

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

- Fiona McDougall
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- Kelly Wyres - Monash
- Wayne Boardman - U Adl
- Paul Thompson - Taronga
- Roy Robbins-Brown- Doherty
- David Gordon - ANU
- Sabine Schiller – Mres

Marine species

- Mariel Fulham – PhD
- 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

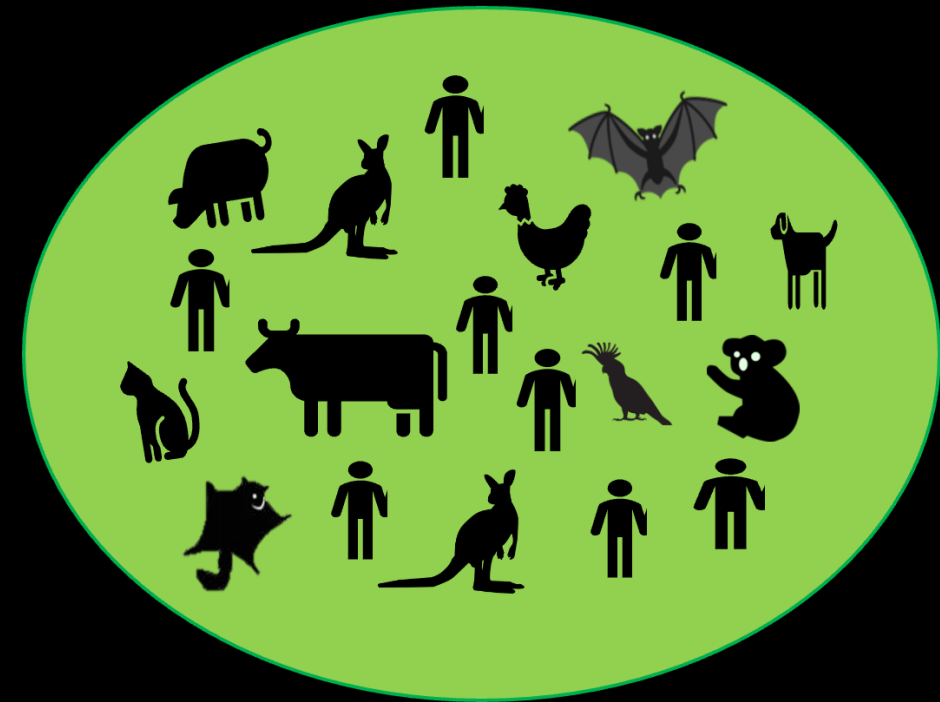
- Nadine Samy (Mres)
- Gilad Bino



Antimicrobial resistance and wildlife

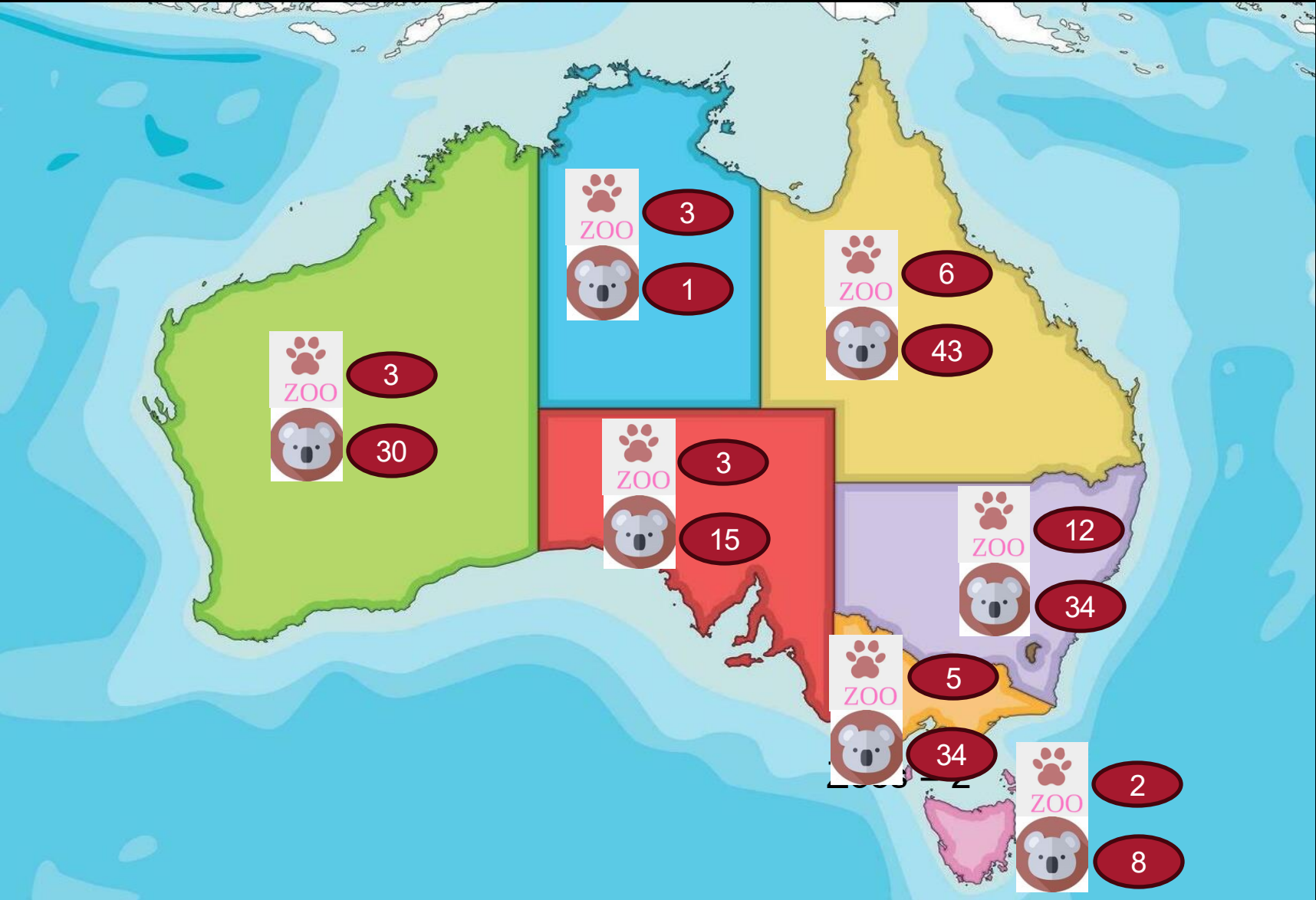
AMR HAS NO SPECIES BOUNDARIES

- 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
<div></div> <div>128,262 native animals across 543 species rescued</div>	<div></div> <div>31,290 (24%) native animals released back to the wild</div>	<div></div> <div>8,621 volunteers</div>

