

The Effect of Morelloflavone on HMG-CoA Reductase Activity

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| ชื่อวิทยานิพนธ์ | ผลของ morelloflavone ต่อเอนไซม์ HMG-CoA reductase |
|-----------------|---|
| ผู้เขียน | นางสาวกูไอดา ต่วนสุหลง |
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บทคัดย่อ

ภาวะคอเลสเทอรอลในเลือดสูงผิดปกติเป็นบ่อเกิดสำคัญของโรคหลอดเลือด หัวใจซึ่งมีอุบัติการเพิ่มขึ้นอย่างต่อเนื่องและยังคงเป็นสาเหตุของการเสียชีวิตอันดับต้น ๆ ของ ประชากรในหลายประเทศ การรักษาด้วยยาลดไขมันในเลือดนั้น ถึงแม้จะให้ผลดี แต่มักเกิด อาการหรือภาวะข้างเคียงบางอย่างที่เป็นอันตรายต่อสุขภาพร่วมด้วย ดังนั้น จึงมีผู้พยายาม ด้นหาสารออกฤทธิ์จากธรรมชาติที่มีศักยภาพในการบำบัดความผิดปกติดังกล่าวซึ่งอาจให้ ผลข้างเคียงที่ไม่พึงประสงค์น้อยกว่ายาแผนปัจจุบันที่ใช้กันอยู่ในขณะนี้ จากการศึกษาที่ผ่าน มาพบว่า morelloflavone ซึ่งเป็นสารไบฟลาโวนอยด์จากใบมะพูดสามารถลดระดับไขมันใน เลือดของกระต่ายที่ได้รับอาหารผสมคอเลสเทอรอลในปริมาณมากติดต่อกันเป็นเวลานาน นอกจากนี้ ยังมีรายงานว่า สารฟลาโวนอยด์จากพืชบางชนิดสามารถยับยั้งการทำงานของ HMG-CoA reductase ซึ่งเป็นเอนไซม์ควบคุมวิถีการสังเคราะห์คอเลสเทอรอลในร่างกาย และ ยังเป็นเป้าหมายหลักในการออกฤทธิ์ของ statins ซึ่งเป็นกลุ่มยาลดไขมันในเลือดที่นิยมใช้กัน อย่างแพร่หลายที่สุดในโลก ดังนั้น เพื่อให้ทราบกลไกการออกฤทธิ์ลดระดับไขมันในเลือดของ morelloflavone จึงได้ศึกษาผลของสารไบฟลาโวนอยด์จังกล่าวต่อเอนไซม์ HMG-CoA reductase ทั้งในหลอดทดลองและในสัตว์ทดลอง

ในการทดสอบกับบริเวณเร่งของ HMG-CoA reductase ซึ่งเตรียมโดยถ่าย ฝากยีน plasmid pET 17b-CD-HMG เข้าสู่แบคทีเรีย *Escherichia coli* สายพันธุ์ BL21 นั้น เอนไซม์ที่ผลิตได้สามารถเร่งปฏิกิริยาในอัตราเร็วสูงสุด (V_{max}) เท่ากับ 5 ± 0.26 μ M NADPH/mg/min โดยมีค่าคงที่ของ Michaelis (K_m) สำหรับ HMG-CoA และ NADPH เป็น 74 ± 0.06 μ M และ 133 ± 0.21 μ M ตามลำดับ ผลการทดสอบกับ morelloflavone พบว่า สารสำคัญชนิดนี้ยับยั้งเอนไซม์แบบแข่งขันกับ HMG-CoA โดยมีค่าคงที่ของการยับยั้ง (K_i) เท่ากับ 80.87 ± 0.06 μ M แต่ยับยั้งแบบไม่แข่งขันกับ NADPH โดยมีค่า K_i เป็น 103 ± 0.07 μ M ในขณะที่ narigenin กับ luteolin ซึ่งเป็นฟลาโวนอยด์หน่วยย่อยของ morelloflavone ยับยั้งเอนไซม์แบบแข่งขันกับ HMG-CoA ด้วยค่า K_i ที่เท่ากัน คือ 83.58 ± 4.37 μ M และ 83.59 ± 0.94 μ M ตามลำดับ ซึ่งใกล้เคียงกับ morelloflavone และสารทั้งสองยังยับยั้งแบบไม่ แข่งขันกับ NADPH โดยมีค่า K_i เท่ากับ 182 ± 0.67 μ M และ 188 ± 0.14 μ M ตามลำดับ การที่ morelloflavone และหน่วยย่อยทั้งสองชนิดสามารถยั้บยั้งเอนไซม์ได้ทัดเทียมกัน แสดง ว่า แต่ละโมเลกุลของ morelloflavone อาจใช้หน่วยย่อยข้างใดข้างหนึ่งเข้าแย่งจับกับเอนไซม์ โดยคาดว่า ตำแหน่งจับที่เหมาะสมน่าจะเป็นวงแหวนฟีนอลิคเดี่ยว (ring B) ซึ่งมีลักษณะคล้าย กับของ HMG และ statin อีกทั้งเป็นโครงสร้างที่ naringenin และ luteolin มีร่วมกัน

สำหรับผลการทดสอบเบื้องต้นในสัตว์ทดลอง พบว่า หนูขาวใหญ่ที่มีภาวะ คอเลสเทอรอลในเลือดสูงจากการบริโภคอาหารไขมัน ซึ่งได้รับ morelloflavone ทุกวันใน ขนาด 0.01% ของน้ำหนักตัว มีการเปลี่ยนแปลงของระดับไขมันในเลือดภายใน 6 สัปดาห์ไม่ แตกต่างจากหนูที่ไม่ได้รับสารทดสอบ ในขณะที่หนูปกติซึ่งได้รับ morelloflavone กลับมีการ เพิ่มขึ้นของปริมาณคอเลสเทอรอลใน HDL (HDL-C) ซึ่งใช้บ่งซี้ระดับไลโปโปรตีนชนิด ดังกล่าวในเลือด นอกจากนี้ mRNA ของ HMG-CoA reductase และตัวรับ LDL ในตับของ หนูทุกกลุ่มที่ได้รับ morelloflavone ยังมีแนวโน้มถูกกระตุ้น ทั้งนี้ อาจจะเป็นผลตอบสนองต่อ กระบวนการสังเคราะห์คอเลสเทอรอลในเซลล์ที่ลดลงนั่นเอง

ผลจากการศึกษานี้แสดงให้เห็นว่า morelloflavone เป็นสารออกฤทธิ์จาก มะพูดที่สามารถยับยั้งเอนไซม์ HMG-CoA reductase ได้ดี ซึ่งควรค่าแก่การค้นคว้าวิจัย เพิ่มเติมเพื่อนำไปสู่การพัฒนาเป็นยาลดระดับคอเลสเทอรอลในเลือดต่อไป Thesis TitleEffect of Morelloflavone on HMG-CoA Reductase ActivityAuthorMiss Ku-Aida TuansulongMajor ProgramBiochemistryAcademic Year2009

ABSTRACT

Hypercholesterolemia is a leading cause in the development of atherosclerosis-associated cardiovascular disease which still occurs increasingly and accounts for the majority of mortality worldwide. Although normalization of plasma cholesterol or LDL concentrations through drug administration is nowadays the most effective way to decrease atherogenic risk, these synthetic lipid-lowering drugs have been shown to cause serious side effects in various clinical settings. As such, there are attempts to find anti-hyperlipidemia agents from natural sources possessing lesser adverse effects. In previous studies, morelloflavone, a biflavonoid abundant in Garcinia dulcis leaves, has exhibited its lipid-lowering property in high-cholesterol fed rabbits. However, a mechanism underlying such beneficial effect of this compound is not known. Several plant derived flavonoids have been reported for their inhibitory activities on HMG-CoA reductase, the key enzyme in the early step of cholesterol biosynthesis pathway and the main target of statins, the most prescribed cholesterol-lowering drugs. Therefore, to investigate whether morelloflavone would reduce blood cholesterol levels primarily through the inhibition of HMG-CoA reductase, morelloflavone from G. dulcis was then examined for its effect on this enzyme both in vitro and in vivo.

