

微小 RNA16 抑制肺癌細胞增生之研究 = Micro RNA16 inhibit lung cancer cell proliferation

石文昭、蔡孟

E-mail: 9806537@mail.dyu.edu.tw

摘要

microRNAs是一群小片段的RNA，它的長度約18~24個核苷酸，不會被轉譯為蛋白質。生物體的運作伴隨而來的是一連串的調控作用，而microRNA更是這其中的要角，像是胚胎的發育、細胞分化、細胞周期的調控等等都可以發現microRNA參與其中。腫瘤的形成就是因為細胞在生物體中並沒有受到正常的調控，所以才會無限制的增生或是轉移；而近年來更陸續的發現乳癌、直腸癌、前列腺癌及神經膠母細胞瘤中都可以發現到microRNA的異常表現，但是在肺癌與microRNA之間的研究文獻則較少。因此本研究要分析與肺癌相關之microRNA，其中包含了miR-16、miR-24、miR-494等等不同的microRNA；進一步分析不同肺癌細胞株的結果顯示，若與正常支氣管上皮細胞相比，miR-16在肺癌細胞中的表現量往往都較正常細胞來的低。我們分別在CL1-5肺腺癌細胞中建立了兩個miR-16表現系統：pSilencerTM3.1-miR16與pcDNATM6.2-miR16。透過已建立完成的表現系統分析，我們發現miR-16大量表現的細胞株pSilencer3.1-miR16-A7、A19與pcDNA6.2-miR16-G16、G17細胞的生長速率會受到抑制達50%；但是對肺腺癌細胞的移動(migration)以及侵入(invasion)能力則不會有明顯的改變。接著透過二維電泳(2-D electrophoresis)的分析以及電腦軟體的預測，我們想找出在肺癌細胞中受到miR-16調控的基因；目前為止我們利用上述兩種方法找到了兩個可能的目標基因，分別是NYGGF4和ALDOA。研究microRNA與癌症將有助於我們瞭解癌症的發生機制及癌症的治療，甚至是作為一個分子標記與預防癌症的再復發。

關鍵詞：微小RNA；微小RNA16；肺癌；癌細胞侵入；癌轉移

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