

# Predictors of Mortality in *Staphylococcus aureus* Bacteremia

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## INTRODUCTION

*Staphylococcus aureus* bacteremia (SAB) is a common and important infection. The exact incidence of SAB is difficult to ascertain, as prospective population-based surveillance studies are infrequently performed. In Scandinavian countries, where data from the nationwide surveillance of SAB are routinely collected, the annual incidence is approximately 26/100,000 population (14, 119, 128). A similar low incidence of 19.7/100,000 population was

reported in a Canadian study in 2008 (160), while in countries with a greater burden of methicillin-resistant *S. aureus* (MRSA),

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incidence rates are generally higher, between 35 and 39/100,000 population (38, 56, 233). In comparison, even higher rates, approximately 50/100,000 population, are inferred from surveillance data from the United States (143, 201). These large geographical discrepancies probably reflect differences in health care systems, infection control practices, and the completeness of surveillance data.

The incidence of SAB increases with advancing age, with the lowest rates observed in pediatric populations, at approximately 8.4/100,000 population per year (71). Similarly, younger adults have lower incidence rates than older adults (14, 160). Other factors associated with higher incidences include male gender, African American ethnicity, community-onset SAB, and specific patient subgroups that have frequent health care contact, including hemodialysis patients (160, 201, 229). Methicillin-sensitive *S. aureus* (MSSA) bacteremia (MSSA-B) episodes generally predominate, especially in countries with a low prevalence of MRSA (14, 38). Infecting clonal patterns vary geographically and are in a constant state of flux, as evidenced by the recent phenomenon, especially in the United States, of hospital strain replacement with typical community MRSA clones such as USA300 (229). Nevertheless, although not as common as other infections, such as tuberculosis or chronic hepatitis, SAB remains an important infection.

## Mortality

Once established, SAB is not a benign condition, resulting in significant morbidity and mortality, especially in patients in intensive care units (ICUs) (156, 265). SAB mortality rates of between 75% and 83% were observed in the preantibiotic era (182, 196). The introduction of antibiotics in the 1940s and 1950s resulted in better outcomes (320). Subsequently, with a greater understanding of SAB management, improved outcomes have continued to be documented throughout the 20th century, with overall mortality rates declining from 36% and 35% in 1981 to 1985 to 21% and 27% in 1996 to 2004 for hospital- and community-onset SAB episodes, respectively ( $P < 0.01$ ) (14). However, recent prospective data (2009) of 1,994 SAB episodes suggest that mortality rates may have stabilized, with a 30-day all-cause mortality rate of 20% (296) and an infection-related mortality rate estimated at approximately 13% (164). Despite these improvements, SAB 30-day all-cause mortality results in approximately 2 to 10 deaths annually per 100,000 population (7, 143, 160). Comparatively, the age-adjusted mortality rates are 3, 0.2, and 2.2 per 100,000 population for AIDS, tuberculosis, and viral hepatitis, respectively, while breast and prostate cancers claim 12.5 and 8.6 lives per 100,000 population annually, respectively (21, 145). Despite these similar mortality rates, the impact and significance of SAB in the community remain underestimated.

Multiple factors influence mortality. These factors include host factors, pathogen-host interactions, and pathogen-specific factors. The purpose of this review is to examine factors that influence mortality and investigate the relative impact of each factor on outcomes for patients with SAB.

## HOST FACTORS

### Age

Age is the most consistent and strongest predictor of all-cause and infection-related 30-day mortality, with the majority of SAB cohort studies using multivariate analysis, confirming age as an in-

dependent predictor of mortality (Table 1). In one of the largest population-based studies ( $n = 9,001$ ), where MRSA-B episodes were linked with death certificates to obtain all-cause 30-day mortality rates (155), the mortality rate was found to increase from 6% in young individuals ( $<15$  years old) to 57% in adults older than 85 years of age. This represents an approximately 1.3-fold increase in mortality from SAB for every 10 years of life (Fig. 1). These data are not explained by other problems that come with aging, which include the presence and number of comorbidities, indwelling medical devices, and health care contact (134). Some of the higher death rates may be attributed to differences in SAB-related investigations and management of elderly patients (15). However, in a case-controlled study, age remained a predictor of mortality despite elderly patients ( $>65$  years) having similar baseline characteristics and SAB management (284). Thus, the increased mortality from SAB associated with aging is directly linked to changes within the host as a consequence of the aging process, and age remains a significant confounder when examining other variables that influence outcomes.

### Gender

The incidence of SAB is generally higher for males than for females (143, 160). Despite this, several studies have shown an increased mortality rate for females. In several retrospective cohort studies of MRSA-B episodes, female gender was an independent predictor of 30-day all-cause mortality. The odds of death was approximately 2-fold higher than that for males in 116 (odds ratio [OR], 3.88; 95% confidence interval [CI], 1.23 to 12.2;  $P = 0.02$ ) (261), 510 (OR, 1.74; 95% CI, 1.13 to 2.68;  $P < 0.001$ ) (220), and 250 (OR, 1.97; 95% CI, 1.06 to 3.64;  $P = 0.017$ ) (185) MRSA-B episodes. Comparable results were seen in two larger studies of 908 (OR, 1.71; 95% CI, 1.02 to 2.84) and 814 (OR, 1.59; 95% CI, 1.09 to 2.34) SAB episodes from Spain (275) and Canada (5), respectively. Possible reasons for why gender may play a role in outcomes include different health-seeking behaviors adopted by women, the infecting MRSA clone, or hormonal differences (289). Alternatively, age differences may explain some of the observed gender differences, as the age-adjusted 30-day mortality rates were similar for males and females in a large retrospective English cohort study of 9,001 MRSA-B episodes (155).

Regardless, these explanations remain largely speculative, and without further evidence, gender is unlikely to be a major factor in SAB mortality, as the majority of studies (only studies with more than 500 SAB cases have been cited) have not detected any gender difference in regard to outcomes (14, 32, 101, 133, 134, 237, 296).

### Ethnicity

In the United States, African American populations have higher rates of SAB (61) and MRSA-B (74, 95, 143) than Caucasian patients. In 2005, in a U.S. population-based surveillance study, predominantly of health care-associated MRSA bacteremic episodes (66%; 5,813/8,792), the annual invasive MRSA infection incidence rate for Caucasians was 27.7/100,000 population (95% CI, 21.9 to 32.4), compared to 66.5/100,000 (95% CI, 43.5 to 63.1) for African Americans (143). Several other studies from New Zealand (109), Australia (96, 289), and the Pacific Islands (98) have documented ethnic differences in SAB incidences, with generally higher incidences in indigenous populations than in nonindigenous populations.

Despite the increased incidences in certain racial groups, the

TABLE 1 Summary of studies that have examined the impact of age on mortality in cases of *Staphylococcus aureus* bacteremia<sup>a</sup>

Study (outcome of interest)	No. of MSSA and MRSA bacteremic episodes	Overall mortality rate (%)	Age variable; OR (95% CI; <i>P</i> )	Reference
Studies that found age to be an independent predictor of mortality (multivariate analysis)				
Single-center retrospective Taiwan study (1990–2004); 30-day all-cause mortality	297 MSSA, 851 MRSA	44.1	Age >65 yr; 1.85 (1.37–2.50; <0.01)	308
Single-center retrospective Spanish study (1991–1998); in-hospital infection-related mortality	683 MSSA, 225 MRSA	12.1	Age per 10 yr; 1.17 (1.05–1.31; ND)	275
Single-center retrospective Canadian study (1991–2005); 30-day mortality	746 MSSA, 96 MRSA	24.0	Age >65 yr; 2.31 (1.43–3.73; <0.01)	5
Single-center prospective Spanish study (1991–2005); 30-day all-cause mortality	0 MSSA, 414 MRSA	28.0	Age per yr; 1.02 (1.00–1.04; 0.013)	273
Single-center retrospective Belgian study (1992–1998); 30-day in-hospital mortality	38 MSSA, 47 MRSA	37.6	Age per yr; 1.02 (1.00–1.02; 0.01)	18
Single-center prospective Danish study (1994–1996); 150-day mortality	275 MSSA, 3 MRSA	34	Age ≥60 yr; 2.4 (1.1–5.3; 0.03)	127
Single-center prospective U.S. study (1994–1998); 90-day all-cause mortality	385 SAB	20.5	Age >65 yr; 2.39 (1.45–3.95; ND)	192
Single-center retrospective U.S. study (1995–1999); 30-day all-cause mortality	204 MSSA, 89 MRSA	23.2	Age >65 yr; 2.0 (1.0–3.8; 0.048)	206
Single-center retrospective Chinese study (2001–2007); 90-day all-cause mortality	0 MSSA, 115 MRSA	21.7	Age per yr; 1.03 (1.01–1.05; 0.019)	32
Multicenter retrospective U.S. study (1995–2003); 90-day infection-related mortality	245 MSSA, 193 MRSA	26.0	Age >65 yr; 2.3 (1.38–3.8; <0.01)	266
Single-center retrospective U.S. study (1996–2006); all-cause mortality	0 MSSA, 489 MRSA	25.0	Age per yr; 1.05 (1.03–1.07; <0.001)	202
Single-center retrospective Taiwan study (1997–2001); 30-day infection-related mortality	0 MSSA, 162 MRSA	47.5	Age ≥60 yr; 2.90 (1.25–6.75; 0.01)	59
Multicenter retrospective United Kingdom study (1997–2004); 30-day mortality	229 MSSA, 232 MRSA	29.0	Age per decade; 1.79 (1.75–1.82; ND)	322
Single-center retrospective Australian study (1997–2008); 30-day all-cause mortality	0 MSSA, 401 MRSA	28.7	Age per yr; 1.03 (1.01–1.05; 0.005)	301
Single-center retrospective Swiss study (1998–2002); in-hospital all-cause mortality	302 MSSA, 6 MRSA	19.5	Age per 10 yr; 1.3 (1.2–1.6; <0.01)	131
Single-center retrospective Taiwan study (2001–2006); in-hospital infection-related mortality	0 MSSA, 177 MRSA	33.3	Age per yr; 1.04 (1.01–1.08; 0.01)	167
Single-center retrospective Taiwan study (2001–2007); 90-day all-cause mortality	0 MSSA, 115 MRSA	21.7	Age per yr; 1.01 (1.0–1.02; <0.05)	33
Single-center retrospective Germany study (2002–2004); in-hospital all-cause mortality	454 MSSA, 67 MRSA	21.7	Age ≥60 yr; 2.4 (1.4–4.2, <0.01)	237
Single-center retrospective Israeli study (2003–2006); in-hospital infection related-mortality	0 MSSA, 250 MRSA	37.2	Age per yr; 1.02 (1.004–1.04; 0.013)	185
Single-center retrospective U.S. study (2003–2008); in-hospital mortality	326 MSSA, 488 MRSA	13.3	Age per yr; 1.04 (1.03–1.05; ND)	257
Single-center prospective U.S. study (2004); in-hospital mortality	0 MSSA, 132 MRSA	22.0	Age >55 yr; 8.09 (2.02–32.5; 0.03)	261
Multicenter prospective <i>post hoc</i> analysis Asian study (2004–2006); 30-day all-cause mortality	380 MSSA, 329 MRSA	24.5	Age ≥65 yr; 1.69 (1.15–2.48; 0.008)	133
Multicenter prospective <i>post hoc</i> analysis Asian study (2004–2006); 30-day all-cause mortality	1,701 MSSA, 2,007 MRSA	13.4	Age per 10 yr; 1.19 (1.12–1.27; <0.01)	134
Single-center retrospective Taiwanese study (2004–2006); 30-day all-cause mortality	186 MSSA, 30 MRSA	12.6	Age per yr; 1.03 (1.01–1.06; <0.01)	310
Single-center retrospective Singaporean study (2005–2006); in-hospital infection-related mortality	100 MSSA, 0 MRSA	23.9	Age >65 yr; 13.7 (2.75–68.3; <0.01)	35
Single-center, prospective U.S. study (2005–2006), 30-day all-cause mortality	90 MSSA, 163 MRSA	Not stated	Age per yr; 1.04 (1.01–1.07)	77
Multicenter retrospective European study (2007); 30-day all-cause mortality	257 MSSA, 77 MRSA	24.0	Age per yr; 1.06 (1.03–1.10; <0.01)	6
Multicenter prospective Australian study (2007–2008); 30-day all-cause mortality	1,415 MSSA, 450 MRSA	20.6	Age >70 yr; ND (ND; <0.001)	296
Multicenter prospective Australian study (2007–2008); 30-day all-cause mortality	324 MSSA, 199 MRSA	17.2	Age per yr; 1.06 (1.04–1.08; <0.01)	101

