

行政院國家科學委員會專題研究計畫 成果報告

以 siRNA 調降 Toll-like 接受器之表達可抑制以 minimally modified LDL 處理之內皮細胞 Monocyte Chemoattractant Protein-1 與 Interle

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執行期間：94 年 08 月 01 日至 95 年 07 月 31 日

執行單位：臺北醫學大學醫學系

計畫主持人：胡朝榮

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以 siRNA 調降 Toll-like 接受器之表達可抑制以 minimally modified LDL 處理之內皮細胞 Monocyte Chemoattractant Protein-1 與 Interleukin-8 表達

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單位名稱：北醫大醫學系神經學科

一、中文摘要

腦中風是台灣的第二死亡原因，腦中風的病因相當分歧，目前的趨勢顯示缺血性中風與出血性中風的比例逐年增加，因此缺血性中風應是腦中風最重要的一部份。同樣的，缺血性中風的病因也相當多，其中包括粥狀動脈硬化（大血管疾病）、小洞性中風（小血管疾病）、心因性、凝血病變、血管炎等，及其他不明病因。粥狀動脈硬化可引起腦中風、冠狀動脈疾病、週邊血管阻塞疾病等為缺血性中風最主要之病因。許多研究指出粥狀動脈硬化是一種慢性發炎反應，在血管內皮細胞受損後，許多發炎介質 (cytokines and chemokines) 均參與其中；吾人在稍早的研究指出阿司匹林 (aspirin)，一種普遍用於缺血性中風預防的藥物，可以抑制發炎介質及單核球對血管內皮細胞的粘黏、穿移。Toll-like receptor 4 (TLR4) 是一種細胞膜上的接受器可啟動固有的免疫反應 (innate immune responses)，以基因技術移除 TLR4 之老鼠較不會罹患粥狀動脈硬化；在白人族群中 TLR4 之基因多型性，Asp299Gly，被證實與粥狀動脈硬化相關，而吾人的研究指出華人雖很少有 Asp299Gly 基因多型性，但在第一內子的一個基因多型性的確與缺血性中風相關；因此可以推論 TLR4 應在粥狀動脈硬化扮演重要角色。流行病學研究發現高血脂症，特別是高低密度脂蛋白 (LDL) 為粥狀動脈硬化之危險因子；最近研究支持脂肪可引起發炎反應，進而引發粥狀動脈硬化。

本研究要以 mmLDL 處理血管內皮細胞引發發炎反應，以 siRNA 抑制 TLR4 之表達，並測量多發炎介質及單核球對血管內皮細胞的粘黏、穿移，並探討 NF- κ B 在此一路

徑之角色，希望可以了解 TLR4 在粥狀動脈硬化的分子機制，未來 TLR4 可作為治療粥狀動脈硬化的一個標第。目前本研究已經建立 siRNA 抑制 TLR4 表達之系統，未來可應用於功能性之評估。

關鍵詞：粥狀動脈硬化、缺血性中風、慢性發炎反應、低密度脂蛋白、分子機制、固有的免疫反應

Abstract

Stroke is the second leading cause of death in Taiwan. The causes of stroke are quite diverse. It's a trend that the ratio of cerebral infarction (CI) and cerebral hemorrhage (CH) is increasing. It implicates that CI plays the most important role in stroke. Many kinds of pathophysiology are attributed to CI, including atherosclerosis (large vessel diseases), lacunar infarction (small vessel diseases), cardiogenic origins, coagulopathy, vasculitis, et al and some of unknown mechanisms. Atherosclerosis resulting in stroke, coronary artery disease (CAD) and peripheral artery occlusion diseases (PAOD) is the major causes of CI. The compelling evidence shows that atherosclerosis is a chronic inflammatory process. Many cytokines and chemokines are involved after endothelial cell damage. In our previous report, we had provided the data the support that Aspirin, the most popular drug for stroke prevention, inhibited monocyte chemoattractant protein-1 (MCP-1) and interleukin-8 (IL-8) expression in TNF-alpha stimulated human umbilical vein endothelial cells. Toll-like receptor 4 (TLR4) is the key

receptor for initiating the innate immune responses. Animals without TLR4 were less susceptible to atherosclerosis even taking high cholesterol diet. There is a genetic polymorphism of TLR4 gene, Asp299Gly, associated with atherogenesis among Caucasian populations. In our another recent study, there is very rare Asp299Gly polymorphism in Chinese population but we still found a polymorphism at intron 1 associated with CI. Based on epidemiological researches, hyperlipidemia, especially high low-density-lipoprotein (LDL) strongly contributes to the atherogenesis. The recent studies have raised the mechanism that lipid plays a role in induction of inflammation, which is one of the most important part of atherogenesis.

