Bangladesh J Pharmacol 2016; 11: 940-943

A Journal of the Bangladesh Pharmacological Society (BDPS)

Journal homepage: www.banglajol.info

Abstracted/indexed in Academic Search Complete, Asia Journals Online, Bangladesh Journals Online, Biological Abstracts, BIOSIS Previews, CAB Abstracts, Current Abstracts, Directory of Open Access Journals, EMBASE/Excerpta Medica, Google Scholar, HINARI (WHO), International Pharmaceutical Abstracts, Open J-gate, Science Citation Index Expanded, SCOPUS and Social Sciences Citation Index; ISSN: 1991-0088

Pattern of skin bacteria in palmar arsenical keratosis

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Article Info

Received: 26 November 2016 Accepted: 10 December 2016 Available Online: 16 December 2016

DOI: 10.3329/bjp.v11i4.30465

Cite this article:

Khalil NB, Misbahuddin M, Sattar ANI. Pattern of skin bacteria in palmar arsenical keratosis. Bangladesh J Pharmacol. 2016; 11: 940-943.

Abstract

The aim of the present study was to examine the involvement of skin bacteria in the pathogenesis of keratosis. Skin swab of five different sites (palm, dorsum of the hand, chest, axilla and alar crease) from arsenic unexposed volunteer (n = 15), arsenic exposed (n = 15) and arsenicosis patients (n = 20) were collected for bacterial culture. The skin of arsenic unexposed volunteers contained Staphylococcus epidermidis and Bacillus spp. Chronic exposure to high concentration of arsenic through drinking water altered the skin bacterial pattern. The growth of Pseudomonas spp. increased in different sites of the skin. But in patient of arsenicosis, the skin of the palm shows the growth of Enterobacter spp.

Introduction

Arsenicosis is a clinical condition that develops following chronic ingestion of high concentration of arsenic through drinking water (Misbahuddin, 2015). Its clinical feature usually manifested first in the skin as black pigmentation (melanosis), rain-drop appearance in the skin of chest and back (leucomelanosis), and presence of nodule/cauliflower growth in the palm and/or sole (keratosis) (Caussy, 2005). One peculiarity of keratosis is that single palm or sole is not affected. Once affected, both palms and/or soles are affected.

Water loss through skin mainly occurs by sweating. In addition, there is transepidermal water loss by diffusion. Compared with that of the back, the transepidermal water loss is four times greater on the forehead, nine times greater on the back of the hand, and one hundred times greater through the palm (Baker and Kligman, 1967). Therefore, large amount of arsenic is excreted through these two sites.

Human skin surface contains around 1 trillion bacteria which survive from the secretory product of skin. There are variations in the type of bacteria (around 1,000 species) in different areas of the skin (Grice et al., 2009). Specific bacteria being associated with moist, dry and sebaceous microenvironments (Grice and Segre, 2013).

The relationship between the bacterial pattern in palm/ sole and the keratosis is not known.

Materials and Methods

This study was conducted in an arsenic endemic area (Comilla District, about 110 km from Dhaka) from October 2014 to January 2016. List of palmar arsenical keratosis patients (n = 105) and their addresses were collected from the Upazilla Health Complex of Muradnagar, Comilla (Figure 1). Detailed history and clinical examination were taken. Photographs of both palms were taken by smart phone. The diagnosis was confirmed by the presence of arsenic in water and nail. Among the 37 patients of moderate arsenical keratosis, 20 were randomly included in this study. Arsenic unexposed healthy volunteers (n=15) and arsenic exposed individuals (n=15; family member of the patient) were included for comparison.

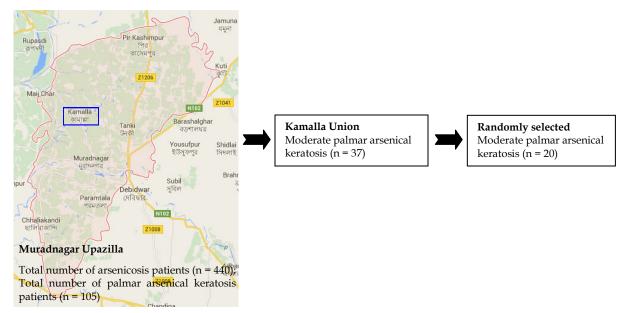


Figure 1: Selection of patients

Inclusion criteria are: a) age (18-60 years), b) sex (male and female), and c) subjects who voluntarily agreed to participate. Exclusion criteria are: a) age (below 18 or above 60 years), b) expectant/lactating mother, c) suffering from chronic diseases (tuberculosis, diabetes, hypertension, hepatitis, arthritis, systemic lupus erythromatosus), d) patient taking vitamin E/selenium, and e) suffering from other skin diseases (atopic dermatitis, psoriasis, eczema, etc).