- ### Wildlife Victoria - 2024
- 97,772 rescues
 - 455 species

- ### WIRES - 2025
- 163,776 rescues
 - 460 carers

Investigating AMR in wildlife systems

URBAN SETTINGS

Flying fox
Ngumuny



Possum
Wali



Macropods *Badagarang*



CONSERVATION

Tasmanian devil
Purinina



Koala
Gula



Platypus
Boondabarra



Echidna
Barrugin



MARINE

Australian sea lion
Bulgurra

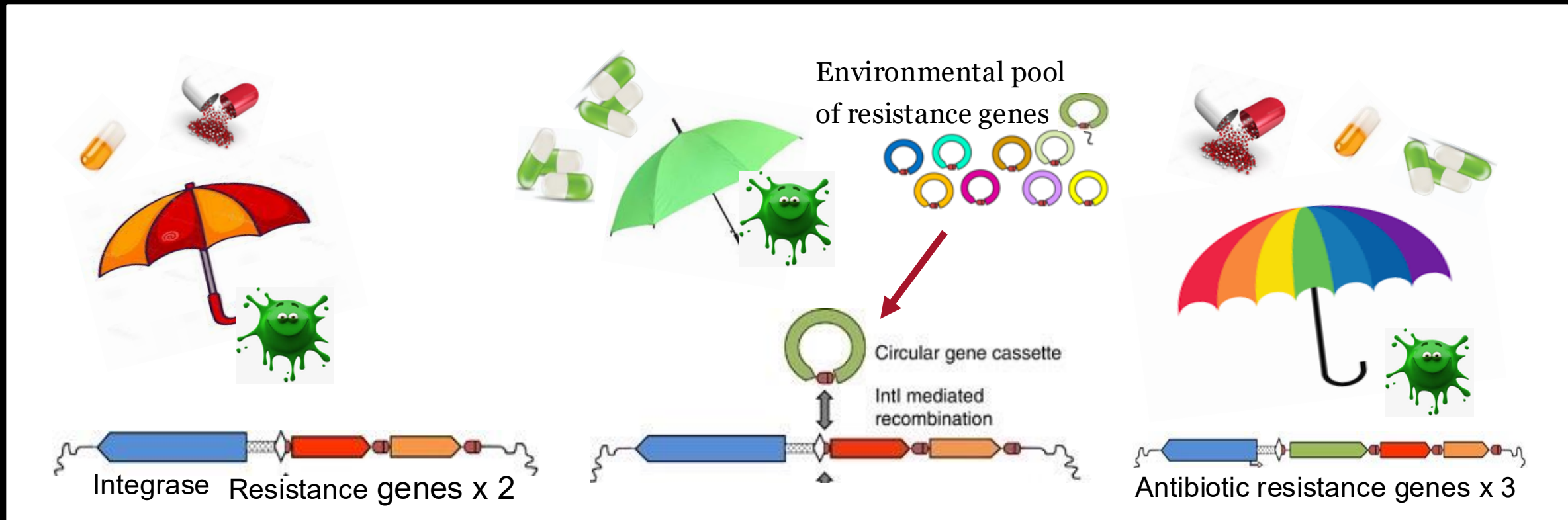


Little penguin
Gur-roo-mul



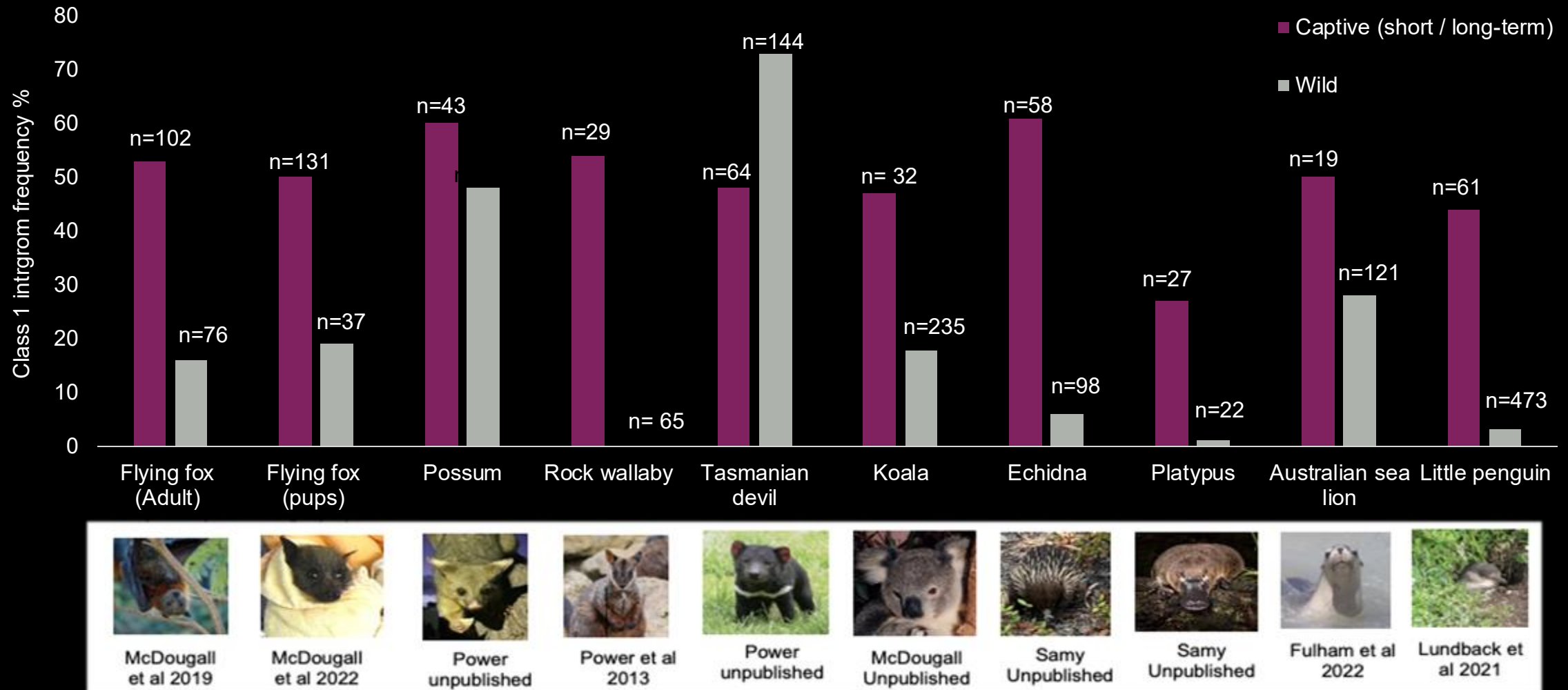
Anthropogenic origins | Clinical class 1 integron

- 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 penicillate*):

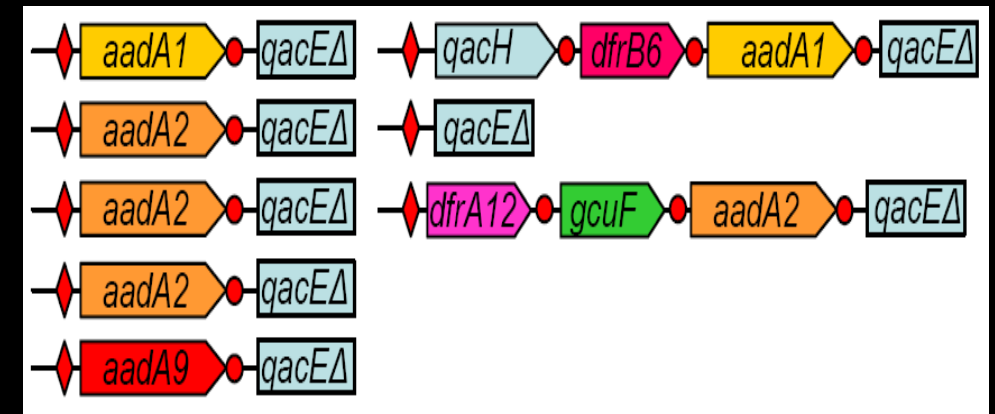
- Class 1 integrons detected in 48% of captive-bred wallabies
- No class 1 integrons were detected in the wild population