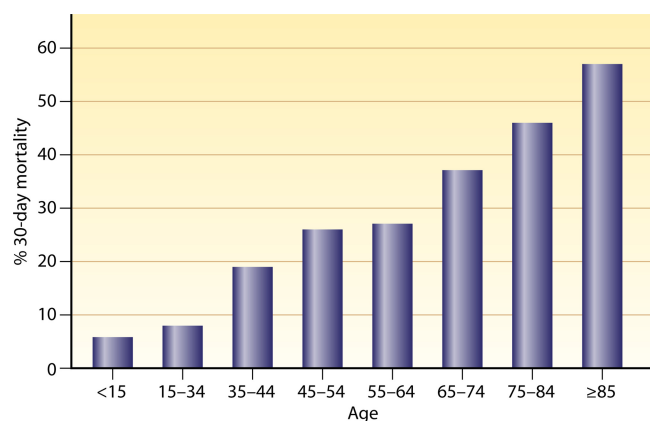
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TABLE 1 (Continued)

Study (outcome of interest)	No. of MSSA and MRSA bacteremic episodes	Overall mortality rate (%)	Age variable; OR (95% CI; <i>P</i> )	Reference
Studies where age was not an independent predictor of mortality				
Single-center prospective Spanish study (1990–1993); infection-related mortality	100 MSSA, 84 MRSA	44.0	Age >65 yr; ND	244
Single-center retrospective Turkish study (1990–1994); infection-related mortality	55 MSSA, 46 MRSA	21.8	ND	291
Single-center retrospective Brazilian study (1991–1992); 14-day all-cause mortality	46 MSSA, 90 MRSA	39.0	Age >60 yr; ND	39
Single-center retrospective French study (1997–1998); infection-related mortality	69 MSSA, 30 MRSA	27.3	Age ≥60 yr; ND	286
Single-center retrospective U.S. study (1997–2000); in-hospital mortality	250 MSSA, 96 MRSA	20.7	Age >65 yr; ND	41
Single-center retrospective Korean study (1998–2001); 60-day infection-related mortality	111 MSSA, 127 MRSA	37.0	Age ≥65 yr; 1.88 (0.97–3.62; ND)	142
Single-center retrospective Korean study (1998–2006); infection-related mortality	294 MSSA, 0 MRSA	19.4	Age ≥65 yr; 1.4 (0.8–2.5; 0.29)	140
Single-center retrospective Brazilian study (2000–2001); infection-related mortality	50 MSSA, 61 MRSA	39.0	Age per yr; 0.98 (0.96–1.00; 0.07)	89
Single-center retrospective Taiwanese study (2000–2008); 30-day all-cause mortality	0 MSSA, 227 MRSA	44.9	Age ≥65 yr; 1.93 (0.98–3.78; ND)	171
Single-center retrospective Belgian study (2002–2004); infection-related in-hospital mortality	88 MSSA, 66 MRSA	40.0	ND	168
Multicenter retrospective U.S. study (2004–2005); in-hospital mortality	32 MSSA, 36 MRSA	19.1	Age per yr; 1.05 (1.0–1.1; 0.08)	184
Single-center retrospective Taiwanese study (2006–2008); 30-day all-cause mortality	0 MSSA, 253 MRSA	30.5	ND	309

<sup>a</sup> Studies employed a multivariate logistic regression model. OR, odds ratio; CI, confidence interval; *P*, probability; ND, not described.

impact of ethnicity on mortality is unclear, primarily because the numbers of deaths are not indicated or are too small to compare (98, 109, 289) and because of multiple confounders associated with indigenous populations (96). These include the prevalence of comorbidities such as diabetes and alcoholism, socioeconomic status (SES), overcrowding, poor infrastructure, limited opportunities, and low income and education (96, 160, 194, 289). There was a suggestion in the literature, however, that mortality may be affected by ethnicity, with higher standardized mortality rates ob-



**FIG 1** Impact of age on overall 30-day mortality from *Staphylococcus aureus* bacteremia. Percentages of patients who succumbed at 30 days following an episode of *Staphylococcus aureus* bacteremia are stratified by 10-year age groups. (Adapted from reference 155 with permission of Elsevier.)

served for African Americans (10/100,000 per annum) than for Caucasians (5.9/100,000) (143). However, unlike the study by Klevens et al. (143), most studies examining the effect of ethnicity showed no difference in outcomes (96) or lower mortality rates in African Americans than in Caucasian patients (26% and 37%, respectively; *P* = 0.04) (243). Similarly, a large multicenter observational prospective study in Australia of 1,865 cases observed a significantly lower 30-day mortality rate for indigenous Australians, 5.7% (3/53), than for people of European descent, 22.2% (350/1,575) (*P* < 0.001) (296). Limitations of these studies include the lack of adjustments for age and comorbidities. Nevertheless, despite these counterintuitive findings, the impact of ethnicity on SAB incidences and mortality remains largely unresolved, with little progress in the understanding of how racial disparities may influence these measures.

### Socioeconomic Status

Socioeconomic status (SES) is known to impact a patient's infection risk (10). For SAB, an inverse relationship exists between incidence and SES, with the lowest rates found for the least deprived economic strata than for the most deprived strata (16 and 21.3/100,000, respectively; *P* < 0.01) (109). In this population-based survey of 779 SAB episodes, SES was measured by using the New Zealand deprivation index, a geographical and census-based measure of income (i.e., place of abode, number of others living in the household, the employment status and qualifications of each person, and access to a telephone and car). Linear regression modeling did not show an effect of the deprivation quintile on mortality rates. Thus, the effect of SES on outcome remains uncertain, as

no other SAB study, to our knowledge, has captured these data. Further study is warranted, however, as defining the role of deprivation in SAB outcomes would allow for the better targeting of preventative strategies and government programs.

### Immune Status

Immunosuppression can be defined as a congenital or acquired quantitative or qualitative deficiency of phagocytic cells, complement, or humoral or cell-mediated immunity. Acquired immune deficiencies predominate especially in adult populations and remain an important area of research with the continuing expansion of stem cell and solid-organ transplantation and the increasing use of immune modulators in general medical therapeutics. Overall, immunosuppression has been observed infrequently as an independent predictor of mortality in SAB studies (98, 131), with the odds of death being approximately 4-fold higher (OR, 4.1; 95% CI, 1.5 to 11.3;  $P = 0.007$ ) than for immunocompetent hosts (131). This may reflect the number of heterogeneous disorders that are lumped together, with very few cohort studies examining the impact of specific immune system components on mortality due to SAB and the lack of functional immune data.

**Innate immunity.** The innate immunity system forms the initial defense against infections and consists of anatomical barriers, complement, and specific cells. The most common acquired defect of the innate immune system is secondary to chemotherapy-induced neutropenia, especially for the treatment of hematological malignancies. Paradoxically, these patients are significantly less likely to die than nonneutropenic cancer patients ( $P = 0.01$ ) (305), which may reflect the higher rates of intravenous line-related SAB and low rates of infective endocarditis (IE) in these patients (270). Congenital disorders are rare, with chronic granulomatous disease (CGD) being the most commonly occurring deficiency. This heterogeneous condition is characterized by the inability of cells to phagocytose catalase-positive microorganisms, resulting in an increased risk of recurrent infections, especially SAB (130). However, no comparative data exist to assess the overall impact of CGD on SAB mortality.

**Humoral immunity.** The protective role of antibody responses to *S. aureus* infections remains to be defined. Similarly, the impact of acquired and congenital B-cell dysfunctions on SAB mortality is unknown.

Individuals with prior *S. aureus* exposure or carriers are known to have lower SAB mortality rates than noncarriers (8% versus 32%;  $P = 0.006$ ) (316). As antibody levels are shaped during *S. aureus* colonization prior to infection (146), it is plausible that antibody levels may provide some protection and improve outcomes for these patients. More direct evidence for protective antibody responses was observed in a single-center prospective Swedish study, which detected a significant ( $P < 0.05$ ) association between 28-day mortality and low levels of antibodies to teichoic acid (relative risk [RR], 4.1; 95% CI, 1.3 to 13.3), lipase (RR, 4.3; 95% CI, 1.3 to 13.1), enterotoxin A (RR, 3.6; 95% CI, 1.1 to 11.9), and scalded skin syndrome toxin (RR, 13.2; 95% CI, 1.7 to 99.7) (118). This study, however, was not limited to bacteremic episodes only and failed to measure other components of the immune system that may impact outcomes.

The failed *S. aureus* vaccine trials (254, 263) provide further evidence that our understanding of the humoral immune system with respect to outcomes of SAB is unclear. Given the impact of

SAB, humoral responses warrant further study, which in turn may enable better vaccination strategies (118, 146, 306).

**Cell-mediated immunity.** AIDS secondary to HIV is one of the most common cell-mediated acquired immune deficiency states worldwide (159). The incidence of SAB among HIV-infected individuals remains higher than that among individuals not infected with HIV (25, 124, 138, 159, 260, 295). This risk is directly associated with the degree of immune deficiency, with the relative risk being 31 times higher (incidence rate ratio [IRR], 31.1; 95% CI, 10.3 to 94.0) for patients with CD4 cell counts of  $<100$  cells/ $\mu$ l than for patients with CD4 cell counts of  $\geq 350$  cells/ $\mu$ l (159). Despite a higher incidence of SAB among HIV-infected individuals, no study to date has detected an increased mortality rate with SAB in HIV-infected cases (227, 260, 261).

Overall, immune deficiency states seem to have a minimal impact on outcomes for patients with SAB. Paradoxically, there is some evidence that immune suppression may actually be protective, as these dampened responses may lead to reduced inflammation, manifesting as less severe disease. However, as comparative data remain limited, the current data should be interpreted with caution.

### Presence of Comorbidities

The presence of one or more comorbidities has been associated, albeit inconsistently, with SAB mortality. These comorbidities include the presence of alcoholism (131), immunosuppression (131), cirrhosis (133, 134, 138, 140), congestive cardiac failure (171), malignancy (33, 77, 133, 134, 140, 308, 309), chronic renal failure requiring hemodialysis (77, 131, 134, 184), and the presence of multiple comorbidities (14).

McCabe and Jackson proposed a system that classifies patients into one of three groups, rapidly fatal, ultimately fatal, and nonfatal, based on the prognosis of the underlying illness (190a). Cohort studies with patients with SAB have found that prognosis as defined by McCabe and Jackson is an independent predictor of mortality (140, 167, 273, 275, 291). Furthermore, an incremental increased risk of death among the three categories has been documented (273, 275), with the odds of death being higher for patients with a rapidly fatal underlying disease (OR, 13.1; 95% CI, 5.7 to 30.9) than for patients with an ultimately fatal underlying disease (OR, 2.02; 95% CI, 1.18 to 3.44) (275). Conversely, patients with nonfatal comorbidities had lower mortality rates (OR, 0.2; 95% CI, 0.1 to 0.4;  $P < 0.01$ ) (237). Cosgrove and colleagues transformed the McCabe-Jackson categories into a continuous variable by assigning a score to each category (fatal disease, 1; nonfatal disease, 3) (41). In this prospective study of 348 SAB episodes, lower scores on multivariate analysis remained an independent predictor of mortality (OR of 38.5 for each incremental decrease;  $P < 0.001$ ).

