

Microbiology Spectrum

The emerging nosocomial pathogen *Klebsiella michiganensis*: genetic analysis of a KPC-3 producing strain isolated from Venus clam

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Corresponding Author(s): Carla Vignaroli, Universita Politecnica delle Marche

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1st Editorial Decision November 23,

20221

November 23, 2022

Dr. Carla Vignaroli Universita Politecnica delle Marche Department of Life and Environmental Sciences via Brecce Bianche Ancona 60131 Italy

Re: Spectrum04235-22 (The emerging nosocomial pathogen *Klebsiella michiganensis*: genetic analysis of a KPC-3 producing strain isolated from Venus clam)

Dear Dr. Carla Vignaroli:

Thank you for submitting your manuscript to Microbiology Spectrum. When submitting the revised version of your paper, please provide (1) point-by-point responses to the issues raised by the reviewers as file type "Response to Reviewers," not in your cover letter, and (2) a PDF file that indicates the changes from the original submission (by highlighting or underlining the changes) as file type "Marked Up Manuscript - For Review Only". Please use this link to submit your revised manuscript - we strongly recommend that you submit your paper within the next 60 days or reach out to me. Detailed instructions on submitting your revised paper are below.

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Sincerely,

Sandeep Tamber

Editor, Microbiology Spectrum

Journals Department American Society for Microbiology 1752 N St., NW Washington, DC 20036 E-mail: spectrum@asmusa.org

Editor comments:

The sequence analysis portion of the method needs more details. Please provide the parameters used for the bioinformatic programs, and specify the type of tree constructed.

Reviewer comments:

Reviewer #1 (Comments for the Author):

Your work is about genome analysis of a K. michiganensis strain, including PCR assays detected resistance genes, WGS, and analysis of plasmid and VFs. However, your article conclusion extend to the whole environment. Maybe your work did not provide sufficient evidence to prove your conclusion.

Reviewer #2 (Comments for the Author):

The authors have presented a relevant and interesting area of research which is currently lacking in the literature. The methods used were appropriate and robust with experimental data alongside bioinformatic review to confirm findings. The experimental data and the bioinformatics support one another. The data presented will further add to our understanding of genome plasticity of Klebsiella sp. and the requirement for surveillance of environmental isolates that have the potential to reach the human food chain to understand transfer of resistance between species and the emergence of multidrug resistant strains.

Suggested edits.

Line 75. In this study, a KPC-producing strain of K. michiganensis isolated from clams collected from natural beds along the Adriatic Sea coast in central Italy was were characterized by WGS.

76 natural beds along the Adriatic Sea coast in central Italy was characterized by WGS.

Line 86. MIC determination for ertapenem, imipenem and meropenem showed that the isolate was resistant to the three carbapenems with MIC values of 256, 16 and 8 µg/ml respectively.

Line 133. These findings also highlighted that pKm_8 could be originated from genetic recombination Requires some editing for grammar.

Staff Comments:

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Thank you for submitting your paper to Microbiology Spectrum.

The emerging nosocomial pathogen *Klebsiella michiganensis*: genetic analysis of a KPC-3 producing strain isolated from Venus clam.

The authors recovered and characterised an isolate of multidrug resistant KPC-3 producing *Klebsiella michiganensis* from a Venus clam mollusc collected from natural beds in the Adriatic Sea coast in central Italy. Using whole genome sequencing genetic elements carrying beta lactam resistance genes were analysed and the ability of the strain to transfer carbapenem resistance by conjugation was investigated. The isolate was identified as being in ST382 with seven plasmid replicons.

The authors have presented a relevant and interesting area of research which is currently lacking in the literature. The methods used were appropriate and robust with experimental data alongside bioinformatic review to confirm findings. The experimental data and the bioinformatics support one another. The data presented will further add to our understanding of genome plasticity of Klebsiella sp. and the requirement for surveillance of environmental isolates that have the potential to reach the human food chain to understand transfer of resistance between species and the emergence of multidrug resistant strains.

Suggested edits.

Line 75. In this study, a KPC-producing strain of K. michiganensis isolated from clams collected from natural beds along the Adriatic Sea coast in central Italy was were characterized by WGS.

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Line 133. These findings also highlighted that pKm_8 could be originated from genetic recombination Requires some editing for grammar.

"Response to Reviewers"

Editor comments:

The sequence analysis portion of the method needs more details. Please provide the parameters used for the bioinformatic programs, and specify the type of tree constructed.

As suggested further details on bioinformatic analysis and the type of phylogenetic tree have been provided.

Reviewer comments:

Reviewer #1 (Comments for the Author):

Your work is about genome analysis of a K. michiganensis strain, including PCR assays detected resistance genes, WGS, and analysis of plasmid and VFs. However, your article conclusion extend to the whole environment. Maybe your work did not provide sufficient evidence to prove your conclusion.

Our conclusions have been suggested not only by WGS analysis of our strain but also from pan-genome analysis of all *K. michiganensis* genomes included in the database.

Reviewer #2 (Comments for the Author):

The authors have presented a relevant and interesting area of research which is currently lacking in the literature. The methods used were appropriate and robust with experimental data alongside bioinformatic review to confirm findings. The experimental data and the bioinformatics support one another. The data presented will further add to our understanding of genome plasticity of Klebsiella sp. and the requirement for surveillance of environmental isolates that have the potential to reach the human food chain to understand transfer of resistance between species and the emergence of multidrug resistant strains. Thank you to the reviewer for the positive comments.

Suggested edits.

Line 75. In this study, a KPC-producing strain of K. michiganensis isolated from clams collected from natural beds along the Adriatic Sea coast in central Italy was were characterized by WGS. 76 natural beds along the Adriatic Sea coast in central Italy was characterized by WGS. The sentence was rephrased.

Line 86. MIC determination for ertapenem, imipenem and meropenem showed that the isolate was resistant to the three carbapenems with MIC values of 256, 16 and 8 μ g/ml respectively. As suggested the sentence has been corrected.

Line 133. These findings also highlighted that pKm_8 could be originated from genetic recombination Requires some editing for grammar.

The sentence has been modified.

November 29, 2022

Dr. Carla Vignaroli Universita Politecnica delle Marche Department of Life and Environmental Sciences via Brecce Bianche Ancona 60131 Italy

Re: Spectrum04235-22R1 (The emerging nosocomial pathogen *Klebsiella michiganensis*: genetic analysis of a KPC-3 producing strain isolated from Venus clam)

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Sincerely,

Sandeep Tamber Editor, Microbiology Spectrum

Journals Department American Society for Microbiology 1752 N St., NW Washington, DC 20036 E-mail: spectrum@asmusa.org

Supplemental Material: Accept