

Supplementary Table 2: Prevalence of *S. aureus* clonal complexes and *tst* in the SHIP-TREND-0 study sample.

CC ¹	Prevalence of CCs among all isolates No. (%)	Prevalence of <i>tst</i> gene within CCs No (%) ²
<i>tst</i> -positive CCs		
CC5	9 (3.75)	1 (1.11)
CC22	15 (6.25)	1 (6.66)
CC30	48 (20.0)	40 (83.33)
CC34	4 (1.66)	2 (50.0)
CC395	5 (2.08)	5 (100.0)
<i>tst</i> -negative CCs		
	159 (66.25)	0 (0.0)
All strains ³	240	49 (20.4)

¹ Clonal complexes (CCs) were determined by *spa* typing (S1).

² *tst* was detected by multiplex PCR (S2).

³ excluding 9 strains that were not available for *spa* typing and virulence gene profiling