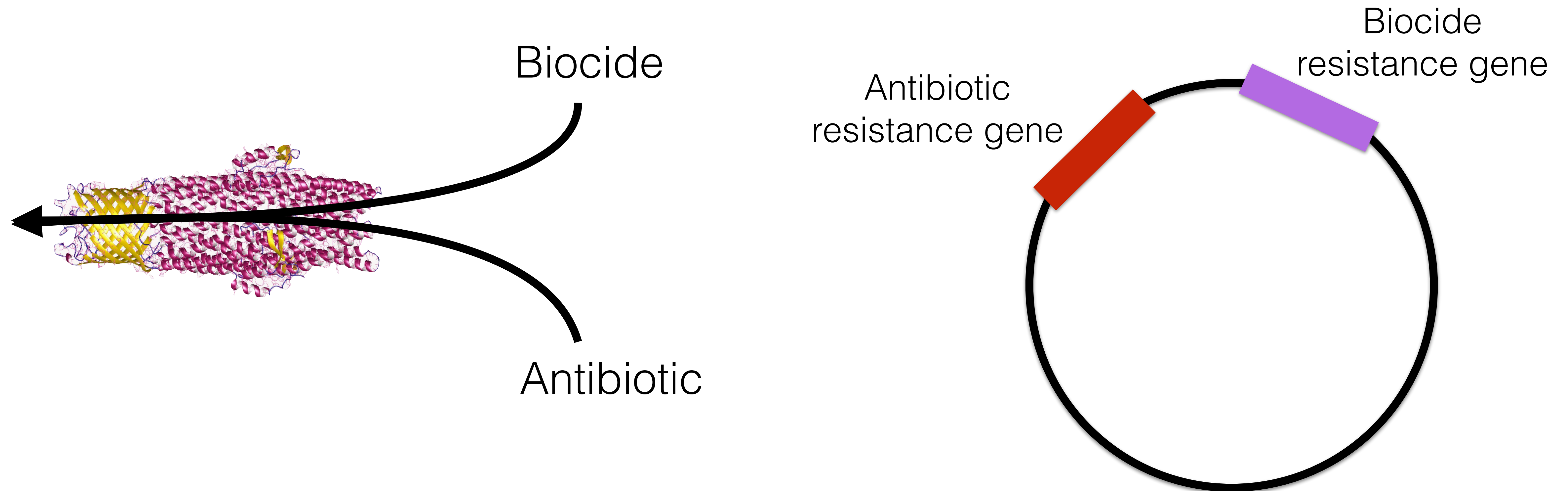


The BacMet database as a tool for biocide resistance risk assessment

Johan Bengtsson-Palme

Biocide-antibiotic co-resistance



Nucleic Acids Research, 2013, 1–7
doi:10.1093/nar/gkt1252

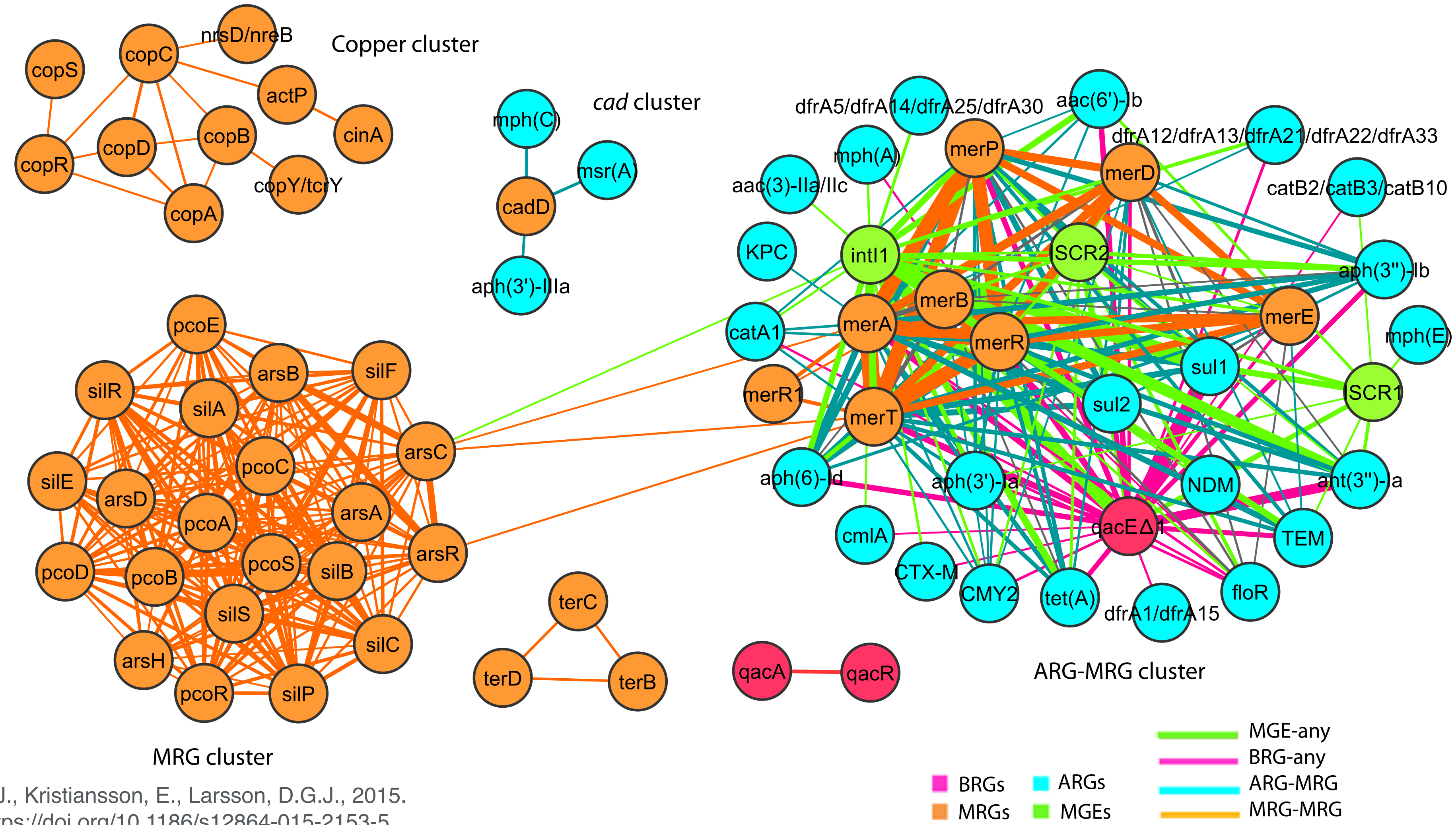
BacMet: antibacterial biocide and metal resistance genes database

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Received August 22, 2013; Revised October 31, 2013; Accepted November 11, 2013

Gene co-occurrences



Pal, C., Bengtsson-Palme, J., Kristiansson, E., Larsson, D.G.J., 2015.
BMC Genomics 16, 964. <https://doi.org/10.1186/s12864-015-2153-5>

BacMet Demo

<http://bacmet.biomedicine.gu.se/>

BacMet – Antibacterial Biocide & Metal Resistance Genes Database

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BacMet is an easy-to-use bioinformatics resource of antibacterial biocide- and metal-resistance genes. BacMet consists of two databases:

- A manually curated database of genes with **experimentally confirmed** resistance function
- A database of **predicted** resistance genes based on sequence similarity to genes with experimentally confirmed function

Database Statistics (version 2.0)

Last updated: 11 March 2018

Predicted Resistance Genes: 155,512

Exp. Confirmed Resistance Genes: 753

Chromosomal-borne: 550

Plasmid/transposon-borne: 203

Biocide resistance genes: 268

Metal resistance genes: 420

Genes with both biocide- and metal-
resistance potential: 65

Total Compounds: 111

Chemical classes: 43

Antibacterial biocides: 58

Metals: 23

'Other compounds' : 30

BacMet provides a high quality, manually curated database of bacterial genes that are experimentally confirmed to confer resistance to metals and/or antibacterial biocides, fully referenced to the scientific literature. BacMet also includes a database of predicted resistance genes, as the resistance genes may differ between species and/or occur in different forms that are not (yet) experimentally investigated. The database of predicted genes is generated by sequence similarity searches in public databases, using a uniform cut-off for genes found on plasmids, and individually set cut-offs for chromosomal genes.

BacMet provides tools for identification of biocide and metal-resistance genes in proteins and DNA sequences including full genomes. The genes in the databases can be accessed either through the browsing option, where one can browse genes by the compounds they confer resistance to or by their name. Alternatively one may use the search function to search for any term in the database, including for example gene name, name of biocide or metal and chemical class. Using the advanced search option, one may search specifically for e.g. plasmid-borne or chromosomal-borne genes. The entire database can also be downloaded for off-line analysis of larger datasets.