Functional status as a surrogate marker for the underlying disease prognosis has likewise been demonstrated to be an independent predictor of mortality, with bedridden patients (185, 204) and individuals from a long-term care facility (133, 134) having higher death rates. Alternatively, 17 preexisting conditions known to be associated with an increased mortality can be scored by using the Charlson weighted index (CWI) (31). Higher scores are associated with an increased 1-year mortality rate compared to lower scores (e.g., 82% for scores of  $>5$ , compared to 12% for scores of 0). Following the publication of the CWI, Lesens et al. were able to validate the utility of this index in a prospective study of 168 patients with SAB (164). Multiple other cohort studies since then

TABLE 2 All-cause mortality for patients with *Staphylococcus aureus* bacteremia stratified by source of infection<sup>a</sup>

Diagnosis	In-hospital mortality (%) (reference[s])	28–30-day mortality (%) (reference[s])	Description (reference[s])
Bacteremia without focus	11–45 (14, 161, 167)	21.9–47.5 (120, 206, 296, 301)	14-day mortality rate, 49% (39); independent predictor of mortality relative to other infections in several studies, with an OR range between 3.88 and 12.3 (15, 39, 206, 301)
Bone and joint infection	0–13.9 (14, 19, 76, 80, 167)	0–29 (120, 296, 301)	14-day mortality rate, 11.7% (296); mortality dependent on site, with higher rates for vertebral osteomyelitis than for nonvertebral infections; independent predictor of reduced mortality relative to other infections in a single study, with an OR of 0.27 (95% CI, 0.08–0.93; <i>P</i> = 0.038) (133)
Prosthetic joint infection		8.3 (296)	
CNS infection	25–56 (14, 126, 223, 228)	10.7 (296)	Independent predictor of mortality relative to other infections in a single study, with an OR of 12.9 (95% CI, 1.1–152.9; <i>P</i> value not stated) (244)
Deep abscess		14.6 (296)	
Intra-abdominal infection		25 (120)	
Intravenous catheter (type not specified or several types)	0–21 (14, 54, 113, 167, 169, 205, 235, 313)	4–32.6 (120, 296, 301)	14-day mortality rate, 12% (39); device with a metastatic infection associated with higher mortality rates (296); independent predictor of reduced mortality relative to other infections in a single study, with an OR of 0.47 (95% CI, 0.31–0.71; <i>P</i> < 0.01) (33)
Central venous catheter	30 (291)	20.5 (296)	
Hickman catheter	22 (53)		
Peripheral venous catheter		16.9 (296)	
Hemodialysis catheter		6.8 (296)	
Infective endocarditis	22.4–66 (8, 15, 66, 106, 107, 167, 237, 242, 291)	25–60 (29, 107, 206, 301)	IVDU-associated IE mortality rate, 11%; non-IVDU IE mortality rate of 21%, vs 29.4% for health care-associated IE (66); infection-related mortality rate, 22% (68); independent predictor of mortality relative to other infections in several studies, with an OR range between 2.8 and 12.1 (77, 168, 237, 296, 301)
Left-sided IE	28.6 (197)	23.9 (296)	Aortic valve IE was an independent predictor of mortality relative to mitral or right-sided IE, with an OR of 1.91 (95% CI, 1.0–3.66; <i>P</i> = 0.05) (197)
Right-sided IE	5.9 (197)	11.8 (296)	
Prosthetic valve IE	30.5–50 (66, 242)		
Cardiac device infection		36.4 (28)	
Pulmonary infection	41.6–62 (14, 79, 167, 200)	39–67 (120, 148, 296, 301)	14-day mortality rate, 65% (39); attributable mortality rate, 46.5% (85); independent predictor of mortality relative to other infections in several studies, with an OR range of between 2.09 and 17.0 (33, 39, 41, 134, 206, 296, 301)
Skin and soft tissue infection	10–18.6 (14, 167)	14.8–17 (120, 296, 301)	
Surgical site infection	0–23 (14, 291)		
Urinary tract infection	9.7 (14)	10 (120)	

<sup>a</sup> For the purpose of this review, no attempt has been made to differentiate between the various study diagnosis definitions, with source of infection, primary focus, or principal diagnosis being used interchangeably. OR, odds ratio; CI, confidence interval; IVDU, intravenous drug user.

have demonstrated that the CWI is an independent predictor of mortality due to SAB (6, 15, 33, 185, 301).

Although several SAB cohort studies have not detected any difference in outcomes for patients with one or more comorbidities (32, 39, 59, 89, 309, 310), this discrepancy probably reflects the small sample size and patient groups selected in these studies. Regardless, overall, the presence of one or more comorbidities generally portends a worse outcome for patients with SAB, and as such, the collection of these data should be encouraged in future SAB studies.

## HOST-PATHOGEN INTERACTIONS

### Principal Diagnosis and Source of Infection

The overall mortality rate from SAB varies depending on the primary focus of infection, with the highest mortality rates occurring in patients with primary bacteremic pulmonary infections (33, 39, 41, 96, 206, 296, 301) and infective endocarditis (IE) (77, 168, 237, 296, 301).

Some of the lowest rates occur in patients with central or peripheral venous catheter-related infections (33) (Table 2).

Within a diagnostic category or SAB source, the site and extent of infection influence outcomes, with higher mortality rates for patients with complicated left-sided IE (e.g., the presence of a paravalvular abscess) than for patients with uncomplicated left-sided IE (197). For SAB secondary to medical device infections, the type of device correlates with outcomes, with higher mortality rates associated with a central line (20.5%) than with a peripheral line (16.9%), a hemodialysis catheter (6.8%), or orthopedic implants (8.3%) (296). Patient characteristics further influence these outcomes, with intravenous drug use (IVDU)-associated IE being associated with improved survival compared to non-IVDU-associated IE (*P* < 0.001) (66). A review of these nuances with respect to presentations and mortality is beyond the scope of this review, as is a review of mortality data for *S. aureus* compared to other microorganisms.

The link between source and outcome was noted and employed by Soriano and colleagues, who stratified SAB episodes into one of three mortality risk groups by the final source of SAB. An incremental increase in the mortality rate was noted, from 5% to 13% and 30% for “low”-risk (intravenous catheter; urinary tract; ear, nose, and throat; and gynecological foci), “intermediate”-risk (bone and joint, soft tissue, and unknown foci), and “high”-risk (endovascular, lower respiratory tract, intra-abdominal, and central nervous system [CNS] foci) groups, respectively (275). In a subsequent prospective study of 414 MRSA-B episodes, the utility of these categories was confirmed, with 2- and 9-fold-increased odds of death for intermediate-risk (OR, 2.29; 95% CI, 1.21 to 4.31) and high-risk (OR, 9.49; 95% CI, 5.1 to 17.6) SAB diagnoses compared to low-risk SAB sources (273).

At presentation, the extent of *S. aureus* infection may not be obvious, with approximately one-third of patients developing a metastatic or complicated SAB infection (67, 161, 238). The clinical implication of the above-mentioned association is that for optimal patient outcomes, defining the extent of SAB is critical (40). Accordingly, several groups have proposed criteria based on clinical features and investigation results to recommend a duration of therapy (69, 125, 207). Of these, the criteria developed by Fowler and colleagues are the most well known and classify SAB episodes into one of three groups: complicated, uncomplicated, and simple SAB (69). Similar to the “risk” groups developed by Soriano et al., complicated SAB episodes have worse outcomes than do uncomplicated and simple episodes (24% versus 40% mortality;  $P < 0.01$ ) (161).

Overall, the source and extent of SAB infection are important and consistent contributors to overall SAB mortality. This association confirms the need to consider SAB as not one single entity but a heterogeneous group of infections in future studies. This may enable the refinement of source-specific management factors to be elucidated.

### Setting of Bacteremia

The setting of SAB onset has traditionally been divided into two categories, health care associated (formerly nosocomial) and community acquired, when subsequent positive *S. aureus* blood culture bottles are obtained  $\geq 48$  h and within 48 h of hospital admission, respectively (102). With changes in the complexity of modern health care, community-onset infections are now further divided into episodes with health care contact (e.g., health care-associated outpatient) and those without (253). The setting of SAB assisted clinicians in predicting the infecting *S. aureus* clonal type and, consequently, antibiotic choice. However, with the advent of community-acquired MRSA (CA-MRSA) (defined by the antibiotic resistance pattern and/or staphylococcal cassette chromosome *mec* [SCC*mec*] type) strains entering the hospital, causing cross infections and replacing common hospital clones, these definitions are becoming less helpful (229).

Most current cohort studies have not found the setting of SAB to influence outcomes (218, 224, 275, 296, 301), with the exception of two studies, which observed increased 30-day mortality rates for hospital-onset SAB episodes (with one study reporting an OR of 1.75, a 95% CI of 1.05 to 2.92, and a  $P$  value of 0.033 [101] and the other reporting an OR of 1.42 and a 95% CI of 1.12 to 1.81 [134]). No typing data were presented in these studies, and therefore, these results should be interpreted with caution, as the setting of bacteremic onset may be a surrogate for a specific *S. aureus*

clone. Regardless, it appears that the setting of SAB has a minimal effect on patient outcomes.

### Persistent Bacteremia

Definitions of persistent bacteremia vary between studies and range from 3 days to up to 7 days following appropriate active antibiotic therapy (172, 220, 302). Regardless, persistent bacteremia is a common manifestation of SAB and occurs for between 6% and 38% of infection episodes (65, 93, 139, 210), with a median time to clearance of 7 to 9 days (65, 139, 210). Risk factors include the source of infection (i.e., infective endocarditis or vertebral osteomyelitis) (66, 163), pathogen phenotypes (vancomycin heteroresistance) (302), antibiotic treatment (139, 219), the presence or retention of prosthetic material (139), and the ability to remove foci of infection (e.g., by surgical drainage) (87, 320). MRSA has been associated with a greater likelihood of persistence (median time to clearance of 8 to 9 days) than MSSA (median time to clearance of 3 days) bacteremia (65, 163). This probably reflects suboptimal activity with vancomycin compared to  $\beta$ -lactams in preventing persistence (267). Alternatively, pathogen-specific factors may be responsible for this persistence (259).

Irrespective of the cause, bacteremic persistence probably is a surrogate for complicated SAB (69), as the likelihood of a metastatic infection increases with an increasing duration of bacteremia, to approximately 45% following  $\geq 10$  days of SAB (139, 210). These complicated infections in turn lead to poorer outcomes (170, 171). Even in the absence of metastatic complications, persistence *per se* portends a worse outcome, with infection-related mortality rates being higher for patients with MRSA-B persistence ( $>3$  days) than for nonpersistent episodes (45.2% and 9.4%, respectively;  $P = 0.002$ ) (323). Similar results were obtained by a larger case-controlled study, where mortality rates for persistent ( $>7$  days) SAB were significantly higher than mortality rates for nonpersistent controls (54.8% and 31.4%, respectively;  $P < 0.01$ ) (93). Only one cohort study of 177 SAB episodes established persistent bacteremia as being an independent predictor of mortality (OR, 17.5; 95% CI, 1.5 to 212;  $P = 0.024$ ) (167).