<https://www.mammalage.com>

Post-release

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



Koala
Phascolarctos cinereas

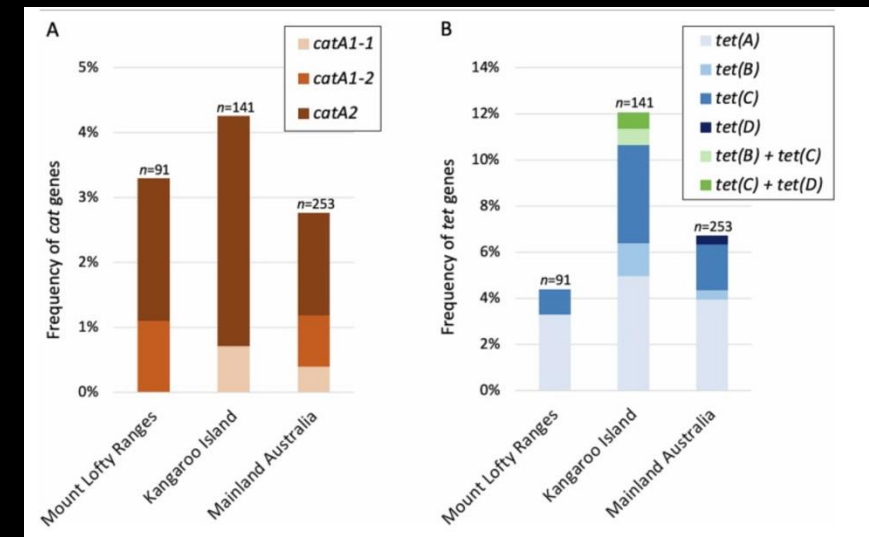
Gula



Conservation | antimicrobial reliance

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



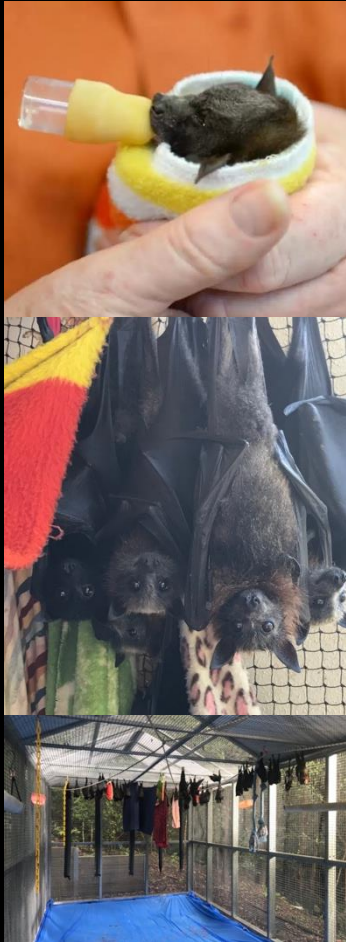
Grey headed flying fox
Pteropus poliocephalus

