



Letter to the Editor

Predictors of mortality of bloodstream infections among internal medicine patients: Mind the complexity of the septic population!



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Sir,

We read with great interest the article by Papadimitriou-Olivgeris and colleagues [1] aiming to evaluate the predictors of mortality among internal medicine patients with bloodstream infections (BSIs). As main result, qSOFA, compared to SIRS and other severity scores, showed better performances in predicting mortality with a high negative predicting value [1].

Our research group has recently conducted a similar evaluation on a cohort of internal medicine patients affected by microbiologically identified BSIs [2]. Despite similarities among the two groups in terms of morbidity and mortality [1,2], the independent predictors of mortality were MEDS scores and vitamin D levels. On the contrary, qSOFA and other severity scores did not show good performances [2].

In our opinion, the reasons for these apparently controversial results could be manifold. First, data on septic patients admitted to internal medicine wards are few with respect to data derived from other contexts (e.g. Emergency Departments and Intensive Care Units) [3–5]. Moreover, the population of septic patients admitted to Internal Medicine wards is heterogeneous [5–8]. With this regard, we agree with the Authors on the complexity of the studied population, and the presence of several comorbidities, in particular malignancies, associated with the worst outcome [1]. All these reasons could favor the presentation variability of septic patients in terms of phenotypic characteristics, host-response, biomarkers profile, and clinical outcomes.

A recently published study by Seymour and colleagues [9] has described the presence of at least four clinical phenotypes of sepsis, which correlated with host-response patterns and prognosis. Probably, in the next future these phenotypes will be of help in understanding the heterogeneity of treatment responses.

Sepsis is a complex syndrome or rather a set of syndromes, characterized by clinical manifestations and evolution that depend on innumerable factors [10]. For this complexity – due to the set of different infections, different individual responses, variegated clinical manifestations and widely varying evolution – it is unlikely to think that a single clinical score or a single laboratory value can give a unique answer in the prognostic field. A combination of multiple predictors (clinical scores and laboratory data) could probably represent the way for a better prognostication. However, future studies are needed to determine the usefulness of sepsis phenotypes in clinical care,

particularly in the context of non-ICU wards where most of septic patients are hospitalized [7].

Declaration of Competing Interest

The authors report no conflicts of interest.

Appendix A. Internal Medicine Sepsis Study Group

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