Pathogen factors such as heteroresistance to vancomycin are associated with persistent bacteremia (OR, 2.37; 95% CI, 1.53 to 3.67;  $P < 0.01$ ) (301). However, heteroresistant vancomycin-intermediate *S. aureus* (hVISA) bacteremic episodes do not lead to poorer outcomes (302) and may be associated with improved overall survival (301) compared to vancomycin-susceptible *S. aureus* (VSSA) bacteremic episodes. These complexities require further study, as it seems that not all persistent episodes translate into poorer outcomes. Nevertheless, overall persistent bacteremia generally results in worse outcomes secondary to the causes and complications that persistence usually indicates.

### Bacteruria

*S. aureus* is an uncommon uropathogen in the absence of bladder catheterization, instrumentation, or surgery and thus typically represents hematogenous spread. The presence of concomitant *S. aureus* bacteruria and bacteremia thus probably represents a higher disease burden or complicated SAB and thus portends a worse outcome. This is supported by a retrospective cohort study of 118 SAB episodes, which detected an increased risk of death for patients with concomitant bacteruria (32%, versus 14% for no *S. aureus* bacteruria;  $P = 0.036$ ) (225). In a subsequent case-controlled study of 308 SAB (42% MRSA) episodes, bacteruria ( $n =$

TABLE 3 Summary of studies that have examined the impact of severity of illness on mortality in *Staphylococcus aureus* bacteremia<sup>a</sup>

Study and outcome of interest	No. of episodes	% of patients with sepsis/shock	Mortality rate for patients with shock (%)	Variable, OR (95% CI; P)	Reference
Studies that found severity of illness to be an independent predictor of mortality (multivariate analysis)					
Single-center retrospective Turkish study (1990–1994); infection-related mortality	101 SAB	23.8	41.7	Septic shock, 5.4 (ND; 0.02)	291
Single-center retrospective Spanish study (1991–1998); in-hospital infection-related mortality	908 SAB	9.7	55.7	Shock, 12.6 (7.2–22.2; ND)	275
Single-center prospective Spanish study (1991–2005); 30-day all-cause mortality	414 MRSA-B	20.3	63.1	Shock, 7.38 (4.11–13.3; <0.001)	273
Single-center retrospective Brazilian study (1991–1992); 14-day all-cause mortality	134 SAB	22.1	73.3	Shock, 8.92 (2.9–27.8; ND)	39
Single-center retrospective Belgian study (1992–1998); 30-day in-hospital mortality	85 SAB	62.2	ND	Hemodynamic instability, 1.76 (1.14–2.71; 0.01); APACHE II per point, 1.04 (1.02–1.06; <0.01)	18
Single-center prospective Danish study (1994–1996); 150-day mortality	278 SAB	25	47	Septic shock, 3.7 (1.5–9.1; 0.004)	127
Single-center retrospective U.S. study (1995–1999); 30-day all-cause mortality	293 SAB	31.4	41.3	Modified APS >60, 15.7 (5.8–49.8; <0.001)	206
Single-center retrospective U.S. study (1996–2006); all-cause mortality	489 MRSA-B	Not applicable	Not applicable	SAPS, 1.51 (1.26–1.80; <0.001)	202
Single-center retrospective Taiwan study (1997–2001); 30-day infection-related mortality	162 MRSA-B	13.6	86.4	Septic shock, 9.31 (2.35–36.8; <0.01)	59
Single-center retrospective Australian study (1997–2008); 30-day all-cause mortality	401 MRSA-B	10.5	52.4	APACHE II per point, 1.11 (1.07–1.15; <0.001)	301
Single-center retrospective Swiss study (1998–2002); in-hospital all-cause mortality	308 SAB	10.7	ND	Septic shock, 14.5 (6.2–61.6; <0.01)	131
Single-center retrospective Brazilian study (2000–2001); infection-related mortality	101 SAB	43.5	68.2	Severe sepsis/shock, 6.68 (3.05–15.4; <0.01)	89
Single-center retrospective Chinese study (2001–2007); 90-day all-cause mortality	115 MRSA-B	11.3	ND	Septic shock, 7.92 (3.64–17.20; <0.001)	32
Single-center retrospective Taiwan study (2001–2006); in-hospital infection-related mortality	177 MRSA-B	Not applicable	Not applicable	Pitt bacteremia score per point, 1.33 (1.04–1.69; 0.024)	167
Single-center retrospective Taiwan study (2001–2007); 90-day all-cause mortality	744 SAB	11.3	ND	Septic shock, 7.92 (3.64–17.20; <0.001)	33
Single-center retrospective Belgian study (2002–2004); infection-related in-hospital mortality	154 SAB	21.4	ND	Septic shock, 10.26 (3.65–28.8; <0.001)	168
Single-center retrospective U.S. study (2003–2008); in-hospital mortality	814 SAB	Not applicable	Not applicable	SAPS, 1.04 (1.03–1.05; not stated)	257
Single-center prospective U.S. study (2005–2006), 30-day all-cause mortality	253 SAB	Not applicable	Not applicable	Illness severity index, <sup>b</sup> 2.78 (1.94–3.99; <0.001)	77
Single-center retrospective Taiwanese study (2006–2008); 30-day all-cause mortality	253 MRSA-B	41.9	ND	Septic shock, 8.11 (4.06–16.19; <0.001)	309
Single-center retrospective Taiwanese study (2004–2006); 30-day all-cause mortality	215 SAB	39.9	ND	Shock, 8.11 (4.06–16.09; <0.001)	310
Multicenter retrospective European study (2007); 30-day all-cause mortality	334 SAB	37.7	38.9	Severe sepsis/shock, 2.68 (1.52–4.75; <0.01)	6
Multicenter prospective Australian study (2007–2008); 30-day all-cause mortality	1,865 SAB	10.8	40.3	Sepsis syndrome, 4.01 (2.40–6.87; <0.001)	296
Studies where severity of illness was not an independent predictor of mortality (multivariate analysis)					
Multicenter prospective Australian study (2007–2008); 30-day all-cause mortality	532 SAB	12.6	ND	Sepsis syndrome, not stated	101

<sup>a</sup> Studies employed a multivariate logistic regression model. OR, odds ratio; CI, confidence interval; P, probability; ND, not described; APACHE, acute physiological and chronic health evaluation; APS, acute physiological score (based on APACHE III); SAPS, simplified acute physiological score.

<sup>b</sup> The illness severity index is a composite scoring system based on the Charlson comorbidity index and a modified acute physiological score.

68 episodes) was an independent predictor of SAB in-hospital mortality (OR, 2.87; 95% CI, 1.4 to 5.9;  $P = 0.004$ ) (36). Similarly, Huggan et al. found that bacteruria was associated with a 2-fold-increased risk of death in SAB episodes (108, 109). Thus, bacteruria is a surrogate for severe disease and, hence, a signal for poorer outcomes.

### Shock/Sepsis and Severity of Illness

The presence of sepsis or shock, although definitions vary slightly between studies, is strongly associated with worse outcomes for

patients with SAB (Table 3), with mortality rates ranging between 38% and 86% (6, 59). The reason why the study by Holmes et al. did not observe this association is unclear (101), as the larger data set from which the data for this study were drawn found sepsis syndrome to be a strong independent predictor of 30-day all-cause mortality (OR, 4.01; 95% CI, 2.34 to 6.87;  $P < 0.001$ ) (296). Possible explanations include differences in sepsis definitions and patient populations studied.

The use of the acute physiological and chronic health evaluation (APACHE) score (144) for SAB has been met with some criticism



(91), as this score was developed and validated for use in intensive care units (ICUs). Nevertheless, the APACHE score (APACHE II or APACHE III), based on the worst parameters within the first 48 h of the positive blood culture, has consistently been established as an independent predictor of mortality, with an incremental increase in mortality rates as scores increase (204, 301, 324). Multiple other scoring systems have been developed, all of which have been shown to be predictors of mortality (167, 202). These scoring systems include the simplified acute physiological score (SAPS), the sequential organ failure assessment score (SOFA), the Pitt bacteremia score, and the multiple organ dysfunction score (MODS).

Single acute organ dysfunction, especially acute renal failure (15, 18, 131); respiratory failure requiring mechanical ventilation (308); altered mental status (82); and hematological dysfunction (33, 134, 310, 312) portend a worse outcome, albeit inconsistently, with SAB. Alternatively, surrogate markers for infection severity such as transit to an ICU (301) or the acquisition of SAB in the ICU (6, 237, 261, 275) have been found to be independent predictors of mortality compared to ward patients. Thus, the severity of SAB, irrespective of the method used to measure this, is one of the more consistent predictors of overall 30-day mortality.

## **PATHOGEN FACTORS**

### **Methicillin Resistance**

Resistance to most  $\beta$ -lactam antibiotics, including the semisynthetic ( $\beta$ -lactamase-resistant) penicillins, such as methicillin and flucloxacillin, is due to the expression of the low-affinity penicillin binding protein PBP2a (92). PBP2a is encoded by the *mecA* gene and is found on an integrated mobile genetic element called the staphylococcal cassette chromosome *mec* (SCC*mec*) element (116, 136, 190, 272).

In many hospital settings over the past 20 years, the incidence of MRSA-B has increased around the world. This increase has not been associated with a corresponding decline in MSSA infections but has added to the overall burden of SAB (21, 322). Numerous studies spanning the last decade have examined the impact of methicillin resistance on mortality in SAB, with conflicting results. Only studies employing multivariate analyses and which were not included in the two meta-analyses mentioned below are summarized in Table 4. Two meta-analyses were performed, by Whitby et al. (317) and Cosgrove et al. (42), in 2001 and 2003, respectively. Although the time periods for study capture were very similar, the numbers of studies included differed greatly due to the fact that Whitby et al. limited their analyses to nosocomial SAB episodes only. Cosgrove et al. examined data from 31 independent studies (1980 to 2000) and showed that mortality rates were significantly higher for MRSA than for MSSA bacteremic episodes (OR, 1.93; 95% CI, 1.54 to 2.42;  $P < 0.001$ ). Similarly, Whitby et al. detected an increased mortality rate with methicillin resistance (OR, 2.12; 95% CI, 1.76 to 2.57;  $P < 0.001$ ) from the nine studies included (1978 to 2000). It was suggested that the major weakness of the studies examined in the meta-analyses by Whitby et al. and Cosgrove et al. is that they ignored the length of hospital stay prior to the onset of bacteremia (110, 111), which leads to a time-dependent bias, the significance of which was demonstrated by a study by Wolkewitz et al. (321), as the MRSA mortality rate was similar to that for MSSA episodes following an adjustment for the length of hospital stay. However, these data may themselves be imperfect secondary to the small numbers of MSSA-B ( $n = 26$ ) and MRSA-B ( $n = 34$ ) cases detected.

Other sources of criticism include the inclusion of studies that failed to adjust for other confounding factors such as comorbidities, age, and severity of illness (224, 276). For example, in a large ( $n = 815$ ) single-center prospective United Kingdom study (1995 to 2000) by Melzer et al. (195), MRSA-B was not associated with increased mortality after adjustments for the above-mentioned host confounders (OR, 1.72; 95% CI, 0.92 to 3.2;  $P = 0.09$ ). Conversely, adjusting for these variables may nullify the true impact of resistance (276). Nevertheless, when Cosgrove et al. analyzed only studies that included adjustments for potential confounders (such as age, gender, and severity of illness), the mortality rate remained higher for MRSA-B episodes (OR, 1.88; 95% CI, 1.33 to 2.69;  $P < 0.001$ ) (42). Similarly, in a cohort study of 753 community-onset SAB episodes (33% MRSA), methicillin resistance remained an independent predictor of mortality when data were stratified by comorbidities or adjusted for confounders in the multivariate logistic regression model (33).

Alternative study designs have attempted to address the significance of methicillin resistance. Harbarth et al. performed a case-controlled study with matching based on previously identified confounders (91), while in a Korean case-controlled study, matching occurred by the use of a propensity score (218). Both studies were unable to detect a difference in mortality for MRSA-B compared to MSSA-B.

