The blood and drainage cultures are suggested for early diagnosis of bloodstream infection (BSI), which are time consuming and laborious. Nasal colonization of bacteria is one of the modalities, occasionally can predict BSI. We hypothesized that nasal culture, as an accessible fluid may be helpful to predict future BSI in hemodialysis patients. The present prospective study evaluated 63 patients undergoing maintenance hemodialysis at the Pars hospital dialysis center, Tehran, Iran, from November 2015 until February 2016. Nasal fluid of patients were collected from the 1–cm internal anterior part of both nostrils of patients by a sterile swab and cultured in Trypticase soy agar. All patients were followed for three months for BSI. The results of first nasal fluid sample revealed that 33.3% in first sampling and 27.0% in sampling had positive nasal fluid culture. The type of bacteria in all positive cases was Staphylococcus aureus. The rate of BSI infection in the patients with positive and negative first nasal fluid culture was 9.5% and 2.4% respectively with no significant difference. We found also no significant association between BSI positivity and nasal culture results so that positive BSI was revealed in 5.9% of patients with positive nasal fluid culture and 4.3% in those with negative nasal fluid culture with no meaningful difference. None of the baseline variables including age and gender, underlying risk factor, access, or duration of dialysis was associated with BSI positivity. In hemodialysis patients, BSI may not be predicted by nasal fluid culture positivity.

Introduction

End stage renal disease (ESRD) unless in renal transplanted patients is associated with lifelong dependence of patients on maintenance dialysis, which can affect patients’ quality of life [1] and is associated with greater risk of morbidity and mortality [2]. One of the most important complications is bloodstream infection (BSI), which is the most common cause of hospitalization and the second most common cause of mortality in hemodialysis patients [3]. The overall incidence of BSI among those patients receiving maintenance hemodialysis due to ESRD remains high [4]. Infection acquisition in hemodialysis facilities is mainly due to non-compliance with health principles and in this regard, all guidelines have emphasized following the principles of infection control for preventing BSI in such vulnerable population [5]. At first attempt, medical directors and clinicians should aware the infection control practice guidelines particularly the ways to colonize and then spread the infection. First, it should be noted that about 70% of BSIs are related to vascular access especially central venous catheters [6,7]. These accesses are responsible for spreading a variety of bacterial and even viral infections [8].

In addition to the access–related infections, infection–related hospitalization of...
hemodialysis patients is associated with low albumin concentrations, immobility, and residence in a day care center [9]. In addition, *Staphylococcus aureus* in blood culture, and antibiotic resistance are important factors associated with greater risk of mortality and morbidity of BSI in hemodialysis patients [10]. Moreover, persistent nasal colonization with resistant *S. aureus* is suggested as an important risk factor associated with increased all-cause mortality rate [11]. Thus, nasal colonization by *S. aureus* in hemodialysis patients is considered as a risk factor associated with bacteremia and general infection, especially for infections of endogenous origin [12] and its eradication may decrease the infection rate [13]. Moreover, coagulase–negative staphylococci and diphteroids may be positive in one–fourth of patients’ nasal vestibules [14]. Therefore, routine nasal fluid culture can be considered an easy access route to predict and prevent future infections. In this regard, some earlier studies could find the evidences on this claim. As shown by Pujol et al. [15], nasal carriers of *S. aureus* are at increased risk for bacteremia induced by *S. aureus* as compared to non-carriers. Another evidence has been also published by Chow et al on utilizing topical and oral therapy for eradication of staphylococcal nasal leading decrease in bloodstream infections [16]. Therefore, we hypothesized that nasal fluid culture, as an accessible fluid may be able to predict which maintenance hemodialysis patients are prone to BSI.

**Methods**

**Study design**

This prospective study evaluated hemodialysis patients regarding the association between nasal colonization and BSI. The protocol of the study was approved by Ethics committee of Iran University of Medical Sciences. The study protocol and objectives were first explained to all participants and patients who were willing to participate in the study were recruited after signing the written informed consent. All study steps were designed and implemented based on principles of Helsinki’s Declaration.

Patients undergoing chronic hemodialysis at dialysis center of Pars hospital, Tehran, Iran, from November 2015 until February 2016, and were currently using no kinds of antibiotics or immunosuppressive medications were included into the study. All patients diagnosed to have cancer or those receiving antibiotics or immunosuppressive therapy during the study period were excluded from the study.

The collected information included patients’ demographics and the underlying disease including diabetes mellitus (DM), hypertension, lupus erythematosus, Chronic tubulointerstitial nephritis (TIN), focal segmental glomerulosclerosis (FSGS), polycystic kidneys, glomerulonephritis, and renal stone with and without UTI. Moreover, the type of hemodialysis access was recorded in all patients.