The samples drinking water (50 mL) and nails (>200 mg) were collected in separate containers to estimate the amount of total arsenic by SDDC method (Bhuiyan et al., 2015). One drop of concentrated hydrochloric acid was added to the container for drinker water as preservative.

Skin swabs were collected from the right palm of the hand for bacterial identification. Other four sites were included (dorsum of the hand, front of the chest, axilla and alar crease) as control. At first rubbed the selected areas eight times by a sterile swab stick separately, and then moistened with a solution of nutrient broth and Tween 80 (10%). Swab stick was then placed in separate screw capped test tube containing 1 mL of the above mixed solution and finally brought to the Department of Microbiology and Immunology. On the next day, inoculation of bacteria was done on blood and chromogenic agar media and finally incubated at 35°C for 16-18 hours.

Bacteria were identified by observing the colony, Gram staining, microscopic examination and biochemical tests.

The study was approved by the University Institutional Review Board (Ref. No. 5102). All participants were informed about the nature and purpose of the study in local language (Bangla) and then written consent was taken from each participant.

Results

Staphylococcus epidermidis and Bacillus spp. were mainly found in the skin of healthy volunteers (Table I). There was no statistically significant difference in pattern of *S. epidermidis, Bacillus spp.* and *Enterococci spp.* between healthy volunteers, arsenic exposed volunteers and patients. When comparison was done between arsenic exposed volunteers and patients, there was a significant difference of *Enterobacter spp.* pattern in the palm of the hand.

There was statistically significant difference of *Pseudomonas spp.* pattern on skin of arsenic exposed volunteers and patients in comparison to healthy volunteers. Statistically significant changes were observed in the axilla and alar crease area of arsenic exposed volunteers and patients in comparison to healthy volunteers. In the dorsum of the hand, there was a significant change of *S. aureus* in patients in comparison to healthy volunteers.

Discussion

The present study shows that the normal human skin in an arsenic affected area contains *Staphylococcus epidermidis* and *Bacillus spp.* Chronic exposure to high concentration of arsenic through drinking water alters the skin bacterial pattern. The growth of *Pseudomonas spp.* Increases in different sites of the skin. But in patient