Despite these potential weaknesses, the majority of studies summarized in Table 4 support an increased mortality rate associated with MRSA-B (in comparison to MSSA-B). Several explanations for this apparent association have been provided. These include pathogen-specific factors such as SCC*mec*-associated virulence factors (3), with SCC*mec* type II being an independent predictor of mortality in a cohort study of 253 SAB (64% MRSA) (77) and 744 SAB (33% MRSA) (33) episodes. Alternatively, differences in empirical prescribing (243) and poor vancomycin efficacy (3, 86, 114) may explain some of these differences. Compared to semisynthetic penicillins, vancomycin has slower bactericidal activity *in vitro*, especially with high-inoculum infections (158, 252), and variable tissue penetration (2, 88, 157). Finally, MRSA infection may just be a surrogate for host factors such as comorbidities rather than methicillin resistance *per se* (91). Ultimately, more research is required to clearly define the reasons, which are likely to be multifactorial, for the observed mortality differences.

### **Vancomycin MIC**

The current Infectious Diseases Society of America MRSA-B guidelines continue to recommend vancomycin as the treatment of choice for susceptible MRSA-B isolates (172), with vancomycin-susceptible breakpoints set by the Clinical and Laboratory Standards Institute (CLSI). These breakpoints were lowered in 2006 from an MIC of 4  $\mu\text{g/ml}$  to an MIC of 2  $\mu\text{g/ml}$  following reports of increased mortality associated with vancomycin-intermediate *S. aureus* (VISA) infections (72, 288).

However, questions remain regarding whether or not these breakpoints should be lowered further, thus limiting the role of vancomycin in the treatment of MRSA-B (47, 198). This controversy is in part maintained by conflicting reports of treatment failure and increased mortality with high- but susceptible-MIC isolates (i.e., an MIC of  $\leq 2$   $\mu\text{g/ml}$ ). For a more complete examination, including other variables that influence decisions about the use of vancomycin in MRSA-B management, readers are directed to the above-mentioned two publications.

TABLE 4 Summary of studies that have examined the impact of methicillin resistance on mortality in *Staphylococcus aureus* bacteremia<sup>a</sup>

Study	No. of MRSA isolates/no. of MSSA isolates	Mortality rate for MRSA vs MSSA (%)	Mortality risk for MRSA vs MSSA	Adjustment(s) for confounders	Reference
Meta-analyses					
Meta-analysis of 9 studies (1990–2000); mortality	778/1,431	29 vs 12	Pooled RR, 2.03 (95% CI, 1.55–2.65; $P < 0.001$ )	None	317
Meta-analysis of 31 studies (1980–2000); mortality	1,360/2,603	Not stated	Pooled OR, 1.93 (95% CI, 1.54–2.42; $P < 0.03$ )	Varied between studies analyzed	42
Studies that found MRSA to be an independent predictor of mortality (multivariate analysis)					
Single-center retrospective Taiwanese study (1990–2004); 30-day mortality	851/297	49.8 vs 27.6	OR, 1.78 (95% CI, 1.3–2.44; $P < 0.001$ )	None	308
Single-center retrospective Belgian study (1992–1998); 30-day mortality	47/38	63 vs 18	HR, 1.93 (95% CI, 1.18–3.18; $P < 0.01$ )	None	18
Multicenter retrospective U.S. study (1995–2003); 90-day infection-related mortality	184/235	34.2 vs 19.6	HR, 1.8 (95% CI, 1.2–3.0; $P < 0.01$ )	Patients with pneumonia	266
Single-center retrospective (2002–2004) and prospective (2005–2007) German study; 90-day all-cause mortality	67/454	42 vs 19	OR, 2.6 (95% CI, 1.4–4.9; $P < 0.01$ )	None	237
Single-center retrospective Belgian study (2002–2004); infection-related in-hospital mortality	66/88	52 vs 32	OR, 3.04 (95% CI, 1.15–8.04; $P = 0.021$ )	None	168
Multicenter prospective Asian study (2004–2006); 30-day mortality	329/380	33.1 vs 17.1	OR, 1.69 (95% CI, 1.15–2.49; $P < 0.01$ )	None	133
Single-center retrospective United Kingdom study (2005–2006); 90-day mortality	34/26	35.3 vs 19.2	Time-averaged HR of 1.69 (95% CI, 0.72–4)	Age, gender, comorbidities, hospitalization (first admission, time averaged)	321
Studies that found MRSA not to be an independent predictor of mortality (upon multivariate analysis)					
Single-center retrospective U.S. study (1991–2000); infection-related mortality	170/183	30.6 vs 15.3	OR, 1.4 (95% CI, 0.7–3; $P = 0.4$ )	Age, site of infection, APACHE score, ICU at onset, hospital onset, and delayed therapy	175
Single-center retrospective Canadian study (1991–2005); 30-day mortality	69/746	33 vs 23	OR, 2.21 (95% CI, 0.99–4.96; $P$ value ND)	Age, gender, comorbidities, residence, absence of treatment, site of infection	5
Single-center prospective United Kingdom study (1995–2000); infection-related mortality	382/433	29.6 vs 13.6	OR, 1.72 (95% CI, 0.92–3.2; $P = 0.09$ )	Age, hospital specialty, primary site of infection	195
Single-center retrospective French study (1997–1998); infection-related mortality	30/69	43.3 vs 20.3	OR, 2.8 (95% CI, 0.99–7.1; $P$ value ND)	Appropriate treatment within the first 72 h	286
Single-center prospective U.S. study (1997–2000); in-hospital mortality	96/252	22.9 vs 19.8	OR, 0.72 (95% CI, 0.39–1.96; $P = 0.45$ )	None	41
Multicenter retrospective United Kingdom study (1997–2004); 30-day mortality	227/214	34.0 vs 27.0	OR, 1.49 (95% CI, 0.99–2.26; $P$ value ND)	None	322
Single-center retrospective Brazilian study (2000–2001); infection-related mortality	61/50	54.9 vs 24.7	HR, 3.52 (95% CI, 0.96–6.60; $P = 0.06$ )	Adequacy of therapy, gender, age, severity of clinical status and underlying illness	89
Single-center prospective Taiwanese study (2001–2006); 30-day all-cause mortality	30/185	10.0 vs 13.2	OR, 0.74 (95% CI, 0.22–2.45; $P$ value ND)	None	310
Multicenter prospective Asian study (2004–2006); 30-day mortality	2007/1701	16.2 vs 10.8	Not stated	Age, comorbidities, source of infection, health care onset	134
Single-center retrospective U.S. study (2004–2008); patients $\geq 80$ yr of age; in-hospital mortality	46/30	34.8 vs 20.0	OR, 2.1 (95% CI, 0.7–6.3; $P = 0.16$ )	None	15
Multicenter prospective Australian study (2007–2008); 30-day all-cause mortality	450/1,415	30.0 vs 17.7	OR, 1.04 (95% CI, 0.58–1.86; $P = 0.89$ )	None	296

<sup>a</sup> Only studies post-2000 and not included in the meta-analyses by Cosgrove et al. (42) and Whitby et al. (317) are represented. MRSA, methicillin-resistant *S. aureus*; MSSA, methicillin-sensitive *S. aureus*; OR, odds ratio; CI, confidence interval;  $P$ , probability; ND, not described.

The first study to demonstrate episodes of vancomycin treatment failure with a high MIC ( $\geq 1.5 \mu\text{g/ml}$  by the Etest) was reported in 2006 by Hidayat et al. (97). Subsequently, two other groups observed similar results (173, 181), despite the differing

treatment failure definitions and MIC testing methodologies used (Etest and Vitek, respectively).

Although a mortality difference with high-MIC (by Etest) compared to low-MIC episodes was observed in a study by Musta et al.,

the significance of these results was questioned by the authors themselves, as this association was not upheld on multivariate analysis (202). Soriano et al., in their single-center study of MRSA-B ( $n = 414$ ), did, however, observe an increased mortality rate with high-MIC ( $2 \mu\text{g/ml}$  by Etest) episodes (OR, 6.39; 95% CI, 1.68 to 24.3;  $P < 0.01$ ) albeit for empirically vancomycin-treated patients only (273). These data were subsequently criticized, as the MIC was an independent predictor of mortality only when shock, which occurred less frequently in high-MIC episodes, was adjusted for (177, 230). However, those authors maintained that shock was a confounding variable rather than an intermediate variable and thus should be corrected for (274). Subsequently, three further studies found a high MIC ( $\geq 1.5 \mu\text{g/ml}$ ) to be an independent predictor of mortality (with one study reporting an OR of 2.39, a 95% CI of 1.2 to 4.8, and a  $P$  value of 0.014 [312], another study reporting an OR of 6.05, a 95% CI of 2.3 to 15.93, and a  $P$  value of 0.001 [285], and a further study reporting an OR of 2.44, a 95% CI of 1.21 to 4.92, and a  $P$  value of 0.012 [101]) compared to low-MIC episodes. Of note, a study by Wang et al. was the first to use broth microdilution to determine MICs (312), while a study by Holmes et al. found that the vancomycin MIC also predicted mortality for flucloxacillin-treated MSSA episodes (101).

However, these findings are by no means universal, as no association of mortality with increasing vancomycin MICs was noted in one of the largest retrospective single-center studies of SAB ( $n = 814$ ) (258). Comparable results were observed by two other retrospective single-center studies (154, 210). The impact of a high MIC is further questioned by the findings of a small study of 49 SAB episodes (MRSA and MSSA), which showed improved survival with high-MIC ( $\geq 1.5 \mu\text{g/ml}$  by the Etest) episodes (232). These studies have likewise come under scrutiny, with contrary findings explained by the inclusion of nontreated patients (258) or vancomycin-treated MSSA episodes in the analysis (232, 258).

Overall, the majority of studies point to a possible increased mortality rate for MRSA-B episodes with high but susceptible vancomycin MICs. As MICs were almost exclusively determined by the Etest, a lowering of the breakpoints is not currently warranted, as the correlation between results of the Etest and the gold standard, broth microdilution (used to set breakpoints), is moderate at best (300). The clinical significance of this association remains unclear, as it may not reflect vancomycin failure *per se* but pathogen-related characteristics (100), especially in light of recent findings of observed increased mortality rates associated with high-vancomycin-MIC, flucloxacillin-treated MSSA episodes (101). Pending further studies, treatment recommendations continue to support vancomycin use irrespective of the MIC, with changes to therapy guided by the clinical responses (172).

### Vancomycin Resistance

**Heteroresistant vancomycin-intermediate *S. aureus*.** Since the first description of hVISA in 1997 by Hiramatsu and colleagues (99), questions remain regarding the best method to detect this phenotype. The current gold standard is the modified population analysis profile-area under the curve (PAP-AUC) method, in which the presence of heteroresistance is confirmed when the ratio of the number of viable colonies plotted against increasing vancomycin concentrations of the test strain to that of the reference strain (Mu3) is  $\geq 0.9$ . For greater details pertaining to various laboratory detection methods and other factors, including hVISA

epidemiology, pathogenesis, and mechanisms of resistance, readers are referred to two recent reviews (104, 302).

Several studies have subsequently examined the impact of hVISA bacteremia compared to vancomycin-susceptible *S. aureus* (VSSA) episodes. The first study reported was from a single Australian center in 2004, which observed no increased mortality with hVISA (by PAP-AUC) episodes (30). Nevertheless, the rate of mortality of hVISA bacteremia cases was found to be high, at approximately 33%, in a retrospective review of 25 confirmed (by PAP-AUC analysis) hVISA episodes (105). This high mortality rate was confirmed by Maor et al., who detected an in-hospital mortality rate of 75% for 16 confirmed (by the macromethod Etest [MET]) hVISA bacteremic infections from 264 screened MRSA-B episodes (186).