In this study, we will down-regulate the TLR4 expression by siRNA and survey the MCP-1, IL-8 expression at mRNA and protein levels, and then test the effects on monocytes adhesion/migration function in the endothelial cells treated by minimally modified LDL (mmLDL). We also will explore the role of NF- κ B in this pathway. Hopefully, we can clarify the molecular mechanism of TLR4 on the atherogenesis and TLR4 might be a target to develop a novel therapy for atherosclerosis.

Key Words: Atherosclerosis, Toll-like receptor 4, MCP-1, IL-8, adhesion, migration, siRNA, mmLDL, endothelial cell, monocyte

二、緣由與目的

1. Stroke is the second leading cause of death in Taiwan. The complications of the survivors include disability in motor, sensory and even mental functions. Every episode of stroke induces huge impacts to the patient and his family. Stroke is not only a common neurological disease but also an urgent issue of public health in

Taiwan. Stroke patients are used to classified according to the causes, in terms of hemorrhage and ischemia [5]. Although the causes of stroke are very diverse, most of stroke patients are classified as ischemic stroke. [6]

- 2. Atherosclerosis resulting in cerebral or carotid arterial stenosis/occlusion plays the most important role in occurrence of ischemic stroke.** Atherosclerosis and its complications such as stroke, myocardial infarction and peripheral vascular disease, remain the major causes of morbidity and mortality in the world. The pathophysiology of atherosclerosis consists of consequent events, including endothelial cell injury, monocytes adhesion, migration, proliferation of smooth muscle cells in sub-endothelial space. [7] In our stroke registration and other series, large vessel disease, associated with atherosclerosis, is the leading cause of Ischemic stroke). [8]
- 3. Atherosclerosis is an inflammatory process.** Inflammatory cells infiltrate in the atheroma, the pathologic presentation of atherosclerosis. Cytokines, chemokines and adhesion molecules are involved in the atherosclerosis. [9] Studies have showed that chemokines and adhesion molecules are involved in causing atherosclerosis by promoting directed migration of inflammatory cells. [7]. There is compelling evidence showing atherosclerosis is a chronic inflammatory disease. Inflammatory processes may be potential targets of therapy in preventing or treating atherosclerosis and its complications. [10]
- 4. Toll-like receptor 4 (TLR4) is one of toll-like receptors (TLRs) family. TLRs are pattern-recognition receptors and they initiate innate immune responses**

after interaction with pattern-specific ligands. [11] Endothelial cells could be activated by microbial ligands through TLRs with resultant expression of inflammatory mediators. [12] (Figure 1) TLR4 was found in human atherosclerotic coronary arteries by immunohistochemistry detection in plaque and adventitia. [13] Genetically knockout mice studies document TLR4 is contributed to atherogenesis via NF-kB signal pathway. [14]

TLR4 genetic polymorphisms are associated with atherogenesis. [8,15] These findings implicate the important role of TLR4 in atherogenesis.

5. **Oxidized LDL (oxLDL) and minimally modified LDL (mmLDL) induce atherosclerosis.** Hyperlipidemia, especially high serum level of LDL, is a risk factor for atherosclerosis. [16] OxLDL might alter TLR4 expression and has been linked to innate immune responses which affect the down-stream pathway by activating NF-kB. This is a relevant mechanism for LDL associated with chronic inflammation and atherosclerosis. [17]
6. **Minimally modified LDL (mmLDL) activates NF-kB pathway and is involved in pathophysiology of atherosclerosis.** MmLDL is a ligand of CD14, which contains no down stream signal pathway but MmLDL, similar to LPS, is a ligand of CD14, which is closely linked to TLR4. The signals from CD14 would be transduced to TLR4 and then affect the NF-kB pathway. [18] (Figure 1) These findings implicate mmLDL could induce an inflammatory response by the CD14-TLR4-NF-kB-inflammation pathway.

7. **siRNA can work in animals by intravenous (IV) administration.** There have been a few reports documenting the systemic (IV) administration of siRNA can alter the gene expression in target organs. [19] These findings have shed light on the new approach of atherosclerosis. Our *in vitro* study might provide a piece of basic information for the further *in vivo* researched on siRNA therapy for atherosclerosis. [20]

8. The specific aims of this study are as below.

Specific aim 1: siRNA down-regulating TLR4 expression in endothelial cells

Specific aim 2: Down-regulation of TLR4 suppresses IL-8, MCP-1 expression

Specific aim 3: Down-regulation of TLR4 suppresses IL-8, MCP-1 expression by inhibition of NF-kB activity

Specific aim 4: Down-regulation of TLR4 interfering the monocytes dhesion/migration to endothelial cells

