

# بسم الله الرحمن الرحيم



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شبكة المعلومات الجامعية التوثيق الالكتروني والميكروفيلم





# جامعة عين شمس

التوثيق الإلكتروني والميكروفيلم

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### DNA PLOIDY IN RELATION TO DIFFERENT PROSTATIC CARCINOMA CLASSIFICATIONS

### Thesis

Submitted for partial fulfillment of Master degree In PATHOLOGY

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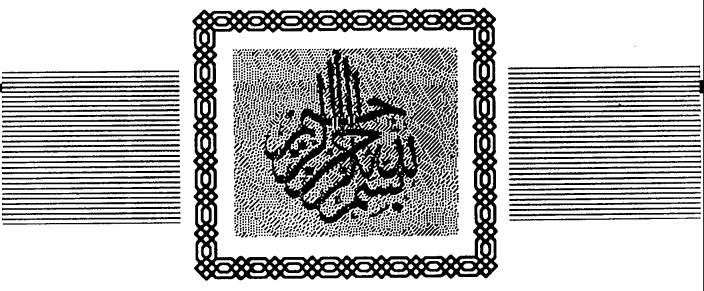
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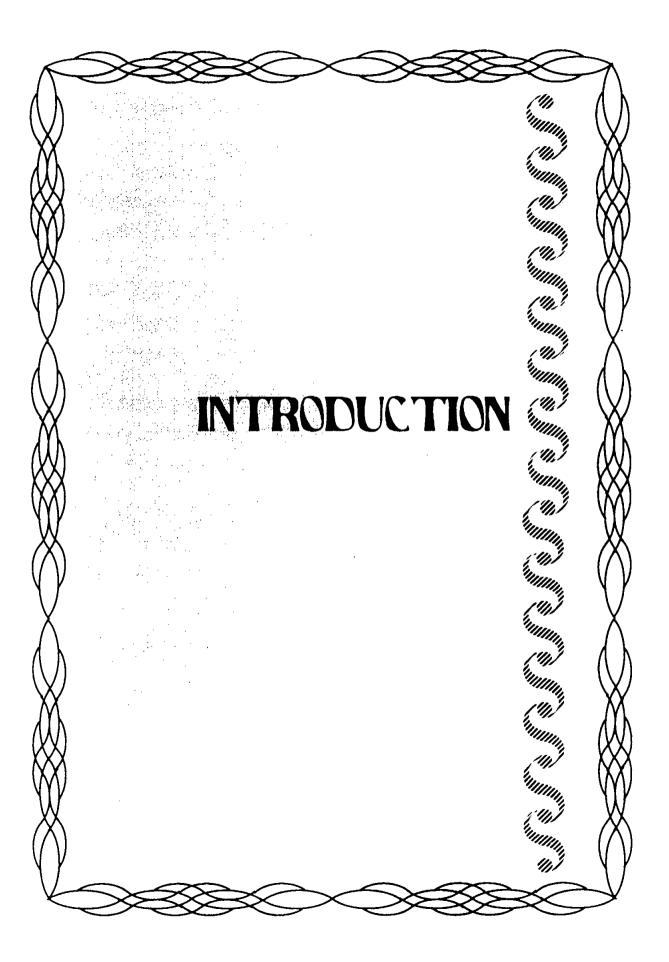
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# INTRODUCTION

In Egypt prostatic carcinoma represent 39.84% of male genital tract malignancy as reported in the Cancer Pathology Registry, National Cancer Institute (Mokhtar, 1991). In U.S.A., it is the second cause of cancer deaths in males. Histologically, most prostatic malignancies are adenocarcinoma of varying degrees of differentiation (Kumar et al., 1992)

Several grading systems have been described based on the degree of nuclear anaplasia (Mostofi, 1975), degree of glandular differentiation and growth pattern of the tumour (Gleason, 1977) or according to the percentage of gland formation (Brawn et al., 1982). In general there is a good correlation between the degree of differentiation and prognosis (Kumar et al., 1992).

Ploidy level, as determined from quantitative cytophotometric DNA analysis of Feulgen stained tumour cell nuclei, is another accurate method, that can be used to discriminate low from high grade malignant tumour. It was found that low grade malignant variant had DNA distribution similar to those of normal or benign prostatic cells, as most of the low grade

malignant nuclei had DNA values in the diploid or in the tetraploid region. On the other hand, most of the tumour cell nuclei of the high grade malignant tumours were outside these regions (aneuploid). However, the degree of prognostic information provided by ploidy level in relation to other prognostic parameters as histologic grade, is still somewhat unclear (Forsslund and Zetterberg, 1990).

# AIM OF THE WORK

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The aim of the present work is to study DNA content by single cell cytophotometry in various histopathologic grades of prostatic adenocarcinoma using different methods of classification. Also, it aims to evaluate the role of this technique in predicting the best method of classification that can give an idea about the ploidy state and the biologic behaviour of this malignancy.