

AUTHOR CORRECTION

Correction for Aamot et al., Genotyping of 353 *Staphylococcus aureus* Bloodstream Isolates Collected between 2004 and 2009 at a Norwegian University Hospital and Potential Associations with Clinical Parameters

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Volume 50, no. 9, p. 3111–3114, 2012. Page 3112, Table 1: For the sixth entry in the “CC” section of the first column, “*spa* C304/024” should read “*spa* CC304/024,” and owing to misclassification of infections as hospital acquired (HA) or community acquired (CA), several entries in the third and fourth columns were wrong. The conclusions of the paper are unchanged, as the misclassifications went in both directions. The first four columns of the table should appear as follows.

<i>spa</i> type, <i>spa</i> CC, PVL, or <i>tst</i>	No. (%) of isolates	No. (%) with:	
		CA infection	HA infection
<i>spa</i> type			
t084	46 (13.0%)	27 (58.7%)	19 (41.3%)
t002	18 (5.1%)	11 (61.1%)	7 (38.9%)
t065	15 (4.2%)	11 (73.3%)	4 (26.7%)
t012	14 (4.0%)	7 (50.0%)	7 (50.0%)
Other <i>spa</i> types ^c	260 (73.7%)	158 (60.8%)	102 (39.2%)
Total	353	214 (60.6%)	139 (39.4%)
CC			
<i>spa</i> CC084	69 (20.7%)	38 (55.1%)	31 (44.9%)
<i>spa</i> CC021	55 (16.5%)	36 (65.5%)	19 (34.5%)
<i>spa</i> CC133	32 (9.6%)	24 (75.0%)	8 (25.0%)
<i>spa</i> CC065	30 (9.0%)	19 (63.3%)	11 (36.7%)
<i>spa</i> CC002	22 (6.6%)	14 (63.6%)	8 (36.4%)
<i>spa</i> CC304/024	22 (6.6%)	15 (68.2%)	7 (31.8%)
<i>spa</i> CC359	20 (6.0%)	12 (60.0%)	8 (40.0%)
Other <i>spa</i> CCs ^e	83 (24.9%)	49 (59.0%)	34 (41.0%)
Total	333	207 (62.2%)	126 (37.8%)
PVL	11 (3.1%) ^f	11 (100%)	0
<i>tst</i>	54 (15.3%)	33 (61.1%)	21 (38.9%)

Page 3112, Table 1: Footnote *f* should read “ $P = 0.004$ for PVL in CA infection versus HA infection: 100% versus 0%.”

Page 3113, Fig. 1, second row: “*spa*-CC304/204” should read “*spa* CC304/024.”

Citation Aamot HV, Blomfeldt A, Eskesen AN. 2016. Correction for Aamot et al., Genotyping of 353 *Staphylococcus aureus* bloodstream isolates collected between 2004 and 2009 at a Norwegian university hospital and potential associations with clinical parameters. *J Clin Microbiol* 54:2403. doi:10.1128/JCM.01434-16.
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