DEDICATION

To my family



The bacteriological work described in this thesis was carried out at the department of Microbiology, Faculty of Veterinary

Medicine University of Khartoum under supervision of Professor

Sulieman Mohamed El Sanousi.

The molecular part of the study was conducted at the Center of excellence, Department of Zoology, Faculty of Science, University of Khartoum under supervision of Dr. Mai Abd Elrahman Mohamed El Masri.

ACKNOWLEDGMENTS

Thanks to ALMIGHTY ALLAH for helping me to complete this work.

I would like to express my sincere gratitude and appreciation to my

supervisor Professor Sulieman Mohamed El Sanousi for his advice, interest, encouragement, leadership, criticism and patient guidance throughout this work. Thanks are also extended to my co-supervisor Dr. Mai Abd Elrahman Mohamed El Masri for leadership and guidance throughout this work particularly the molecular part and for providing me with primers; PCR kits and electrophoreses dye used in this work. Special regards are extended to Professor Mansour Faris Hussein and Dr Abdelkhalig Babiker for providing me the Lancefield grouping kits from Saudia Arabia. Thanks are extended to the staff of the Department of Microbiology Faculty of Veterinary Medicine, U. of K. for their cooperation and assistance with special regards to Mrs. Fawzia M. Hussein, Mrs. Mona Mutassim, Mr. Elshaarani Omer, Mr. A/Moniem Ramadan, Mr. Murwan Alamin, Mr. Elrasheed Omer, Mr. Hassan A. Hassan and Miss.

Mawahib Awad.

Special thanks are due to the staff of the Pathology Department, Faculty of Veterinary Medicine, U. of K. for providing a separate laboratory for me to do my work; in addition to some chemicals and reagents with special thanks to Professor A. A. Gameel, Professor A. M. Saad and Professor Afaf I. Abu Elgasim for continuous encouragement, and to Mr. S. Farah, Mr. H. Abdelwahab, and Mr. Y. Jubarra for technical help.

I'm also grateful to my colleagues Mr. M. Suleiman, Faculty of Medicine U. of Khartoum and Mr. Eltayib Abbas, Faculty of Vet.

Medicine, U. of Khartoum for providing me some reagents and glassware.

I wish to express my sincere gratitude to Professor Yousif Fadlalla for his unlimited help, cooperation and advice.

Thanks are extended to the administration of Khartoum E. N. T. Teaching Hospital, Saudi Maternal Hospital, Omdurman Teaching Hospital and Khartoum North Teaching Hospital and their medical staff, laboratory technicians and matrons for helping in samples collection; with special regards to my colleague Mrs. Manal Kamal. Thanks are also extended to Matron Suha Abu Elgasim (Saudi maternal hospital) for collection the vaginal swab samples.

Thanks are extended to Mrs. Rimaz Mohamed Ahmed and Miss.

Roua Mohamed Elnour who assist me in reading DNA concentration in the Nano-drop apparatus. Thanks are extended also to Miss Shaza El zain Hussein, Miss Salma Ahmed Ramadan and Miss Lymia Izzeldin for assisting me in loading DNA with special thanks to my colleague Mr. Ahmed Guzailan (Yemen) who assists also in loading DNA. Special thanks to Miss Amal Jaffar Taha for providing me with sterilized PCR tubes and tips.

Finally I would like to thank the administration of Khartoum
University farm from whom I obtained the sheep blood and cow serum
used in this work for preparing some types of media used with special
regard to my colleague Mr. M. Awadelkariem.

Abstract

A total of 500 samples were collected from patients with suspected Streptococcal infection. These included throat swabs (150 samples) collected from patients who attended Khartoum Ear Nose Throat (ENT) Teaching Hospital, suffering from sore throat, vaginal swabs (125 samples) collected from pregnant women (35-37 weeks of gestation) who attended the Saudi Maternal Hospital, Urine (125 samples) and blood (100 samples) samples collected from patients who attended Omdurman and Khartoum North Teaching Hospitals. All samples were collected from suspected patients regardless of their age.

The samples were cultured on 5% sheep blood agar. The type of haemolysis and the colony morphology were recorded as part of the diagnosis and the Gram's stain was done from overnight cultures using the standard technique.

Suspected Streptococcus isolates were subjected to different biochemical reactions to confirm the diagnosis. Commercial Lancefield grouping kits were then used to confirm the group identification.

Finally Polymerase Chain Reaction (PCR) was done to confirm the results of traditional diagnosis. Three types of primers were used: genus specific primer to confirm that the organism isolated was Streptococcus and the other two were used for sequencing in an attempt to identify isolates at the level of species

Antimicrobial sensitivity was done to find out the susceptibility of these organisms to antibiotics.

The results showed that group F Streptococci was isolated from swabs of patients suffering from sore throat (3.48%) beside group A Streptococci (96.52%).

Out of 125 samples of vaginal swabs, 4 samples (3.2%) were diagnosed as group B Streptococci, while 8 samples were isolated from 125 samples of urine and 5 samples were isolated from 100 samples of blood cultures.

Out of the 35 samples selected for sequencing, 28 samples were identified to the level of species. PCR results using genus specific primer indicated that ~ 100% of the isolates were of Streptococcus species. Sequencing results indicated that isolates were similar to the GenBank data base.

بسم الله الرحمن الرحيم

ملخص الاطروحة

تم جمع 500 عينة من المرضى منها مائة وخمسون مسحة حلقية من المرضى المترددين على مستشفى الآذن والآنف والحنجرة التعليمي بالخرطوم المصابين بالتهاب الحلق بغرض عزل ومعرفة نوع الباكتيريا المسببة للمرض من النوع المكورات السبحية (استربتوكوكس). ومنها مائة خمسة وعشرون مسحة مهبلية من النساء الحوامل المترددات على المستشفى السعودي بامدرمان في عمر 35-7 اسبوع من الحمل ومائة خمسة وعشرون عينة من البول ومائة عينة من الدم من المرضى المترددين على مستشفى امدرمان التعليمي و مستشفى بحرى التعليمي.

تم جمع هذا العدد من المرضى المترددين من مختلف الاعمار وتمت زراعة جميع العينات في الاوساط الملائمة لنمو هذا النوع من الباكتريا لمدة اربعة وعشرين ساعة ولمدة ثمانية واربعون ساعة اخرى في حالة عدم نموها قبل التخلص منها. وتم ملاحظة الشكل العام للباكتيريا ونوع التخثير الذي تفرزه على كرويات الدم الحمراء كجزء من التشخيص. وتم صبغ العينات بصبغة جرام وبعد ذلك تم اجراء كل الاختبارات الكيميائية التي تمكن تشخيص هذا النوع من الباكتيريا ومعرفة مجموعاتها. كما تم استخدام الامصال التجارية (لانسفيلد) التي تحدد ايضاً انواع المجموعات لهذا النوع من الباكتيريا. كما تم اختبار جميع العينات للمضادات الحيوية.

وقد اثبتت النتائج ان هناك مجموعات اخرى غير النوع المعروف للاصابة بالتهاب الحلق (أ) حيث ظهرت المجموعة (ف) ايضاً. كما تم اجراء التحليل الكهربى للعينات واختبار البلمرة المتسلسل كوسيلة تشخيصية حديثة للمرض بدلاً عن الوسائل التقليدية. واخيراً تمت قراءة شريط الحامض النووى للتأكد من التشخيص ومقارنة النتائج مع بنك الجينات.

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