Presence of carbapenem-resistant bacteria in soils affected by illegal waste dumps

Tomislav Ivankovic^a, Jasna Hrenovic^a, Goran Durn^b, Svjetlana Dekic^a, Snjezana Kazazic^c Ivica Kisic^d

^a University of Zagreb, Faculty of Science, Zagreb, Croatia; ^b University of Zagreb, Faculty of Mining, Geology and Petroleum Engineering, Zagreb, Croatia; ^c Ruder Bošković Institute, Division of Physical

Chemistry Zagreb, Croatia; ^d University of Zagreb, Faculty of Agriculture, Zagreb, Croatia

ABSTRACT

carbapenem-resistant bacteria (CRB) are The currently at the top of the WHO priority list of bacteria that pose the greatest threat to human health. Considering that soil is one of the important environments for the emergence of antibiotic-resistant bacteria, we isolated and quantified cultivable CRB in soils across Croatia, including ones affected by illegal dumps. We cultivated CRB at two temperatures, distinguishing between the intrinsically resistant CRB (37°C, mostly *Stenotrophomonas* spp.) and the ones that are presumably human-associated and clinically relevant (42°C, Acinetobacter sp, Enterobacteriaceae, Burkholderia spp.).

Qur study demonstrated that distinguishing between the two offers a better insight into the diversity of CRB in the environment. The ones cultivated at 37 °C were found in almost all soil samples, while the presumably clinically relevant ones were absent from uncontaminated pasture and grassland, indicating that human-associated CRB are unlikely to be found in soils spared from anthropogenic influence.



MATERIALS AND METHODS

Soil samples were taken at 14 locations (eight illegal dump sites, one coke factory, one thermal power plant, two arable lands, one pasture, and a grassland). Used methodology is based on cultivation of environmental samples on 37 and 42°C in parallel. The latter supresses the growth of Stenotrophomonas sp. which are intrinsically carbapenem-resistant and ubiquitous in soil, and therefore dominant in environmental/soil samples cultivated at 37°C. Cultivation at 42°C allows the detection of other CRB that are usually present in far smaller numbers than Stenotrophomonas sp.. This methodology has been FIND OUT MORE... successfully used in our previous studies where CRB were monitored in wastewater treatment plant and now it was, to best of our knowledg implemented for soil samples for the first time.

Incubating the samples at 37 and 42°C, respectively, enabled us to identify CRB with acquired resistance, that are normally present in smaller numbers and shaded by large numbers of intrinsically resistant Stenotrophomonas sp. and/or other species.

Intrinsically resistant CRB were found in every soil sample, including the uncontaminated pasture and grassland (log 2.4-5.3 CFU g⁻¹) confirming that CRB with intrinsic resistance are abundant in soil. Presumably clinically relevant CRB, with acquired resistance, were found in

The CRB were cultivated on selective CHROMagar[™] Acinetobacter medium (CHROMagar, Paris, France). With adition of CR102 supplement, the growth of carbapenem-resistant Acinetobacter sp. and other carbapenem-resistant Gram-negative bacteria, belonging mostly to the Enterobacteriaceae, Pseudomonas spp., and Stenotrophomonas genera is allowed.

most dump sites (log 1.2 - 4.7 CFU g⁻¹), but not in uncontaminated pasture and grassland, recognizing illegal dump sites as potential sites of dissemination of antibiotic-resistant bacteria to the environment.

	No of		No		No
Species	isolates				
Acinetobacter baumannii	3	Enterobacter asburiae	1	Pediococcus sp.	1
Acinetobacter sp.	6	Enterobacter cloacae	3	Providencia sp.	1
Burkholderia ambifaria	12	Enterobacter ludwigii	1	Providencia stuartii	1
Burkholderia multivorans	2	Enterobacter sp.	2	Pseudomonas putida	1
Burkholderia sp.	2	Escherichia coli	3	Sphingobacterium thalpophilum	1
Cupriavidus gilardii	3	Escherichia sp.	1		
Cupriavidus respiraculi	5	Ochrobactrum intermediu	<i>m</i> 3		

List of bacteria with acquired resistance to carbapenems randomly isolated after cultivation at 42°C. Bacteria isolated after cultivation at 37°C were intrinsically resistant to carbapenems with 10 isolates of Stenotrophomonas sp. and one isolate of Elizabethkingia meningoseptica.

An example of CHROMagar AcinetobacterTM plate (without CR102 supplement for cultivation of carbapenem-resistant bacteria) inoculated with wastewater sample after incubation at 37°C for 48 hours. The blue

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colonies are usually *Enterobacteriaceae* and red colonies are

Stenotrophomonas sp., Acinetobacter sp., Pseudomonas sp. or other

Gram-negative bacteria.



by Prof.dr.sc. Jasna Hrenovic.