For the kinetics study, the catalytic domain of house mouse HMG-CoA reductase prepared by transforming plasmid pET 17b-CD-HMG to *Escherichia coli* of BL21 strain, catalyzed the reaction at a maximum velocity (V_{max}) of 5 ± 0.26 μ M NADPH/mg/min, with the Michaelis constant (K_m) values for HMG-CoA and NADPH calculated to be 74 ± 0.06 μ M and 133 ± 0.21 μ M, respectively. Morelloflavone inhibited the enzyme by competing with HMG-CoA resulting in the inhibition constant (K_i) of 80.87 ± 0.06 μ M but exhibited noncompetitive inhibition towards NADPH with K_i of 103 ± 0.07 μ M. In accordance with morelloflavone, both

flavonoid subunits, narigenin and luteolin, competed with HMG-CoA with similar K_i values of 83.58 ± 4.37 µM and 83.59 ± 0.94 µM, respectively. These isoflavone compounds were also noncompetitive with NADPH and gave K_i of 182 ± 0.67 µM and 188 ± 0.14 µM, respectively. Due to the finding that morelloflavone and its subunits were equally effective at inhibiting HMG-CoA reductase, it was likely that this biflavonoid would bind with the enzyme active site *via* either naringenin or luteolin subunits. The moiety favorable for such binding could be the phenolic structure (ring B), commonly existing in both flavonoids and having HMG-like feature.

In animal experiments, treatment with morelloflavone at a dose of 0.01% (w/w/day) for 6 weeks showed no effect on plasma lipids in diet-induced hypercholesterolemic rats. In normal rats receiving the compound, however, their HDL-cholesterol levels became elevated indicating the stimulation of circulating HDL particles in these animals. A tendency for enhanced hepatic mRNA levels of HMG-CoA reductase and LDL receptor in response to the suppression of cellular cholesterol synthesis was also detected upon morelloflavone administration.

The findings from this study thus indicate a potentiality of morelloflavone isolated from *G. dulcis* as an effective HMG-CoA reductase inhibitor that might be developed into a new hypocholesterolemic agent in the future.

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Ku-aida Tuansulong

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LIST OF ABBREVIATIONS

| cDNA | = | Complementary deoxynucleic acid |
|------------------|---|---|
| DMSO | = | Dimethyl sulfoxide |
| DTT | = | Dithiotreitol |
| EDTA | = | Ethylenediamine tetraacetic acid |
| E. coli | = | Escherichia coli |
| g | = | Gravity force |
| g | = | Gram |
| h | = | Hour |
| HDL-C | = | High density lipoprotein cholesterol |
| HMG-CoA | = | 3-Hydroxy-3-methylglutaryl coenzyme A |
| kDa | = | KiloDalton |
| K_m | = | Michalis Menten constant |
| LB broth | = | Luria-Bertani broth |
| Μ | = | Molar |
| min | = | Minute |
| ml | = | Milliliter |
| mM | = | Millimolar |
| NADPH | = | β -nicotinamide adenine dinucleotide |
| | | phosphate, reduced form |
| nm | = | Nanometer |
| OD | = | Optical density |
| pH | = | -Log hydrogen ion concentration |
| RT-PCR | = | Reverse transcriptase polymerase chain reaction |
| SDS-PAGE | = | Sodium dodecylsulfate polyacrylamide |
| | | gel electrophoresis |
| sec | = | Second |
| V _{max} | = | Maximum rate of reaction |
| % | = | Percent |
| °C | = | Degree Celsius |
| | | |

LIST OF ABBREVIATIONS (CONTINUED)

| μg | = | Microgram |
|-----|---|------------------|
| μl | = | Microliter |
| μm | = | Micrometer |
| v/v | = | Volume by volume |
| w/w | = | Weight by weight |

CHAPTER 1

INTRODUCTION

1.1 Background and Rationale

Epidemiological studies have indicated that high blood cholesterol or hypercholesterolemia confers high risk for atherosclerosis and occlusive related vascular disorders (Levy and Brink, 2005). According to the world health report 2002 by the World Health Organization (WHO), hypercholesterolemia causes 18% of cerebrovascular disease and 56% of ischemic heart disease among populations all over the world (WHO, 2002). Although therapeutic life style changes including lowfat diet, weight management, and increased physical activity are important in a regulation of blood cholesterol, but in many cases, pharmaceutical lipid regulating interventions are also required. HMG-CoA reductase or 3-hydroxy-3-methylglutaryl coenzyme A reductase (EC 1.1.1.34) catalyzes the rate-limiting step in cholesterol biosynthesis. Therefore, inhibition of this enzyme is the major target for lowering of blood cholesterol level. A class of HMG-CoA reductase inhibitors collectively called statins have been developed since the first compound was discovered by Endo and Kuroda in 1970s (Endo, 2004). They remain the drug of choice for blood cholesterol lowering in people with or at risk of cardiovascular disease until now. Although, these drugs exert health benefits, they also cause adverse effects in various clinical settings (Wagstaff et al., 2003). Therefore, a search for naturally occurring compounds as regulators of cholesterol metabolism has been focused. Flavonoids are a large group of natural polyphenolic compounds ubiquitous in the plant kingdom. They exert a broad range of biological and pharmacological activities (Havsteen, Flavonoids from various sources have been shown to cause significant 2002). reduction of hepatic HMG-CoA reductase in experimental animals (Lee et al., 1999; Choi et al., 2001; Bok et al., 2002; Ohtsuki et al., 2003) and to inhibit the activity of this enzyme in vitro (Sung et al., 2004).

Garcinia dulcis (Roxb.) Kurz, commonly known as Maphuut in Thai,

is a tropical fruit tree belonging to the Clusiaceae (Guttiferae) family that grows mainly in Southeast Asia (Yupwattanaphun *et. al.*, 2002). All parts of this plant have been reported to contain a biflavonoid, morelloflavone (5, 7, 4', 5", 7", 3", 4"'-heptahydroxy-[3, 8"]-flavonylflavanone) (Ansari *et al.*, 1976; Deachathai *et al.*, 2005; 2006; 2008). This compound possesses many remarkable biological activities such as anti-HIV (Lin *et al.*, 1997), anti-inflammation (Gil *et al.*, 1997), anti-oxidant (Hutadilok-Towatana *et al.*, 2007), anti-malaria (Ngouamegne *et al.*, 2008), and anti-tumor (Pang *et al.*, 2009). In the previous study, morelloflavone from *G. dulcis* leaf has been found to lower blood cholesterol level in diet-induced hypercholesterolemic rabbits (Decha-Dier *et al.*, 2008). Therefore, to elucidate the mechanism underlying that hypocholesterolemic action of morelloflavone, the current study was then undertaken both *in vitro* and *in vivo*. In Part 1, the inhibitory effect of morelloflavone on HMG-CoA reductase activity was determined. Its effect on hepatic HMG-CoA reductase expression and activity in hypercholesterolemic rats was also studied in second part.

1.2 Objectives

The general aim of this thesis is:

To investigate whether morelloflavone from *G. dulcis* could inhibit HMG-CoA reductase *in vitro* and *in vivo*.

