

# One Health and Antimicrobial Stewardship – Why wildlife matter

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I acknowledge the Traditional Custodians of the land where Macquarie University is located, the Wallumattagal Clan of the Dharug nation, and the First Peoples of the lands where our studies were undertaken.

I appreciate that we can meet today on lands of the Wurundjeri, Woi-wurrung and Bunurong Boon Wurrung peoples of the East Kulan.



# Acknowledgements

## Bats

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- Paul Thompson - Taronga
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- David Gordon - ANU
- Sabine Schiller – Mres

## Marine species

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- Rachael Gray – U Syd
- Ida Lundback – Mres

## Tasmanian devils

- Samantha Fox - DPIPWE
- Carolyn Hogg - U Syd

## Koala

- Fiona McDougall
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- Natasha Speight - U Adl
- Tamsyn Stephenson - U Adl
- Wayne Boardman - U Adl
- Jen McLelland – Zoos SA
- Oliver Funnell – Zoos SA
- Ian Smith – Zoos SA
- KI Vet Hospital

## Platypus

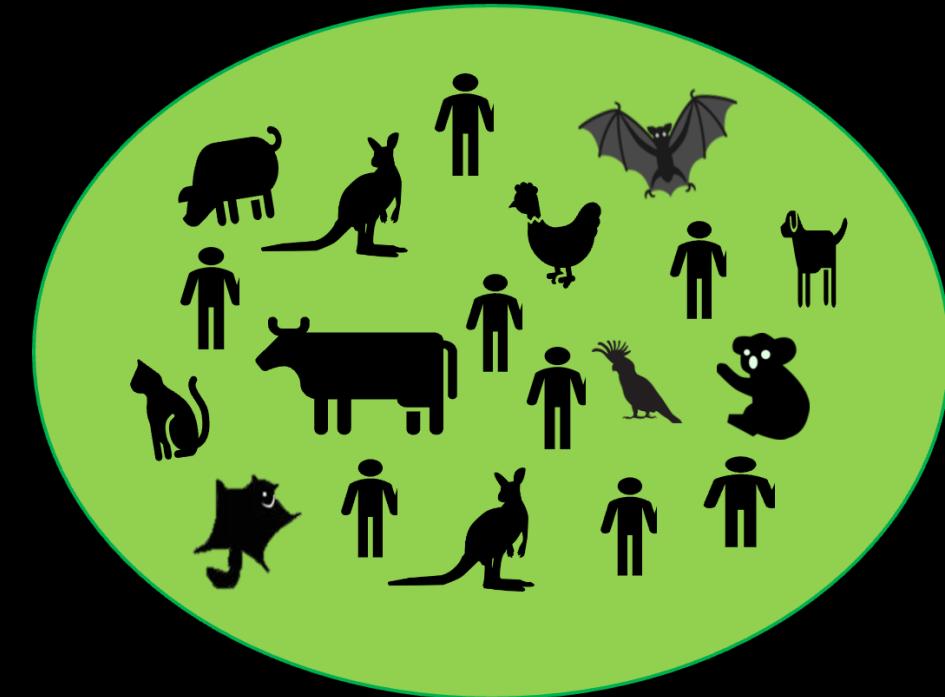
- Nadine Samy (Mres)
- Gilad Bino



# Antimicrobial resistance and wildlife

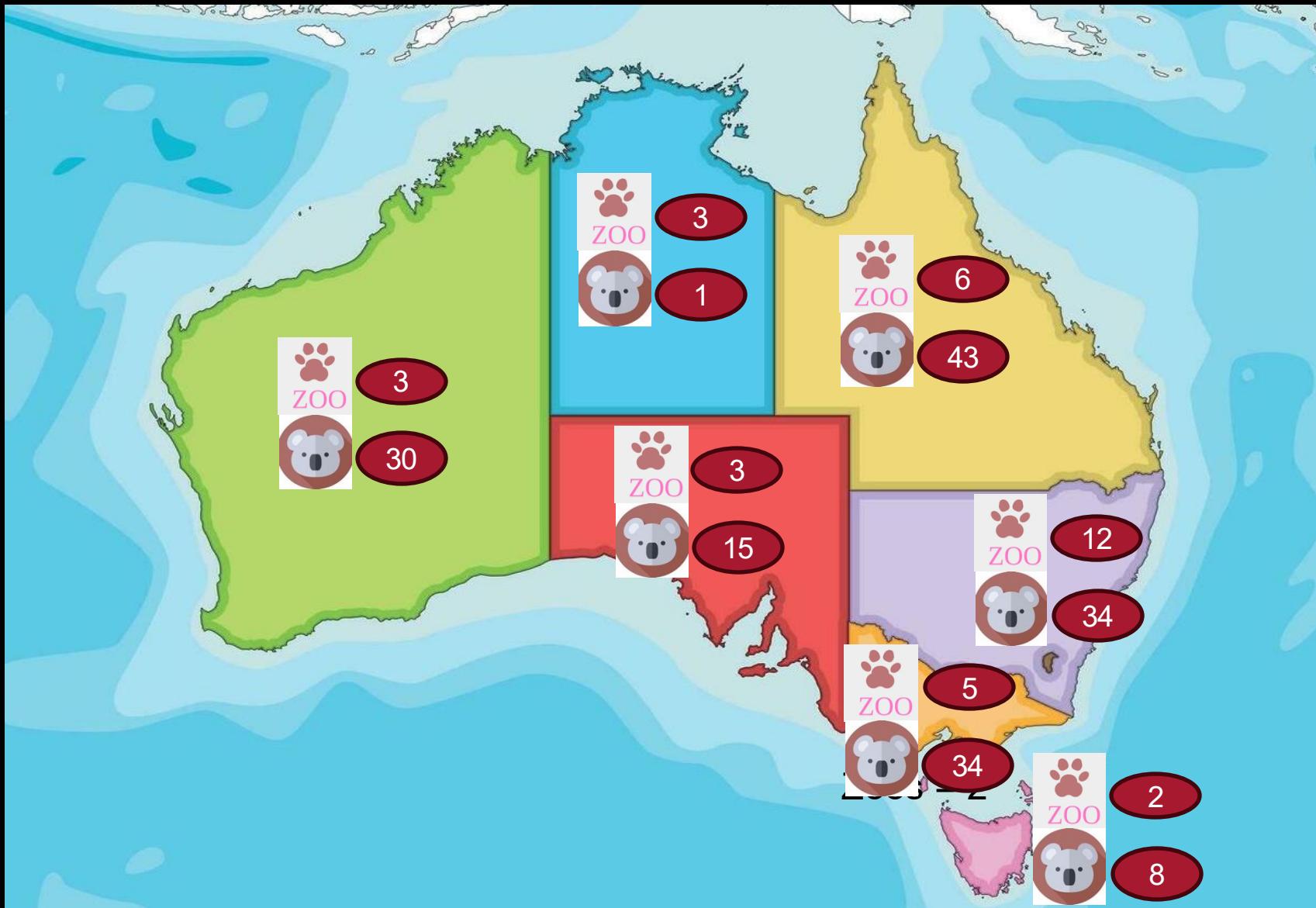
## AMR HAS NO SPECIES BOUNDARIES

- Detected in diverse wildlife
  - Species representative of all major phyla
  - Land to the sea
  - Remote and urban areas
- Sentiments of AMR in wildlife
  - Naturally occurring
  - Antibiotics not used to treat wild animals
  - No impact for wildlife



**Overlooks the significance of wildlife in AMR ecology, antimicrobial stewardship, and One Health**

# Australia's wildlife system – a snapshot!



## NSW 2021-2022

NSW Department of Environment report

Wildlife rescues	A second chance	People power
 <b>128,262</b> native animals across <b>543</b> species rescued	 <b>31,290</b> (24%) native animals released back to the wild	 <b>8,621</b> volunteers