Ngununy

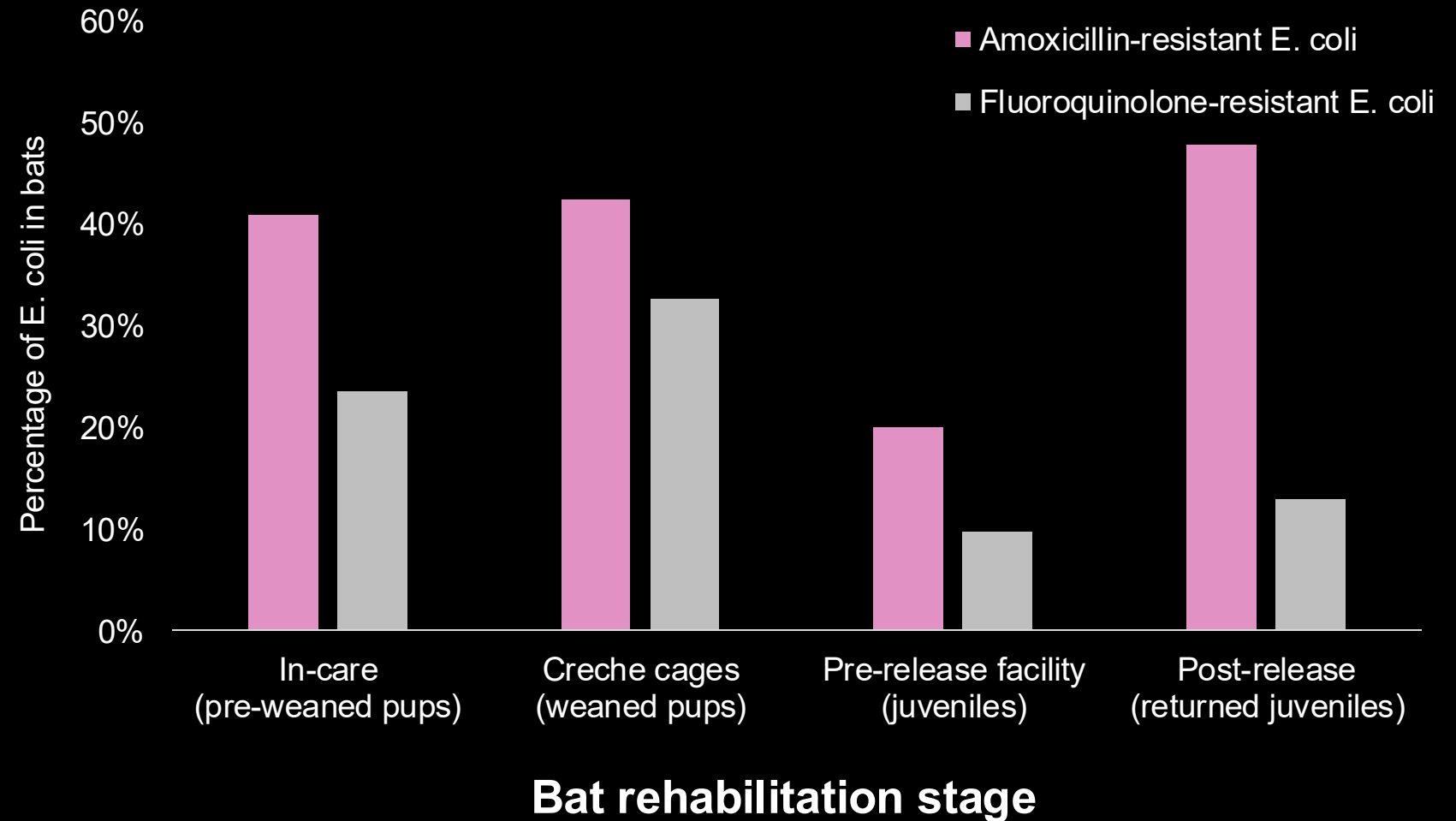


Image: Adam Mckeown

Wildlife rehabilitation | Antibiotic-resistant *E. coli*



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



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



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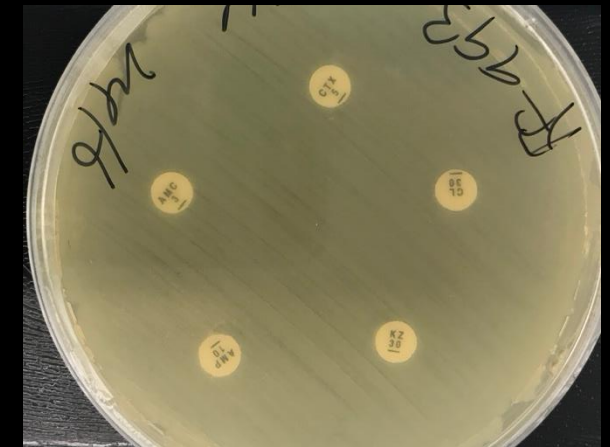
48% *E. coli* isolates from bats multidrug-resistant

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

aph(3')-Ia, *aph(6)-Id*, *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 (*blaCTX-M-15*)

4% Carbapenem-resistant (*blaOXA-23*)

Anthropogenic origins | AMR *E. coli* in bats

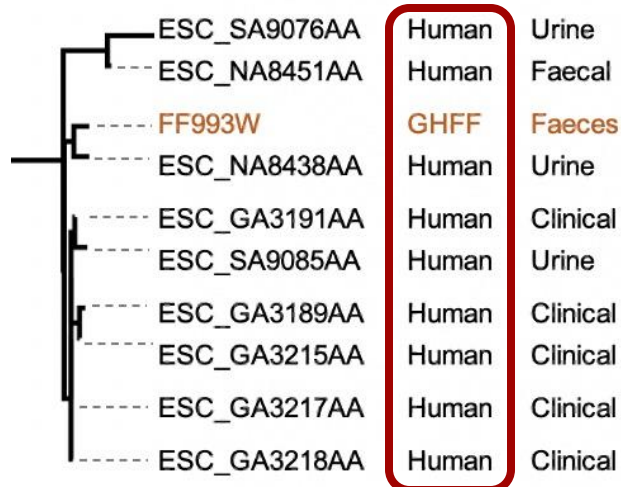
>95% antibiotic-resistant *E. coli* strains from bats are lineages associated with humans and/or domestic animals

