Supplementary Materials

In-hospital mortality from healthcare-associated infection by multidrug-resistant *Pseudomonas aeruginosa*: a competing risks analysis of a 4-year propensity-matched cohort study in southern China

Zhou M, Kritsotakis EI, Xu B, Guo Z, Zeng Y, Zhou B, Brinks R, Wang J

Table of Contents

R code for main analyses	2
Table S1. Multicollinearity analysis	
Table S2. STROBE Statement—Checklist of items that should be included in reports of cohort studies	6
Figure S1. Flow chart of data collection and patient selection	8
Figure S2. Length of hospitalisation of patients with healthcare-associated infections by <i>P. aeruginosa</i>	9
Figure S3. Incidence rate trends for healthcare-associated infections by non-multidrug-resistant <i>P. aeruginosa</i> (non-MDRPA), 2018-2021	11
Table S3. Results of multivariable competing risks survival analysis in the matched samples	13
Table S4. Results of multivariable competing risks analysis in the unmatched samples	14

R code for main analyses

library("MatchIt")

Matching based on the estimated propensity scores

```
formula <- mdr ~ age + sex + admission_department + infection_site + diabetes + immunocompromised + COVID_period matchit_model <- matchit(formula, data=df, distance='logit', method='nearest', replace=FALSE, caliper=0.2, discard=, ratio=1)
```

Check the balance after matching

summary(matchit_model, standardize = TRUE)

Competing risks 14-day survival analysis after matching

library(survival)

survival analysis variables

```
# define time and event variables (from infection onset to death/dicharge)

df2$dthtm= as.numeric(round(df2$discharge_date - df2$infection_date))

df2$dth <- ifelse(df2$alive mortality == 1, 1, 0) # alive mortality == 1 = died [dth==1 = died]
```

admin right censoring at 14 days after infection

```
df2$dthtm14 <- df2$dthtm

df2$dthtm14[df2$dthtm14 > 14] <- 14

df2$dth14 <- ifelse(df2$dthtm<=14 & df2$dth ==1, 2, 1)

df2$dth14[df2$dthtm>14] <- 0
```

#14d in-hospital death

```
# bivariable analysis
```

```
fgmodel\_mtch\_univr1\_d14 <- FGR(Hist(dthtm14, dth14) \sim mdr \ , \ data = df2, \ cause = 2) \\ \qquad \# \ dth14=2 \ when \ inpatient \ death print(fgmodel\_mtch\_univr1\_d14)
```

Attachment 1 to: Zhou M, Kritsotakis EI, Xu B, Guo Z, Zeng Y, Zhou B, Brinks R, Wang J. In-hospital mortality from healthcare-associated infection by multidrug-resistant Pseudomonas aeruginosa: a competing risks analysis of a 4-year propensity-matched cohort study in southern China. GMS Hyg Infect Control. 2025;20:Doc68. DOI: 10.3205/ddgkh000597

```
# multivariable analysis
```

#14d discharge alive

bivariable analysis

fgmodel_mtch_univr2_d14 <- FGR(Hist(dthtm14, dth14) ~ mdr , data = df2, cause = 1) # dth14=1 is discharge alive print(fgmodel_mtch_univr2_d14)

multivariable analysis

fgmodel_mtch_multivar2_d14 <- FGR(Hist(dthtm14, dth14) ~ mdr + infection_site + empiric_therapy

+ age65 + sex + admission department + admission diagnosis + diabetes + immunocompromised

+ COVID_period , data = df2, cause = 1) # dth14=1 is discharge alive

print(fgmodel_mtch_multivar2_d14)

Table S1. Multicollinearity analysis

Variable	Generalised Variance Inflation Factor	Degrees of Freedom	Adjusted Generalized Variance Inflation Factor			
Age ≥65 years	1.160	1	1.077			
Sex = Male	1.045	1	1.022			
Admission Department	1.950	3	1.118			
Admission Diagnosis	2.123	5	1.078			
Infection Site	1.220	3	1.034			
Receipt of empiric therapy	1.287	1	1.134			
Diabetes = Yes	1.056	1	1.028			
Immunocompromised = Yes	1.150	1	1.073			
COVID Period	1.228	2	1.053			

Note: Generalised Variance Inflation Factor (GVIF) analysis for the included variables showed no evidence of multicollinearity. The adjusted GVIF values for all variables were close to 1, indicating low correlation between predictors and suggesting that multicollinearity is not an issue for this set of covariates.