## Wildlife Victoria - 2024

- 97,772 rescues
- 455 species

## WIRES - 2025

- 163,776 rescues
- 460 carers

# Investigating AMR in wildlife systems

## URBAN SETTINGS

Flying fox  
*Ngununy*



Possum  
*Wali*



Macropods *Badagarang*



## CONSERVATION

Tasmanian devil  
*Purinina*



Platypus  
*Boondabarra*



Koala  
*Gula*



Echidna  
*Barrugin*



## MARINE

Australian sea lion  
*Bulgurra*

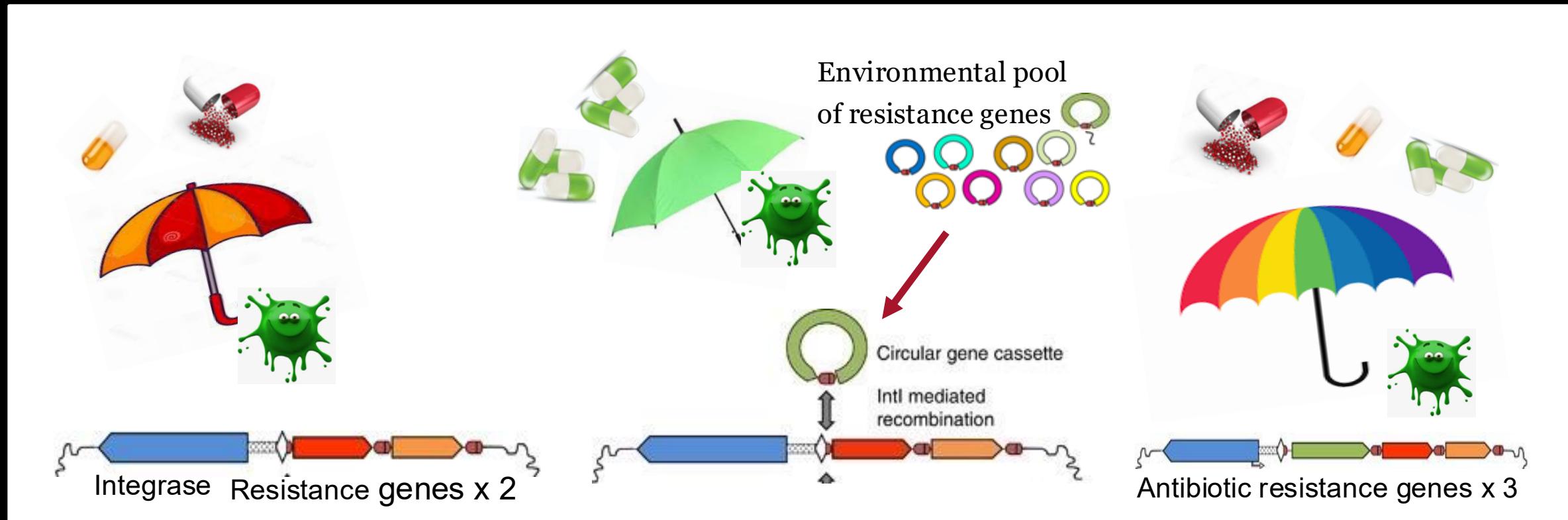


Little penguin  
*Gur-roo-mul*



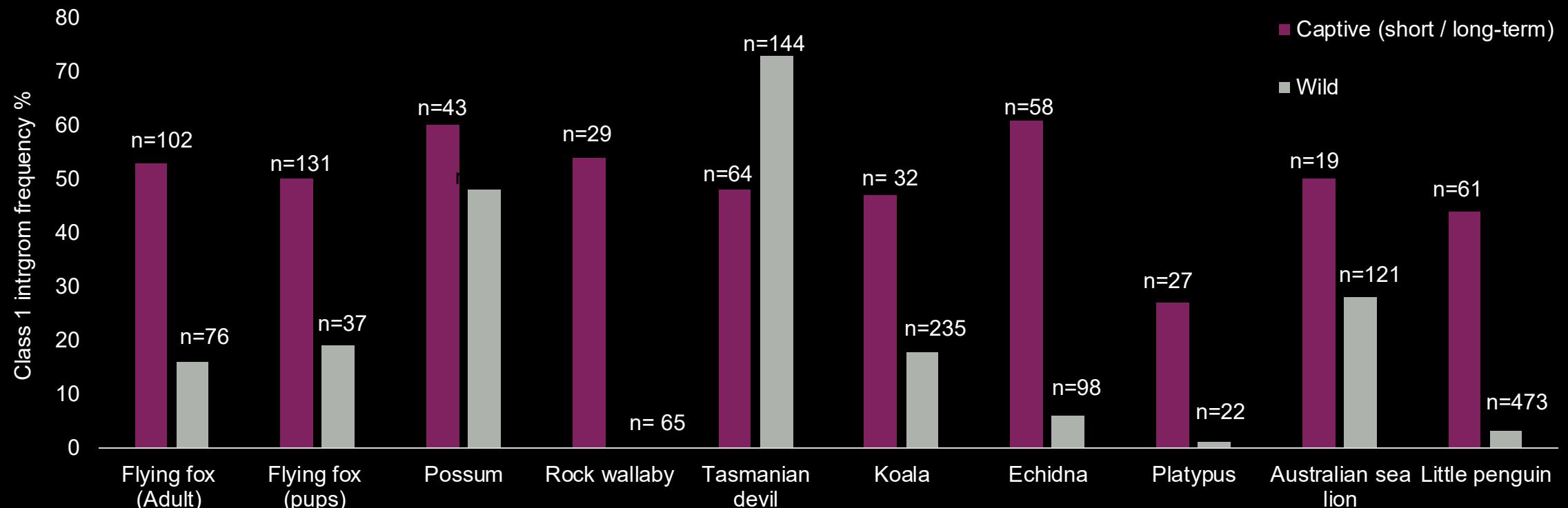
# Anthropogenic origins | Clinical class 1 integron

- Assembled under antibiotic selection pressure 100 years ago
- Present in >80 clinically important bacterial species
- Proposed as an indicator of human pollution (Gillings 2015)



# Anthropogenic origins | Clinical class 1 integron

## Clinical class 1 integrons - more frequent in wildlife rehabilitation and in close-contact to people



# Brushtail rock wallaby

*Petrogale penicillata*