That same group, however, did not detect a 30-day overall mortality difference in their later retrospective comparative study of 250 MRSA-B episodes (51% for hVISA versus 46% for VSSA;  $P = 0.48$ ) (185). Similar results were seen in a small single-center retrospective review of 56 persistently bacteremic (for more than 7 days) MRSA episodes (63).

In 2009, Musta et al. reported the largest study of hVISA bacteremic episodes (202). In this single-center retrospective study of 489 MRSA-B episodes, 71 (17%) hVISA (confirmed by MET) episodes were detected. No outcome differences were observed between the two groups. Similarly, no increased mortality rate was detected with hVISA (confirmed by PAP-AUC analysis) episodes in a retrospective review of 65 MRSA infective endocarditis patients by the International Collaboration of Endocarditis (ICE) (8).

In contrast, an Australian retrospective study detected a reduced mortality rate with hVISA (confirmed by PAP-AUC analysis) for 401 consecutive MRSA-B episodes over a 12-year period (301). The hVISA phenotype was an independent predictor of reduced mortality (OR, 0.27; 95% CI, 0.09 to 0.83;  $P = 0.02$ ). Possible explanations for these conflicting findings include the detection of hVISA exclusively in ST239-like MRSA clones and the higher number of hemodialysis patients in the hVISA group.

Overall, hVISA bacteremia is not associated with an increased mortality rate (OR, 1.18; 95% CI, 0.80 to 1.72;  $P = 0.4$ ) (302) despite hVISA being associated with persistent bacteremia (8, 30, 209, 302), high-bacterial-load infections (e.g., infective endocarditis) (30), and high vancomycin MICs (104, 303), as the likelihood of detection of hVISA increases with rising MICs, with the greatest chance occurring with isolates exhibiting an MIC of  $2 \mu\text{g/ml}$ . The failure to detect a mortality difference may reflect the reduced virulence that was demonstrated *in vitro* for these isolates (191). Clinical data are lacking but are suggested by the lower rates of shock with infections with high-vancomycin-MIC isolates (154, 273), the lower risk of acquiring hVISA infections than VSSA infections (103), and the lower mortality rates detected in one study (301).

**Vancomycin-intermediate *S. aureus*.** An MRSA isolate is currently defined as being a VISA isolate if the vancomycin MIC is between 4 and  $8 \mu\text{g/ml}$ . These changes occurred in 2006, when the vancomycin breakpoints were lowered from 8 to 16 to 4 to  $8 \mu\text{g/ml}$  for "intermediate" resistance (288). Part of the rationale for doing so was secondary to the findings of a case-controlled study by Fridkin et al. in 2003. In that study, *S. aureus* isolates with reduced vancomycin susceptibility (RVS) or VISA isolates by the current breakpoints were significantly associated with increased mortality rates compared to vancomycin-susceptible infections

(63% versus 12%; OR, 12.7; 95% CI, 3.4 to 48) (72, 73). Despite changes to the breakpoints, VISA remains uncommon (72), and thus, no other large-scale comparative data with respect to VISA outcomes have been reported. Regardless, mortality is generally considered to be increased secondary to vancomycin failure, and as such, alternative antibiotics are recommended for the treatment of VISA infections (172).

**Vancomycin-resistant *S. aureus*.** Vancomycin-resistant *S. aureus* (VRSA) infections have been limited to a few cases (4, 62, 249, 268). Most have occurred in the United States and have not been associated with bloodstream infections. Thus, the impact of VRSA on mortality in SAB is unknown.

### Exotoxins

Secreted toxins (exotoxins) produced by *S. aureus* contribute to pathogenicity and the ability to colonize the host. These toxins can broadly be categorized as either superantigens (SAGs) or cytotoxic exotoxins. Although there are few studies that have directly examined the impact of exotoxin production on mortality in bacteremic patients, clues to the potential role that exotoxins play in mortality can be garnered from virulence studies and animal models.

**Superantigens.** Staphylococcal SAGs include the staphylococcal enterotoxin serotypes (e.g., SEA to SEE, SEG, and SEI), the staphylococcal enterotoxin-like serotypes (e.g., SEI-H and SEI-J to SEI-V), and the distantly related toxic shock syndrome toxin 1 (TSST-1) (formerly serotype SEF) (255). The majority of isolates do not produce SAGs (132), while 90% of isolates that do generally produce more than one enterotoxin (48). Combinations observed include SEA and TSST-1 in 12% of strains, and SEG, SEI, SEN, SEO, and SEM occur together in 100% of strains, as these are carried on the same gene cluster. Conversely, certain toxin combinations remain rare (e.g., SEA and SEB) (123, 132).

SAGs represent a large family of biologically and genetically related toxins that are pyrogenic, cause T-cell proliferation, and induce the release of proinflammatory cytokines (189, 193). Unlike conventional antigens, which stimulate <0.01% of the T-cell population, picomolar concentrations of SAGs are able to stimulate 5 to 30% of the T-cell population (147). The magnitude and speed with which T-cell activation occurs and the accompanying cytokine induction can lead to shock (329).

As shock is a strong independent predictor of mortality in patients with SAB, it is reasonable to assume that isolates expressing these exotoxins would result in greater mortality. However, studies of patients with SAB found similar distributions of enterotoxins (SEA to SEG, SEH, SEI, and TSST-1) in strains isolated from patients who did survive and those who did not survive (48, 55, 241). Furthermore, a prospective study by Desachy et al. found no significant difference (OR, 1.60; 95% CI, 0.34 to 7.59;  $P = 0.55$ ) between the numbers of enterotoxin-producing strains isolated from surviving (70%; 16/23) and deceased (79%; 11/14) community-onset MSSA-B patients (48). Additional studies using a mouse septic model found no correlation between strains producing SEA, SEB, SEC, SED, or TSST-1 and mouse lethality (287).

Although enterotoxins play an important role in staphylococcal virulence, clinical data do not indicate that the presence of one or more of these factors impacts outcomes.

**Cytotoxic exotoxins.** Almost all *S. aureus* strains produce cytotoxic exotoxins. These toxins are able to disrupt host cells and evade host immune responses. Of the wide variety of staphylococ-

cal cytotoxic exotoxins produced, the most clinically important ones are the hemolysins (alpha-, beta-, gamma-, and delta-hemolysins), which have hemolytic activity toward human erythrocytes and have both dermonecrotic and neurotoxic effects (52).

**(i) Alpha-hemolysins.** The presence of alpha-hemolysin, measured by a zone of hemolysis using supplemented nutrient agar, was significantly associated with complicated bacteremia ( $P = 0.024$ ) but not mortality in a prospective study of 96 SAB infections (118).

**(ii) Beta-hemolysins.** A human study by Jin found that bacteremias caused by *S. aureus* strains with a disrupted beta-hemolysin (*hlyB*) gene (93/161; 58%), due to an insertion of a phage, were associated with a 4.3-fold decrease in mortality rates (129). While this may suggest that beta-hemolysin production increases the risk of mortality, previous mouse and rabbit models of ocular infections and septic arthritis have shown that beta-hemolysin plays a negligible role in *S. aureus* virulence (44, 213).

**(iii) Gamma-hemolysins.** Gamma-hemolysin is a unique three-gene (*hlgABC*) two-component hemolysin (60, 183). A mouse bacteremic model showed a 20% decrease in mortality, on average, in mice infected with an *hlgABC* deletion mutant of a USA300 strain, LAC (LAC  $\Delta hlgABC$ ), compared to the wild-type strain at 72 h (183). However, the survival rates after LAC  $\Delta hlgABC$  infection following phagocytosis did not differ compared to those after infection with the wild-type strain. In addition, the ability of gamma-hemolysin to cause pore formation in human neutrophils was significantly inhibited in the presence of 20 to 50% human serum or plasma. As *hlgABC* expression is dependent on specific host environments, growth conditions, and/or the disease state, the authors suggested that although gamma-hemolysin may contribute initially to immune evasion or survival in the blood, it plays a minor role in virulence and overall mortality.

**(iv) Delta-hemolysin and the accessory gene regulator.** The accessory gene regulator (*agr*) is a global regulon comprised of two transcripts, RNA II and RNA III. RNA II encodes *agrDBCA*, which is involved in generating autoinducing peptides (AIPs) for quorum sensing (178). AIPs also induce the expression of the other *agr* components. RNA III has multiple functions, including coding for delta-hemolysin and controlling the expressions of various exotoxins (e.g., hemolysins, enterotoxins, SEB, SEC, and TSST-1), virulence factors, and housekeeping genes (34). The production of delta-hemolysin can thus be used as a semiquantitative assay for *agr* function (292).

*agr* dysfunction has been associated with reduced killing by innate host defense cationic peptides (70, 250) and persistent (duration of  $\geq 7$  days) rather than resolving bacteremia (71.4% and 38.9%, respectively;  $P = 0.057$ ) (70). In addition, antibiotic efficacy, especially vancomycin bactericidal activity, is reduced with *agr*-dysfunctional isolates (245, 293, 294). All these factors combined probably explain the findings of a recent study by Schweizer et al., who detected an increased 30-day in-hospital mortality rate (RR, 1.52; 95% CI, 1.05 to 2.21;  $P = 0.03$ ) with *agr*-dysfunctional SAB (60% MRSA-B) episodes (258). This association was no longer significant following adjustments for age, comorbidities, severity of illness, and methicillin resistance but remained an independent predictor of 30-day mortality in severely ill patients when data were stratified by severity of illness (modified APACHE III score of  $>28$ ), thus suggesting that *agr* dysfunction is likely to

impact patients who most require an intact immune system and optimal antibiotic efficacy.

Based on the sequence diversity at this locus, 4 *agr* groups (groups I to IV) have been recognized (179). Several studies have examined the associations between the *agr* group and mortality, with conflicting results. Increased mortality was noted for *agr* non-group II (199) and group II (258) isolates, while no mortality difference between *agr* groups was noted in two studies (49, 120). These discrepancies may be explained in part by the considerable geographical variation in the *agr* group distribution (37 to 92% for group I, 4 to 50% for group II, 0 to 34% for group III, and 0 to 5% for group IV) among *S. aureus* isolates obtained from colonized and infected individuals (120, 251, 258, 264, 304).

(v) **Panton-Valentine leukocidin.** Like gamma-hemolysin, Panton-Valentine leukocidin (PVL) is a two-component pore-forming toxin (231) with cytolytic activity toward human neutrophils, macrophages, and monocytes (117). The genes encoding PVL, *lukS-PV* and *lukF-PV*, are known to be harbored by seven distinct temperate bacteriophages which can be acquired by *S. aureus* (20). The prevalence of PVL in *S. aureus* strains varies widely across the world (0 to 50%), and PVL is typically found in community-associated MRSA and MSSA isolates (16, 57, 121, 149, 153).

*In vitro* studies exposing human neutrophils to the supernatants of isogenic PVL-positive strains and also a PVL-negative strain, grown under conditions that promoted PVL upregulation (i.e., casein-hydrolysate-yeast extract [CCY] medium) (183), have provided conflicting evidence for the role that PVL plays. Neutrophil pore formation was observed in both samples treated with the supernatants from the PVL-positive and PVL-negative strains albeit 25 min later. This suggests that in some strains, there may be functional redundancies that exist, whereby other hemolysins (e.g., alpha-hemolysin) may be involved with disease progression (183). PVL did not affect the survival time (3 to 4 days) in a rabbit bacteremia model (51) despite rabbit neutrophils having similar susceptibilities to the effects of PVL as human neutrophils (217).

