COMMENTARY

New Delhi metallo- β -lactamase-1: detection and prevention

Timothy R. Walsh PhD

See related research article by Kus and colleagues on page 1257 and at www.cmaj.ca/lookup/doi/10.1503/cmaj.110477.

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Correspondence to: Timothy R. Walsh, t.r.walsh@uq.edu.au

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The advent of New Delhi metallo-βlactamase-1 (NDM-1) as a new mechanism of microbial resistance went mostly unmarked, despite it being plasmidmediated, highly transferable and able to confer resistance to most antibiotics (with the exceptions of colistin and tigecycline).^{1,2} However, after the Indian subcontinent was linked to the spread of organisms producing NDM-1, and in response to the resulting reaction from India's Government, NDM-1 became an overnight global phenomenon.^{3,4} Thus far, most cases of infection or colonization with NDM-1producing bacteria have originated in southern Asia, most often in India. It is now clear that NDM-1 is firmly established in India, both in the hospitals and in the community. At the community level, the situation is exacerbated by the poor sanitary conditions with which many Indians must live on a daily basis.5

From a diagnostic and microbiologic perspective, NDM-1 is not easily identified in clinical bacteria. Some methods of detection lack the sensitivity necessary for the rapid and assured detection of NDM-1 that facilitates antibiotic stewardship and the introduction of measures to control the spread of infection.⁶

The story of NDM-1 is still in its infancy; however, it has already eclipsed the *Klebsiella pneumoniae* carbapenemases in terms of the number of countries in which it is found. In addition, because of the promiscuous nature of the plasmids containing the NDM-1 gene, the enzyme has been found in many more species than have the *K. pneumoniae* carbapenemases. Furthermore, our

Key points

- Most cases of infection or colonization with bacteria producing New Delhi Metalloß-lactamase-1 (NDM-1) have originated in southern Asia, most often in India.
- The spread of such bacteria to the international community such that direct links to southern Asia cannot be confirmed is inevitable.
- The rise of NDM-1 is not only due to its high rate of genetic transfer among unrelated bacterial species, but also to human factors such as travel, sanitation and food production and preparation.

knowledge of NDM-1 is compromised by poor case studies; additional information, such as carriage and persistence, is required.

In their article for the *CMAJ*, Kus and colleagues⁷ present two important but contrasting case studies. The first patient, an 86-year-old man, had no contact with India or southern Asia. The *Morganella morganii* isolated from a culture of the patient's urine showed sensitivity to meropenem based on results from the Vitek II automated susceptibility testing system (bio-Mérieux, France), but carbapenemases were not detected by the Modified Hodge Test. However, despite these results, the organism was NDM-1 positive. These data throw into considerable doubt the validity of some of the methods heavily relied upon by diagnostic laboratories for the detection of antibiotic resistance.⁶

That the patient had no primary contact with southern Asia (indeed, he had not travelled outside of Canada in the previous 10 years) may be touted as evidence by the people who wish to propagate the myth that NDM-1 is everywhere.^{8.9} However, the link between southern Asia and most NDM-1–producing isolates is overwhelming, and the spread of such bacteria to the international community, such that direct links to southern Asia cannot be confirmed, is inevitable.

Another important feature of this particular case is that the isolate could be found in the patient's urine five months after the original diagnosis, an unusual show of persistence. Clearly, the isolate is not overtly pathogenic; however, it is able to persist in unusual environments, indicating a possible link between antibiotic resistance and persistence rather than pathogenicity. Unfortunately, samples of the patient's stool were not taken, which may have shown the presence of the *M. morganii* strain or other isolates that could have conferred the NDM-1 gene to the patient's natural flora.

In the second case, an NDM-1–producing isolate of *Providencia rettgeri* caused a urinary tract infection in a 71-year-old woman who had recently travelled to India. For this patient, the link to southern Asia is indisputable. The patient

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presented to hospital in Toronto, where samples of her urine were taken in which the NDM-1– producing *P. rettgeri* was detected. Again, a follow-up culture of the patient's urine several months later showed that the organism was persisting, serving as another example of antibiotic resistance being linked to persistence rather than to pathogenicity.¹⁰ As in the previous case, because samples of this patient's stool were not taken, fecal carriage and in situ transfer of the NDM-1 gene could not be investigated.

Both patients were colonized by or infected with bacteria not usually associated with NDM-1, suggesting that the NDM-1 gene pool has now established itself in species other than *Escherichia coli* and *K. pneumoniae*. NDM-1– producing bacteria typically show a high level of multidrug resistance or near pan-drug resistance. Although the organisms isolated from both of these patients were resistant to colistin, treatment with that antibiotic would not have been appropriate anyway. In fact, the only drug available to treat infections caused by NDM-1–producing organisms is tigecycline, which could not have been used because it is not indicated for the treatment of urinary tract infections.

Both of the patients described by Kus and colleagues7 were asymptomatic, so treatment of their infections was not necessary. However, if the patients had been symptomatic, their urinary tract infections would have been untreatable - Providencia and Morganella species are intrinsically resistant to colistin. Thankfully, neither of these organisms are prominent pathogens in the West, although we have very little data on their prevalences in southern Asia. A previous study by Walsh and colleagues⁵ suggests that the NDM-1 gene pool in southern Asia (i.e., carriage in normal flora) could be as high as 20%. It is therefore possible that, owing to poor sanitation, NDM-1producing organisms could further spread throughout southern Asia, causing an increase in the prevalence of conditions such as cystitis.4,5

The examples reported by Kus and colleagues⁷ are but two of several cases of colonization or infection with NDM-1–producing organisms known to have occurred in Canada; for the foreseeable future, the Canadian medical community can brace itself for many more — whether they are linked to India or not. The spread of NDM-1 is not only due to its high rate of genetic transfer among unrelated bacterial species, but also to human factors such as travel, sanitation and food production and preparation. The rise of NDM-1 has caused us to rethink antibiotic resistance and the global spread of resistance mechanisms. As such, increased vigilance and appropriate surveillance are mandatory.

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Affiliations: Timothy R. Walsh is with the Section of Medical Microbiology, School of Medicine, Cardiff University, Heath Park, Cardiff, Wales; and with the Clinical Centre for Research, University of Queensland, Herston, Queensland, Australia.