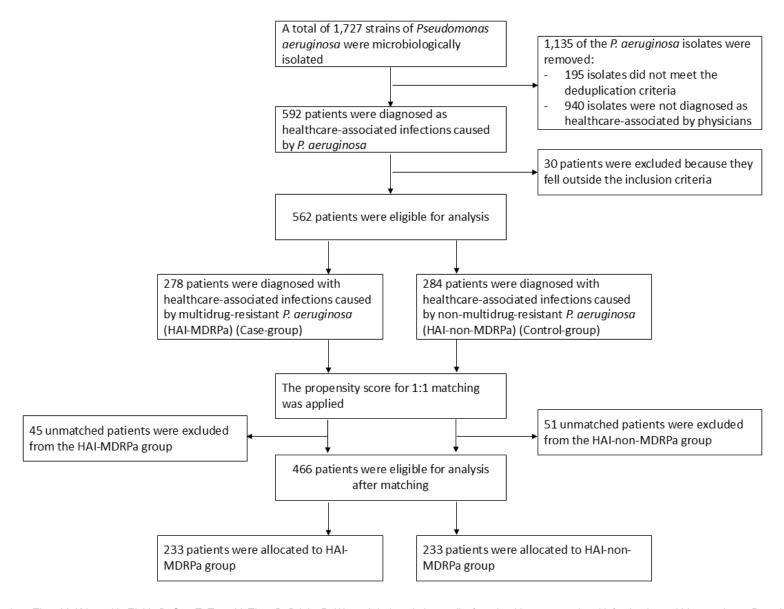
Table S2. STROBE Statement—Checklist of items that should be included in reports of cohort studies

	No	Recommendation	Page(s)
		(a) Indicate the study's design with a commonly used term in the title or the abstract	1
Title and abstract	1	(b) Provide in the abstract an informative and balanced summary of what was done and what was found	3
Introduction	.		
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	4-5
Objectives	3	State specific objectives, including any prespecified hypotheses	5
Methods	•		
Study design	4	Present key elements of study design early in the paper	5
Setting	5	Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection	5
Darticipanto	6	(a) Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up	5-6
Participants	0	(b) For matched studies, give matching criteria and number of exposed and unexposed	7
Variables	7	Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable	5
Data sources/ measurement	8	For each variable of interest, give sources of data and details of methods of assessment (measurement). Describe comparability of assessment methods if there is more than one group	5-6
Bias	9	Describe any efforts to address potential sources of bias	N/A
Study size	10	Explain how the study size was arrived at	N/A
Quantitative variables	11	Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why	5-6
		(a) Describe all statistical methods, including those used to control for confounding	6-8
Statistical methods	12	(b) Describe any methods used to examine subgroups and interactions	Subgroup modelling of incidence trends for major types of HAIs-MDRPa, Figure 1 and Figure S3
		(c) Explain how missing data were addressed	8
		(d) If applicable, explain how loss to follow-up was addressed	N/A
		(e) Describe any sensitivity analyses	Table 2
Results			