The specific aims are:

1. To determine the effects of morelloflavone on HMG-CoA reductase kinetics in comparison with those of the enzyme inhibitors, by using HMG-CoA reductase produced from recombinant DNA as a study model.

2. To preliminarily investigate the regulatory effects of this compound on both mRNA expression and activity of hepatic HMG-CoA reductase in high-fat diet induced hypercholesterolemic rats.

1.3 Benefits Gained

Basically, *G. dulcis* or Ma-phuut will be more well-known and become more valuable. Some insights into the mechanism underlying hypolipidemic/antiatherosclerotic action of its chemical constituent, morelloflavone, will be obtained. Taken together, all results from the recent past and present studies will substantiate the potential of this natural compound to be medicinally applicable for the management of atherosclerosis associated hypercholesterolemia in the future.

The publication arisen from this thesis is:

Tuansulong, K., Hutadilok-Towatana, N., Sotthibhandhu, R., Mahabusarakam, W., and Pinkaew, D. (2009) Morelloflavone inhibits HMG-CoA reductase activity. In: Proceedings for the 2nd Biochemistry and Molecular Biology Conference, 7-8 May 2009, Khon Kaen, Thailand, pp. 62-66.



CHAPTER 2

REVIEW OF LITERATURE

2.1 Blood cholesterol

2.1.1 Hypercholesterolemia

Cholesterol is the most abundant steroid in our body. It serves as a membrane structural component and a precursor of adrenal and gonadal hormones, vitamin D, and bile acids. Over 70% of total cholesterol in individuals arises from endogenous biosynthesis mainly in liver and intestine, and the remainder is provided by the diet (Grundy, 1988). In blood stream, cholesterol resides within two kinds of lipoproteins: low density lipoprotein (LDL) which carries this lipid of both dietary and *de novo* synthesis origins from liver to peripheral tissues for cellular uptake and utilization, and high density lipoprotein (HDL) which transports cholesterol from tissues to liver for degradation and excretion. Since LDL has the cholesterol content about 45% whereas HDL contains only 18%, thus any changes in LDL level unquestionably impacts on the total amount of cholesterol in the circulation (Bettelheim and March, 1991). At the present time, a high LDL level together with a low HDL level is well accepted as a symptom of faulty cholesterol transport and also a warning sign for possible 'atherosclerosis' or the hardening and narrowing of the arteries, a major risk factor for cardiovascular diseases (CVD) (Myant, 1990). Therefore, the significance of any particular cholesterol level cannot be assessed without taking into account either the total cholesterol (TC)/HDL ratio or LDL/HDL ratio. According to the blood lipid guidelines shown below, an acceptable TC/HDL ratio for a low risk of CVD is 5 or less for men, and women is 4.5 or less. The ability of HDL to predict the development of coronary atherosclerosis has been estimated to be about four times greater than LDL (www.exrx.net/Testing/LDL&HDL.html).

| TC/HDL ratios | |
|---------------|---|
| Men | Women |
| <3.4 | <3.3 |
| 4.0 | 3.8 |
| 5.0 | 4.5 |
| 9.5 | 7.0 |
| >23 | >11 |
| | <i>Men</i> <3.4 4.0 5.0 9.5 |

| | LDL/H | DL ratios |
|-------------|-------|-----------|
| Risk Levels | Men | Women |
| Very low | 1.0 | 1.5 |
| Average | 3.6 | 3.2 |
| Moderate | 6.3 | 5.0 |
| High | 8.0 | 6.1 |

Source: www.exrx.net/Testing/LDL&HDL.html.

In certain cases, however, when the blood cholesterol level rises above 7-8 mmol/L or 240 mg/dL and it is known as hypercholesterolemia (Chait *et al.*, 1993). This abnormal condition is not a disease in itself but has a good correlation with various circulatory diseases (Levy and Brink, 2005). It plays a crucial part in atherogenesis, along with accompanying high blood pressure, which may lead to heart attack or stroke. Hypercholesterolemia has been estimated to cause nearly 8% of total deaths worldwide (World Health Organization, 2002).

2.1.2 Causes of hypercholesterolemia

Both the dietary and environmental factors determine individual's circulating cholesterol concentration. Based on data from a number of survey studies, the populations in western countries have their average blood cholesterol levels higher than those in Asia (Sung *et al.*, 2002). The high blood cholesterol among western people is in connection with their lifestyles and likely to be influenced by their dietary

intakes which are rich in saturated fats (Mirmiran *et al.*, 2008). In comparison to unsaturated fat-fed animals, the conversion of cholesterol to bile acids in those fed with saturated fat is lower (Bravo *et al.*, 1997). Down-regulation of LDL receptor by saturated fatty acids has also been evident (Bucci *et al.*, 1998).

High cholesterol levels can also run in families which is called 'familial hypercholesterolemia'. Familial hypercholesterolemia is an autosomal codominant inherited disorder of lipoprotein metabolism. It is clinically characterized by dramatically elevated levels of total and LDL-cholesterol in the circulation as well as the presence of tendon xanthomas and premature atherosclerosis (Artieda et al., 2005). The genetic basis of familial hypercholesterolemia resulted from the LDL receptor gene (LDLR) deficiency is well-described. A large array of mutations in LDLR with varying frequency in the population, leads to a lack of functional receptors for LDL on the cell surface and thus a reduced rate of plasma LDL As a result, the elevated LDL level causes hypercholesterolemia clearance. (Goldstein et al., 2001). Similarly, mutations in the gene encoding for apolipoprotein B (APOB) also reduce LDL clearance, resulting in the familial defective B. which is clinically indistinguishable apolipoprotein from familial hypercholesterolemia (Defesche et al., 1993; Boren et al., 2001). Recently, a third putative familial hypercholesterolemia locus located on chromosome 1, encoding proprotein convertase subtilisin/kexin 9 (PCSK9) was identified (Abifadel et al., 2003). The clinical phenotype caused by mutations in LDLR, APOB or PCSK9, characterized by elevated levels of plasma LDL-cholesterol (LDL-C) is nowadays referred to as autosomal dominant hypercholesterolemia (Alonso et al., 2009). Homozygosity for the apolipoprotein E allele which does not bind to the LDL receptor also has severe effects on familial hypercholesterolemia phenotype (Nagasaka et al., 2008).

2.1.3 Hypercholesterolemia treatments

The formation of foam cells induced by oxidative modification of LDL is the hall mark of atherosclerosis (Shafi *et al.*, 2000). Accumulation of LDL particles in the circulation confers higher risk for these lipoproteins to be oxidized and taken up

by macrophages to become foam cells. For this reason, LDL-C has been recognized as an important determinant of clinical atherosclerotic diseases and thus treatment for hypercholesterolemia is usually focused on management of high LDL-C levels. In some cases, LDL-C can be lowered without medication (nonpharmacological therapy), and in other cases, medication is necessary.

2.1.3.1 Nonpharmacological therapy

Standard nonpharmacological therapy consists primarily of modifying diet and lifestyle. This therapy may modestly reduce LDL-C, but is not likely to lower the LDL-C level more than about 30 mg/dl (Sloane *et al.*, 2007).

Therapeutic lifestyle features that may lower LDL-C levels as recommended by the National Cholesterol Education Program (Expert Panel on Detection, Evaluation, and Treatment of High Blood Cholesterol in Adults, 2001) include:

Diet change: Minimizing cholesterol and fat intake, especially saturated fat which raises cholesterol levels, together with increasing dietary fiber and plant stanols/sterols.

Weight management: Losing a modest amount of weight can reduce in LDL levels achieved through an improved diet.

Increased physical activity: Exercise can decrease LDL and increase HDL levels.

