Screening cultures for detection of methicillin-resistant *Staphylococcus aureus* in a population at high risk for MRSA colonisation: identification of optimal combinations of anatomical sites

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This retrospective study analysed the diagnostic yield of single-site, two-site, and three-site anatomical surveillance cultures in a population of 4,769 patients at high risk for methicillin-resistant *Staphylococcus aureus* (MRSA) colonisation. Cultures of seven anatomical sites were used as the gold standard against which to measure the sensitivity of MRSA detection. Detection rates for the seven single-sites, 21 two-site, and 35 three-site combinations are presented. Single-site swabbing only detected 50.5% (nose) of total cases, while three-site surveillance achieved a 92% (groin + nose + throat) sensitivity of detection at best. It is recommended that at least three anatomical sites should be screened for MRSA colonisation in these high-risk patients.

Keywords: MRSA screening; optimal sensitivity; infection control

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Infections arising from methicillin-resistant *Staphylococcus aureus* (MRSA) are an important cause of healthcare-associated infections (HCAI) throughout the world. They are associated with an increase in morbidity, mortality, and an increase in healthcare costs and there are indications that MRSA infections are increasing worldwide (1). Individuals colonised with MRSA are at a greater risk of developing infections with this organism (2). The aim of this retrospective study was to determine the optimal body sites and combinations of sites for detection of MRSA carriage in a high-risk population using sampling from seven anatomical sites as a gold standard. Single-site swabbing has a low sensitivity of detection despite it being recommended in some guidelines (3). Using culture on selective media, we determined the sensitivity of a single anatomical site sampling in this high-risk population as well as all possible combinations of two and three anatomical sites to find the highest detection rates.

**Materials and methods**

Between January 1 2010 and November 30 2012, 4,769 sets of body swabs were taken from adult patients admitted to the Abertawe Bro-Morgannwg University Hospital in Swansea, Wales. The body sites swabbed in all patients were as follows: axilla, hairline, groin, nose, perineum, throat, and umbilicus. Only patients where all these anatomical sites were swabbed simultaneously were accepted to this retrospective study. These were adult patients who were classed as being at high risk for MRSA colonisation because of their frequent re-admissions to healthcare facilities; direct inter-hospital transfers; recent admissions at a hospital known or likely to have a high prevalence of MRSA; residents of nursing or residential care homes; patients being admitted to high-risk areas; transfers from outside the United Kingdom, or patients with a history of colonisation with MRSA in the past. Upon admission, swabs were taken from the above-mentioned body sites, transported in charcoal transport medium (Amies) and plated separately onto chromogenic MRSA medium (P&O Laboratories). Suspect colonies were subcultured onto Columbia blood agar (Oxoid Ltd) and identified with a latex agglutination kit (Pro-Lab) and each new isolate further tested on panel PMIC/ID-67 of the Becton-Dickenson Phoenix ID system for
confirmation of identity and provision of an antimicrobial susceptibility profile. *S. aureus* isolates were also identified using mass spectrometry, MALDI-TOF (Bruker) (4, 5). Patients were considered colonised if MRSA was grown from any of the body sites tested. The seven-site body screen was regarded as the reference standard against which the sensitivity of single and combined swabs was measured.

**Statistics**

The number of different combinations for the two-site and the three-site determinations were calculated to be seven for the single-site swabs, 21 for the two-site swabs, and 35 for the three-site swabs. Confidence intervals were calculated using the normal approximation method of the binomial confidence interval.

Ethical approval was deemed unnecessary by the local ethics committee, as this was a retrospective study.

**Results**

The number of seven-site anatomical sampling screens taken from 4,769 individuals that detected MRSA was 925 (19.4%). The patient was considered colonised with MRSA if any one of the swabs was found positive for MRSA. Single swabs detected MRSA colonisation at rates between 18% (hairline) and 50.5% (nose) as shown in Table 1. The ability of two anatomical site swabs to improve the detection of MRSA colonisation was determined by using all possible combinations from the original set of seven sites sampled. There were 21 combinations possible and these are shown in Table 1. The two-site swabs improved the detection rate of MRSA by up to 50% with the best combination being the groin and throat where 74.5% of colonisations were detected. These were closely followed by the groin/nose, nose/perineum, and the nose/throat combinations. Of note is that some of the two-swab combinations, such as the axilla/hairline and the hairline/umbilicus combinations, had poorer detection rates than single swabs of the nose, groin, perineum, or throat. However, 14 other two-site combinations were more sensitive at detecting MRSA colonisation than the single nasal swab (Table 1).

Three anatomical site combinations yielded even better results than the two-site anatomical combinations when compared to the seven-site anatomical gold standard. Table 1 shows all 35 different combinations of three anatomical site sampling with the best detection rate accounted for by a combination of groin+nasal+throat, which detected 92% of colonisations closely followed by perineum+nose+throat combinations (91%). It can be noted from Table 1 that all but one of the triple-site sampling had detection rates superior to that of the single nose swab. The only combination with inferior results was the axilla+hairline+umbilicus combination. Sampling of the groin and throat together improved MRSA detection by 47% and sampling of the groin, nose and throat together improved MRSA detection by 82% when compared to single-site sampling from the nose.