三、研究方法

Cell culture

We purchased human umbilical vein endothelial cells (HUVECs) and endothelial cell growth medium (EGM-2, CC3156), from Clonetics (San Diego, CA), which contains 10% fetal bovine serum, hydrocortisone, hFGF-B, vEGF, R3-IGF-I, ascorbic acid, hEGF, GA-1000 and heparine. HUVEC which was used between passages 1 and 6, were maintained in EGM-2 medium in a humidified chamber containing 5% CO₂ at 37 °C. Cells were cultured in 6-well or 24-well plates until confluent, and were washed twice and incubated with serum-free medium for 12 h before different concentrations of ticlopidine (0.1, 1, 10, 30 µg/ml) were added. After being incubated for 12 h, cells were stimulated with TNF-α (10 ng/ml) (R&D Systems; Minneapolis, MN)

for 24 h for MCP-1 and IL-8 expression. After incubation, the supernatants were collected for ELISA analysis, and the cells were used for RNA isolation.

siRNA for Toll-like receptor 4

1. Design siRNA according to the TLR4 mRNA (BC025294)
2. Four siRNA sequences were chosen, the sequences as following
 siRNA 1: Sense strand:
 AUCCAGACAAUUGAAGAUGtt
 Antisense strand:
 CAUCUUCAAUUGUCUGGAUtt
 siRNA 2: Sense strand:
 GCCACCUCUCUACCUUAAUtt
 Antisense strand:
 AUUAAGGUAGAGAGGUGGCtt
 siRNA 3: Sense strand:
 ACCCAUCCAGAGUUUAGCtt
 Antisense strand:
 GCUAAACUCUGGAUGGGGUtt
 siRNA 4: Sense strand:
 CUUCCCCAUUGGACAUCUCtt
 Antisense strand:
 GAGAUGUCCAAUGGGGAAGtt
3. The siRNA was synthesized by in vitro transcription by *Silencer*TM siRNA Construction Kit (Ambion). The procedure will be performed according to the manufacture guide.
4. The siRNA was transfected to endothelial cells by *Silencer*TM siRNA Transfection Kit (Ambion). The procedure will be performed according to the manufacture guide.
5. The efficiency of siRNA will be determined by western blot analysis to assay the amount of TLR4 protein level at time point 24, 48 and 72 hours after

transfection.

6. Optimal time point will be chosen according to the western blot data.

RNA isolation and reverse transcription-polymerase chain reaction (RT-PCR) analysis

Total RNA was extracted from 1×10^6 cells according to manufacturer's instructions. Briefly, Trizol (1 ml) (Life Technologies, Grand Island, NY) was added to the plates to lyse the cells, and then the cells were transferred to the microfuge tube. Chloroform was added and total RNA was collected in the aqueous phase after centrifugation. Finally, RNA was precipitated by isopropyl alcohol, then washed and re-dissolved in DEPC treated water. The concentrations of RNA samples were measured with a spectrophotometer (GeneQuant II, Pharmacia Biotech) to determine the OD 260 and OD260/280 values. In total, 5 μ g of RNA samples was reverse-transcribed with an oligo-dT primer to synthesize first-strand cDNA consecutively under 65 °C for 15 min, 25 °C for 10 min, 42 °C for 60 min, 95 °C for 10 min, using SuperScript reverse transcriptase (Life Technologies, Grand Island, NY) and followed by storage of products at 4 °C²⁴.

四、結果及討論

(1) 結果

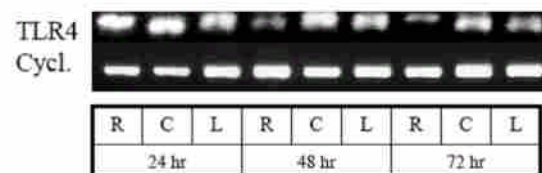


Figure 1. RT-PCR of TLR4. The mRNA of TLR4 was down-regulated by siRNA in 48 and 72 hours. R=siRNA, C=control, L=lipofitamin, Cycl.=cyclophyline (as internal control)

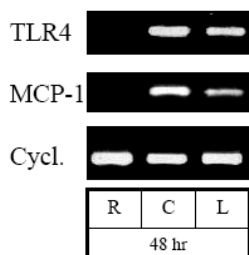


Figure 2. RT-PCR of TLR4 and MCP-1. MRNA of both TLR4 and MCP-1 is down-regulated by siRNA at 48 hour. R=siNNA, C=control, L=lipofitamin, Cycl.=cyclophyline (as internal control)

(2) 討論

At present, we have established an *in vitro* system of cerebral endothelial cells for down-regulation of TLR4. Suppression of TLR4 might result in decrease of MCP-1 expression. We will use this system to investigate the consequences, especially on the adhesion function, after down-regulation of TLR4.

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