2.1.3.2 Pharmacological therapy (Medication)

Hypocholesterolemic agents are divided into five major classes.

1) Bile acid sequestrants

Bile acid sequestrants are cross-linked polymeric cationic gels (anionexchange resins) that bind to the negatively charged bile acids in the intestine. Because these sorbents are large-sized, they are not absorbed into the body and thus the bound bile acids are excreted with the stool (Stedronsky, 1994). Consequently, the liver's pool of bile acids is depleted, leading to increased conversion of cholesterol to bile acids in hepatocytes. Decline in hepatic cholesterol content stimulates the production of LDL receptors causing increased LDL clearance and a net decrease in blood cholesterol due to lower LDL-C level (Jain *et al.*, 2007). However, this effect is partially offset by the enhanced cholesterol synthesis caused by the up-regulation of HMG-CoA reductase (Jain *et al.*, 2007).

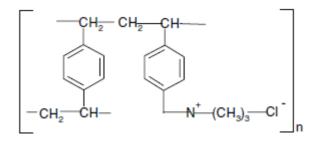


Figure 1 Structure of cholestyramine, a typical bile acid sequestrant (Jain *et al.*, 2007).

With over 30 years of clinical experience, the use of cationic polymers for sequestration of bile acids has become an established approach for treating elevated blood cholesterol levels (Zarras and Vogl, 1999). Cholestyramine, a quaternary amine, and colestipol, a mixture of tertiary and quaternary diamines, are the two widely used bile acid sequestrants (Huval *et al.*, 2001). Cholestyramine is a polymer of styrene and divinylbenzene with active sites formed from trimethylbenzylammonium groups. Colestipol is a copolymer of diethylenetriamine and chloro-2, 3-epoxypropane. These two first generation bile acid sequestrants, however, have low clinical potency, considered to be due to a competition between the polymeric sequestrant and active bile acid reuptake transporter system of the gastrointestinal tract (Huval *et al.*, 2004). Recently, a more potent bile acid sequestrant, colesevelam, has been launched (Donovan *et al.*, 2000). This drug has high binding capacity, strong binding strength, and selectivity towards bile acids in the presence of competing desorbing forces of the gastrointestinal tract (Huval *et al.*, 2004). Despite the appeal of their safety profiles, bile acid sequestering agents may contribute to calcium loss and therefore increase the risk for osteoporosis. Overtime deficiencies of vitamin A, D, E, and K may occur also (Jain *et al.*, 2007).

2) Niacin

Niacin (nicotinic acid) is a water-soluble vitamin of B type. It has been widely used clinically to regulate abnormalities in lipid/lipoprotein metabolism and in the treatment of atherosclerotic coronary heart disease (Meyers et al., 2004). It inhibits the lipolysis of triglycerides by hormone-sensitive lipase, which reduces transport of free fatty acids to liver and decreases hepatic triglyceride synthesis by inhibiting both the synthesis and esterification of fatty acids that increases apolipoprotein (apo) B degradation (Aithul et al., 1955 as cited by Jain et al., 2007). Reduction of triglyceride synthesis reduces hepatic VLDL production, which accounts for the reduced LDL levels. Niacin enhances lipoprotein lipase activity, which promotes the clearance of chylomicrons and VLDL-triglycerides. It also raises HDL-C levels by decreasing the fractional clearance of apo A-1 in HDL, rather than by enhancing HDL synthesis (Knopp et al., 1985). These unique beneficial effects of niacin on lipoproteins have been assumed to contribute to its anti-atherosclerotic properties. Clinical studies have demonstrated that niacin alone or in combination can slow or reverse the progression of atherosclerosis, and reduce cardiovascular event rates and total mortality in patients with hypercholesterolemia and established atherosclerotic cardiovascular disease (Meyers et al., 2004). In recent trial studies, a combination of niacin with simvastatin has shown more protective effects than simvastatin monotherapy in patients with elevated non-HDL-C by increasing more plasma HDL-C levels as well as lowering more triglyceride levels (Sanford and Curran, 2008).

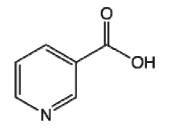
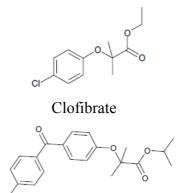


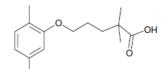
Figure 2 Structure of niacin (Jain *et al.*, 2007)

3) Fibrates (fibric acid derivatives) or lipoprotein lipase stimulants

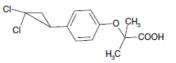
The primary effect of fibrates is a significant reduction in triglyceride levels. Additionally, increased HDL-C and decreased LDL-C levels may also be obtained (Birjmohun *et al.*, 2005). Major anti-hypertriglyceridemic actions of fibrates are increased lipolysis, increased hepatic fatty acid uptake, and reduced hepatic triglyceride production (Steals *et al.*, 1998). Fibrates seem to reduce plasma cholesterol to a variable extent from their ability to reduce intestinal cholesterol absorption, and more importantly, to increase the biliary output of cholesterol and phospholipids while decreasing the biliary secretion of bile acids by hepatic cholesterol 7 α -hydroxylase inhibition (Sirtori *et al.*, 1992; Catapano *et al.*, 1992; Bertolotti *et al.*, 1995). Since the discovery of clofibrate (ethyl ester of *p*chlorophenoxyisobutyrate) as an effective anti-lipidemic agent in the mid-1960s (Gennes *et al.*, 1965), four more fibrate derivatives, gemfibrozil, fenofibrate, ciprofibrate, and benzafibrate, are now prescribed.



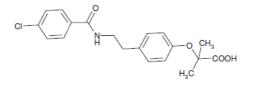
Fenofibrate



Gemfibrozil







Benzafibrate

Figure 3 Structure of the five different fibrates (Jain *et al.*, 2007).

These fibrate agents are agonists of the peroxisome proliferatoractivated receptor alpha (PPAR α) group of nuclear receptors, which control the expression of key elements in cholesterol transport and lipid metabolism, as well as endothelial function, vascular inflammation and remodeling (Steals *et al.*, 1998; Lock *et al.*, 1989; Duval *et al.*, 2007; Zandbergen and Plutzky, 2007). Their agonistic effects on PPAR α result in the peripheral lipolysis of triglyceride rich lipoproteins through the hepatic stimulation of the lipoprotein lipase gene, a reduction in apo C-III, resuting in a stimulation of the catabolism of triglyceride-rich lipoproteins, mainly VLDL and IDL (Shepherd, 1995). Fibrate-mediated PPAR α stimulation of apo A-I and apo A-II expression also increases HDL-cholesterol levels (Steals *et al.*, 1998). An increased hepatic expression of sterol regulatory element binding protein-2 (SREBP-2), responsible for higher HMG-CoA reductase and phosphatidate phosphohydrolase activities in gemfibrozil administered rats, has been reported (Roglans *et al.*, 2001)

4) Ezetimibe

Ezetimibe is also known as a cholesterol absorption inhibitor. This drug localizes at the brush border of the small intestine, where it inhibits the absorption of cholesterol from the diet. Specifically, ezetimibe appears to bind to a critical mediator of cholesterol absorption, the Niemann-Pick C1-Like 1 (NPC1L1) protein on the gastrointestinal tract epithelial cells (Garcia-Calvo et al., 2005) as well as in hepatocytes (Temel et al., 2007). In addition to this direct effect, decreased cholesterol absorption leads to an up-regulation of LDL-receptors on the cell surface and thus decreases LDL levels in the blood (Repa et al., 2005). In clinical practice, ezetimibe replaces bile acid sequestrants as second-line therapy for the treatment of familial hyperlipidemias and in most drug-intolerant patients. In comparison to the resins, ezetimibe shows significantly lower gastrointestinal side-effects (Sweeney et al., 2007; Florentin, 2008). It is able to reduce LDL-C by 15-25% from baseline (Mikhailidis et al., 2005; Capps, 2006). Besides monotherapy, ezetimibe is commonly used in combination with other agents (Toth and Davidson, 2005). Coadministration of ezetimibe and fenofibrate is more efficacious than fenofibrate alone.

