



Zdunya, 17th–21st September 2024

INVESTIGATION OF THE EFFECTIVENESS OF PREVENTION STRATEGIES AGAINST DRUG-RESISTANT BACTERIA IN THE HEALTHCARE NETWORK

Monika J. Piotrowska^{1,6}, Konrad Sakowski^{1,2,7}, Agata Lonc^{1,8}, Mirjam E. Kretzschmar^{3,9}, Rafael Mikolajczyk^{4,10}, Johannes Horn^{5,11} and Andre Karch^{4,12}

¹Institute of Applied Mathematics and Mechanics, University of Warsaw
Banacha 2, 02-097 Warszawa,

²Institute of High Pressure Physics, Polish Academy of Sciences,
Sokolowska 29/37, 01-142 Warszawa,

³University Medical Center Utrecht, Utrecht University
3584 CG, Utrecht, The Netherlands,

⁴Institute for Medical Epidemiology, University Halle-Wittenberg
06108, Halle (Saale), Germany

⁵ Institute for Epidemiology and Social Medicine, University of Münster
48149, Münster, Germany

⁶monika@mimuw.edu.pl, ⁷konrad@mimuw.edu.pl, ⁸a.lonc@uw.edu.pl,

⁹M.E.E.Kretzschmar@umcutrecht.nl, ¹⁰rafael.mikolajczyk@uk-halle.de,

¹¹johannes.horn@uk-halle.de, ¹²andre.karch@ukmuenster.de

ABSTRACT

Multidrug-resistant bacterial infections are a problematic kind of healthcare-associated infections (HAI), which spread within the healthcare system due to favorable conditions to develop. Each year over 3.8 million patients acquire HAIs and over 90,000 people lose their lives due to HAIs in the European Economic Area only. Thus, our aim is to propose HAI preventive strategies and evaluate them by the means of numerical simulations. It is the extension of our previous work [1]–[5].

In this study, a few qualitatively different strategies are considered to determine whether it is beneficial to target specific groups of patients in specific hospitals, like patients highly susceptible to HAIs, from the standpoint of lowering the prevalence in the whole healthcare system. This is a theoretical research, based on numerical simulations of a regional network of healthcare facilities. The simulations are based on the deterministic mathematical model of a pathogen spread within a healthcare network. The model includes patient transfers, patient readmissions and stratification of patients into risk groups. Simulations are performed with an open-source library developed within this study, and they use a computer model of Lower Saxony healthcare network, created by the analysis of anonymized insurance records. The simulations are performed under the assumption that individual hospitals have preventive means to reduce HAI transmission, which are not

explicitly stated, but the available funds are insufficient to introduce those means at a sufficient scale to simply eradicate HAIs.

The considered strategies aim at selecting individual hospitals and risk groups to impose those means to effectively reduce healthcare system-level prevalence for the smallest possible cost. If additional funds are low, the efficiency of many strategies is similar, with risk group stratification or not. If funds are very high, then even basic strategies are sufficient to lower the prevalence to a negligible level. Targeting individual groups is more sophisticated, and if the additional costs are moderate, then it is possible to obtain a considerably larger decrease in prevalence for the same cost in comparison to targeting all groups. In addition, targeting only highly susceptible patients does not allow for reducing prevalence to negligible levels. Other groups must also be targeted to reach such a goal.

ACKNOWLEDGEMENTS

The work was carried out within the Excellence Initiative – Research University, University of Warsaw, New Ideas in Priority Research Area III “Analysis of coordinated countermeasures against spread of hospital-acquired infections within healthcare systems by numerical simulations based on mathematical models” (AL, MJP, KS) and it was a continuation of the work conducted within the 3rd JPI AMR framework cofound grant no 681055 for the consortium EMerGE-Net (Effectiveness of infection control strategies against intra- and inter-hospital transmission of Multidrug-resistant Enterobacteriaceae) funded by National Science Centre, Poland, Grant Number 2016/22/Z/ST1/00690 (MJP, KS, AL); Netherlands ZonMw, Grant Number 547001005 (MEK); Bundesministerium für Bildung und Forschung, Grant Number 01KI1704C (JH, AK, RM).

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** (2023), 18593.
- [2] M.J. Piotrowska, A. Puchalska, K. Sakowski, *On the network suppression of the pathogen spread within the healthcare system*, Appl. Math. Comput. **457** (2023), 128169.
- [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*, Clinical Microbiology and Infection **29**(1) (2023), P109.E1-109.E7.
- [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) (2020), e1008442.
- [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** (2020), 100408.