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Research Article

DISTRIBUTION OF MICROORGANISMS IN THE ORAL CAVITY OF CHILDREN WITH LEUKEMIA BEFORE AND AFTER CHEMOTHERAPY

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Abstract

The oral cavity is a unique environment, that gives a critical protective interface between the external and internal environment because it consists of oral mucosa which serves as a barrier to a large number of microbial species present in this moist and warm. So, the hard tissue breaks through the epithelial surface of the oral cavity due to invading by the different microorganisms (bacteria, yeasts), which regarded commensal microorganisms in the oral cavity. Impaired in the immune system in some cases as in patients with leukemia, these microorganisms become pathogens. Current study shows more prevalence for bacteria in oral cavity to patients before receive chemotherapy about 46 (53.4 %) and after take chemotherapy reach to 40 (46.5 %) additional to bacterial prevalence we observed yeasts in oral cavity for children's patients especially *Candida* sp. with increased number of *Candida* sp. After chemotherapy more than before therapy reach to 19 (59.4 %) and 13 (40.6 %) respectively.

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1. Introduction

Leukemia is a group of malignant hematologic diseases with mesenchymal (myeloid or lymphoid) organ originating from the bone marrow, which generates a high number of abnormal hematopoietic cells concerning their proliferation, differentiation, and programmed cell death (apoptosis) (Chandran *et al.*, 2015). The most common type of leukemia among children is (ALL) Acute lymphoblastic leukemia, about 75 % of all childhood leukemia and 25 % of all malignancy in childhood (Wang *et al.*, 2014; Babu *et al.*, 2016). Among some individuals, leukemia **Key words:** Oral cavity, Leukemia patients, *Candida* sp. and Commensal microorganisms.

first manifests in the oral cavity and the frequent manifestations that occur in leukemia patients include gingival bleeding, oral ulceration, gingival enlargement. candidiasis. and periodontitis (Chamilos et al., 2006; Garrett et al., 2007; Javed et al., 2012). Most children with leukemia suffering from neutropenia that means decreased in absolute neutrophils count less than 500 cell/ mm³ that play important role in regulating some microbial flora in the oral cavity including bacteria, Staphylococcus spp., Lactobacillus spp., Streptococcus spp., Klebsiella spp., Escherichia *Enterobacter* spp., *Citrobacter* coli, spp., Acinetobacter spp. and yeasts (Zuckermann et al., 2012). Candida sp. are commensal yeasts with more abundance and colonization in mucus membranes surface in the oral cavity for a patient

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with leukemia (Alnuaimi *et al.*, 2015). Generally, the first manifestation in patients with leukemia occurs in the oral cavity and observed the prevalence *of Candida* spp. and cause candidiasis, pseudomembrane *Candida*, and erythematous (Mushi *et al.*, 2016; Javed *et al.*, 2009; Scully, 2004). Previous studies found that oral bacteria are responsible for 25 % to 50 % of systemic infections in neutropenic patients (Wingard, 2001).

2. Materials and Methods

Patients and Specimens

This study was conducted at Basra children's specialty hospital, Basra province, Iraq from August 2016 to June 2017. Fifty specimens were taken from the oral cavity by using sterile cotton swabs from each patient before receiving chemotherapy and lately 4 - 6 weeks after receiving chemotherapy.

Identification Study

Bacteria and yeasts were identification by used traditional methods that including Macroscopic and Microscopic examination, Routine media (Blood agar, Chocolate agar, MacConkey agar of bacteria, Sabouraud's dextrose agar for fungi and Selective medium (CHROM agar Candida medium) of yeasts, all these media were prepared accordance the instructions for manufacture protocol of company procedure. In addition to biochemical test of bacteria (Catalase, Coagulase, Oxidase and API 20 E, API Staph, API 20 Strep from BioMerieux SA, France) and Yeast tests (Germ tube, API Candida from BioMerieux SA, France). Molecular studies including DNA Extraction for veasts by Commercial kit (from Favorgen company, Canada) was also included in the present research. PCR amplification was performed in a final volume of 25 µl. And each reaction consists of 2.5 µl of Green Master Mix, 0.5 µl MgCl₂, 0.5 µl of each forward (ITS1, F-5-TCC GTA GGT GAA CCT GCG G -3) and revers (ITS4, R-5- TCC TCC GCT TAT TGA TAT GC-3), 1.5 µl of template DNA, finally added about 9.5 µl DDH₂O to complete volume into 25 µl. An initial