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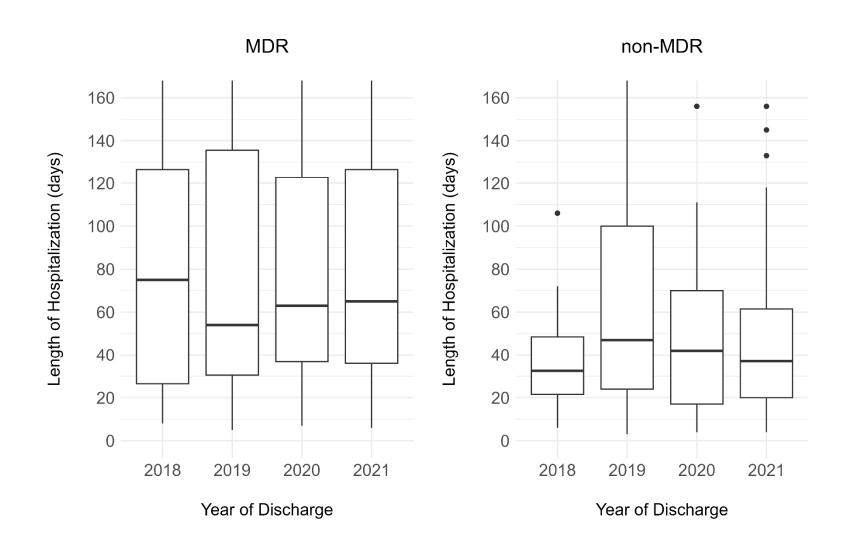
Participants	13	(a) Report numbers of individuals at each stage of study—e.g., numbers potentially eligible, examined for eligibility, confirmed eligible, included in the study, completing follow-up, and analysed	8, Table 1		
		(b) Give reasons for non-participation at each stage	Supplementary Figure S1		
		(c) Consider use of a flow diagram	Supplementary Figure S1		
December date	4.4	(a) Give characteristics of study participants (e.g., demographic, clinical, social) and information on exposures and potential confounders	8		
Descriptive data	14	(b) Indicate number of participants with missing data for each variable of interest	N/A		
		(c) Summarise follow-up time (e.g., average and total amount)	N/A		
Outcome data	15	Report numbers of outcome events or summary measures over time	8-9		
		(a) Give unadjusted estimates and, if applicable, confounder-adjusted estimates and their precision (e.g., 95% confidence interval). Make clear which confounders were adjusted for and why they were included	Table 2 and Supplementary Tables S3 and S4.		
Main results	16	(b) Report category boundaries when continuous variables were categorized	N/A		
		(c) If relevant, consider translating estimates of relative risk into absolute risk for a meaningful time period	We reported subdistribution hazard ratio		
Other analyses	Report other analyses done—e.g. analyses of subgroups and interactions, and				
Discussion					
Key results	18	Summarise key results with reference to study objectives	10-11		
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias or imprecision. Discuss both direction and magnitude of any potential bias	12		
Interpretation	20	Give a cautious overall interpretation of results considering objectives, limitations, multiplicity of analyses, results from similar studies, and other relevant evidence	10-11		
Generalisability	21	Discuss the generalisability (external validity) of the study results	N/A		
Other information					
Funding	22	Give the source of funding and the role of the funders for the present study and, if applicable, for the original study on which the present article is based	13		

Figure S1. Flow chart of data collection and patient selection



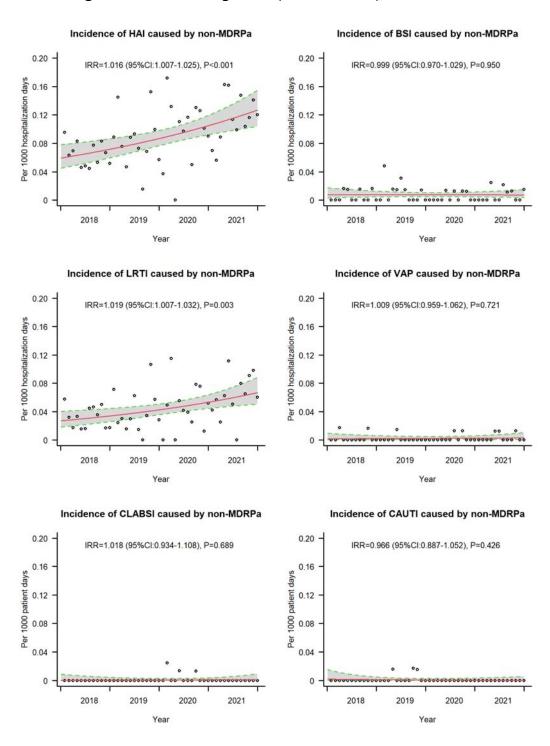
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Figure S2. Length of hospitalisation of patients with healthcare-associated infections by P. aeruginosa



MDR, multidrug resistance

Figure S3. Incidence rate trends for healthcare-associated infections by non-multidrug-resistant *P. aeruginosa* (non-MDRPA), 2018-2021



BSI, Bloodstream infection; CAUTI, catheter-associated urinary tract infection; CI, confidence Interval; CLABSI, central line-associated bloodstream infection; HAI, healthcare-associated

infection; IRR, mean monthly incidence rate ratio; LRTI, lower respiratory tract infection; MDRPa,
multidrug-resistant Pseudomonas aeruginosa; VAP, ventilator-associated pneumonia.