Table I

healthy volunteers	Isolation of different bacteria from the five anatomical sites of skin					
Palm						
Palm 11 7 14 Dorsum 10 8 17 Chest 10 8 13 Axilla 10 10 18 Axilla 10 10 18 Axilla 10 10 18 Alar crease 9 9 17 Palm 9 7 7 Dorsum 8 5 14 Chest 8 6 10 Axilla 6 7 8 Alar crease 7 8 7 Palm 4 4 5 Dorsum 2 5 10 0.04* Axilla 1 4 5 Alar crease 1 4 5 Alar crease 1 5 4 Dorsum 3 0 5 9 Palm 3 0 5 9 Chest 3	Area	healthy volunteers	volunteer		p value	
Dorsum	Staphylococcus epidermidis					
Chest	Palm	11	7	14		
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Axilla 10 10 18 18 17 18 18 18 19 19 17 18 18 18 18 18 18 18 18 18 18 18 18 18	Chest	10	8	13		
Alar crease 9 9 9 17			10	18		
Palm 9 7 7 7 Dorsum 8 5 5 14 Chest 8 6 6 10 Axilla 6 7 8 8 Alar crease 7 8 7 8 7 Palm 4 4 4 5 Dorsum 2 5 10 0.04* Chest 2 4 9 Chest 2 4 9 Axilla 1 4 5 Alar crease 1 5 4 Chest 2 5 10 0.04* Chest 2 4 9 Axilla 1 4 5 Alar crease 1 5 4 Dorsum 3 0 5 Chest 3 4 2 Chest 3 4 2 Chest 3 4 2 Chest 3 1 5 Chest 3 1 1 5 Chest 3 1 1 5 Chest 3 1 1 1 1 1 Alar crease 3 0 2 2 Interobacter spp. Palm 2 0 7 0.029* Dorsum 2 5 8 8 Chest 5 2 7 Axilla 6 1 7 Alar crease 1 1 1 5 Chest 5 2 7 Axilla 6 1 7 Axilla 6 1 7 Alar crease 1 1 1 5 Chest 5 2 7 Axilla 6 1 1 7 Alar crease 1 1 1 5 Chest 5 1 0 0.029* Chest 5 2 7 Chest 5 1 0 0.014* Chest 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Alar crease		9	17		
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Palm 2 0 7 0.029* Dorsum 2 5 8 Chest 5 2 7 Axilla 6 1 7 Alar crease 1 1 5 Pseudomonas spp. 2 7 0.015†, 0.003& Palm 1 10 13 0.015†, 0.003& Chest 1 12 16 0.005†, 0.001& Axilla 2 10 15 0.043†, 0.009& Scherichia coli 9 10 0.007†, 0.004& Palm 2 7 9 Dorsum 2 7 9 Dorsum 2 7 2 Chest 4 11 5 Axilla 3 13 10 0.038†; 0.043*		3	0	2		
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Axilla 2 10 15 0.043†, 0.009& Alar crease 0 9 10 0.007†, 0.004& Escherichia coli Palm 2 7 9 Dorsum 2 7 2 Chest 4 11 5 Axilla 3 13 10 0.038†; 0.043*	Dorsum	1	8	20	0.045^{\dagger} , $0.001^{\&}$	
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Sscherichia coli Palm 2 7 9 Dorsum 2 7 2 Chest 4 11 5 Axilla 3 13 10 0.038†; 0.043*	Axilla	2	10	15	0.043†, 0.009&	
Scherichia coli Palm 2 7 9 Dorsum 2 7 2 Chest 4 11 5 Axilla 3 13 10 0.038†; 0.043*	Alar crease	0	9	10	0.007†, 0.004&	
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Dorsum 2 7 2 Chest 4 11 5 Axilla 3 13 10 0.038†; 0.043*	Palm	2	7	9		
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Axilla 3 13 10 0.038†; 0.043*						
					0.038†; 0.043*	
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Samples were collected from the right palm, right dorsum of the hand, right axilla and right alar crease and front of the chest; McNemar's test was done for statistical analysis. †arsenic unexposed vs arsenic exposed; *arsenic exposed vs arsenicosis; &arsenic unexposed vs arsenicosis

of arsenicosis, the skin of the palm shows the growth of *Enterobacter spp*.

The most common is the *Staphylococcus epidermidis* which is occasionally pathogenic. The highest incidence of *S. aureus* is present in the groin and toe web (Aly and Maibach, 1977). Diphtheroids (lipophilic and nonlipophilic), together with coagulase-negative staphylococci and micrococci, are the dominant aerobic and facultative residents of the normal skin flora (Kligman, 1965).

One study shows that total microbial count was higher in arsenicosis (forearm> finger web > upper chest) than the control (Jyoti et al., 2011). *E. coli* and *Pseudomonas spp.* are the most predominant candidates. Other candidates like *Acinetobacter spp., Moraxella spp., Klebsiella spp.* and *Neisseria spp.* are found in arsenicosis patients.

Our focus is on the palm of arsenicosis. It is the site where keratosis develops in addition to sole (Caussy, 2005). Our data shows that *Enterobacter spp* is significantly present within it. As the patient of arsenicosis is exposed to high concentration of arsenic and its metabolites which might be involved in the growth of *Enterobacter spp*. But the growth of *Enterobacter spp* is not significantly increased in individual with arsenic exposure. We can predict that cauliflower growth may enhance the growth of *Enterobacter spp* in the palm.

It is apparent that chronic arsenic exposure increases the growth of *Pseudomonas spp* in all the five sites of skin examined. That is, arsenic and its metabolites create a good environment in the skin of arsenic exposed individual and patient for the growth of *Pseudomonas spp*.

Conclusion

Arsenic exposure increases the growth of *Pseudomonas spp.* in the skin surface whereas the palm of arsenical

keratosis shows, in addition, the presence of *Enterobacter spp*.

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