The likelihood of developing bacteremia is not affected by the presence of PVL (121, 203), with MRSA-B rates being similar in patients with intranasal MRSA, irrespective of PVL carriage (16). PVL is associated with diseases such as skin and soft tissue infections (22), osteoarticular infections (137), and necrotizing pneumonia (78). With the exception of necrotizing pneumonia (79), PVL has not been found to be associated with increased mortality rates, including SAB (16, 212, 311, 314). Potential reasons for this include the high likelihood of a skin and soft tissue source for SAB (203, 212, 290, 314) and the predominance of PVL-associated disease occurring in younger patients (9, 203). PVL does not seem to augment already-established infections, including bacteremias (153), but is considered important in the early stages of establishing infections (16, 50, 51) or evading an initial innate immune response by lysing neutrophils (22, 51) and is therefore unlikely *per se* to influence mortality. Nevertheless, the role and contribution of PVL to pathogenesis, morbidity, and mortality remain a controversial area subject to much study.

Overall, the data point to a possible role for various exotoxins in influencing outcomes in SAB. These effects probably result from a complex interplay between toxin expression (concentration and duration) and host immune responses (287), which in turn influence clinical manifestations and subsequent outcomes. However, these complexities are unlikely to be teased apart in the

foreseeable future secondary to the multiple confounders (246), and thus, the impact of toxins remains largely speculative based on *in vitro* and animal data.

### **S. aureus Clonal Types**

The application of various typing methodologies, such as pulsed-field gel electrophoresis (PFGE) and multilocus sequence typing, has shown that *S. aureus* has a varied clonal population structure. Few studies have examined the impact of different clonal types on mortality in SAB. A single-institution prospective U.S. study of 116 MRSA-B episodes found that USA300 isolates typed by PFGE (34%) were associated with a reduced mortality rate compared to rates associated with USA100 and USA500 isolates upon a univariate analysis only (8%, 29%, and 36%, respectively) (261). The most likely explanation for this finding is that USA300 isolates infected younger patients, with skin and soft tissue infections as the predominant source of SAB. This may change over time as USA300 becomes more established in health care settings (229). A more recent retrospective multicenter U.S. study (2005 to 2008) examined case data for 1,104 typed MRSA-B isolates (138). In contrast to data from the previous study, a multivariate analysis revealed that the USA300 type (37.5% of 1,104 isolates) was an independent predictor of mortality compared to non-USA300 PFGE types after adjustment for age (hazard ratio [HR], 1.63; 95% CI, 1.19 to 2.23;  $P = 0.002$ ). Although animal data support increased virulence with USA300 strains (166), clinical data remain limited. Clearly, more research is required in this area to determine if various *S. aureus* clones have a definitive association with increased mortality in SAB.

## **MANAGEMENT**

### **Antibiotic Choice**

The choice of antibiotic is important, especially for the treatment of MSSA-B episodes. Cefazolin, a narrow-spectrum cephalosporin, has been used for the treatment of MSSA-B since the 1970s. The role of this agent was questioned, however, following case reports of treatment failures, suggesting a reduced efficacy compared to the efficacy of antistaphylococcal penicillins (23, 234). Potential explanations for this included an increased susceptibility of cefazolin to the inoculum effect (208) and to staphylococcal  $\beta$ -lactamases (248). No randomized control trial comparing these agents has been performed to definitively answer questions regarding relative efficacies. A single-center cohort study comparing cloxacillin ( $n = 281$ ) with cefazolin ( $n = 71$ ) for MSSA-B infections detected no mortality or outcome difference between the two agents (222). Baseline patient characteristics differed significantly between the two groups, and this probably influenced the clinicians' initial antibiotic selections, thus limiting possible conclusions. In a recent propensity-score-matched, case-control study, where the nafcillin supply was interrupted for a 2-year period (i.e., cefazolin period), correction for clinicians' antibiotic choices was able to occur (162). Although no mortality difference or treatment failure was documented at the end of therapy (4 weeks), the low number of episodes (41 in each arm) and the infrequent high-inoculum infections (infective endocarditis,  $n = 2$ ) likewise limit the conclusions of this study. Thus, questions remain about the comparative efficacies of these agents.

For the treatment of MSSA-B infections, the 30-day mortality rate was higher for patients receiving empirical treatment (i.e.,

commenced within the first 48 h) with a third-generation cephalosporin (OR, 2.24; 95% CI, 1.23 to 4.08;  $P = 0.008$ ) or  $\beta$ -lactam- $\beta$ -lactamase inhibitor combinations (OR, 2.68; 95% CI, 1.23 to 5.85;  $P = 0.013$ ) than for patients receiving cloxacillin or cefazolin (222). In that same study, no conclusions could be made for definitive MSSA-B therapy (given from days 3 to 9, i.e., the first week after the receipt of blood culture results), as the numbers of patients treated with broad-spectrum  $\beta$ -lactam agents were too small compared to the numbers of patients treated with cloxacillin or cefazolin.

Vancomycin has consistently been associated with increased rates of treatment failure and high mortality rates compared to  $\beta$ -lactams when used for the management of MSSA-B (29, 85, 139, 140, 174, 281). In two cohort studies, vancomycin treatment was significantly associated with infection-related mortality upon a multivariate analysis (with one study reporting an OR of 14.5 and a 95% CI of 1.43 to 146.64 [85] and the other study reporting an OR of 3.3, a 95% CI of 1.2 to 9.5, and a  $P$  value of 0.02 [140]). This association remained in the latter study, even after adjusting for other variables, including the likelihood of receiving vancomycin. Further evidence for suboptimal vancomycin treatment responses came from a large prospective cohort study of 1,865 SAB episodes, where glycopeptide use and not antibiotic resistance (i.e., MRSA) was an independent predictor of mortality (OR, 1.64; 95% CI, 1.08 to 2.48;  $P = 0.02$ ) (296). This difference in outcomes may persist even after replacing vancomycin treatment once susceptibility results become available for patients with endocarditis (174). All groups, including hemodialysis patients, seem equally susceptible to treatment failures with vancomycin (281). Possible reasons for these poor outcomes with vancomycin include slower bactericidal activity, especially in high-inoculum infections (26, 158, 252, 271), and variable tissue penetration (2, 88, 157). Vancomycin is thus considered a suboptimal therapy for MSSA-B compared to semisynthetic penicillins.

For the treatment of MRSA-B, vancomycin remains the treatment of choice, with daptomycin as an alternative, except for patients with left-sided endocarditis (172). This recommendation is based on an open-label, randomized, controlled noninferiority SAB study, where daptomycin was as effective as standard therapy including antistaphylococcal penicillins for MSSA-B (65).

There seems to be little difference between the glycopeptides provided that teicoplanin is dosed appropriately, especially for cases of endocarditis (319). In a recent systematic review and meta-analysis, all-cause mortality rates were similar between teicoplanin and vancomycin treatments, with significantly greater toxicity when vancomycin was prescribed (283).

A direct comparison of treatment outcomes with linezolid compared to vancomycin in SAB is lacking. Although not aimed at investigating the efficacy of linezolid for SAB, three multicenter open-label randomized trials included SAB episodes. For the treatment of suspected *S. aureus* infections (18% SAB) (278) and Gram-positive infections in pediatric patients (80 SAB episodes) (135), the efficacy of linezolid was comparable to that of vancomycin. In contrast, linezolid was associated with an excess of deaths compared to the number of deaths associated with vancomycin treatment in a trial of catheter-related episodes (142 SAB episodes in the intention-to-treat analysis) (318). Further analyses suggested that these deaths occurred in patients with either Gram-negative pathogens or no pathogens isolated at baseline and not as a result of SAB. In a systematic review and meta-analysis, overall,

linezolid was equivalent in efficacy to vancomycin (OR, 0.49 for mortality; 95% CI, 0.49 to 1.58;  $P = 0.46$ ) for the treatment of SAB (12).

In a randomized controlled trial comparing cotrimoxazole (trimethoprim-sulfamethoxazole) to vancomycin in 101 injection drug users with *S. aureus* infections (66% SAB), vancomycin was found to be superior with regard to the duration of bacteremia and clinical cure rates (188). As all MRSA infections were cured, the outcome differences were explained by cotrimoxazole treatment failure in MSSA episodes. Although the numbers of patients were small, a recent retrospective case-controlled study observed no mortality or treatment failure difference between patients treated with cotrimoxazole ( $n = 38$ ) and those treated with vancomycin ( $n = 76$ ) for MRSA-B (81). These data suggest a possible role for cotrimoxazole in the management of MRSA-B. However, without a randomized controlled trial, these findings remain speculative.

Theoretical considerations suggest a potential role of adjunctive rifampin in the treatment of SAB, as rifampin penetrates biofilms (11, 327) and is rapidly bactericidal (45). Several small clinical studies have been reported, with conflicting results (165, 298, 299). Thus, without a clear benefit (226) and possibly increased hepatic adverse events (236), adjunctive rifampin therapy is generally not recommended (172). The addition of gentamicin has similarly been associated with greater toxicity for no mortality benefit (43, 122) and likewise is no longer recommended (172). With the exception of adjunctive levofloxacin, which did not result in improved outcomes for the treatment of quinolone-sensitive SAB episodes in a randomized trial (247), other combinations remain limited to *in vitro* data and case reports, with variable successes (211).

For the management of SAB, with the exception of the definitive inferiority of glycopeptides compared to  $\beta$ -lactams for the treatment of MSSA-B, no alternative agent or combination has been shown to result in better patient outcomes than current standard therapy. This may in part reflect antibiotic registration study design, and thus, further research aimed at determining better treatment options is urgently required, especially for MRSA-B episodes.

### Empirical Antibiotics/Antibiotic Timing

The harmful effects of delayed appropriate empirical therapy, defined as *in vitro* antibiotic activity against the infecting pathogen, have been well documented for patients with sepsis (150, 221) and bacteremia (163). Multiple studies have confirmed these observations for the treatment of SAB (MSSA-B or MRSA-B) (139, 176, 256, 275), with a recent meta-analysis observing an overall 2-fold-increased survival benefit (OR, 1.84; 95% CI, 1.25 to 2.71;  $P < 0.01$ ) with the administration of appropriate empirical therapy for MRSA-B episodes (220).

However, unlike studies of sepsis, no “time response curve” (increasing mortality for every hour of delay in antibiotic administration) has been detected for SAB treatment; rather, time cutoffs have been detected, after which mortality is increased. In a single-center retrospective study of 167 hospital-acquired SAB (67% MRSA) episodes, Lodise et al., using classification regression tree analysis, identified an empirical/delayed antibiotic “breakpoint” of 44.75 h (176). This time cutoff has, however, never been replicated and has ranged between studies from 24 h up to more than 72 h (139, 176, 256, 275). The importance of

appropriate empirical antibiotics (within 24 h) remained, even after adjusting for other risk factors of mortality, in a 2010 study by Paul et al. of 510 health care-onset MRSA-B episodes (220). Poorer outcomes have also been demonstrated using a case-controlled study design, with a 2.35-fold increase in mortality in MRSA-B patients whose therapy was delayed by more than 2 days (OR, 2.35; 95% CI, 1.01 to 5.44;  $P = 0.05$ ) (187).

Provided that the antibiotics prescribed have some activity, even if they are not considered standard therapy, for example, gentamicin alone or in combination with other agents (e.g., ciprofloxacin), mortality may be improved (240). Antibiotic dosing is critical, however, as Shime et al. demonstrated an incremental survival benefit if patients received empirical antibiotics (within 48 h) and attained a vancomycin target of  $>400 \mu\text{g} \cdot \text{h/ml}$  compared to those who did not achieve appropriate targets (262).

