markov model
- Calibrate to current observed levels of mortality and morbidity
- Project varying resistance over time and calculate the change in mortality and morbidity



Data sources - incidence

Incidence rates for bacteraemia







Limitations

- Limited data. E. coli monitoring in England goes back to 2013.
- Limited evidence for how resistance interacts with incidence.
- Bias? Monitoring is of HCAIs.



Data sources – mortality

Death rates for bacteraemia

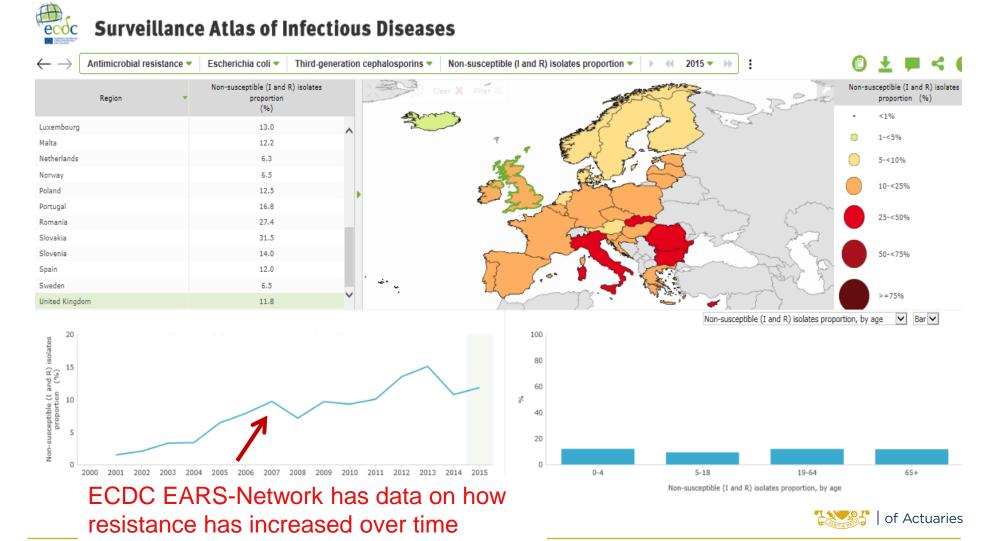


Limitations

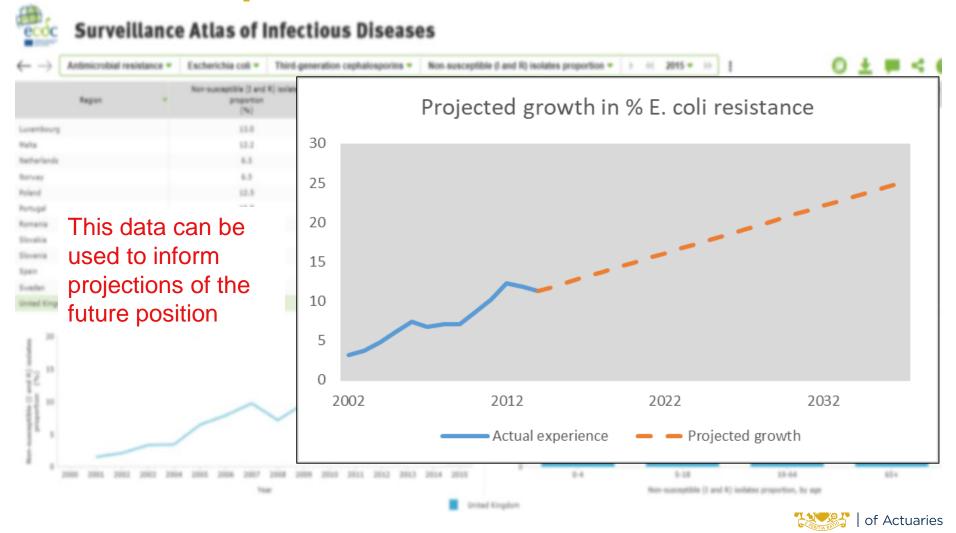
- Granularity of data:
 - Confounding causes of death?
 - Academic literature is helpful here.
- Large error bounds around estimates of the relative virulence of resistant and susceptible strains.
- Bias? The most ill are more likely to be sampled.



