

شبكة المعلومات الجامعية التوثيق الإلكتروني والميكروفيلو

بسم الله الرحمن الرحيم





MONA MAGHRABY



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جامعة عين شمس التوثيق الإلكتروني والميكروفيلم قسم

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Nested Multiplex PCR Assay for Simultaneous Detection of Bacterial and Fungal Blood Stream Infections in Patients with Hematological Malignancies

Thesis

Submitted for Partial Fulfilment of MD Degree in Medical Microbiology and Ommunology

 \mathfrak{B} y

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Tist of Contents

Title Po	age No.
List of Tables	5
List of Figures	7
List of Abbreviations	8
Abstract	10
Introduction	i
Aim of the Work	14
Review of Literature	15
I. Hematological Malignancies	15
II. Neutropenic Fever in Hematological Malignancies	18
III. Blood-Stream Infections in Hematological Malignan	cies25
IV. Diagnosis of Blood Stream Infections	38
V. Management of Neutropenic Fever	62
Patients and Methods	70
Results	83
Discussion	98
Summary	108
Conclusion	110
Recommendations	111
References	112
Arabic Summary	

Tist of Tables

Table No	o. Title	Page No.
Table 1:	Classification of infection risk with NF .	21
Table 2:	Summary of LCBI categories	27
Table 3:	Summary of MBI-LCBI categories	28
Table 4:	Commercially available molecular assay the diagnosis of BSIs	•
Table 5:	New drugs for MDR bacteria and characteristics	
Table 6:	The main characteristics of escalation de-escalation strategy	
Table 7:	Sequence of primers utilized in the n multiplex PCR assay	
Table 8:	The clinical and laboratory data of the patients.	
Table 9:	Results of conventional blood cutechnique.	
Table 10:	Gender distribution among cases positive & negative blood cultures	
Table 11:	Age distribution among cases with posit negative blood cultures	
Table 12:	Distribution of isolated pathogens a positive blood cultures.	
Table 13:	Distribution of poly-microbial positive cultures among cases	
Table 14:	Detailed results of nested multiplex analysis of blood samples	

Tist of Tables cont...

Table No	o. Title	Page No.
Table 15:	Results of nested multiplex PCR compared to conventional blood culture.	•
Table 16:	Relation between the median CRP valuations & the results of conventional culture & the nested multiplex PCR ass	blood
Table 17:	Relation between the median ANC valuations & the results of conventional culture & the nested multiplex PCR ass	blood

Tist of Figures

Fig. No.	Title Page	No.
Figure 1:	Spectrum of Gram-positive bacteremia in patients with cancer	
Figure 2:	Time trend of bacterial etiology in neutropenic patients BSI	34
Figure 3:	Different molecular methods for the diagnosis of sepsis	
Figure 4:	Time to result of selected blood culture- dependent and blood culture-independent technologies for the diagnosis of sepsis	
Figure 5:	Hi-safe dual performance blood culture bottle	
Figure 6:	Results of conventional blood culture technique	85
Figure 7:	Gender distribution among cases with positive & negative blood cultures	86
Figure 8:	Age distribution among cases with positive & negative blood cultures	87
Figure 9:	Distribution of poly-microbial positive blood cultures among cases	89
Figure 10:	Detailed results of nested multiplex PCR analysis of blood samples.	90
Figure 11:	Agarose gel electrophoresis of the amplified products of the 1 st run PCR	91
Figure 12: Figure 13:	Amplification plots of positive samples Melt curve analysis	92
Figure 14:	8	95
Figure 15:	Range & median of CRP in positive & negative PCR.	95
Figure 16:	Range & median of ANC in positive & negative BC	

Tist of Abbreviations

Abb.	Full term
ANC	Absolute Neutrophil Count
	American Society of Clinical Oncology
BC	
	Blood Stream Infection
	Centers for Disease Control & Prevention
	Central Line Associated Blood Stream
02220	Infection
<i>CRP</i>	C-Reactive Protein
CoNS	Coagulase-Negative Staphylococci
DNA	Deoxyribonucleic Acid
	Extended Spectrum Beta Lactamase
	Fluorescence in Situ Hybridization
	Gram-Negative Bacilli
	Hematopoietic Cell Transplantation
	Hematological Malignancy
	Intensive Care Unit
	Infectious Disease Society of America
	Laboratory Confirmed Blood Stream Infection
MALDI-TOF MS	Matrix-Assisted Laser Desorption / Ionization Time of Flight Mass Spectrometry
<i>MBI</i>	Mucosal Barrier Injury
	Multi-Drug Resistant
	Methicillin-Resistant Staphylococcus
	aureus
MSSA	Methicillin-Sensitive Staphylococcus aureus

Tist of Abbreviations cont...