Table S3. Results of multivariable competing risks survival analysis in the matched samples

		ays		ys	Overall							
	In-hospital death Discharge alive			live	In-hospital de	Discharge alive		In-hospital death		Discharge alive		
Variable	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р
Multidrug-resistant	1.07 (0.52 - 2.19)	0.850	0.44 (0.31 - 0.63)	<0.001	0.95 (0.51 - 1.77)	0.860	0.56 (0.43 - 0.74)	<0.001	1.37 (0.78 - 2.39)	0.280	0.66 (0.54 - 0.82)	<0.001
Age > 65 years	1.34 (0.67 - 2.72)	0.410	0.55 (0.38 - 0.82)	0.003	1.39 (0.74 - 2.60)	0.300	0.68 (0.51 - 0.91)	0.010	1.64 (0.94 - 2.88)	0.082	0.70 (0.55 - 0.89)	0.004
Male Sex	1.21 (0.49 - 3.00)	0.680	0.87 (0.58 - 1.31)	0.510	1.35 (0.60 - 3.00)	0.470	0.90 (0.66 - 1.22)	0.480	1.19 (0.63 - 2.22)	0.600	0.89 (0.69 - 1.14)	0.350
Admission Department												
Internal Medicine	0.04 (0.00 - 0.35)	0.004	1.01 (0.58 - 1.75)	0.980	0.05 (0.01 - 0.20)	<0.001	1.21 (0.81 - 1.82)	0.350	0.11 (0.04 - 0.30)	<0.001	1.93 (1.30 - 2.87)	0.001
Surgery	0.09 (0.03 - 0.31)	<0.001	0.75 (0.45 - 1.26)	0.270	0.08 (0.03 - 0.23)	<0.001	0.76 (0.50 - 1.16)	0.210	0.11 (0.05 - 0.25)	<0.001	1.51 (1.04 - 2.19)	0.030
Other	0.00 (0.00 - 0.00)	<0.001	0.61 (0.34 - 1.11)	0.110	0.00 (0.00 - 0.00)	<0.001	0.69 (0.44 - 1.07)	0.098	0.06 (0.01 - 0.28)	<0.001	1.47 (1.02 - 2.10)	0.036
Admission Diagnosis												
Neurological disorder	1.65 (0.32 - 8.41)	0.550	0.63 (0.28 - 1.42)	0.270	1.18 (0.34 - 4.08)	0.790	0.64 (0.35 - 1.17)	0.150	1.07 (0.36 - 3.16)	0.910	0.76 (0.44 - 1.31)	0.320
Respiratory disease	0.84 (0.18 - 4.01)	0.830	1.72 (0.78 - 3.78)	0.180	0.55 (0.16 - 1.85)	0.330	1.33 (0.72 - 2.48)	0.360	0.54 (0.19 - 1.58)	0.260	1.29 (0.71 - 2.34)	0.400
Injury or orthopaedic condition	0.39 (0.03 - 5.74)	0.490	0.65 (0.28 - 1.53)	0.320	0.48 (0.07 - 3.37)	0.460	0.70 (0.36 - 1.35)	0.290	0.26 (0.04 - 1.68)	0.160	0.91 (0.52 - 1.60)	0.750
Cancer or related disorder	2.44 (0.38 - 15.76)	0.350	0.71 (0.29 - 1.73)	0.450	1.59 (0.36 - 7.06)	0.540	0.68 (0.34 - 1.37)	0.280	1.43 (0.38 - 5.39)	0.590	0.65 (0.34 - 1.23)	0.190
Other diagnosis	1.71 (0.32 - 9.19)	0.530	1.23 (0.58 - 2.60)	0.590	0.97 (0.26 - 3.66)	0.970	1.30 (0.74 - 2.29)	0.360	1.14 (0.38 - 3.47)	0.810	1.24 (0.73 - 2.12)	0.420
Infection Site												
Lower Respiratory Tract	1.09 (0.28 - 4.23)	0.900	0.55 (0.35 - 0.86)	0.008	0.99 (0.36 - 2.72)	0.990	0.61 (0.43 - 0.88)	0.008	1.08 (0.42 - 2.82)	0.870	0.69 (0.51 - 0.95)	0.021
Urinary Tract	1.80 (0.37 - 8.74)	0.470	1.01 (0.55 - 1.85)	0.990	1.00 (0.25 - 4.02)	1.000	0.73 (0.43 - 1.26)	0.260	1.59 (0.50 - 5.13)	0.430	0.64 (0.41 - 0.99)	0.043
Other Site	2.35 (0.55 - 10.07)	0.250	0.97 (0.60 - 1.58)	0.900	1.86 (0.58 - 5.95)	0.300	0.90 (0.60 - 1.37)	0.640	1.96 (0.67 - 5.77)	0.220	0.79 (0.55 - 1.14)	0.210
Receipt of empiric therapy	0.65 (0.23 - 1.84)	0.420	1.29 (0.73 - 2.25)	0.380	0.79 (0.33 - 1.90)	0.600	1.11 (0.74 - 1.67)	0.610	0.63 (0.31 - 1.28)	0.200	1.22 (0.93 - 1.61)	0.160
Diabetes	1.46 (0.34 - 6.26)	0.610	0.81 (0.22 - 2.97)	0.750	1.97 (0.67 - 5.79)	0.220	1.06 (0.45 - 2.49)	0.900	2.11 (0.66 - 6.73)	0.210	0.93 (0.44 - 1.94)	0.840
Immunocompromised	1.97 (0.45 - 8.67)	0.370	1.23 (0.59 - 2.57)	0.580	2.98 (0.76 - 11.76)	0.120	1.14 (0.67 - 1.95)	0.640	1.32 (0.34 - 5.09)	0.690	1.16 (0.78 - 1.73)	0.450
Year												
2020	0.50 (0.20 - 1.22)	0.130	0.89 (0.56 - 1.39)	0.600	0.54 (0.24 - 1.22)	0.140	0.95 (0.67 - 1.36)	0.790	0.60 (0.29 - 1.25)	0.170	1.24 (0.95 - 1.61)	0.120
2021	0.46 (0.18 - 1.15)	0.098	1.27 (0.85 - 1.89)	0.250	0.44 (0.20 - 0.95)	0.036	1.43 (1.04 - 1.97)	0.030	0.46 (0.23 - 0.92)	0.028	1.64 (1.25 - 2.15)	<0.001