Photo Credit: Hugh McGregor

# Conservation | dissemination risks

Endangered brush-tail rock wallabies (*Petrogale penicillata*):

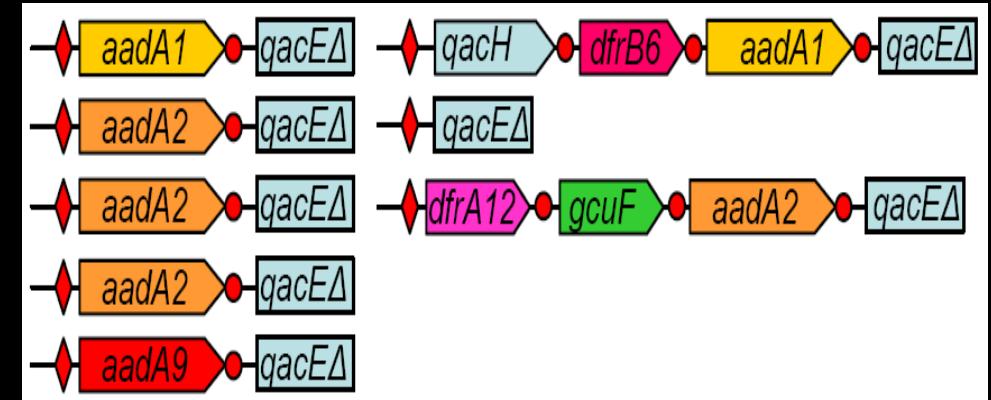
- Class 1 integrons detected in 48% of captive-bred wallabies
- No class 1 integrons were detected in the wild population



<https://www.mammalage.com>

## Post-release

- Risk spread of resistant bacteria to wild wallabies
- Dissemination into new environments



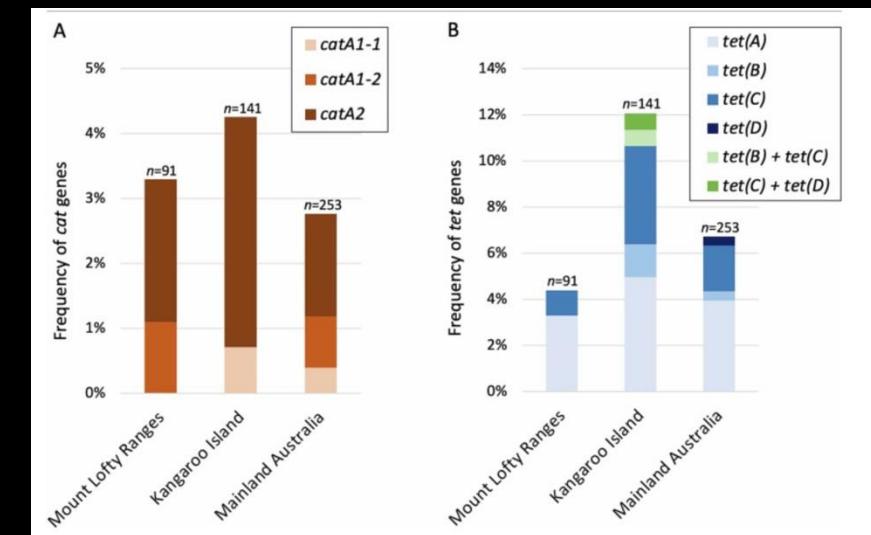
Koala  
*Phascolarctos cinereas*

Gula



## Koala chlamydial disease treatments - Chloramphenicol and Doxycycline LA

- Resistance genes detected in koalas
- Co-infections with *Chlamydia pecorum* and resistant bacterial strains
- Potential for emergence of resistant *Chlamydia pecorum* strains - Non-resolving chlamydial disease and treatment failure
- Risk of secondary infections