Bat *E. coli* isolate ST10 O89:H9

Bat *E. coli* ST963 ONT:H18

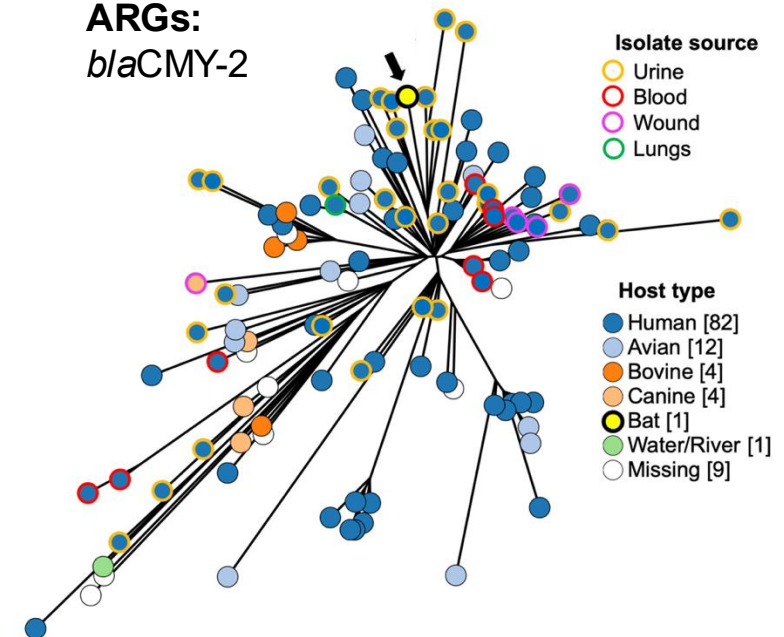
ARGs:

*bla*NDM-5
*bla*CTX-27
*bla*TEM-1B
*cat*A2
*dfr*A14
gyrA/parC/parE
*sul*2
Tet(A)



ARGs:

*bla*CMY-2



Emergence risks | within host evolution

STRAIN EMERGENCE

E. coli ST38

Penicillins, chloramphenicol



Penicillins, fluoroquinolones, chloramphenicol

E. coli ST131 Cx

Fluoroquinolones, trimethoprim



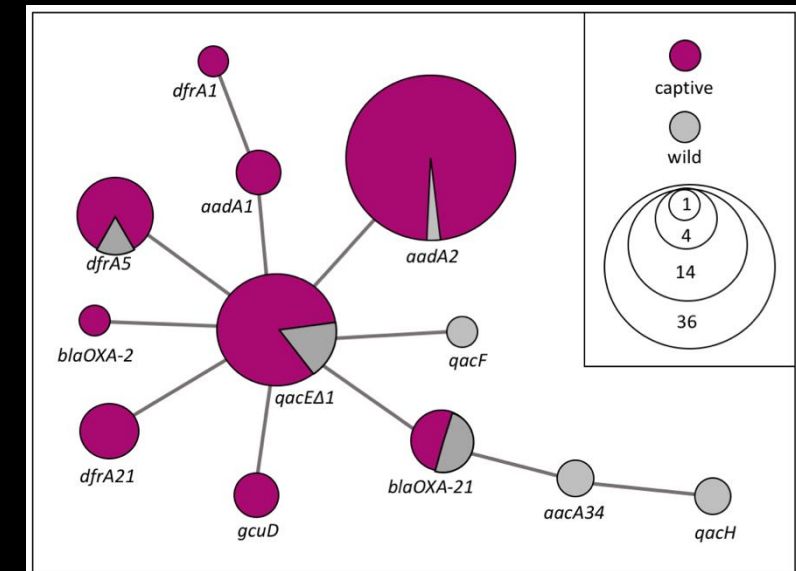
Penicillins, ESBL, fluoroquinolones,
trimethoprim/sulfonamides, aminoglycosides