It greatly reduces LDL-C, as well as significantly improves triglycerides, HDL-C total cholesterol, non-HDL-C, and apo B levels in patients with mixed hyperlipidemia (McKenney et al., 2006), while the addition of ezetimibe to niacin-based regimens lowers LDL-C level by 25% but does not change the level of HDL-C (Jelesoff, 2006). Its concurrent use with simvastatin represents the most common combined therapy, due to the fact that ezetimibe can add an extra 20% reduction in LDL-cholesterol to that produced by statins alone (Wierzbicki et al., 2005). The molecular mechanism for this enhanced efficacy has not been identified. Apo B-100 kinetics in pigs treated with ezetimibe plus simvastatin have revealed that such combination increases NPC1L1 expression in liver and intestine, consistent with increased SREBP2 expression, and also decreases VLDL and LDL apo B-100 concentrations through reduced VLDL production and up-regulation of LDL receptor-mediated LDL clearance (Telford et al., 2007). LDL receptor protein expression as well as LDL receptor and HMG-CoA reductase gene expression upon their combination therapy have been studied in healthy men. The results demonstrate that the co-administration abrogates the ezetimibe-induced increase in cholesterol synthesis and up-regulates the LDLR gene but not protein expression, possibly through a parallel up-regulation of PCSK9 expression (Gouni-Berthold et al., 2009).

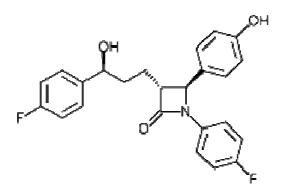


Figure 4 Structure of ezetimibe (http://www.biochemsoctrans.org/bst).

5) Statins or HMG-CoA reductase inhibitors

In the early 1970s, Dr. Akira Endo and his research team at Sankyo Co. Ltd. in Tokyo, isolated and studied various fungal metabolites from Penicillium *citrinium* for their hypocholesterolemic activities through the inhibition of HMG-CoA reductase which catalyzes the first committed step of the endogenous cholesterol biosynthetic pathway. The discovery of a potent inhibitor namely 'mevastatin' (formerly called ML-226B or compactin) in 1973 was a key step toward attaining an effective means of lowering blood cholesterol levels in humans (Endo, 2004). In 1976, lovastatin (mevinolin or monacolin K from Monascus ruber) produced from Aspergillus terreus, became the first statin to be commercially marketed. Since then, a number of statin derivatives have been developed. This class of agents, fermentation-derived and synthetic ones with varying potency, becomes the most commonly prescribed medications for blood cholesterol reduction in people with or at cardiovascular risk nowadays (Kapur *et al.*, 2008). Statins also have a wide range of 'pleiotropic effects' independent of cholesterol-lowering (Tandon et al., 2005). These include improvement of endothelial dysfunction, increased nitric oxide bioavailability, antioxidant properties, inhibition of inflammatory responses, stabilization of atherosclerotic plaques, recruit endothelial progenitor cell, а putative immunosuppressive activity, and inhibition of cardiac hypertrophy (Davignon, 2009). Such beneficial effects of statins have been known to be mediated through inhibition of small GTP-binding proteins (Turner et al., 2008; Rashid et al., 2009).

In 1980, Drs. Michael Brown and Joseph Goldstein at University of Texas Southwestern in Dallas, USA established that statins act by competitively inhibiting HMG-CoA reductase which is the rate-limiting enzyme of the mevalonate pathway of cholesterol synthesis in mammals (Brown and Goldstein, 1980). Due to the fact that they possess a side group that is structurally similar to HMG-CoA, statins then act by competitively inhibiting HMG-CoA reductase activity in the liver which results in decreased cholesterol synthesis (Istvan, 2003). Inside the liver cell, other enzymes of the protease class sense the reduced intracellular cholesterol content. In response, they cleave a group of proteins called 'membrane-bound sterol regulatory element binding proteins', which then migrate to the nucleus and up-regulate

production of various proteins and enzymes, including LDL receptor. This results in an enhanced clearance of LDL particles from the bloodstream and eventually lower circulating cholesterol level (Ma *et al.*, 1986).

Although statins are the most potent cholesterol-lowering drugs available, their treatments are associated with major side effects including increased liver enzymes and skeletal muscle pain and/or damage due to creatine kinase inhibition (Bybee *et al.*, 2004), and induced memory loss (Endres *et al.*, 1998). More serious but rare reactions include myositis and myopathy, with the potential for rhabdomyolysis leading to acute renal failure (Graham *et al.*, 2004). A common variation in the SLCO1B1 gene, which participates in the absorption of statins, has been shown to significantly increase the risk of myopathy (The research collaborative group, 2008). Among currently used statins, the risk of myopathy is lowest with pravastatin and fluvastatin probably because they are more hydrophilic and as a result have less muscle penetration (Pierno *et al.*, 1995). Lovastatin induces the expression of gene atrogin-1, which is believed to be responsible in promoting muscle fiber damage (Hanai *et al.*, 2007).

Besides inhibiting cholesterolgenesis, statin therapy also limits availability of mevalonate, the ubiquinone precursor, leading to lower levels of coenzyme Q10 (Ghirlanda *et al.*, 1993). The deficiency of coenzyme Q10 which is an essential cofactor in the mitochondrial electron transport and also a lipid-soluble antioxidant, has been implicated in several clinical disorders, including heart failure, hypertension, Parkinson's disease, and malignancy (Molyneux *et al.*, 2008). Despite initial concerns that statins might increase the risk of cancer, a systematic review and meta-analysis of statin trials and observational studies have concluded later that statins have no influence on cancer risk (Browning and Martin, 2007). Though, a 2005 trial has revealed that patients taking statins for over 5 years reduced their risk of colorectal cancer by 50% (Poynter *et al.*, 2005). A secondary analysis of data performed in large colorectal adenoma chemoprevention trials has found no association between statin use and reduced risk of recurrence of colorectal carcinomas (Wei *et al.*, 2005). As statins inhibit the acetyl CoA pathway of mevalonate, resulting in the decreased formation of higher isoprenoids which are necessary for post-

translational processing, anti-neoplastic effects of these drugs through inhibition of protein prenylation have been demonstrated in a variety of cancer cell lines (van de Donk *et al.*, 2002; Otsuki et al., 2004; Shellman *et al.*, 2005; Glynn *et al.*, 2008; Maksimova *et al.*, 2008; Nonaka *et al.*, 2009).

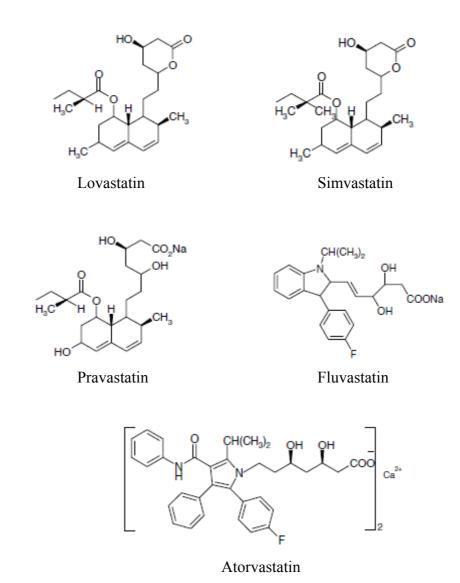


Figure 5 Structure of commonly used statins (Jain *et al.*, 2007).