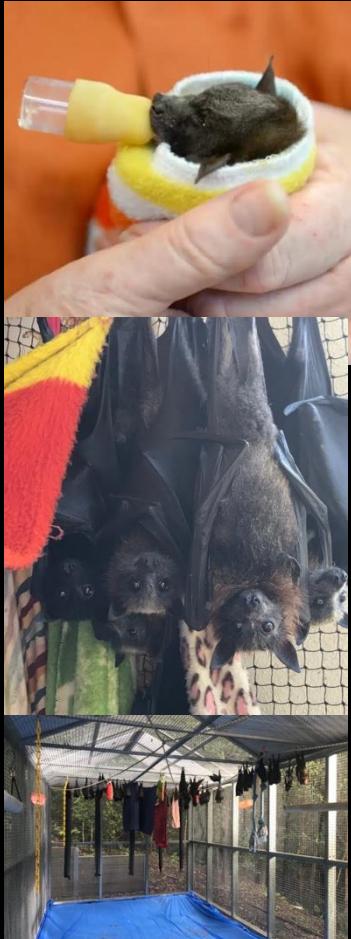
# Grey headed flying fox

*Pteropus poliocephalus*

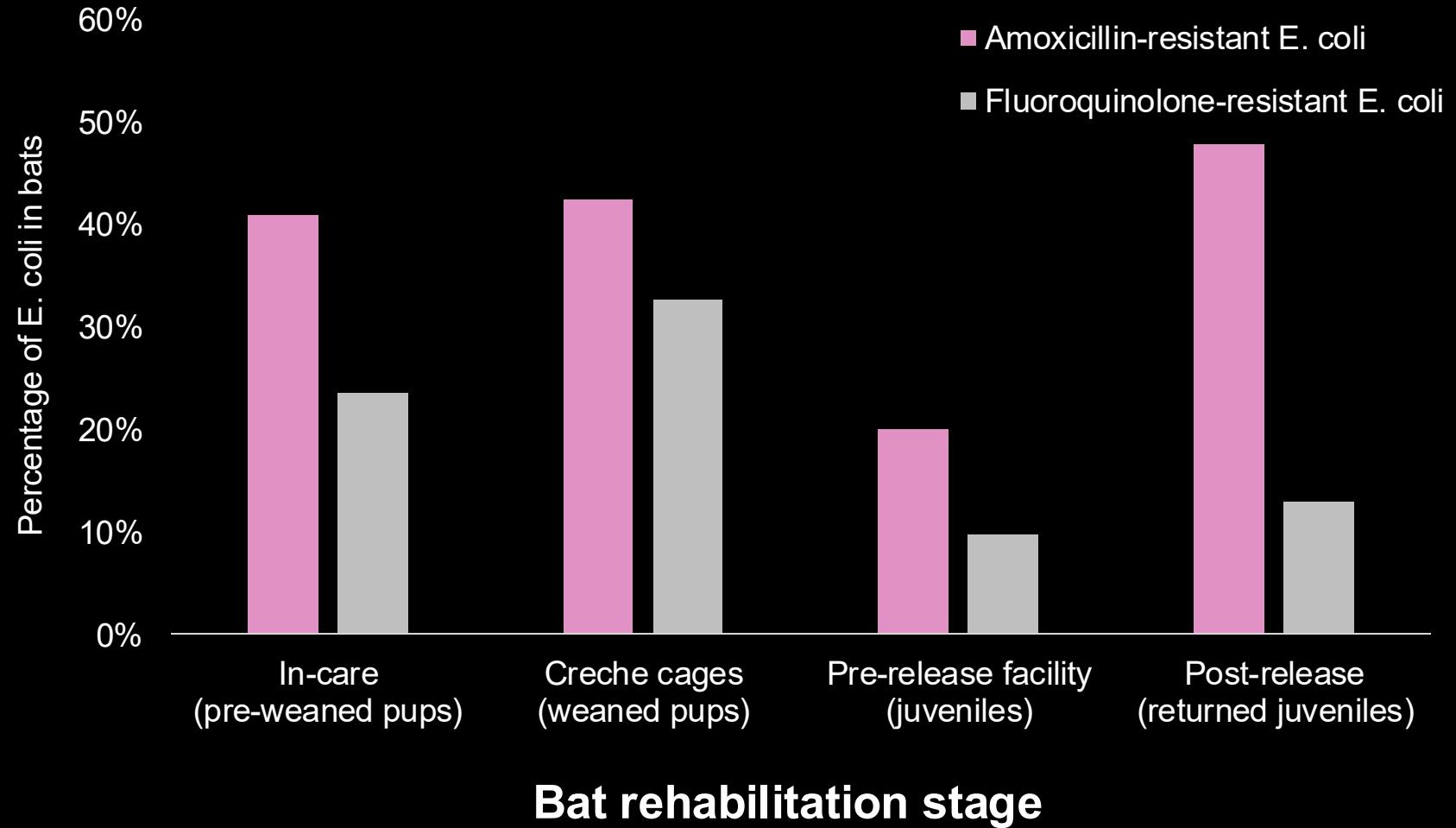
Ngununy



Image: Adam McKeown



Detected in 35.3% bat faecal samples ( $n=568$ )