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Confirmed Database](#)[Predicted Database](#)

- A manually curated database of experimentally confirmed resistance function
- A database of predicted resistance genes based on sequence similarity to genes with experimentally confirmed function

Database Statistics

Last updated: 11/11/2011

Predicted Resistance Genes

Exp. Confirmed Resistance Genes

Chromosomal-borne: 550
Plasmid/transposon-borne: 203

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[By Biocides](#)[By Metals](#)[By other compounds](#)[By Biocide Resistance
Genes](#)[By Metal Resistance
Genes](#)[By Genes that have
both Biocide and Metal
Resistance Potential](#)[Total BacMet
Experimentally
Confirmed Database](#)

Database of antibacterial biocide- and metal-resistance genes. BacMet consists of

Experimentally confirmed resistance function
based on sequence similarity to genes with experimentally confirmed function

BacMet provides a high quality, manually curated database of bacterial genes that are experimentally confirmed to confer resistance to metals and/or antibacterial biocides, fully referenced to the scientific literature. BacMet also includes a database of predicted resistance genes, as the resistance genes may differ between species and/or occur in different forms that are not (yet) experimentally investigated. The database of predicted genes is generated by sequence similarity searches in public databases, using an uniform cut-off for genes found on plasmids, and individually set cut-offs for chromosomal genes.

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Browsing Experimental Confirmed database by biocides

Biocide Name	Chemical Class	Resistance Gene(s)
2-nitroimidazole	Imidazole	Click here
Acriflavine	Acridine	Click here
Alexidine	Biguanides	Click here
Benzylkonium Chloride (BAC)	Quaternary Ammonium Compounds (QACs)	Click here
Cetrimide (CTM)	Quaternary Ammonium Compounds (QACs)	Click here
Cetylpyridinium Chloride (CPC)	Quaternary Ammonium Compounds (QACs)	Click here
Chlorhexidine	Biguanides	Click here
Chlorine Dioxide (ClO₂)	Halogens	Click here
Crystal Violet	Triarylmethane	Click here
Dequalinium chloride	Quaternary Ammonium Compounds (QACs)	Click here
Dibrompropamide	Diamidine	Click here
Dodine	Acetate	Click here
Diphenyl Ether	Phenyl	Click here
Empigen	Amine betaine	Click here
Ethidium Bromide	Phenanthridine	Click here
Ethylbenzene	Aromatic hydrocarbons	Click here
Ethylmercury chloride	Organo-mercury	Click here
Glycerol	Alcohol	Click here
Hexamidine	Diamidine	Click here
Hydrochloric Acid (HCl)	Acid	Click here
Hydrogen Peroxide (H₂O₂)	Peroxides	Click here
Linoleic acid	Free Fatty Acids	Click here
m-xylene	Aromatic hydrocarbons	Click here
Methylene Blue	Thiazinium	Click here

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Browsing BacMet compounds

Information of 2-nitroimidazole from BacMet

Compound Information

- **Compound name:** 2-nitroimidazole
- **Chemical Class:** Imidazole
- **Description:** It is an industrial biocide. It is only effective against anaerobic organisms and it is effective against sulfate-reducing bacteria (SRB) and compatible with other chemicals. Used to prepare nitroimidazole substituted boronic acids as precursors for imaging hypoxic tissue. Also used to prepare potential site-selective radiosensitizers for estrogen receptor-rich tumors.
- **Database Links:** [CHEBI:67135](#)

BacMet database and website is designed and maintained by [Chandan Pal](#)

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Browsing Experimentally Confirmed database

Result: Your query found 1 resistance gene(s) from the BacMet Experimentally Confirmed database

Gene_name	Experimentally Verified Resistance Gene Information	Predicted Resistance Gene Information
ruvB	<ul style="list-style-type: none">• BacMet ID: BAC0293• Code for: Enzyme• Family: Malic enzymes family• Sequence: FASTA• Cross-database IDs: Link• Organism: <i>Pseudomonas corrugata</i> 28• Location: Chromosome• Compound: Chromium (Cr), Cetylpyridinium Chloride (CPC) [class: Quaternary Ammonium Compounds (QACs)], Dodine [class: Acetate], 2-nitroimidazole [class: imidazole]• Description: It is a fragment of ruvB gene product. It is a oxidoreductase from malic enzyme family.• Length (amino acid): 379• Reference: Decorosi et al. 2009; Pubmed- 19768364	Similar resistance genes in 'BacMet Predicted database'