AMR DETRMINANT EMERGENCE

Novel clinical class 1 integrons

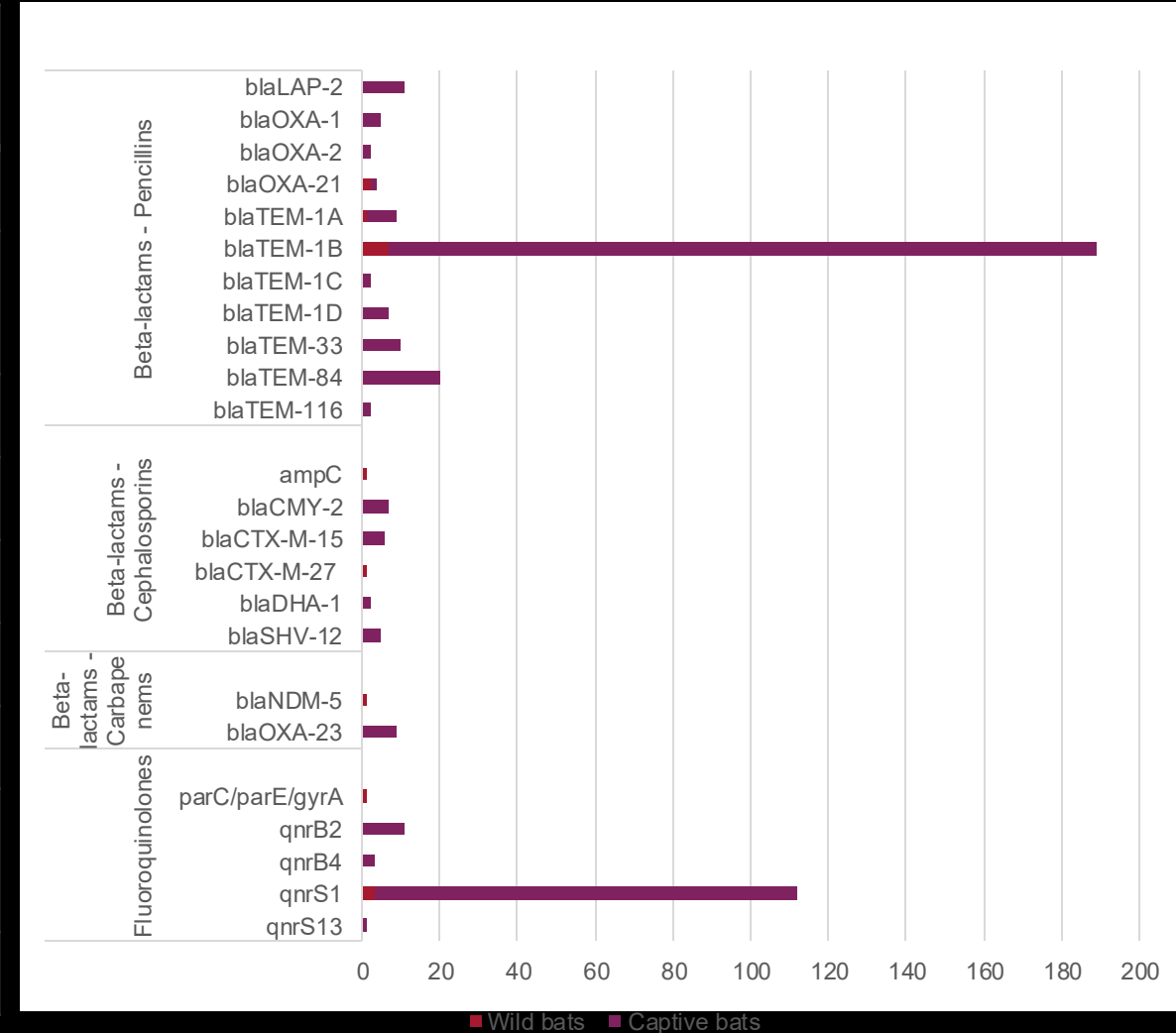
Novel clinical class 1 integrons and genes

- 5' CS-*qacH*-*aacA34*-*bla*_{OXA-21}-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 - Carbapenems	2
Aminoglycosides	9
Chloramphenicol	2
Tetracyclines	3
Fluoroquinolones	5
Sulfonamides/ trimethoprim	11
TOTAL	49