# Wildlife rehabilitation | MDR, ESBL and carbapenem-resistant *E. coli*

48% *E. coli* isolates from bats multidrug-resistant

***E. coli* isolate FF1992A:**

*aph(3')-Ia, aph(6)-IId, aadA2, blaSHV-12, blaTEM-1B, dfrA14, dfrA19, qnrB2, qnrS1, sul1, sul2*

***E. coli* isolate FF2394:**

*aph(3')-Ib, blaCTX-M-15, dfrA1, qnrS1, sul2, tet(A)*



4% ESBL-producing (*bla*CTX-M-15)

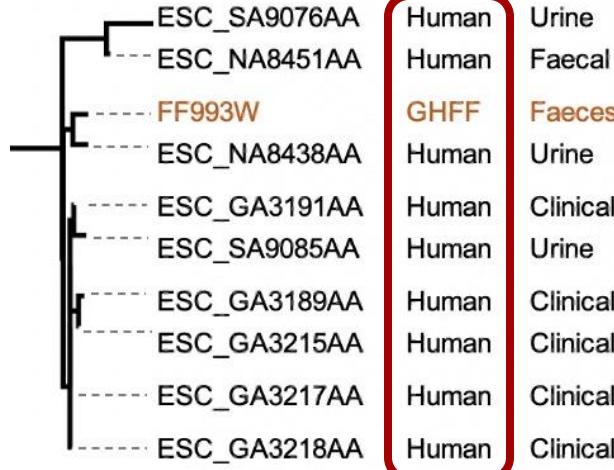
4% Carbapenem-resistant (*bla*OXA-23)

>95% antibiotic-resistant *E. coli* strains from bats are lineages associated with humans and/or domestic animals

Bat *E. coli* isolate ST10 O89:H9

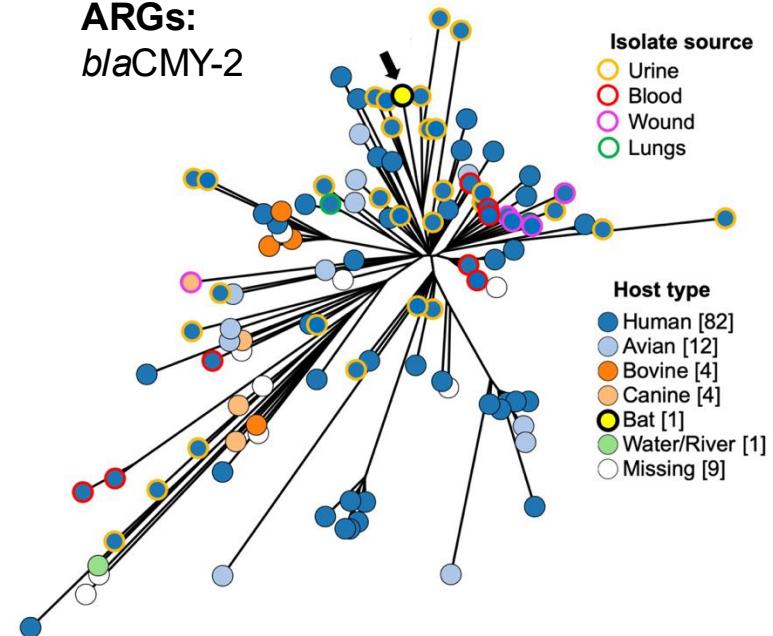
**ARGs:**

- blaNDM-5*
- blaCTX-27*
- blaTEM-1B*
- catA2*
- dfrA14*
- gyrA/parC/parE*
- sul2*
- Tet(A)*



Bat *E. coli* ST963 ONT:H18

**ARGs:**  
*blaCMY-2*



# Emergence risks | within host evolution

## STRAIN EMERGENCE

### *E. coli* ST38

Penicillins, chloramphenicol



Penicillins, fluoroquinolones, chloramphenicol

### *E. coli* ST131 Cx

Fluoroquinolones, trimethoprim



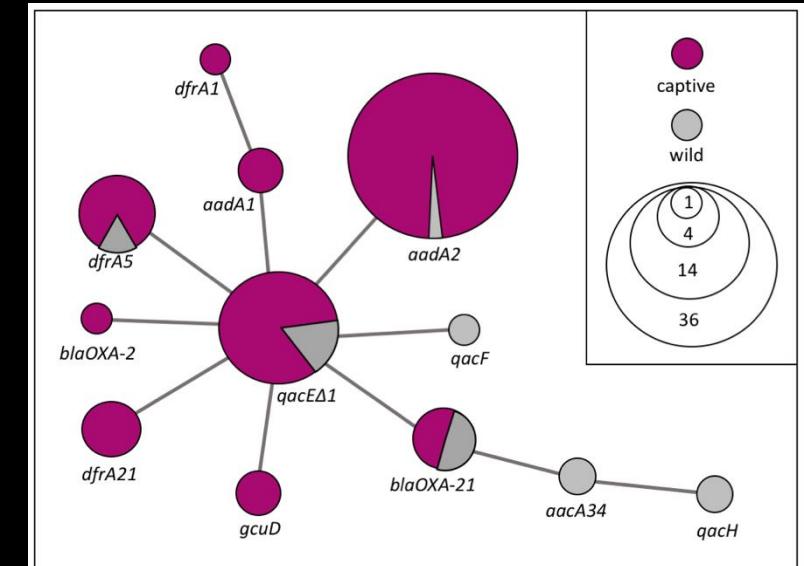
Penicillins, ESBL, fluoroquinolones,  
trimethoprim/sulfonamides, aminoglycosides