denaturation step at 94 °C for 5 min then the second step was 25 cycles of denaturation at 94 °C for 30 seconds. Annealing at 56 °C for 45 seconds and extension at 72 °C for 1 min, the last final step of 72 °C for 7 min. After that, the amplified products were visualized by 0.8 % agarose gel electrophoresis about (0.2 gm) dissolved in 25 ml of TBE buffer, then stained with 0.2 μ l of Ethidium bromide and photographed (Mirhendi *et al.*, 2006). After the finished amplification process, the samples of PCR products were sent into sequenced by micro gen company in Korea.

Statistical analysis

Statistical analysis of data was carried out by using the t-test sample (paired samples test and paired samples correlation) with differences at P<0.05 which is considered to be statistically significant. This calculation was carried out according to the Statistical Package for Social Science (SPSS version 20) and the least significant difference at a level less than (0.05).

3. Results and Discussion

The oral cavity is a unique environment where contains multiple types of microorganism, whether bacteria or fungi that coexist with the state of ecological balance in healthy individuals (Dale et al., 2005; Kimball et al., 2006). Exposure diseases that cause weakness of the to immunodeficiency, leukemia, and other cancers will lead to the imbalance of the oral cavity environment, as there was a decrease in the richness and less diversity of oral microbiota compared to healthy controls (Wang et al., 2014). The patients who use chemotherapy that cause neutropenia and another antimicrobial agent like aminoglycoside agent (gentamycin, amikacin and netilmicin) which used as prophylaxis drug receive to the febrile patient with leukemia to reduce risk of bacterial that invading the body when granulocyte white blood cells(neutrophils) are less than 1500 cell/mm³ (Andrews et al., 2003). In the current study, the results for 50 specimens throat swabs which taken from 25 children with leukemia before and after chemotherapy as shown in Table - 1, where a total of 86 bacterial isolates were obtained 46 (53.4 %) were isolated before chemotherapy and 40 (46.5 %) were isolated after 4 - 6 weeks of chemotherapy, using the above-mentioned diagnostic methods for diagnosis of bacteria. This result agreed with Wang *et al.* (2014) who proposed that the state number and diversity of bacterial isolation before chemotherapy more than after chemotherapy. As for yeast species that coexisted with bacteria in the oral cavity were all

of *Candida* species 32 isolates was obtained 13 (40.6 %) before chemotherapy and about 19 (59.4 %) were isolated after 4 - 6 weeks of chemotherapy, since most children patients in both cases before and after chemotherapy suffering from neutropenia, so this result agreed with Netea *et al.* (2015) who showed the increase in invading opportunistic infections in immune suppression patients, as *Candida* spp. particularly *Candida albicans.*

Bacterial species	No. % of isolates		No. % of isolates	
	be	efore	after	
	cheme	otherapy	chemotherapy	
Viridans group of Streptococci	(7)	15.2%	(5)	12.5%
Streptococcus mitis	(4)	8.7%	(4)	10%
Streptococcus trails	(3)	6.5%	(3)	7.5%
Streptococcus spp.	(2)	4.4%	(2)	5%
Globicatelia sanguinis	(2)	4.4%	(1)	2.5%
Enterococcus spp.	(3)	6.5%	(4)	10%
Staphylococcus aureus	(2)	4.4%	(2)	5%
Staphylococcus xylosus	(3)	6.5%	(2)	5%
Staphylococcus spp.	(4)	8.7%	(3)	7.5%
Staphylococcus lugdunesis	(0)	0%	(1)	2.5%
Staphylococcus sciuri	(0)	0%	(2)	5%
Staphylococcus cohnii	(0)	0%	(1)	2.5%
Leuconostac spp.	(0)	0%	(1)	2.5%
Lactobacillus	(1)	2.2%	(0)	0%
Diphtheroid	(3)	6.5%	(1)	2.5%
Klebsiella pneumoniae	(2)	4.4%	(3)	7.5%
Klebsiella spp.	(3)	6.5%	(1)	2.5%
Escherichia coli	(1)	2.2%	(1)	2.5%
Pseudomonas oryzihabstans	(1)	2.2%	(0)	0%
Pseudomonas luteola	(1)	2.2%	(0)	0%
Enterobacter cloacae	(1)	2.2%	(1)	2.5%
Acinetobacter lwoffii	(2)	4.4%	(0)	0%
Ochrabacterium anthropi	(1)	2.2%	(0)	0%
Serratia marcesence	(0)	0%	(1)	2.5%
Enterococcus sakazakii	(0)	0%	(1)	2.5%
Total = 86 99.9 %	46	53.4%	40	46.5%