**Laboratory methods**

Swabs were taken from the anterior nares of the nose by the Medical Laboratory Technologist. The nasal culture was performed at a recent referring the hospital for hemodialysis (immediately after hemodialysis procedure). After transfer to the laboratory, swabs were placed on blood agar base plates (Biolife Italiana, Milano, Italy) supplemented with 5% defibrinated sheep blood (Darvash Co., Tehran, Iran) and incubated at 35°C ± 2 for 48 hours. Suspected colonies were identified as *Staphylococcus aureus* by colony morphology, catalase production, and DNase test. *S. aureus* isolates were screened for methicillin resistance (MRSA) following the Clinical Laboratory Standards Institute (CLSI) disk-diffusion method. Overnight cultures from sheep blood agar plates were plated on Mueller-Hinton agar and a 30 μg cefoxitin disk (surrogate test for oxacillin) was placed on the inoculated plate. Zone diameters were measured and recorded after 24 hour incubation at 35°C ± 2 as susceptible (greater than 22 mm) or resistant (less than 21 mm).
Sampling was performed by one trained nurse for all participants by the same protocol. The sampling was performed at the two time points as baseline as well as after a three-month follow-up time.

After the results of nasal fluid culture were proven, patients were divided into two groups of negative and positive culture results, and were both followed for 3 months regarding BSI. The results of nasal fluid cultures were then compared with the etiology of BSI in follow-up.

**Statistical analysis**

The collected data were reported by mean ± standard deviation (SD), and percentage. Kolmogorov–Smirnov test was used to assess the normal distribution of data, which showed normal distribution of all data. Thus, continuous variables were compared using T test and categorical variables using chi-square test. The association of variables were tested by odds ratio and 95% confidence interval (CI). For the statistical analysis, the statistical software IBM SPSS Statistics for Windows, Version 21.0. Armonk, NY: IBM Corp. was used. P values less than 0.05 were considered statistically significant.

**Results**

A total of 63 patients entered analysis. The mean age of participants as was 62.79±16.18 (range: 24–86) years; 32 cases (43.8%) were female and 31 (42.5%) were male. The most common underlying disease was diabetes mellitus in 22 cases (34.9%) (Table 1).

Mean duration of hemodialysis was 54.89 months (median: 32, and range of 10–348) months. Hemodialysis access route was AVF or AVG in 44 patients (69.8%), and catheter in 19 patients (30.2%).

The results of first nasal fluid sample revealed that 21 patients (33.3%) in first sampling and 17 patients (27.0%) in second sampling had positive nasal fluid culture. The type of bacteria in all positive cases was *Staphylococcus aureus*. At 3-month following up, the BSI positivity was revealed to be 4.8%. The rate of BSI in the patients with positive and negative first nasal fluid culture was 9.5% and 2.4% respectively indicating no significant association (p = 0.256). The main source for BSI in both groups with and without BSI positivity was CVCs. In total, two out of three cases with positive BSI were infected with methicillin resistant strains. Similarly, we found no significant association between BSI positivity and nasal fluid culture results because of positive BSI revealed in 5.9% of patients with positive nasal fluid culture and 4.3% in those with negative nasal fluid culture with no meaningful difference (p = 0.800). In univariate analysis, none of the baseline variables including gender (p = 0.573), underlying risk factor (p = 0.555), access (p = 0.214), age (p = 0.353), or duration of dialysis (p = 0.682) was associated with BSI positivity (Table 2).

Figure 1 shows the frequency of bloodstream infection in the patients with nasal culture in two steps.

<table>
<thead>
<tr>
<th>Table 1: Baseline characteristics of study subjects (n = 63).</th>
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<tbody>
<tr>
<td>Male gender</td>
</tr>
<tr>
<td>Mean age, year</td>
</tr>
<tr>
<td>Mean duration of dialysis (months)</td>
</tr>
<tr>
<td>Route of access</td>
</tr>
<tr>
<td>AVF or AVG</td>
</tr>
<tr>
<td>Catheter</td>
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<tr>
<td>Causes of ESRD</td>
</tr>
<tr>
<td>Diabetes mellitus</td>
</tr>
<tr>
<td>Glomerulonephritis</td>
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<tr>
<td>Poly kidney disease</td>
</tr>
<tr>
<td>Renal stone</td>
</tr>
<tr>
<td>Hypertension</td>
</tr>
<tr>
<td>ATN</td>
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</tbody>
</table>
Can bloodstream infection be predicted by nasal culture in hemodialysis patients?

