An Introduction to CARD & Biocuration

Brian Alcock May 25, 2022

A Bit About Me...

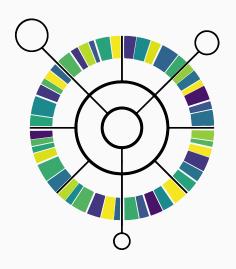


- Born and raised in Newfoundland
- McMaster M.Sc. (Biology) -2013-2015
- McArthur Lab December 2015

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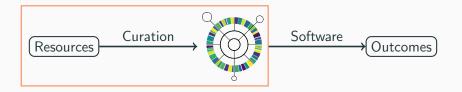
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- No background in CompSci prior to my MSc!



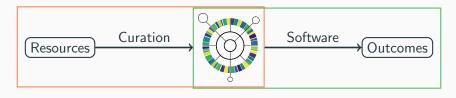
- The Comprehensive Antibiotic Resistance Database
- Antimicrobial surveillance and stewardship
- Biocuration and software development
- Data standardization and harmonization



Brian: Curation, Ontologies, AMR detection models Amos: Software Engineering, RGI, AMR prediction



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Art: How you represent it to the world

Everyday examples: museums; radio stations; Twitter; TikTok

What Gets Included?

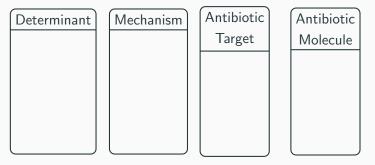
CARD's Golden Rule: to be included in CARD an AMR determinant must be described in a peer-reviewed scientific publication, with its DNA sequence available in GenBank, including clear experimental evidence of elevated minimum inhibitory concentration (MIC) over controls. (CARD 2020)

(Over time we've added some exceptions but let's not worry about those right now...)

 CARD is built around ontologies — think of these like a Word Bank where each term is connected — mainly the Antibiotic Resistance Ontology (ARO)

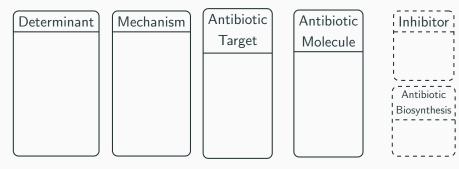
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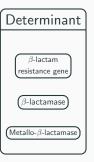
NDM β -lactamase

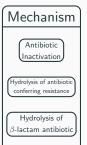
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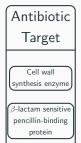


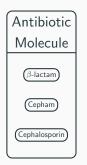
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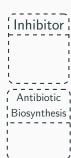
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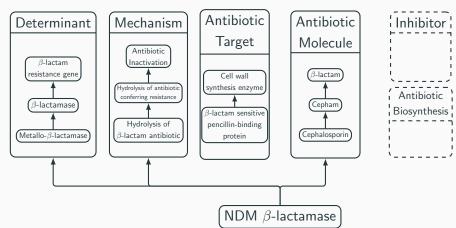




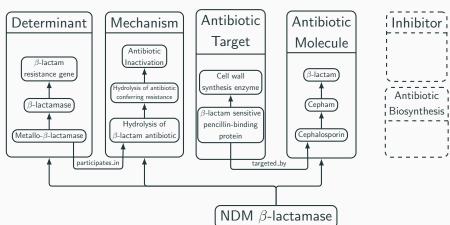


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NDM on CARD

NDM beta-lactamase [AMR Gene Family] **Download Sequences** ARO:3000057 Accession Definition NDM beta-lactamases or New Delhi metallo-beta-lactamases are class B beta-lactamases that confer resistance to a broad range of antibiotics including carbapenems, cephalosporins and penicillins. Drug Class carbapenem, cephalosporin, cephamycin, penam Resistance Mechanism antibiotic inactivation Classification 13 ontology terms | Hide + process or component of antibiotic biology or chemistry + antibiotic molecule + mechanism of antibiotic resistance + antibiotic inactivation [Resistance Mechanism] + beta-lactam antibiotic + determinant of antibiotic resistance + determinant of beta-lactam resistance + hydrolysis of antibiotic conferring resistance + antibiotic inactivation enzyme + hydrolysis of beta-lactam antibiotic by metallo-beta-lactamase + beta-lactamase + cephem + class B (metallo-) beta-lactamase Parent Term(s) 5 ontology terms | Hide + confers resistance to carbapenem [Drug Class] + confers_resistance_to cephalosporin [Drug Class] + confers_resistance_to cephamycin [Drug Class] + confers resistance to penam [Drug Class] + subclass B1 (metallo-) beta-lactamase Sub-Term(s) 17 ontology terms | Hide + NDM-1

Adding a New Resistance Gene

One incredibly useful aspect of ontologies is we do not need to record redundant information for new genes:



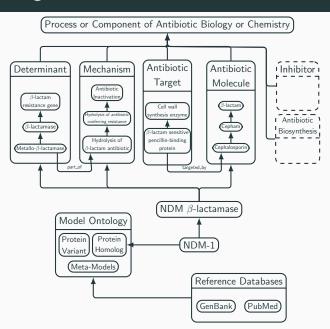
AMR Detection Models

We append "resistance genes" with genetic sequence, mutation and bit-score information to create an AMR Detection Model — a bioinformatic tool to detect resistance genes from genomes:



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Putting it Together



Core takeaways from this brief introduction:

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- 1. CARD is a platform and bioresource for amalgamating AMR data from peer-reviewed sources
- 2. ARO is critical for providing a controlled vocabulary universally understood AMR language
- Biocuration is both the collection and presentation of these data for improved user experience
- 4. Our software uses these data structures and AMR detection models for computational AMR prediction

Thanks for listening!

(Questions are welcome!)