Abb.	Full term
NF	Neutropenic Fever
	National Healthcare Safety Network
PCR	Polymerase Chain Reaction
PMNLs	Poly-morpho-nuclear Leucocytes
<i>rDNA</i>	Ribosomal Deoxyribonucleic Acid
<i>TTP</i>	Time to Positivity
WBCs	White Blood Cells

Abstract

Introduction: Bloodstream Infections (BSIs) are a main cause of life-threatening complications among patients with cancer.

Methodology: This study aimed to identify microbial pathogens causing BSI in febrile neutropenic patients with hematologic malignancy and compare the results of conventional blood culture with a nested multiplex real time PCR assay done directly on whole blood samples. The nested multiplex PCR was based on *16S rDNA* and *18S rDNA* sequence-specific primers; hence, it allowed the identification of most species of bacteria and fungi.

Results: Forty adult patients with febrile neutropenia, admitted at Hematology ward of Ain Shams University Hospitals, were included in this study. Each patient was subjected to conventional blood culture and nested multiplex PCR. Blood culture was positive in 19 patients (47.5%). About 68.4% of the positive cultures were monomicrobial, while 31.6% were polymicrobial. A total number of 26 isolates were grown from positive cultures; *Staphylococcus aureus* was the most common (30.8%), followed by *Klebsiella pneumoniae* (19.2%). Regarding nested PCR, positive results were detected in 37/40 patients (92.5%) which was statistically significantly higher than that of blood culture. Eighteen samples that tested negative by culture were positive using the molecular approach. The agreement between the two approaches was 55%.

Conclusion: Nested multiplex real time PCR can be a promising tool in order to achieve rapid diagnosis in cancer patients clinically suspected of BSIs. Its utilization could affect the choice of antimicrobial treatment whether bacterial or fungal and, therefore avoid unnecessary use of antimicrobials.

Key words: Bloodstream Infections; Nested Multiplex PCR; Hematological Malignancy; Diagnosis.

Introduction

loodstream infections (BSIs) are life-threatening especially in patients with haematological malignancies (HMs). These patients are at increased risk of infections, not only because of the malignancy itself, but also because of neutropenia induced by intensive chemotherapy & its cytotoxic effect on the cells that line the gastrointestinal tract. Neutropenia in these patients is considered an important factor in assessment of infection risk and treatment choice. According to Infectious Diseases Society of America (IDSA) guidelines, the infection risk is described as high risk if neutropenia is prolonged; >7 days, and neutrophil count is $\leq 100 / \text{mm}^3$ (*Habip et al.*, 2014).

Bacteria are incriminated in most cases of these infections: coagulase-negative staphylococci (CoNS), Streptococcus viridans and Escherichia coli have been found to be the most commonly isolated pathogens. Risk factors for bacteraemia in patients with haematological malignancies were found to be use of a central venous catheter and neutropenia for more than 6 days. Fungi play an important role as well. In fact, fungal BSIs are rapidly increasing over the last decades mostly due to increased cases of immune suppression. Candida and Aspergillus spp. are the most commonly isolated fungal pathogens and Candida is considered the 4th most common cause of BSI in patients with haematological malignancies (Chen et al., 2017).



Mixed fungal/bacterial infections are also not uncommon: as combined Candida and bacterial bloodstream infections have been reported in as many as 23% of all episodes of candidaemia (Gosiewski et al., 2014b).

The early stages of BSI are associated with nonspecific clinical symptoms leading to difficult and delayed diagnosis. It's of utmost importance to rapidly diagnose such infections to swiftly initiate the proper antimicrobial therapy with the least side effects. The diagnostic "gold standard" nowadays for bloodstream infections is still blood culture (BC) techniques, performed either in the conventional specialized media or in automated culture systems. Blood culture is advantageously of low cost but hampered by the long period it takes for the results to properly appear, delaying the initiation of appropriate treatment. It's also unfortunately associated with low sensitivity as positive results are given in less than 50% of cases in spite of the clear signs of sepsis in patients. Moreover, subjecting the patients to empiric antimicrobial therapy before collection of blood samples for culture might inhibit the growth of microorganisms, especially if they are of fastidious nature (*Habip et al.*, 2014).

Molecular methods are a promising reliable alternative to BCs for the detection of these life-threatening infections. These techniques showed much higher sensitivity and less affection by the implementation of empirical antimicrobial therapy (Lamoth et al., 2010).