## AMR DETERMINANT EMERGENCE

### Novel clinical class 1 integrons

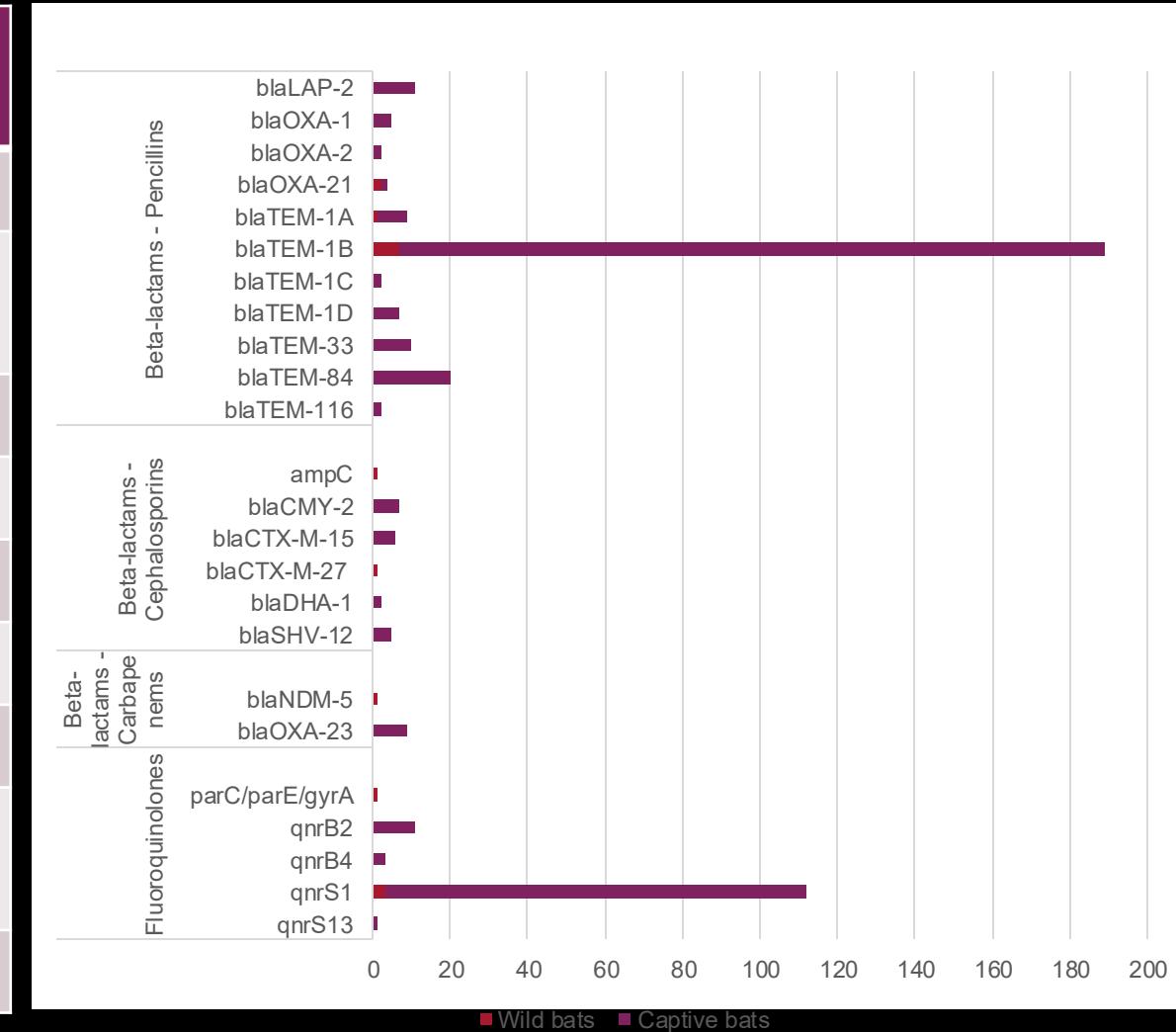
Novel clinical class 1 integrons and genes

- 5' CS-*qacH*-*aacA34*-*bla*<sub>OXA-21</sub>3'CS
- 5'CS-*qacF*-3'CS

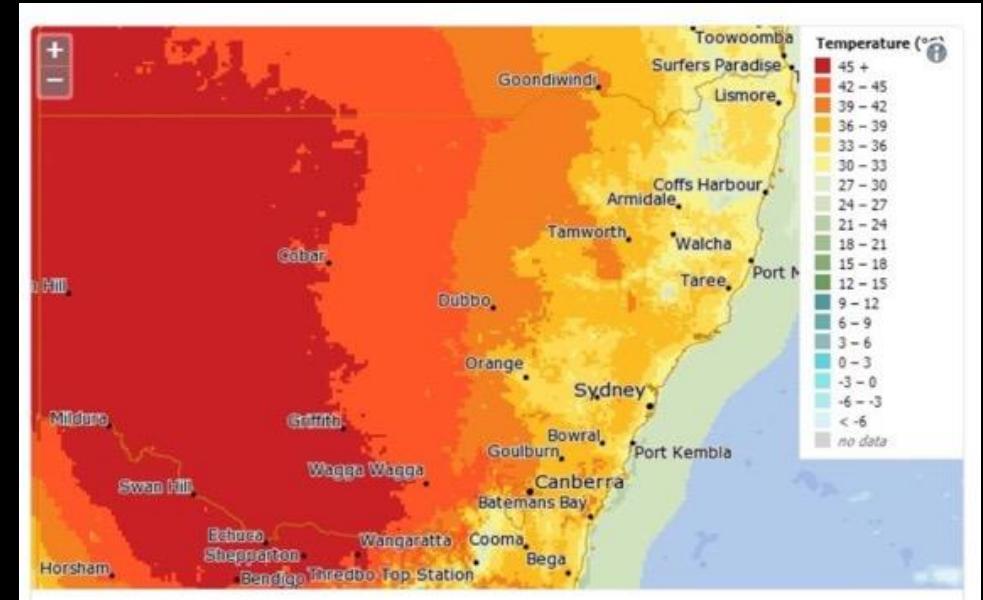


# 49 different antibiotic resistance genes in Grey-headed flying fox

Antibiotic class	No. different ARGs
Beta lactams (penicillin ARGs)	11
Beta lactams – Cephalosporin ARGs	6
Beta lactams - Carbepenems	2
Aminoglycosides	9
Chloramphenicol	2
Tetracyclines	3
Fluoroquinolones	5
Sulfonamides/ trimethoprim	11
<b>TOTAL</b>	<b>49</b>