2.2 HMG-CoA reductase

2.2.1 Characteristics

Basically, all body tissues containing nucleated cells are capable of synthesizing cholesterol but about two-third of endogenously produced cholesterol comes from liver alone (Williams *et al.*, 2002). The cholesterol biosynthesis pathway begins with conversion of acetyl CoA to HMG-CoA as shown in Figure 6. HMG-CoA reductase (3-hydroxy-3-methyl-glutaryl-CoA reductase, EC 1.1.1.88), which catalyzes the reduction of HMG-CoA to mevalonate, is the rate limiting enzyme of this pathway and provides feedback regulation of cholesterol synthesis in cells (Brown and Goldstein, 1980).

In humans, the gene for HMG-CoA reductase is located on the long arm of the fifth chromosome (5q13.3-14) (Cherng *et al.*, 2008). The enzyme has a molecular weight of about 97 kD and contains 887 amino acids in total. There are 39 out of 449 residues at the amino terminal region which act as a signal sequence for specific targeting to endoplasmic reticulum (ER) (Olender and Simoni, 1992). It is an intrinsic membrane-bound protein of the ER, containing eight transmembrane domains with the active site located in a long carboxyl terminal domain in the cytosol (Roitelman. *et al.*, 1992). The enzyme normally forms a tightly associated tetramer, but it is only the r of the molecule that is crucial for catalysis (Istvan, 2003).

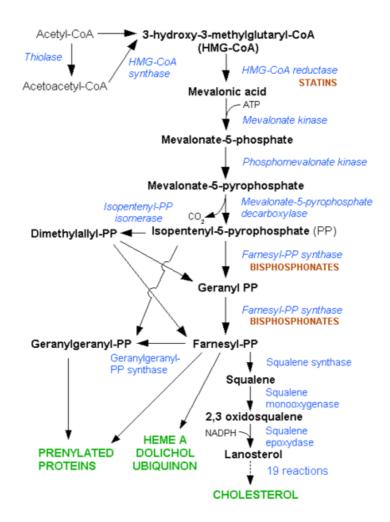


Figure 6 The mevalonate pathway of cholesterol synthesis (http://www.smbrower.com/mediawiki/images/8/8f/Cholesterol_Synthesis _Pathway.png).

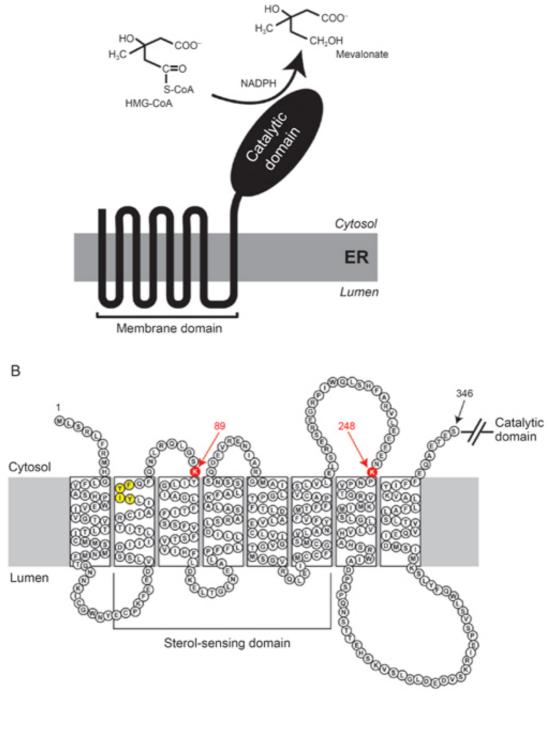
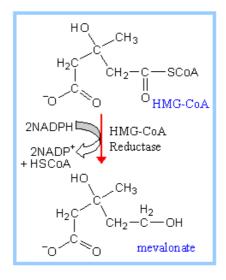


Figure 7 Structure of membrane-bound HMG-CoA reductase (Roitelman *et al.*, 1992)

А

HMG-CoA reductase is a NADPH dependent enzyme which catalyzes the reduction of HMG-CoA to mevalonate in two steps, as follows.



Source: www.rpi.edu/dept/bcbp/molbiochem/MBWeb/mb2/part1/cholesterol.htm.

This reaction involves the transfer of 4 electrons *via* 2 molecules of NADPH cofactor to a substrate that has been activated for reaction with the sulfhydryl (-SH) containing coenzyme A (CoASH). Some of the key amino acid residues at the active site of this allosteric enzyme that bind to HMG-CoA have been identified (Jain *et al.*, 2007).

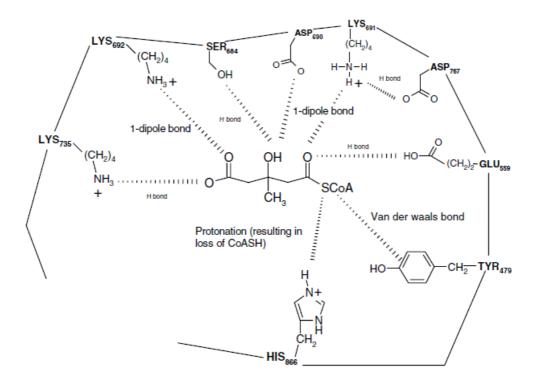


Figure 8 The substrate binding site of HMG-CoA reductase (Jain et al., 2007).

According to Jain and co-workers (2007), the binding interactions of HMG-CoA reductase and its substrate include:

• Lys735 (in cationic conjugate acid form), which anchors the substrate to the enzyme through an ion-ion bond with the C5 anionic carboxylate group of HMG-CoA.

• Lys692, also in the cationic form, stabilizes the carbonyl oxygen of which group through an ion-dipole bond.

• The C3-OH group of the substrate, which is stabilized by two residues, Ser684 and Asp690. Serine acts as a H-donor to form a H-bond with the oxygen of the OH group, and anionic Asp forms an ion-dipole bond with the alcoholic hydrogen.

• Lys691 (in cationic conjugate acid form), engages in an ion-dipole bond with the carbonyl oxygen of C1. It is this carbonyl group that gets reduced to

the primary alcohol through the action of the HMGR enzyme. Lys691 is found in the region (or domain) of the receptor referred to as the *cis* loop.

• An anionic Asp767, promotes this important ion-dipole interaction by forming its own ion-ion bond with Lys residue. This interaction stabilizes the action for interaction with the HMG-CoA substrate.

• Glu559 (in unionized acid form), also forms hydrogen bonds to the same C1 carbonyl oxygen. By forming these two important bonds, this carbonyl group is held tightly to the enzyme, and is properly oriented for the reduction to occur. In addition to these interactions, Tyr479 engages through Van der Waals bond with the adenine base of the CoA portion of the substrate, forming a kind of 'hydrophobic shield' that closes down the binding pocket for effective reduction by the NADPH cofactor. During the first NADPH reduction, the doubly bonded nitrogen of His866 (in cationic acid form) acts as a hydrogen donor to the sulfur atom of the thioester (SCoA), to liberate CoASH from the substrate. This, in turn, produces the mevaldehyde intermediate.

2.2.2 Regulation of HMG-CoA reductase

The liver is the main organ responsible for controlling cholesterol homeostasis in the body. In mammalian cells, the intracellular concentration of cholesterol is regulated by the balance between endogenous synthesis and exogenous uptake). Endogenous synthesis is subjected to feedback control by hepatic HMG-CoA reductase activity, while the exogenous supply is mainly controlled by the modulation of LDL receptor (Brown and Goldstein, 1980; 1986). The feedback control of HMG-CoA reductase and LDL receptor occurs through regulation of gene transcription or by regulation of protein level through translation or degradation (Reynolds *et al.*, 1984). Regulation of HMG-CoA reductase is multi-valent as shown in Figure 7, being achieved at several levels; transcription, translation, degradation and phosphorylation (Cherng *et al.*, 2008).