Note: The reference group is the intensive care unit for the admission department, cardiovascular disease for the admission diagnosis, bloodstream infection for the infection site, and 2018-2019 for the year

sHR: subdistribution hazard ratio from the Fine-Gray model; 95% CI: 95% confidence interval.

Table S4. Results of multivariable competing risks analysis in the unmatched samples

		ays		ıys	Overall							
	In-hospital de	Discharge alive		In-hospital death		Discharge alive		In-hospital death		Discharge a	live	
Variable	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р	sHR (95%CI)	Р
Multidrug-resistant	1.20 (0.62 - 2.35)	0.590	0.48 (0.34 - 0.68)	<0.001	1.14 (0.65 - 2.01)	0.640	0.61 (0.47 - 0.79)	<0.001	1.48 (0.89 - 2.49)	0.130	0.68 (0.56 - 0.83)	<0.001
Age > 65 years	1.14 (0.60 - 2.19)	0.690	0.68 (0.48 - 0.97)	0.033	1.33 (0.75 - 2.35)	0.330	0.80 (0.61 - 1.04)	0.100	1.58 (0.94 - 2.66)	0.087	0.75 (0.61 - 0.94)	0.011
Male Sex	1.50 (0.64 - 3.53)	0.350	0.80 (0.56 - 1.13)	0.210	1.78 (0.84 - 3.78)	0.140	0.80 (0.61 - 1.04)	0.096	1.53 (0.85 - 2.76)	0.160	0.83 (0.66 - 1.03)	0.097
Admission Department												
Internal Medicine	0.04 (0.01 - 0.24)	<0.001	0.95 (0.58 - 1.57)	0.850	0.05 (0.01 - 0.18)	<0.001	1.03 (0.70 - 1.50)	0.890	0.14 (0.06 - 0.34)	<0.001	1.75 (1.22 - 2.53)	0.002
Surgery	0.07 (0.02 - 0.23)	<0.001	0.76 (0.47 - 1.23)	0.260	0.07 (0.02 - 0.19)	<0.001	0.83 (0.57 - 1.21)	0.330	0.10 (0.04 - 0.22)	<0.001	1.51 (1.08 - 2.12)	0.017
Other	0.00 (0.00 - 0.00)	<0.001	0.69 (0.41 - 1.14)	0.150	0.02 (0.00 - 0.20)	<0.001	0.78 (0.53 - 1.15)	0.210	0.06 (0.02 - 0.21)	<0.001	1.58 (1.14 - 2.19)	0.007
Admission Diagnosis												
Neurological disorder	1.00 (0.25 - 3.95)	1.000	0.66 (0.31 - 1.39)	0.270	0.85 (0.28 - 2.58)	0.780	0.70 (0.39 - 1.24)	0.220	0.68 (0.27 - 1.74)	0.420	0.86 (0.52 - 1.41)	0.540
Respiratory disease	0.65 (0.18 - 2.34)	0.510	1.78 (0.85 - 3.75)	0.130	0.58 (0.20 - 1.69)	0.320	1.39 (0.77 - 2.53)	0.270	0.49 (0.20 - 1.21)	0.120	1.30 (0.74 - 2.28)	0.360
Injury or orthopaedic condition	0.25 (0.02 - 3.17)	0.290	0.62 (0.27 - 1.44)	0.270	0.34 (0.05 - 2.15)	0.250	0.74 (0.39 - 1.39)	0.350	0.17 (0.03 - 1.02)	0.053	0.96 (0.57 - 1.63)	0.890
