

Reducing the Spread of Drug-resistant Bacteria in the Healthcare Network Using Mathematical Modelling Approach

Monika J. Piotrowska, Konrad Sakowski, Agata Lonc, Mirjam E. Kretzschmar, Rafael Mikolajczyk, Johannes Horn and Andre Karch

We compare the effectiveness and estimate the costs of selected intervention strategies designed to reduce the spread of multidrug-resistant bacteria in a regional network of healthcare facilities. The proposed model is based on our previous research [1]–[5] and combines intra-hospital pathogen transmission with direct and indirect inter-hospital transfers and readmissions of patients. Additionally, we distinguish three patient groups: a class of patients with a single hospital stay and two classes with low and high patient susceptibilities to colonisation with multidrug-resistant bacteria. The considered facility-level intervention strategies are applied to either all patients or to selected patient groups. While it is just the individual facilities or patient groups in facilities that undergo interventions, they are selected specifically to lower the prevalence in the whole network. The simulations, based on German anonymised insurance data, suggest that strategies aimed at selected patient groups may significantly reduce the system-wide prevalence and be more cost-effective than those aimed at the whole hospital population, up to a certain cost.

References

- [1] P. Brachaczek, A. Lonc, M.E. Kretzschmar, R.T. Mikolajczyk, J. Horn, A. Karch, K. Sakowski, M.J. Piotrowska, *Transmission of drug-resistant bacteria in a hospital-community model stratified by patient risk*, Sci. Rep. 13, 18593 (2023)
- [2] M.J. Piotrowska, A. Puchalska, K. Sakowski, *On the network suppression of the pathogen spread within the healthcare system*, Appl. Math. Comput. 457, 128169 (2023)
- [3] M.J. Piotrowska, K. Sakowski, J. Horn, R. Mikolajczyk, A. Karch, *The effect of re-directed patient flow in combination with targeted infection control measures on the spread of multi-drug-resistant Enterobacteriaceae in the German health-care system: a mathematical modelling approach*, CMI 29(1), P109.E1-109.E7 (2023)
- [4] M.J. Piotrowska, K. Sakowski, A. Karch, H. Tahir, J. Horn, M.E. Kretzschmar, R.T. Mikolajczyk, *Modelling pathogen spread in a healthcare network: indirect patient movements*, PLoS Comput. Biol. 16 (11), e1008442 (2020)
- [5] M.J. Piotrowska, K. Sakowski, A. Lonc, H. Tahir, M.E. Kretzschmar, *Impact of inter-hospital transfers on the prevalence of resistant pathogens in a hospital-community system*, Epidemics 33, 100408 (2020)

- First Author:** Monika J., Piotrowska
Affiliation: *Institute of Applied Mathematics and Mechanics, University of Warsaw
02-097, Poland*
e-mail: monika@mimuw.edu.pl
- Second Author:** Konrad, Sakowski
Affiliation: *Institute of Applied Mathematics and Mechanics, University of Warsaw
02-097, Poland*
e-mail: konrad@mimuw.edu.pl
- Third Author:** Agata, Lonc
Affiliation: *Institute of Applied Mathematics and Mechanics, University of Warsaw
02-097, Poland*
e-mail: a.lonc@uw.edu.pl
- Fourth Author:** Mirjam E., Kretzschmar
Affiliation: *University Medical Center Utrecht, Utrecht University
3584 CG, The Netherlands*
e-mail: M.E.E.Kretzschmar@umcutrecht.nl
- Fifth Author:** Rafael, Mikolajczyk
Affiliation: *Institute for Medical Epidemiology, University Halle-Wittenberg
06108, Germany*
e-mail: rafael.mikolajczyk@uk-halle.de
- Sixth Author:** Johannes, Horn
Affiliation: *Institute for Medical Epidemiology, University Halle-Wittenberg
06108, Germany*
e-mail: johannes.horn@uk-halle.de
- Seventh Author:** Andre, Karch
Affiliation: *Institute for Epidemiology and Social Medicine, University of Münster
48149, Germany*
e-mail: andre.karch@ukmuenster.de