**Discussion**

In this study, we assessed the role of MRSA screening in patients at high risk for MRSA colonisation. Sampling from seven anatomic body sites was used as the reference gold standard against which to measure single-site, two-site, and three-site sampling combinations. This was to determine the smallest combination of surface body sites that can give the highest detection rates of MRSA colonisation. Colonisation with MRSA is associated with an increased likelihood of clinical infection with MRSA (6). Screening patients for MRSA has been advocated to detect carriers and enforce decolonisation procedures which can reduce infections and prevent transmission to others. Different countries have differing policies on the screening procedures. US guidelines suggest only that nasal swabbing may be sufficient (3), whereas many in the United Kingdom recommend at least two sites for screening (7, 8). In areas of low endemicity, single-site screening may be sufficient to detect colonisation. However, in patients at high risk for MRSA colonisation, the single-site nasal swab is likely to miss a significant number of colonisations and allow spread of the organism in healthcare establishments hence putting patients at risk of developing infections with MRSA. In this study, we found that swabs of the anterior nares alone detected only about half (50.5%) of all possible cases whereas a triple swab combination of nose, groin, and throat detected 92% of colonisations. This contrasts with the Scottish results where swabs of the nares detected 66.4% of MRSA colonisations (9). Throat swabs alone performed less well (10). Table 1 shows the results of two-site swabbing, whereby groin and throat combinations improved sensitivity of detection to 74.5%. It can also be seen that 10 of the 21 two-site combinations gave superior results to the nose swab alone. Certain anatomical site combinations lacked sensitivity and gave poor results. Axilla, hairline, and umbilicus combinations could not achieve the sensitivity of single nose or throat swabs (Table 1). However, triple swab combinations of most sites were as sensitive as or of higher sensitivity than nasal, throat, or groin swabs individually. From Table 1, it can be seen that only one of the 35 different combinations failed to achieve the sensitivity of the single nasal swab. Optimal two-site screening is likely to miss up to 25% of MRSA colonisations. This study is unique in that it calculates all of the different combinations of anatomic sites including all 21 possible two-site and all 35 three-site combinations. This allows clinicians to predict the relative accuracy of combination swab results in relation to the commonly
Table 1. Numbers of MRSA colonisations detected at different anatomical sites and combinations of two and three sites

