



Rapid Identification of *bla*_{IMP-1} and *bla*_{IMP-6} by Multiplex Amplification Refractory Mutation System PCR

Akiyo Nakano, M.S.^{1,*}, Ryuichi Nakano, Ph.D.^{1,*}, Yuki Suzuki, M.S.¹, Kyoichi Saito, M.D.¹, Kei Kasahara, M.D.², Shiro Endo, M.D.³, and Hisakazu Yano, M.D.¹

Department of Microbiology and Infectious Diseases¹ and Center for Infectious Diseases², Nara Medical University, Nara, Japan; International University of Health and Welfare³, Shioya Hospital, Tochigi, Japan

Dear Editor,

Carbapenem-resistant Enterobacteriaceae have acquired carbapenemase genes [1], which differ substantially across countries [2]. Transferable carbapenemase IMP-type metallo-β-lactamases, particularly IMP-1 and IMP-6, are commonly identified in the clinical setting in Japan [3, 4] and exhibit different substrate specificity despite having a difference of only one amino acid (IMP-6: Ser214Gly). IMP-1 producers are more resistant to imipenem than to meropenem, whereas IMP-6 producers are more resistant to meropenem [5]. We previously found that the susceptibility rate of IMP-6-positive *Escherichia coli* was higher for imipenem than for meropenem [3]. Thus, IMP-6-producing isolates may be erroneously categorized as imipenem-susceptible, which could lead to treatment failure in patients.

It is important to distinguish between the genes *bla*_{IMP-1} (encoding IMP-1) and *bla*_{IMP-6} (encoding IMP-6) because of the differences in substrate specificity; these variants can be identified by either DNA sequencing or amplification refractory mutation system (ARMS) PCR [6]. However, these methods are costly and time-consuming, and ARMS PCR requires multiple reaction tubes per sample. Therefore, we developed a simple, rapid multiplex ARMS PCR assay to discriminate between IMP-1 and IMP-6. We designed two sets of Multiplex ARMS PCR primers

based on the nucleotide sequences of *bla*_{IMP-1} and *bla*_{IMP-6}. Primers IMP-f and IMP-r were designed to recognize the conserved sequences of *bla*_{IMP-1} and *bla*_{IMP-6}, while primers IMP-1f and IMP-6r were designed to recognize specific nucleotide sequences of *bla*_{IMP-1} and *bla*_{IMP-6}, respectively [7].

PCR amplification using the primer pair IMP-1f (5'-AAGGC-AAAAGTGGTTGTTCCCTA-3') and IMP-r (5'-CGACTTGTAGAA-ATTTAGTTGC-3') results in a 120-bp *bla*_{IMP-1} gene-specific fragment, and PCR amplification using the primer pair IMP-f (5'-CTCGATCTATCCCCACGTATG-3') and IMP-6r (5'-TCGTCTCCAAC-TTCACTGTGAGC-3') generates a 350-bp *bla*_{IMP-6} gene-specific fragment (underlined sequences indicate nucleotide alternations for allele-specific primers). The 3' termini of the primers correspond to the specific nucleotide sequence at position 640 of IMP-1 (AGT, Ser) and IMP-6 (GGT, Gly), respectively. We performed multiplex ARMS PCR amplification using the Qiagen Multiplex PCR Master Mix (Qiagen, Venlo, The Netherlands) with the following conditions: denaturation for 15 minutes at 95°C and 32 cycles of 10 seconds at 95°C, 30 seconds at 50°C, and 20 seconds at 72°C. DNA fragments were analyzed by electrophoresis. We used IMP-1-producing *E. coli* (NR2406) and IMP-6-producing *E. coli* (NR2407) from Japanese clinical isolates as positive controls. The bands, 120- or 350-bp in size, were obtained

Received: September 5, 2017

Revision received: October 10, 2017

Accepted: February 7, 2018

Corresponding author: Ryuichi Nakano

Department of Microbiology and Infectious Diseases, Nara Medical University, 840 Shijo-chou, Kashihara, Nara 634-8521, Japan
Tel & Fax: +81-744-29-8839, E-mail: rnakano@naramed-u.ac.jp

*These authors contributed equally to this article.

© Korean Society for Laboratory Medicine

This is an Open Access article distributed under the terms of the Creative Commons Attribution Non-Commercial License (<http://creativecommons.org/licenses/by-nc/4.0>) which permits unrestricted non-commercial use, distribution, and reproduction in any medium, provided the original work is properly cited.

within 90 minutes or less (Fig. 1).

We performed this method on 350 carbapenemase-producing clinical isolates collected from general hospitals across Japan from 2012 to 2016 and archived in our laboratory [4, 8].

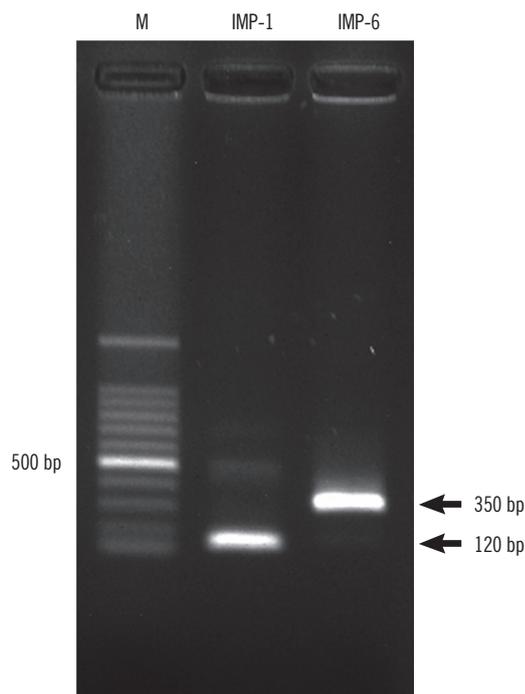


Fig. 1. Representative multiplex ARMS PCR patterns of IMP-1- and IMP-6-producing strains.
Abbreviations: ARMS, amplification refractory mutation system; Lane M, 100-bp DNA ladder marker.