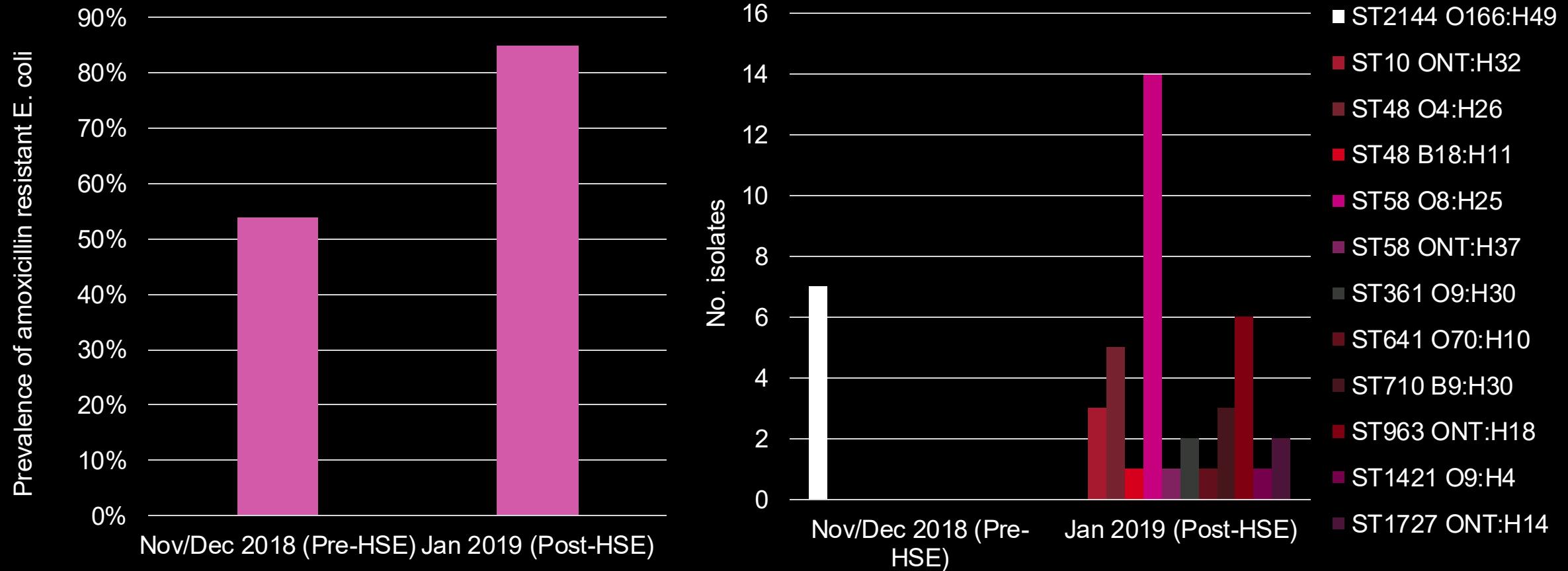


# Wildlife emergencies



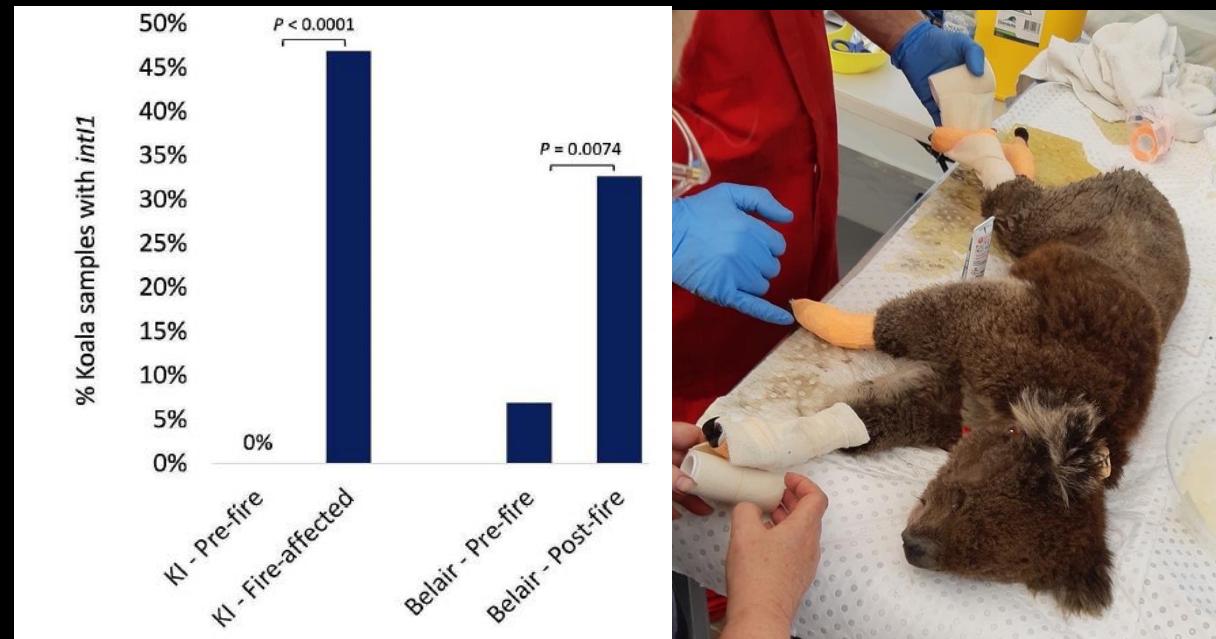
# Anthropogenic impacts | AMR dynamics and heat stress in bats