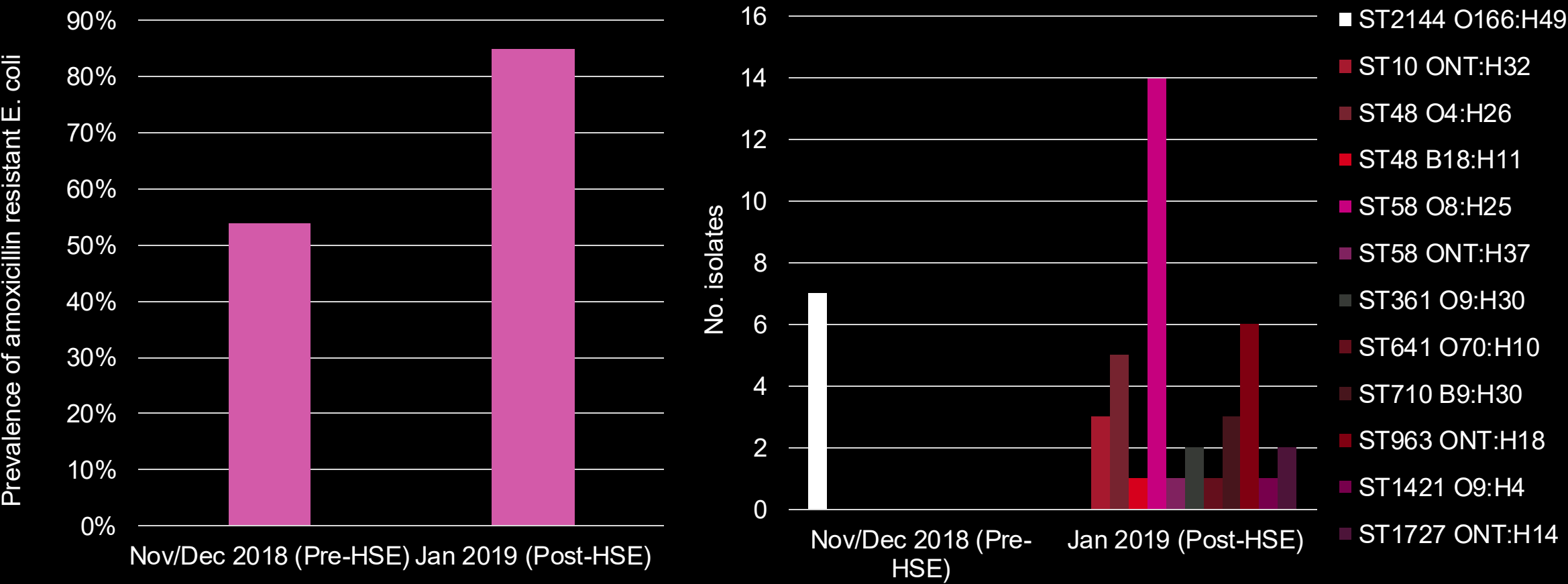


Wildlife emergencies



Anthropogenic impacts | AMR dynamics and heat stress in bats

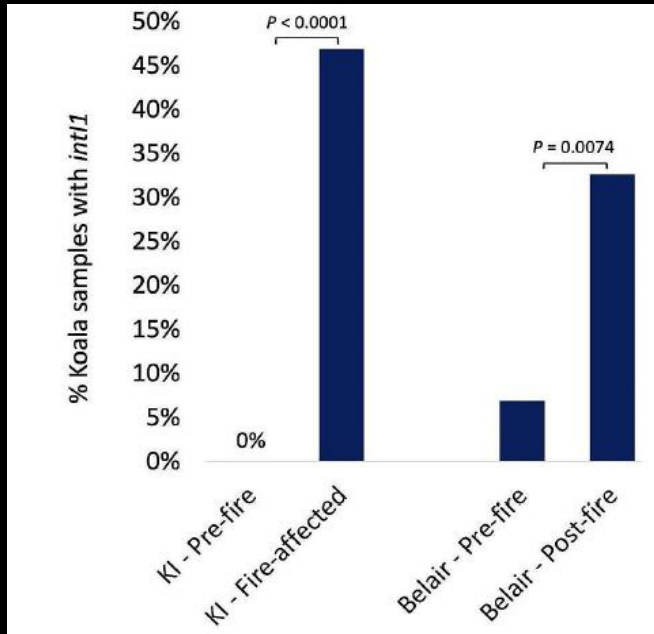
Post heat stress bats - Increase in AMR *E. coli* and in diversity of strain types



Anthropogenic impacts | AMR dynamics and fire

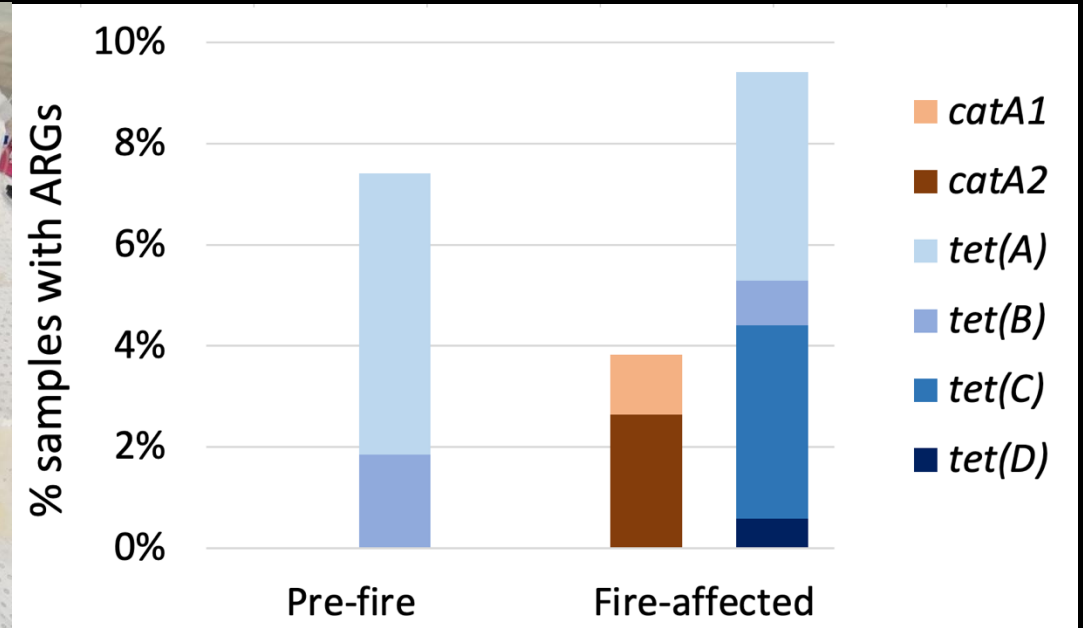
Class 1 integrons

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)

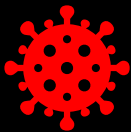
Anthropogenic impacts | ongoing selection



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^a, Jackie E. Mahar^a, Jane Hall^b, Hannah Bender^b, Ayda Susana Ortiz-Baez^a, Wei-Shan Chang^a, Edward C. Holmes^{a,*}, Karrie Rose^{b,**}

^a Sydney Institute for Infectious Diseases, School of Medical Sciences, The University of Sydney, NSW, 2006, Australia

^b 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

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



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QUESTIONS



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