Table - 1: Distribution of Bacteria species in Oral cavity for Children with Leukemia in Study Group

	No. of isolates %	No. of isolates %
Candida Species	Before chemotherapy	After chemotherapy
Candida albicans	10 31.3%	16 50%
Candida africana	1 3.1%	2 6.3%
Candida krusei	1 3.1%	1 3.1%
Candida glabrata	1 3.1%	0 0%
Total=32 100%	13 40.6%	19 59.4%

Table - 2: Distribution of Candida Species in Oral Cavity for Children with Leukemia in Study Group

 Table - 3: Statistical Analysis of Cultured of Candida species for Children That Calculate Twice Before and After Chemotherapy

No.	Type sample	Correlation coefficient	P-value
1	<i>Candida</i> spp.	0.658	0.491





Figure - 1: Types of *Candida* species identified by API system (A) before chemotherapy and (B) after chemotherapy in a patient with leukemia



Figure - 2: Results of the gene sequence of Candida species and their strains before and after chemotherapy

	Befor	re chemotherapy			After chemotherapy
No.	Results of API	Results of Sequence	No.	Results of API	Results of Sequence
1	Candida albicans	Candida albicans strain h3b	2	Candida albicans	Candida albicans strain CBS:2735
3	Candida albicans	Candida albicans strainATCC18804	4	Trichosporon spp	Candida albicans strain h10a
5	Candida krusei	Pichia kudriavzevii CBS:5147	6	Candida krusei	Pichia kudriavzevii CBS:5147
7	Trichosporon spp	Candida albicans isolate 79209	8	Candida albicans	Candida albicans isolate LMICRO 142
9	Candida albicans	Candida aldicans strain d9b	10	Candida albicans	Candida albicans strain d9b
11	Candida albicans	Candida albicans strain ATCC 752	12	Candida albicans	Candida albicans strain ATCC 752
	1		13	Candida albicans	Candida albicans strain Hb13
14	Candida albicans	Candida albicans strain H245B	15	Candida albicans	Candida albicans strain ATCC 18804
16	Candida glabrata	Candida glabrata CBS:138	17	Candida albicans	Candida albicans strain ATCC 18804
	1		18	Candida albicans	Candida albicans strain ATCC 18804
19	Candida albicans	Candida albicans strain ATCC 18804	20	Candida albicans	Candida albicans strain ATCC 18804
	1		21	Candida albicans	Candida africana CBS 8781
22	Candida albicans	Candida albicans strain ATCC 18804	23	Candida albicans	Candida albicans strain ATCC 18804
	1		24	Candida albicans	Candida albicans strain ATCC 18804
25	Candida albicans	Candida albicans strain ATCC 18804	26	Candida albicans	Candida africana CBS 8781
27	Candida albicans	Candida africana CBS 8781	28	Candida albicans	Candida albicans strain ATCC 18804
29	Candida albicans	Candida albicans strain ATCC 18804	30	Candida albicans	Candida albicans strain ATCC 18804
	1		31	Candida albicans	Candida albicans strain ATCC 18804
	1		32	Candida albicans	Candida albicans strain ATCC 18804
	*Total No. for paties	at = 25 , % for patient before chemotherapy	= (13)52	1% , % for patient af	ter chemotherapy= (19)79%

Table – 4: Results of API Candida and Result Data for *Candida* Species Sequence Before and After Chemotherapy

* Total No. for isolate =32 , % for isolates before chemotherapy= (13) 40.6% , % for isolates after chemotherapy= (19)59.37%

