## 中文摘要

目前研究發現,人類 hMYH 雙等位基因突變(biallelic germline) mutations,  $G: C \rightarrow T: A$ )與大腸直腸癌及其它癌症的產生有著密切的 關係。hMYH 為腺嘌呤醣基解? , 它是一種基因修復蛋白。本研究是 以 allele-specific polymerase chain reaction (AS-PCR)方法來進行 hMYH 基因中單核? 酸多態性(single nucleotide polymorphism, 簡稱 SNP)分 析。一開始先以質體 DNA(含 hMYH 基因)作為模板, 建立 AS-PCR 的反應條件,並且發現適當地調整 DNA 模板濃度、引子接合溫度或 改變 dNTP 混合比例,將可增加 AS-PCR 檢測 hMYH 基因中 SNP 的 可信度。此外,我們也證明了在對偶特異性引子的3'末端倒數第二位 置增加鹼基錯配,可提升 AS-PCR 方法的特異性。最後對 30 位已被 台中榮民總醫院診斷患有大腸直腸癌的患者,篩選 hMYH 基因上的突 變熱點 G382D、Y165C 及 V232F。雖然沒有患者的 hMYH 基因被檢 測出有突變熱點存在,但此實驗結果與先前由韓國科學家所報導之結 果相似。在大腸直腸癌病理學上,本實驗結果暗示 hMYH 突變發生率 會隨著種族的差異而有所不同。

關鍵字:對偶特異性聚合?連鎖反應(AS-PCR), hMYH 基因,單核? 酸多態性(SNP)、大腸直腸癌(colorectal cancer)

III

## Abstract

Biallelic germline mutations in hMYH, a gene for adenine DNA glycosylase, have been reported to be associated with colorectal cancer. In this study, we have modified the allele-specific polymerase chain reaction (AS-PCR) for single nucleotide polymorphism (SNP) genotyping. In order to make the AS-PCR more practical and to assess its ability in distinguishing the SNPs of *hMYH*, we employed the plasmid DNA containing *hMYH* as the DNA template to establish the AS-PCR conditions, and we found that the reliability of AS-PCR might be improved by optimizing the concentration of DNA template, the annealing temperature, as well as the ratio of dNTP. In addition, we demonstrated that the incorporation of an additional mismatch at the penultimate position near the 3' of allele specific primer could further enhance the specificity of AS-PCR. Finally, we screened for the mutations in hMYH, such as G382D, Y165C and V232F, from the genome of 30 Taiwanese patients who had been diagnosed as colorectal cancer by the Veterans General Hospital at Taichung. However, none of patients were detected with these reported mutations in hMYH, in which was similar to that reported for Korean population. Our results further indicated that the frequencies of hMYH mutations depended on the pathology of colorectal cancer and might also vary among different ethnic groups.

Keywords: allele-specific polymerase chain reaction (AS-PCR), *hMYH* gene, single nucleotide polymorphism (SNP), colorectal cancer