## Post heat stress bats - Increase in AMR *E. coli* and in diversity of strain types



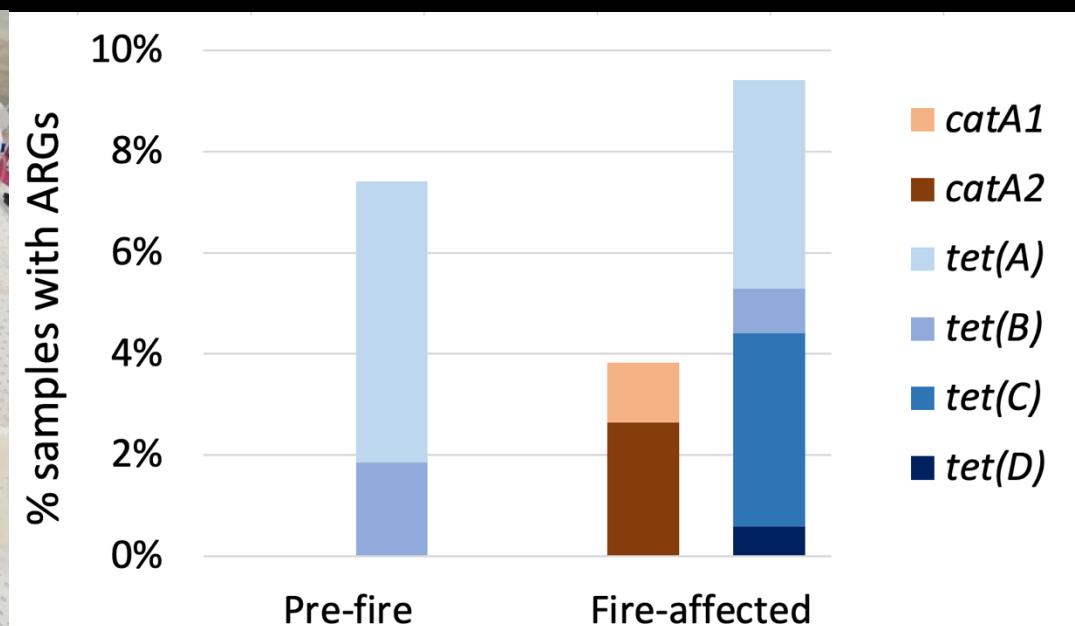
## Class 1 integrons

Wild koalas sampled pre-fire (KI = 36, Bel = 32),  
Post-fire (KI = 329 Bel = 98),



## Chloramphenicol and tetracycline

Wild koalas sampled pre-fire (n= 36)  
Rehabilitation post-fire (n=95)



# Anthropogenic impacts | Antimicrobial stewardship during wildlife emergencies

- Examined treatment records of 355 koalas admitted to wildlife hospitals during the 2019-2020 bushfires



## Koala Hospital A



**40.3%** koalas received antibiotics



75.7% prophylactic treatment of non-infected cutaneous burns  
(penicillins, enrofloxacin)



23.6% for treatment of infections  
(penicillins, enrofloxacin)

## Koala Hospital B



**35.0%** koalas received antibiotics

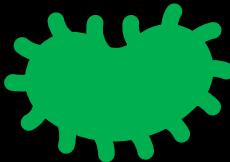


18.8% prophylactic treatment of non-infected cutaneous burns  
(penicillins)



81.2% for chlamydial disease & other infections  
(chloramphenicol, enrofloxacin and amoxicillin)

# Anthropogenic impacts | ongoing selection



Good gut  
bacteria



Resistant  
bacteria



Resistant  
secondary  
infections



Gammaretroviruses, novel viruses and pathogenic bacteria in Australian bats with neurological signs, pneumonia and skin lesions

Kate Van Brussel<sup>a</sup>, Jackie E. Mahar<sup>a</sup>, Jane Hall<sup>b</sup>, Hannah Bender<sup>b</sup>, Ayda Susana Ortiz-Baez<sup>a</sup>, Wei-Shan Chang<sup>a</sup>, Edward C. Holmes<sup>a,\*</sup>, Karrie Rose<sup>b, \*\*</sup>

<sup>a</sup> Sydney Institute for Infectious Diseases, School of Medical Sciences, The University of Sydney, NSW, 2006, Australia

<sup>b</sup> Australian Registry of Wildlife Health, Taronga Conservation Society Australia, Mosman, New South Wales, Australia

## Nadine Samy - Master research

- Data mining from wildlife pathogen database
- 55% of the *Pseudomonas aeruginosa* strains from these bats were resistant to enrofloxacin
- We do not know if the bats received enrofloxacin.

# Antimicrobial Stewardship and wildlife

AMR HAS NO SPECIES BOUNDARIES

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## Key points

- Wildlife species have acquired AMR strains from anthropogenic sources
- Antibiotics are used to treat wild animals
- Knowledge of AMR ecology, drivers, and stewardship required for the wildlife health sector

## Translation of research findings

- Using wildlife AMR data to educate wildlife care groups on AMR and AMS
- Establishing an AMS Action plan for wildlife (CSIRO, WHA and ZAA, wildlife experts representing One Health)



# QUESTIONS



Image N. Edards, link CC-BY-NC-ND