Not all patients may benefit, however, from appropriately timed empirical therapy, with several studies, irrespective of design, being unable to detect a mortality difference (5, 6, 59, 141, 168, 243). An indication as to which patient subgroups would benefit emerged from a study by Lodise et al., who observed that the greatest impact of empirical antibiotics (within 44.75 h) occurred in patients with severe disease (APACHE II score of  $>15.5$ ) or complicated SAB (176). However, a larger single-center study of 814 SAB episodes detected a reduced mortality rate only for empirically (24 h before to 24 h after the index blood culture) treated patients with the lowest severity-of-illness score (modified APACHE III score) (HR, 2.42; 95% CI, 1.7 to 3.44) (257), whereas a study by Kim et al. noted that empirical antibiotics (within 48 h) resulted in outcome differences only for patients with a noneradicable focus compared to eradicable foci of SAB (142). Conversely, some patients will improve irrespective of inadequate empirical or definitive SAB therapy (6). Theoretically, the greatest benefit is likely to occur when antibiotics are still able to affect the progression of infection (141) and thus impact infection-related mortality (257). This suggests, therefore, that ill but less severely ill patients are most likely to benefit (167).

Potential explanations for these discrepancies include differing empirical therapy or appropriate antibiotic therapy (including vancomycin for MSSA episodes) definitions, patient groups studied, and possible confounders (e.g., initial treatment selection bias). Several studies did not differentiate between empirical and definitive therapies (131, 176, 184), while the need to adjust for confounders, especially shock, remains debated (257, 273). Regardless, earlier appropriate antibiotic prescribing is likely to be of benefit, albeit not for all patients, and is generally recommended by most experts.

### Surgery

The impact of surgery on mortality is unclear. Surrogate markers of source control suggest that surgery may improve outcomes. Jensen et al., in a prospective study of 278 SAB cases, demonstrated that eradicable foci were associated with a reduced mortality rate compared to noneradicable foci of infection (13% and 31%, respectively;  $P = 0.002$ ) (127). Similarly, in a study of 294 MSSA-B episodes, the eradication of infection was an independent predictor of reduced mortality (OR, 0.3; 95% CI, 0.1 to 0.7;  $P = 0.006$ ) (140). Furthermore, the intervention *per se* seems to be responsible for better outcomes, as an increased mortality rate was detected for patients with an eradicable focus that was not removed or drained (OR, 4.17; 95% CI, 1.09 to 3.62;  $P = 0.04$ )

(142). Fowler et al., in a series of 244 infected intravascular device infections, found comparable results, with higher rates of relapse and mortality if the device was not removed (OR, 6.5; 95% CI, 2.1 to 20.2) (69). Only one study to date has found infection-related surgery with respect to SAB to be an independent predictor of reduced mortality (OR, 0.27; 95% CI, 0.09 to 0.83;  $P = 0.022$ ) (301). This observation is strongly confounded, however, by the patient's ability to undergo surgery or the surgeon's willingness to perform the procedure. As the eradication of infection sources improves outcomes, it would make intuitive sense that surgical source control would have a similar impact.

Unlike uncomplicated SAB diagnoses, there is definitive evidence that surgery improves outcomes compared to medical therapy alone, although this is limited to patients with complicated infective endocarditis. In a multinational propensity-matched prospective cohort study, the adjusted in-hospital mortality rate was lower for patients with paravalvular complications, systemic embolization, and *S. aureus* native valve IE (152). Other studies have shown improved outcomes with surgery for patients with congestive heart failure and perivalvular complications (1, 307).

For patients with prosthetic joint infections, reduced cure and retention rates are associated with an increased duration of symptoms and suggest that earlier surgical intervention is one of the factors that may result in improved outcomes (328). Besides these data, which provide possible indirect evidence for the efficacy of early intervention, the optimal timing of surgery for the management of SAB is unknown.

### Role of Infectious Disease Consultation

Infectious disease (ID) consultation for SAB episodes generally results in a more detailed evaluation, a greater detection of endocarditis or metastatic complications, and improved adherence to treatment guidelines (69, 125). Fowler et al. showed that an acceptance of ID management resulted in significantly fewer relapses, while Jenkins et al. observed a trend toward fewer treatment failures and reduced mortality following the implementation of an ID consultation service for all SAB patients. In two separate studies, of 308 SAB episodes (82% received an ID consult) (131) and 293 SAB episodes (36% received an ID consult) (206), ID consultation was associated with better 30-day mortality rates by a univariate analysis, while in a German study of 521 SAB episodes, ID consultation was an independent predictor of reduced mortality (OR, 0.6; 95% CI, 0.4 to 1.0;  $P = 0.045$ ) (237). These results were obtained even though sicker patients and patients with more complicated diseases, such as infective endocarditis, were referred. Differences in these results are probably related to study design, patient populations, baseline institutional expertise in SAB management, and adherence to ID advice.

### Special Populations

**Neonates.** *S. aureus* is among the most important causes of bacteremia in neonatal intensive care units (NICUs) (115, 279), with specific *S. aureus* clonal outbreaks (e.g., phage type 86 [112] or USA300 [64]) being documented in newborn nurseries. Similar to adults, SAB is an important cause of both hospital-acquired (325) and community-onset (214, 269, 326) infections in neonates from developing countries.

Data from the U.S. NICU Network (279) and the Danish SAB Registry ( $n = 300$ ) (58) suggest that overall mortality rates are similar to those for adults (17.2% and 23%, respectively). Identifi-

fied risk factors for mortality include low birth weight (<1,000 g;  $P < 0.01$ ) (58) and age at the onset of infection, with early-onset MSSA sepsis ( $\leq 48$  h of age) being associated with a significantly higher mortality rate than that associated with late-onset MSSA infections (39% versus 7.3%;  $P < 0.01$ ) (115).

The impact of methicillin resistance is unclear, with MRSA mortality rates being significantly higher than those associated with MSSA episodes (24.6%, versus 9.9% for MSSA;  $P < 0.001$ ) in an Australian and New Zealand NICU study by Isaacs et al. (115). However, MRSA episodes occurred in infants who were more premature and of lower birth weights. In contrast, at the Duke University Medical Center (37), NICU patient characteristics were similar, with no significant differences in mortality rates (19% for MRSA versus 6% for MSSA infections), length of stay, or later neurodevelopmental impairment between the two groups.

A report of CA-MRSA bacteremia in neonates from Texas (94) noted a high mortality rate of 38%. However, comparative data with other SAB clonal episodes were not documented. In a neonatal intensive care unit study by Kuint et al. (149), of 11 CA-MRSA (*pvl*-negative), 20 multidrug-resistant MRSA, and 12 MSSA bacteremic episodes, mortality (9%) was not dependent on the *S. aureus* subtype despite CA-MRSA episodes occurring in younger neonates.

In summary, the data on SAB mortality in neonates are limited. There is a suggestion, however, that patient factors such as the degree of prematurity and birth weight may influence outcomes, whereas the effects of methicillin resistance or *S. aureus* clonal types cannot be definitively determined.

**Children.** Age remains one of the most consistent predictors of mortality, with children generally having lower SAB mortality rates than adults. In one of the largest prospective cohort studies to date that included children, the mortality rate was the lowest for individuals aged between 10 and 19 years (296). Although rates tend to be higher in children aged between 0 and 9 years, this probably reflects mortality in neonates rather than children older than 1 year of age, with mortality being 10- to 17-fold higher in <1-year-olds than in older children (71).

Overall, recent mortality rates vary widely in developing countries, at 35 to 40% in Thailand (0- to 10-year-olds) (212), 25% in Kenya (151), and 15% in South Africa (75), with lower rates documented in Tanzania (7.1%) (17) and Mozambique (6%) (269). These results are difficult to interpret, as they may represent access to and the level of health care as well as the impact of high HIV coinfection rates in these countries. In contrast, overall mortality rates are generally much lower in series from developed countries: 0.7% (infection-related mortality) in Australia (282), 3% in New Zealand (109), 1.5% in Finland (180), and 2% in the United Kingdom (46). In the United States, mortality rates for children with SAB range between 3.2% and 18% (13, 90, 297). The wide range of mortality rates may be explained in part by the associated comorbidities, with a higher 1-year mortality rate (18%) seen in a cohort study including many children with congenital heart disease (297). This is supported by a Danish study of 2,468 pediatric SAB episodes in patients aged between 1 and 20 years, where the presence of comorbidities was an independent risk factor for death (OR, 2.49; 95% CI, 1.71 to 3.62) (71).

The place of SAB onset influences outcomes, with community-onset episodes having a lower mortality rate than health care-acquired episodes (0.6% and 3.9%, respectively) (71), probably secondary to the predominance of skin and soft tissue infections

and bone and joint infections in community-onset episodes. Conversely, the presence of comorbidities and, hence, the need for hospitalization may explain the higher mortality rates for children aged between 11 and 20 years with nosocomial SAB episodes. As with adults, the source of infection influences outcomes, with no mortality occurring in oncology patients with line-associated SAB (239, 277). Infective endocarditis, although rare in children, is associated with an increased 1-year mortality rate (40%, versus 12% for children with no IE) (297). Both IE (patients 0 to 10 years old) and pulmonary infections (patients <1 and 11 to 20 years old) were independent predictors of mortality (71). Although small case series have documented high mortality rates associated with *pvl*-positive and CA-MRSA infections (27, 83, 84), a lack of comparative data and the small sample size limit drawing conclusions about the impact of these factors on outcomes.

The role of antibiotic resistance is likewise unclear. Burke et al., in a retrospective cohort study of 164 U.S. children (24), reported a difference in mortality rates for MRSA-B compared to MSSA-B (10% versus 4%). In contrast, a study of 69 U.S. children, when adjusted for underlying diseases and delays in appropriate antibiotic therapy, found MRSA not to be associated with increased mortality (280). The impact of treatment is similarly unclear, with no large comparative studies of children being reported. Although vancomycin treatment failure was documented in a small pediatric series ( $n = 22$ ) of MRSA-B, the mortality rate seemed similar to those of other reported data, at 13.6%. Treatment failure was associated with prematurity and *pvl*-positive infections but not a high vancomycin MIC (by either the Etest or broth microdilution) (315).

In summary, pediatric SAB mortality rates were low in recent series from developed countries and, similarly to adults, are declining with better management in children, from 20% (1990 to 1994) to 11% (1993 to 2007) and from 19.5% (1971 to 1976) to 2.5% (1996 to 2000), based on data from Argentina (215, 216) and Denmark (71), respectively. Although outcomes are influenced by the presence of comorbidities and infection manifestations, the impact of other variables remains unclear and warrants further study.

## CLINICAL IMPLICATIONS, FUTURE RESEARCH, AND CONCLUSIONS

The most consistent predictor of mortality in patients with SAB is age. The impacts of other host factors, except for the presence of comorbidities, are unclear. Pathogen-host interactions, especially the presence of shock, and the source of SAB are strong predictors of outcome. Although antibiotic resistance may be associated with increased mortality, questions remain as to whether this reflects pathogen-specific factors or poorer responses to antibiotic therapy, namely, vancomycin. Optimal management relies on starting appropriate antibiotics in a timely fashion, with outcomes being improved in certain patient subgroups.

Of all these factors, the treating clinician can only influence patient management. These remain uncertain, however, with no real indication of superior outcomes with any alternative antibiotic, especially for the treatment of MRSA-B. With the advent of "personalized" care, hopefully, better treatment algorithms can be developed (e.g., the need for early surgery or a certain antibiotic for line-related SAB). For this to occur, a greater understanding of predictors of mortality is required, which possibly necessitates the inclusion of genomic approaches in future studies. Furthermore,



more international collaborative studies should be encouraged, similar to the International Collaboration of Infective Endocarditis or HIV cohort studies. This would allow the standardization of data collection and adequate power to tease out some of the controversies in SAB management.

In conclusion, SAB remains a common infection with significant associated mortality. Its impact and significance remain underestimated in the community. A greater recognition of the impact of SAB will hopefully lead to more research on and a better understanding of predictors of mortality, which subsequently would assist SAB management and optimize patient outcomes.

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