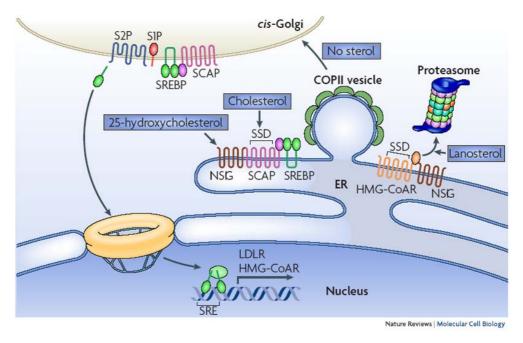


Figure 9 Regulation of intracellular cholesterol metabolism (http://www.nature.com/nrm/journal/v9/n2/fig_tab/nrm2336_F2.html).

1) Transcription of the gene

In 1993, Brown and Goldstein discovered that sterol regulatory element binding proteins (SREBPs) at ER membrane are responsible for the endproduct feedback regulation of gene transcription factors controlling intracellular cholesterol and lipid metabolism (Bengoechea-Alonso and Ericsson, 2007). The SREBPs are synthesized as inactive proteins that are inserted into ER membrane through two membrane-spanning domains. In ER, the C terminus of SREBP interacts with a protein called 'SCAP' (SREBP-cleavage-activating protein), which functions as a sterol sensor. In sterol-depleted cells, SCAP escorts SREBPs as SREBP-SCAP complex from ER to Golgi body, where they are cleaved by two membrane-associated proteases, the site 1 (S1P) and site 2 (S2P) proteases. The active fragments of the SREBPs are translocated to the nucleus, where they bind to the promoters of SREBP target genes, including HMG-CoA reductase and LDL receptor genes. Under conditions of ample sterol, the ER retention protein 'INSIG' prevents entry of the SREBP-SCAP complex to Golgi via COPII-coated vesicles. HMG-CoA reductase is also post-transcriptionally regulated by lanosterol, the first sterol intermediate in the

cholesterol pathway, with INSIG binding of the enzyme molecule leading to its proteasomal degradation (Bengoechea-Alonso and Ericsson, 2007).

2) Translation of mRNA

It has been believed that mevalonate, the product of HMG-CoA reductase, regulates the reductase translation through its conversion to a non-sterol substance by a mechanism that is still to be defined. This proposal comes from the findings that Chinese hamster ovary cells develop high levels of both the reductase protein and activity when they are deprived of mevalonate. In the presence of sterols, the mRNA is partially suppressed whereas the level of reductase remains relatively high. When mevalonate is added, the mRNA level does not decline further but the level of the enzyme declines owing to both reduced translation and accelerated degradation of the enzyme protein (Brown and Goldstein, 1988).

3) Degradation of the reductase

Rising levels of intracellular sterols increase the susceptibility of the reductase enzyme to proteolysis. Sterol-sensing domain (SSD) at helices 2-6 of transmembrane HMG-CoA reductase senses the higher levels of lanosterol through its C4-dimethyl group, leading to ubiquitination of Lys248. This ubiquitinyl residue then serves as a signal for proteolytic degradation of the enzyme (Cherng *et al.*, 2008).

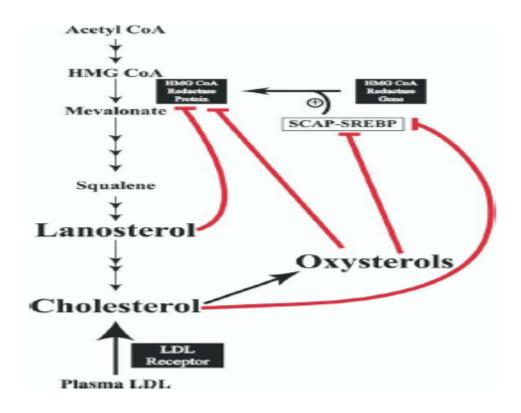


Figure 10 Model for sterol-mediated regulation of HMG CoA reductase (Song *et al.*, 2005)

Feedback control of cholesterol synthesis is also mediated in part by cholesterol and its oxy derivative (25-hydroxycholesterol) levels which induce binding of HMG CoA reductase to INSIG in ER. Such binding suppresses the activation of SREBPs, leads to ubiquitination, and accelerates proteasomal degradation of the reductase. Through these reactions, cholesterol synthesis is maintained such that important by-products of the synthetic pathway are continually provided, while at the same time avoiding the accumulation of cholesterol (Song *et al.*, 2005).

4) **Phosphorylation of the reductase**

Short-term regulation of the reductase is operated *via* phosphorylation and dephosphorylation processes. The principal enzymes involved

are AMP-activated kinase α (Hardie, 1992) and protein phosphatase 2A (Gaussin *et al.*, 1997). The phosphorylation site on HMG-CoA reductase has been identified at the C-terminal domain as Ser871 in rodents (Sato *et al.*, 1993) and Ser872 in humans (Istvan *et al.*, 2000). The phosphorylated form of HMG-CoA reductase is inactive, while dephosphorylation activates it (Hardie, 1992).

According to a 2007 review (King, 2007), HMG-CoA reductase is phosphorylated by AMP-activated protein kinase α , (AMPK α). AMPK itself is activated *via* phosphorylation by at least 2 enzymes. The primary kinase sensitive to rising AMP levels is LKB1. The second AMPK phosphorylating enzyme is calmodulin-dependent protein kinase kinase (CaMKK). CaMKK induces phosphorylation of AMPK in response to increasing intracellular Ca²⁺.

Additional regulation of HMG-CoA reductase occurs through the cAMP signaling pathway. An increase in cAMP leads to activation of cAMPdependent protein kinase (PKA). PKA phosphorylates phosphoprotein phosphatase inhibitor-1 (PPI-1) leading to an increase in PPI-1 activity. PPI-1 then inhibits the activity of numerous phosphatases including protein phosphatase 2C and HMG-CoA reductase phosphatase which remove phosphates from AMPK and HMG-CoA reductase, respectively. This maintains AMPK in the phosphorylated and active state, and HMG-CoA reductase in the phosphorylated and inactive state. Conversely to glucagon and epinephrine which increase cAMP production, insulin stimulates the removal of phosphates, and thereby activates the enzyme activity.

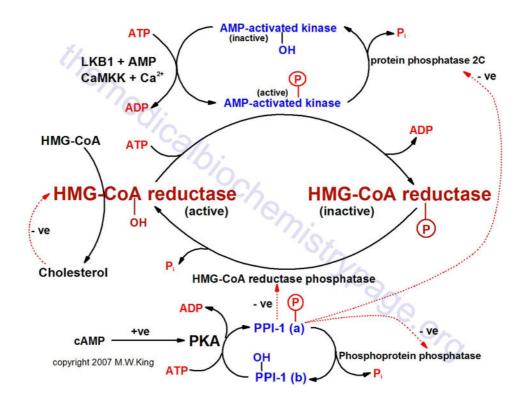


Figure 11 Regulation of HMG-CoA reductase by covalent modification as a result of phosphorylation and dephosphorylation (King, 2007).

2.2.3 Inhibitors of HMG-CoA reductase

Natural products from many sources have been reported to possess HMG-CoA reductase inhibitory properties.