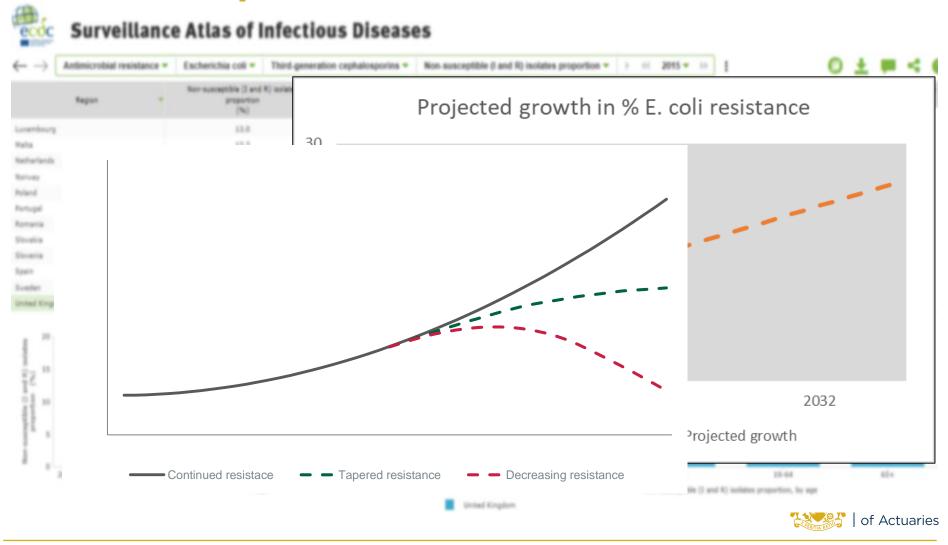
Trends in resistance can be observed...



...and extrapolated forwards

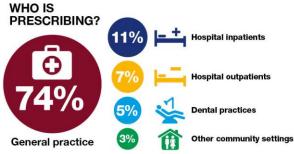


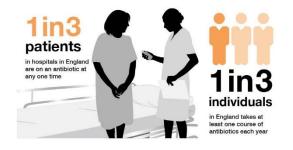
...and extrapolated forwards



Future of resistance?







30 years since a new class of antibiotics was last introduced....

Barriers to R&D Investment

Cautious optimism in 2 new compounds



Infographics sourced from "Review on Antimicrobial Resistance" 2014





Initial Results: E. coli resistance

Parametrisation based on:

- Growth in E. coli bacteria resistant to 3rd generation cephalosporin antibiotics
- Ages 19-64, i.e. working age population
- Projected position in 2037, i.e. 20 years' time

Results:

Central scenario

1% increase in mortality rate (qx) from one strain

Perhaps ~0.2% / 0.25% pa reduction to CMI model LTR?

Allowing for all main strains of bacteria

In a bad scenario (95% confidence level not 1-in-200), there could be a 10-20% increase in overall mortality (with all main strains)



Working party – next steps

Sessional meeting February 2019

Model development

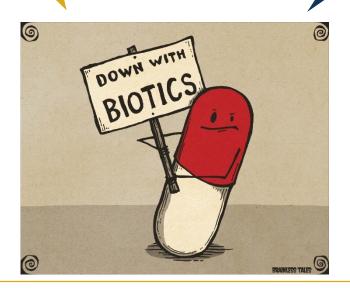
- Parameterisation other main bacteria (5)
- Interactions between pathogens
- Validation / Documentation

- Full model release
- Suggested parameterisation based on UK data
- Associated paper main issues relating to sources of ABR, mitigation actions, recent trends, other projection results / methodologies, and background to our model and results from the model



Questions

Comments



Expressions of individual views by members of the Institute and Faculty of Actuaries and its staff are encouraged.

The views expressed in this presentation are those of the presenter.