| Anatomical site | No. of patients MRSA detected (total = 925) | Sensitivity (%) of detection (95% CI) |
|-----------------|--------------------------------------------|--------------------------------------|
| **Single-site** |                                            |                                      |
| Nose            | 467                                        | 50.5% (47.3–53.7)                    |
| Throat          | 444                                        | 48% (44.8–51.2)                      |
| Groin           | 408                                        | 44.2% (40.9–47.3)                    |
| Perineum        | 365                                        | 39.5% (36.3–42.6)                    |
| Umbilicus       | 214                                        | 23.1% (20.4–25.9)                    |
| Axilla          | 171                                        | 18.5% (16.2–21)                      |
| Hairline        | 167                                        | 18% (15.6–20.5)                      |
| **Two-site**    |                                            |                                      |
| Groin + throat  | 689                                        | 74.5% (71.7–77.3)                    |
| Groin + nose    | 667                                        | 72.1% (69.2–75)                      |
| Nose + perineum | 664                                        | 71.8% (68.9–74.7)                    |
| Nose + throat   | 662                                        | 71.6% (68.7–74.5)                    |
| Perineum + throat| 582                                         | 63% (59.8–66)                        |
| Throat + umbilicus | 553                                 | 59.8% (56.6–62.9)                    |
| Nose + umbilicus | 549                                        | 59.35% (56.2–62.5)                   |
| Axilla + nose   | 524                                        | 56.6% (53.5–59.8)                    |
| Axilla + throat | 521                                        | 56.3% (53.1–59.5)                    |
| Groin + perineum | 518                                        | 56% (52.8–59.2)                      |
| Hairline + throat | 513                                     | 55.46% (52.3–58.7)                   |
| Hairline + nose | 501                                        | 54.1% (51–57.4)                      |
| Groin + umbilicus | 482                                         | 52.1% (48.9–55.3)                    |
| Groin + hairline | 476                                        | 51.45% (48.2–54.7)                   |
| Axilla + groin  | 457                                        | 49.4% (46.2–52.6)                    |
| Perineum + umbilicus | 455                                        | 49.2% (46–52.4)                      |
| Axilla + perineum | 435                                         | 47% (43.8–50.2)                      |
| Hairline + perineum | 432                                         | 46.7% (43.5–49.9)                    |
| Axilla + umbilicus | 300                                         | 32.4% (29.4–35.4)                    |
| Hairline + umbilicus | 293                                         | 31.7% (28.7–34.7)                    |
| Axilla + hairline | 267                                         | 28.8% (25.9–31.8)                    |
| **Three-site**  |                                            |                                      |
| Groin + nose + throat | 850                                         | 92% (90.1–93.6)                      |
| Nose + perineum + throat | 840                                         | 91% (88.9–92.7)                      |
| Groin + perineum + throat | 783                                         | 84.6% (82.3–87)                      |
| Groin + nose + perineum | 769                                         | 83.1% (80.7–85.5)                    |
| Groin + throat + umbilicus | 752                                         | 81.3% (78.8–83.8)                    |
| Nose + throat + umbilicus | 742                                         | 80.2% (77.6–82.8)                    |
| Axilla + groin + throat | 732                                         | 79.1% (76.5–81.7)                    |
| Groin + hairline + throat | 726                                         | 78.5% (75.8–81.1)                    |
| Groin + nose + umbilicus | 724                                         | 78.3% (75.6–81)                      |
| Perineum + throat + umbilicus | 720                                         | 77.8% (75.2–80.5)                    |
| Nose + perineum + umbilicus | 719                                         | 77.7% (75–80.4)                      |
| Axilla + nose + throat | 717                                         | 77.5% (74.8–80.2)                    |
| Groin + hairline + nose | 708                                         | 76.5% (73.8–79.3)                    |
| Hairline + perineum + throat | 706                                         | 76.3% (73.6–79)                      |
| Axilla + perineum + throat | 703                                         | 76% (73.2–78.7)                      |
| Hairline + nose + perineum | 698                                         | 75.5% (72.7–78.2)                    |
| Axilla + nose + perineum | 694                                         | 75% (72.2–77.8)                      |
| Axilla + groin + nose | 693                                         | 74.9% (72.1–77.7)                    |
| Hairline + nose + throat | 690                                         | 74.6% (71.8–77.4)                    |
| Axilla + nose + umbilicus | 604                                         | 65.3% (62.2–68.4)                    |
There were 925 patients found to be colonised with MRSA in total using the seven-site body screen gold standard. One in high-risk patients using rapid molecular methodologies. These molecular methods can be commercial or in-house (11–13). This would allow a faster turn-around time of results and allow appropriate decolonisation procedures and infection control measures to be implemented more rapidly. Although there have been studies showing the merits of universal decolonisation in intensive care units, these cannot be used for all areas of healthcare and it remains important to screen for MRSA in most institutions because of the risks of infection and transmission (14–19). Because screening is relatively expensive, it is important to determine the optimal and most cost-effective combination of anatomical sites that are tested. Our results indicate that a three-site screen, preferably including groin, nose, and throat will detect the vast majority of cases. The sensitivity of all other possible combinations is also presented. We feel that the results of this study will help healthcare managers to decide the optimal strategy that can be adopted to optimise the ability to detect MRSA colonisation and prevent infection and nosocomial transmission in this at-risk population.

### Table 1 (Continued)

| Anatomical site                  | No. of patients MRSA detected (total = 925) | Sensitivity (%) of detection (95% CI) |
|----------------------------------|---------------------------------------------|--------------------------------------|
| Axilla + throat + umbilicus      | 597                                         | 64.5% (61.5 - 67.6)                  |
| Hairline + throat + umbilicus    | 596                                         | 64.4% (61.3 - 67.5)                  |
| Groin + perineum + umbilicus     | 583                                         | 63% (59.9 - 66.1)                    |
| Groin + hairline + perineum      | 582                                         | 62.9% (59.8 - 66)                    |
| Axilla + hairline + throat       | 581                                         | 62.8% (59.7 - 65.9)                  |
| Hairline + nose + umbilicus      | 581                                         | 62.8% (59.7 - 65.9)                  |
| Axilla + groin + perineum        | 577                                         | 62.4% (59.2 - 65.5)                  |
| Groin + hairline + umbilicus     | 543                                         | 58.7% (55.5 - 61.9)                  |
| Axilla + hairline + nose         | 535                                         | 57.8% (54.6 - 61)                    |
| Axilla + groin + umbilicus       | 534                                         | 57.7% (54.5 - 60.9)                  |
| Axilla + groin + hairline        | 522                                         | 56.4% (53.2 - 59.6)                  |
| Axilla + perineum + umbilicus    | 516                                         | 55.8% (52.6 - 59)                    |
| Hairline + perineum + umbilicus  | 506                                         | 54.7% (51.5 - 57.9)                  |
| Axilla + hairline + perineum     | 481                                         | 52% (48.8 - 55.2)                    |
| Axilla + hairline + umbilicus    | 367                                         | 39.6% (36.5 - 42.8)                  |

There were 925 patients found to be colonised with MRSA in total using the seven-site body screen gold standard.

CI, confidence interval; MRSA, methicillin-resistant *Staphylococcus aureus.*

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