We also determined the antibiotic susceptibility of the isolates by the agar dilution method in accordance with CLSI recommendations [9]. The multiplex ARMS PCR results and imipenem and meropenem minimum inhibitory concentration (MIC) ranges of the isolates are shown in Table 1. The IMP-1 and IMP-6 producers yielded PCR products of the expected size. We confirmed the accuracy of this method by DNA sequencing of the PCR products (120-bp and 350-bp bands) of three representative strains. Cross-reactivity for other types of carbapenemase producers was not detected, and false-positive or false-negative results were not observed (Table 1).

This multiplex ARMS PCR assay successfully discriminated between *bla*_{IMP-1} and *bla*_{IMP-6}. Thus, this method could serve as a specific, rapid, and simple alternative for the detection of IMP-1 or IMP-6 producers and could facilitate clinical treatment, infection control, and large-scale IMP producer screening.

Authors' Disclosures of Potential Conflicts of Interest

No potential conflicts of interest relevant to this article were reported.

Acknowledgment

This study was supported by JSPS KAKENHI (Grant no. 17K16-228 and 17K10027).

Table 1. Characteristics of carbapenemase-producing strains used in this study and the multiplex ARMS PCR results

Species	Strains (N)	β -lactamase	MIC range (μ g/mL)		Specific band* (N)
			IPM	MEPM	
<i>Klebsiella pneumoniae</i>	31	IMP-1	0.125–64	0.125–64	120 bp (31)
<i>Enterobacter cloacae</i>	18	IMP-1	0.125–4	\leq 0.06–4	120 bp (18)
<i>Klebsiella oxytoca</i>	9	IMP-1	0.125–1	0.125–2	120 bp (9)
<i>Escherichia coli</i>	5	IMP-1	0.125–1	0.5–2	120 bp (5)
<i>Citrobacter freundii</i>	2	IMP-1	0.25–1	0.125–0.5	120 bp (2)
<i>Escherichia coli</i>	170	IMP-6	\leq 0.06–1	0.125–32	350 bp (170)
<i>Klebsiella pneumoniae</i>	87	IMP-6	\leq 0.06–0.5	0.25–16	350 bp (87)
<i>Enterobacter cloacae</i>	1	IMP-6	0.125	0.25	350 bp (1)
<i>Klebsiella pneumoniae</i>	12	KPC	2–32	2–64	None
<i>Escherichia coli</i>	12	NDM	1–16	0.5–32	None
<i>Klebsiella pneumoniae</i>	2	OXA-48 like	0.5–1	0.5	None
<i>Citrobacter freundii</i>	1	VIM	1	1	None

*120-bp band, IMP-1-specific band; 350-bp band, IMP-6-specific band.

Abbreviations: MIC, minimum inhibitory concentration; IPM, imipenem; MEPM, meropenem; ARMS, amplification refractory mutation system.

REFERENCES

1. Gupta N, Limbago BM, Patel JB, Kallen AJ. Carbapenem-resistant Enterobacteriaceae: epidemiology and prevention. *Clin Infect Dis* 2011;53:60-7.
2. Nordmann P, Naas T, Poirel L. Global spread of carbapenemase-producing Enterobacteriaceae. *Emerg Infect Dis* 2011;17:1791-8.
3. Yano H, Ogawa M, Endo S, Kakuta R, Kanamori H, Inomata S, et al. High frequency of IMP-6 among clinical isolates of metallo- β -lactamase-producing *Escherichia coli* in Japan. *Antimicrob Agents Chemother* 2012;56:4554-5.
4. Saito K, Nakano R, Suzuki Y, Nakano A, Ogawa Y, Yonekawa S, et al. Suitability of carbapenem inactivation method (CIM) for detection of IMP metallo- β -lactamase-producing Enterobacteriaceae. *J Clin Microbiol* 2017;55:1220-2.
5. Yano H, Kuga A, Okamoto R, Kitasato H, Kobayashi T, Inoue M. Plasmid-encoded metallo- β -lactamase (IMP-6) conferring resistance to carbapenems, especially meropenem. *Antimicrob Agents Chemother* 2001;45:1343-8.
6. Kayama S, Shigemoto N, Kuwahara R, Onodera M, Yokozaki M, Ohge H, et al. Rapid detection of *bla*_{IMP-6} by amplification refractory mutation system. *J Microbiol Methods* 2012;88:182-4.
7. Newton CR, Graham A, Heptinstall LE, Powell SJ, Summers C, Kalsheker N, et al. Analysis of any point mutation in DNA. The amplification refractory mutation system (ARMS). *Nucleic Acids Res* 1989;17:2503-16.
8. Nakano R, Nakano A, Hikosaka K, Kawakami S, Matsunaga N, Asahara M, et al. First report of metallo- β -lactamase NDM-5-producing *Escherichia coli* in Japan. *Antimicrob Agents Chemother* 2014;58:7611-2.
9. CLSI. Performance standards for antimicrobial susceptibility testing, 22nd ed. CLSI supplement M100-S22. Wayne, PA: Clinical and Laboratory Standards Institute; 2012.