BacMet database and website is designed and maintained by [Chandan Pal](#)

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BacMet Predicted Database

Result: Your query found 495 resistance gene(s) from the BacMet Predicted database

Gene name	GI number	GenBank ID	Sequence	Organism	Compound	NCBI annotation
ruvB	15596164	NP_249658.1	FASTA	<i>Pseudomonas aeruginosa</i> PAO1	Chromium (Cr), Tellurium (Te), Selenium (Se)	ruvB gene product
ruvB	7443650	PIR: JC5477	FASTA	<i>Pseudomonas aeruginosa</i>	Chromium (Cr), Tellurium (Te), Selenium (Se)	DNA-binding protein ruvB - <i>Pseudomonas aeruginosa</i>
ruvB	1183842	BAA11819.1	FASTA	<i>Pseudomonas aeruginosa</i>	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction specific DNA helicase
ruvB	152988470	YP_001349890.1	FASTA	<i>Pseudomonas aeruginosa</i> PA7	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction DNA helicase RuvB
ruvB	289674672	ZP_06495562.1	FASTA	<i>Pseudomonas syringae</i> pv. <i>syringae</i> FF5	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction DNA helicase RuvB
ruvB	28871122	NP_793741.1	FASTA	<i>Pseudomonas syringae</i> pv. <i>tomato</i> str. DC3000	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction DNA helicase RuvB
ruvB	289626497	ZP_06459451.1	FASTA	<i>Pseudomonas syringae</i> pv. <i>aesculi</i> str. NCPPB 3681	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction DNA helicase RuvB
ruvB	237800219	ZP_04588680.1	FASTA	<i>Pseudomonas syringae</i> pv. <i>oryzae</i> str. 1_6	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction DNA helicase RuvB
ruvB	71736556	YP_275912.1	FASTA	<i>Pseudomonas syringae</i> pv. <i>phaseolicola</i> 1448A	Chromium (Cr), Tellurium (Te), Selenium (Se)	Holliday junction DNA helicase RuvB
				<i>Pseudomonas</i>	Chromium (Cr), Tellurium	

Query BacMet databases

1. Quick Search:

You can search the BacMet database using any full term or partial term with a wildcard, including for example gene names (e.g. copA or cop*), name of biocides (e.g. Triclosan or Triclo*), metals (e.g. Arsenic or Arsen*) or chemical classes (e.g. Acridine or Acrid*). You will get an output in a table format with the resistance gene information that matches to your search term/pattern.

2. Advanced Search:

You can use the advanced search option if you are especially interested in resistance genes that confer resistance to certain antibacterial compounds or chemical classes, and plasmid- or chromosomal-borne resistance genes. You can use full term or partial term with a wildcard in description section (e.g. resistance or resis*). You will get an output in a table format with the resistance gene information that matches to your search criteria.

Select 'chemical class'/ 'compound' (resistant to):

Select location: ☒ Any ☐ Chromosome ☐ Plasmid (for EXP confirmed database only)

Protein description contains text (e.g. resistance)

Peptide sequence length greater than (for EXP confirmed database only)

Peptide sequence length less than (for EXP confirmed database only)

Select database: ☐ BacMet Predicted database ☒ BacMet Experimentally Confirmed database

Moving towards a risk assessment tool

- Incorporate data on **selective concentrations** and **co-selection opportunities**
- Make BacMet useful for practical risk assessment and management

BacMet Update

- Current data will remain and be expanded upon
- **Substance-based** rather than gene-based focus – commonly the starting point for risk assessment
- **MICs and LOECs** for different species, PNECs for resistance selection and HGT
- **New genes** will be added (from literature and this project)
- **Context information** (hosts, mobility potential, co-location with other resistance genes), based on available genome data
- Relative **abundance in different environment types**

Acknowledgement

- Chandan Pal
- Joakim Larsson
- Erik Kristiansson
- Christopher Rensing
- Funding from FORMAS

Discussion tasks

Short introduction round (3-4 sentences per person)

<http://bacmet.biomedicine.gu.se/>

Who takes the minutes?

How can the BacMet database support biocide resistance risk assessment during approval/authorization of biocides?

What needs to be changed to be able to use the BacMet database for the regulatory process?

Proposed structural changes?

Proposed new categories?

Proposed links with other relevant databases?

BacMet currently is a substance-based database. How to incorporate results for products?

Who would like to present the results?