Discussion

As the main point in our study, we firstly suggested that the presence of positive nasal fluid culture in hemodialysis patients could predict which patients are prone to future positive bloodstream infection as an important adverse events of dialysis in such patients, but our suggestion failed and therefore, bloodstream infection could not be predicted by nasal fluid culture positivity, at least within three months after primary evaluation. However, some earlier studies had demonstrated our suggestion. As previously pointed, Pujol et al. [15], could show that nasal carriers of S. aureus are at increased risk for bacteremia. One of the main causes for this contradictory results may be sourced from our small sample sizes and thus presence of a low numbers of patients with positive BSI. Another issue may refer to the differences in applying different techniques for detection of bacterimia or to the difference in exclusion or eligibility criteria for inclusion of patients into the study.

The estimated prevalence of BSI in hemodialysis patients varies from country to country, and even from one hospital to another, but the reports of other studies, especially in developing countries indicate higher rates of BSI than the present study. In the United States, patients on hemodialysis had higher rates of bacteremia than peritoneal dialysis and 50% of patients experienced at least one episode of BSI [17]. Fram and colleagues reported an incidence of 42% for BSI in hemodialysis patients during six months (10) and other researchers reported the BSI rate at 0.52 per 1000 patient–days [18], which is much higher than the incidence of BSI in our study, which can reflect the appropriateness of preventive health measures taken in Pars hospital's

Table 2: Comparing baseline variables between the groups with and without BSI (n = 63).

<table>
<thead>
<tr>
<th>Item</th>
<th>BSI (+) (n = 3)</th>
<th>BSI (-) (n = 60)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Male gender</td>
<td>1 (33.3)</td>
<td>30 (50.0)</td>
<td>0.998</td>
</tr>
<tr>
<td>Mean age, year</td>
<td>71.33 ± 11.85</td>
<td>62.37 ± 16.33</td>
<td>0.319</td>
</tr>
<tr>
<td>Mean duration of dialysis (months)</td>
<td>39.00 ± 24.64</td>
<td>55.69 ± 69.42</td>
<td>0.379</td>
</tr>
<tr>
<td>Route of access</td>
<td>0</td>
<td></td>
<td>0.214</td>
</tr>
<tr>
<td>AVF or AVG</td>
<td>1 (33.3)</td>
<td>43 (71.7)</td>
<td></td>
</tr>
<tr>
<td>Catheter</td>
<td>2 (66.7)</td>
<td>17 (28.3)</td>
<td></td>
</tr>
<tr>
<td>History of diabetes mellitus</td>
<td>3 (100)</td>
<td>19 (31.7)</td>
<td>0.176</td>
</tr>
</tbody>
</table>

Figure 1: The frequency of bloodstream infection in the patients with nasal culture.
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In the present study, the results of the first nasal culture revealed positive culture in 33% of subjects, and in the second culture in nearly 27% of subjects. The discrepancy in the reported rate of positive nasal canal culture in the present study can be explained by the fact that only 20% of the patients who are carriers of *S. aureus* are persistent carriers, 30% are intermittent carriers, and 50% are non-carriers [22].

In present study, in all positive cases of the first nasal culture and nearly 95% of the second culture, the type of infection was *S. aureus*, which is similar to previous studies indicating as prevalent causes of bacteremia to be Gram positive micro-organism, responsible for 96 episodes (65%), with *S. aureus* as the most frequent (55%), followed by *S. epidermidis* (26%) and Gram-negative for 36 (23%), with *E. coli* (39%) as the most frequent agent, while 9.5% were polymicrobial infection [16]. Therefore, it can be concluded that Gram positive micro-organisms, especially *S. aureus* the most important organism, which call for adequate preventive and treatment measures.

Although the present prospective study addressed an important issue and prediction of future BSI by nasal fluid culture can help decrease the rate of mortality and morbidity of hemodialysis patients, the present study had several limitations, including small sample size. In addition, the study recruited patients from one center that could clearly show the prophylactic policy of that center, but it could limit the generalizability of the results. Therefore, it is suggested that future multicentric studies address this issue with larger sample size, and longer follow-up. Due to this fact that small sample size is a potential limitation of the study because of no more patients available, implementing further studies with larger sizes is recommended. Also, due to the considerable difference in the sensitivity of culture and the PCR technique in detection of bacterial strains, applying both techniques especially the PCR to identify bacterial genome is recommended in future studies.
Conclusion

In conclusion, the results of the present study indicated BSI in 3 out of 63 hemodialysis patients in a three–month’ follow-up, while nasal canal culture was not significantly associated with it, which was possibly due to the small number of BSI cases. According to our result, BSI may not be predicted by nasal fluid culture positivity particularly in hemodialysis patients.

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References

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