Cancer or related disorder	1.78 (0.39 - 8.12)	0.450	0.70 (0.31 - 1.61)	0.410	1.34 (0.37 - 4.90)	0.660	0.85 (0.45 - 1.60)	0.610	0.91 (0.29 - 2.91)	0.880	0.81 (0.47 - 1.42)	0.470
Other diagnosis	1.06 (0.28 - 4.01)	0.930	1.15 (0.57 - 2.35)	0.690	0.79 (0.26 - 2.46)	0.690	1.35 (0.79 - 2.32)	0.280	0.79 (0.31 - 2.00)	0.620	1.32 (0.81 - 2.17)	0.270
Infection Site												
Lower Respiratory Tract	0.81 (0.29 - 2.26)	0.690	0.57 (0.37 - 0.87)	0.010	0.79 (0.34 - 1.82)	0.580	0.66 (0.46 - 0.94)	0.020	0.89 (0.40 - 1.96)	0.760	0.74 (0.54 - 1.00)	0.053
Urinary Tract	1.06 (0.29 - 3.90)	0.930	1.01 (0.58 - 1.77)	0.970	0.80 (0.26 - 2.51)	0.700	0.79 (0.48 - 1.29)	0.350	1.25 (0.48 - 3.26)	0.650	0.72 (0.48 - 1.07)	0.110
Other Site	1.56 (0.49 - 4.98)	0.460	0.80 (0.50 - 1.29)	0.370	1.17 (0.43 - 3.19)	0.760	0.85 (0.57 - 1.27)	0.440	1.36 (0.55 - 3.40)	0.510	0.84 (0.60 - 1.20)	0.340
Receipt of empiric therapy	0.59 (0.24 - 1.45)	0.250	1.35 (0.81 - 2.25)	0.250	0.72 (0.33 - 1.55)	0.400	1.27 (0.88 - 1.84)	0.210	0.59 (0.30 - 1.14)	0.120	1.32 (1.03 - 1.68)	0.029
Diabetes	0.78 (0.18 - 3.35)	0.740	0.45 (0.13 - 1.51)	0.200	1.19 (0.40 - 3.51)	0.760	0.72 (0.34 - 1.50)	0.370	1.76 (0.61 - 5.02)	0.290	0.72 (0.39 - 1.34)	0.300
Immunocompromised	5.16 (1.74 - 15.27)	0.003	1.06 (0.55 - 2.04)	0.850	4.46 (1.59 - 12.5)	0.004	0.96 (0.58 - 1.60)	0.880	2.20 (0.78 - 6.22)	0.140	1.00 (0.70 - 1.44)	1.000
Year												
2020	0.52 (0.24 - 1.12)	0.094	0.94 (0.62 - 1.44)	0.790	0.63 (0.33 - 1.21)	0.170	0.97 (0.69 - 1.35)	0.850	0.59 (0.33 - 1.08)	0.086	1.25 (0.97 - 1.60)	0.080
2021	0.36 (0.15 - 0.86)	0.022	1.25 (0.86 - 1.81)	0.240	0.40 (0.19 - 0.83)	0.014	1.43 (1.07 - 1.92)	0.015	0.38 (0.20 - 0.74)	0.004	1.62 (1.27 - 2.07)	<0.001

Note: The reference group is the intensive care unit for the admission department, cardiovascular disease for the admission diagnosis, bloodstream infection for the infection site, and 2018-2019 for the year

sHR: subdistribution hazard ratio from the Fine-Gray model; 95% CI: 95% confidence interval.