Result of API Candida system that performed for 32 isolates that recovered from 25 patients. These include 13 (40.6 %) isolates before chemotherapy and 19 (59.4 %) isolates after chemotherapy showed a positive result. All the results of the API Candida system were compatible with sequence results proportion 84.4 % except 5 isolates about 15.6 % was not compatible with sequence result as shown in Table - 3. The result before chemotherapy showed that (12) 48 % patients with leukemia did not have an infection with *Candida* spp., and (10) 40 % patients were infected with Candida albicans and (1) 4 % patients for each Candida krusei, Candida glabrata, and Trichosporon spp. as shown in Figure - 1. While the result after chemotherapy showed that (8) 32 % for patients with leukemia did not have an infection with Candida sp., (15) 60 % for Candida albicans, (1) 4 % for Candida krusei and Trichosporon spp. as shown in Figure -2. Only the isolates 4 and 7 showed discrepancies between Api and the sequence results for the previous isolates were identified as Trichosporon spp. with API Candida system, while by using sequence techniques they identified as Candida albicans of strains h10a and 79209 respectively, both isolates have the same characteristic morphology on SDA and reaction on CHROM agar medium. Also, there were some differences among the isolates 21, 26, 27 respectively when API Candida system identified as Candida

albicans and sequence results was *Candida* africana CBS 8781, both isolates have the same characteristic features on SDA and CHROM agar medium. Also, in this study revealed disappeared *Candida glabrata* strain CBS:138 of the isolate (16) was not recovered before chemotherapy while *Candida albicans* strain ATCC 18804 of isolate (17) reported after chemotherapy for the same patient as shown in Table - 4, this difference may be due to chemotherapy effects on *Candida* spp. that agrees with (Teoh and Pavelka, 2016).

The patient's chance of survival was increased through clinicians administer better treatment decisions by used early diagnosis of invasive fungal infections such as candidiasis. The molecular biology methods are more confident than traditional phenotyping methods (Shokohi et al., 2010; Shokohi et al., 2011). For substance, Candida species were identified by Molecular biology methods. Such as multiplex PCR, standard PCR, PCR with species-specific probes, PCR-RFLP, real-time PCR (Mirhendi et al., 2008). The current study using the universal primers ITS1 and ITS4, which amplified between 510 - 870 bp of the ITS1 - 5.8 S-ITS2 region. As well as the result for Candida species sequence before and after chemotherapy in this study has been shown that Candida albicans reach 81.25 % including 26 strains were 15. Candida albicans strain ATCC 18804, 2 Candida albicans strain ATCC 752, 2

Candida albicans strain d9b and 1 number for each next strain H245B 1, 79209 1, LMICRO142, h10a, Hb13, h3b, and CBS:2735). While Candida africana reaches to (3) 9.30 % consists of one type of strain CBS 8781, Candida krusei (Pichia kudriavzevii) reach to (2) 6.30 % in one type of strain CBS:5147 and the last strain was Candida glabrata (1) 3.10 % CBS:138. This study showed that the proportion of infected children with Candida species increased from 52 % before to 76 % after chemotherapy. Candida albicans represent about 40 % - 60 % proportion of infected children with Candida species before and after chemotherapy respectively. This result is confirmed by the results sequence that refers to 81.25 % belong to Candida albicans with different strains, the ATCC 18804 strain is the most common among strains, the second species identified by the results sequence technique was Candida africana which reached 9.37 %. This species did not identify by the Api Candida system which was diagnosed as Candida albicans in isolates (21, 26, 27) respectively, while the results sequence for Candida krusei (teleomorph Pichia kudriavzevii) reach 6.25 %. The results of the current study revealed a high correlation between Candida species, as well as Candida albicans more prevalence from other non -Candida albicans in the oral cavity of children with leukemia before and after chemotherapy. This finding agrees with Subramaniam et al. (2008), Badiee et al. (2009), Mokaddas et al. (2010) and Darbandi *et al.* (2014) who demonstrated that the most often Candida spp. that is associated with oral lesions was Candida albicans, and other Candida spp. have also been isolated from the oral cavity. fungi were caused about 40 % to 50 % of fatal infections among cancer patients. This study shows that a correlation between Candida spp. before and after chemotherapy reach to 0.658 with no significance different P> 0.05.

4. Conclusion

The current study observed that the increased infection with opportunistic microorganisms in patients under immune suppression such as different species of bacteria

and *Candida* spp. particularly *Candida albicans*, result from the effect of chemotherapy in an environmental oral cavity so the most common species of *Candida* identified was *Candida albicans* with different strains and the most common strain was ATCC 18804. Other species were *Candida africana*, *Candida krusei* (*Pichia kudriavzevi*), and *Candida glabrata* respectively.

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