In different animal studies, garlic supplemented diets decrease the hepatic cholesterol synthesis resulting from an inhibition of HMG-CoA reductase (Qureshi *et al.*, 1983a; 1983b; 1987). Organosulfur compounds present in garlic (*Allium sativum*), diallyl disulfide, allicin, allyl mercaptan, and ajoene, suppress HMG-CoA reductase in both rat hepatocytes and HepG2 cells (Gebhardt *et al.*, 1994; Gebhardt and Beck, 1996). Rai and co-workers have shown that diallyl disulfide analogs decrease the enzyme activity by reducing the mRNA level in hypercholesterolemic rat livers (Rai *et al.*, 2009). Qureshi and co-workers have found that α -tocotrienol, a vitamin E analog, from barley potently inhibits HMG-CoA reductase in both rat and chick livers (Qureshi *et al.*, 1986), whereas δ - and γ -

to cotrienols show a greater effect on the enzyme inhibition than α -to cotrienol in endothelial cells (Chao *et al.*, 2002). β -sitosterol which is a major phytosterol present in seed oil such as black cumin seed, cashew, rice bran etc., decreases hepatic HMG-CoA reductase activity in rats (Gylling and Miettinen, 2005). The reduction of HMG-CoA reductase in rat livers have been detected following intake of a polyunsaturated fatty acid, docosahexaenoic acid (DHA), which is abundant in marine fish oil (Froyland *et al.*, 1996: Shirai and Suzuki, 2007). Dietary α -linolenic acid also shows the similar effect (Ihara-Watanabe et al., 1999; 2000). Curcumin significantly lowers this liver enzyme in cholesterol fed rats (Murugan and Pari, 2006). Citrus flavonoids, hesperitin, hesperidin, naringenin and naringin, have been found to inhibit hepatic HMG-CoA reductase in several hypercholesterolemic animal models (Lee et al. 1999; Kim et al., 2003; 2004). Quercetin, a bioflavonoid found in the skins of apples and red onions, significantly lowers this liver enzyme in high cholesterol fed rats (Bok et al., 2002). Gugulipid from the resin of gugul tree (Commiphora mukul) also elicits its inhibitory effects against both liver and intestinal HMG-CoA reductase in cholesterolfed rats (Kumari and Augusti, 2007). Resveratrol, a phytoalexin primarily found in grapes, significantly inhibits the hepatic HMG-CoA reductase activity in apo Edeficient mice (Do et al., 2008) and attenuates the expression of this enzyme in hamsters (Cho et al., 2008). The hydroalcoholic extract of watercress (Nasturtium officinale) leaves attributes to a significantly decreased hepatic HMG CoA reductase activity in hypercholesterolemic rats (Bahramikia and Yazdanparast, 2008).

 β -carotene and its precursor 'lycopene' from higher plants, inhibit this enzyme at a post-translational level, as demonstrated in rat livers (Moreno *et al.*, 1995) and in macrophages (Fuhrman *et al.*, 1997). Artichoke (*Cynara scolymus*) leaf extracts also inhibits HMG-CoA reductase activity in rat hepatocytes, and its constituents, cynaroside and luteolin, have been identified to be mainly responsible for the inhibition (Gebhardt, 1998). A flavonoid compound 'astilbin' (Chen *et al.*, 2001) and tannin derivatives (Chang *et al.*, 2001) isolated from different traditional Chinese herbs as well as their crude extracts (Liu *et al.*, 2002), show the inhibitory effects on HMG-CoA reductase in Vero cells. Policosanol, a defined mixture of high molecular weight aliphatic alcohols purified from sugar cane (*Saccharum officinarum*) wax, suppresses an up-regulation of this enzyme in lipid-depleted cultured Vero fibroblasts (Menendez *et al.*, 2001). FR901512, a novel compound with unique tetraline ring structure, has been isolated from an agonomycetus fungus, and strongly inhibits the enzyme activity in human hepatoma HepG2 cells with IC₅₀ value of 1.0 nM (Hatori *et al.*, 2004). Soy isoflavone, genistein, and omega-3 fatty acids, eicosapentaenoic acid (EPA) and DHA, down-regulate HMG-CoA reductase activity, primarily through post-transcriptional effects in MCF-7 human breast cancer cells (Duncan *et al.*, 2005). Diosgenin, a natural furostanol saponin, decreases HMG-CoA reductase expression in human colon carcinoma cells (Raju and Bird, 2007). Green and black tea extracts also inhibit HMG-CoA reductase activity and thus decrease cholesterol synthesis in hepatoma cells (Singh *et al.*, 2008).

Extracts of medicinal plants such as *Typha augustifolia*, *Polygonum cuspidatum*, *Crataegus pinnatifida*, *Polygonum multiforum*, and *Pueraria thunbergiana*, as well as *Scutellariae radix*, *Coptidis rhizoma*, *Phellodendri cortex*, *Gardeniae fructus*, and *Rhei rhizoma*, contained in two oriental herbal medicines with anti-hyperlipidemic activity, 'orengedokuto' and 'daio-orengedokuto', show their high HMG-CoA reductase inhibitory activities *in vitro* (Lee and Choi., 1999; Kim *et al.* 2000; 2005). A stigmasterol from the mushroom *Pholiota adipose*, having an inhibitory activity of IC₅₀ 6.8 µg, has been isolated from its fruiting body (Yu *et al.*, 2007). Vitisin A and vitisin B from *Vitis vinifera* bark exhibit a remarkable inhibitory activity against HMG-CoA reductase *in vitro* with IC₅₀ values of 42.1 µM and 23.9 µM, respectively (Koo *et al.*, 2008).

In addition to statins originated from molds, the three isoflavones obtained from soybeans, diadzein, genistein, and glycitein, also act as competitive inhibitors of this enzyme with K_i values of 27.7, 49.5, and 94.7 μ M, respectively (Sung *et al.*, 2004b).

2.3 Garcinia dulcis

2.3.1 General description

Garcinia dulcis (Roxb.) Kurz is a sub-woody plant belonging to the Clusiaceae (Guttiferae) family that grows mainly in Southeast Asia (Yupwattanaphun *et. al.*, 2002). It is called Mundu in Indonesia and Malaysia, Baniti in the Philippines, and also known as Ma-phuut in Thailand. *G. dulcis* is a medium sized tree with short trunks and brown bark with white latex. Its leaves are opposite, lanceolate-shaped, 10-30 cm long and 3-5 cm wide. They are pale green when young and become dark green and shiny on the upper surface at maturity. The lower leaves are often hairy. The midrib is prominent with numerous veinlets arranged in parallel. The thick petiole is short being only 2 cm long. Flowers are borne in the axil. They are yellowish white with a sour smell. Fruits are globose, 5-8 cm wide with slightly pointed ends, often rather compressed and crowned by the persistent stigma. The fruit is soft with a thin skin and has a light yellow color, which turns to orange when ripe. The seeds are enveloped in an edible pulp of a darker color than the skin and have a pleasant taste (Subhadrabandhu, 2001).

2.3.2 Uses

The fruits can be eaten fresh, but they are sour and can be made into an excellent jam. Its leaves and seeds have been used in traditional medicine against lymphatitis, parotitis and struma in Indonesia (Kasahara and Henmi, 1986). Its stem bark has been used in Thai folk medicine as antiseptic, the fruit juice as anti-scurvy and expectorant for the relief of cough and sore throat, as mild laxative, and as decongestant, the root extract is also used as antipyretic and antitoxin (Wuttidhammavej, 1997; Bunyapraphatsara and Chokechaichareonporn, 1999).

Accumulated data from phytochemical studies have revealed that various parts of this plant contain, in abundant, a broad range of phenolic compounds (Ansari *et al.*, 1976; Harrison *et al.*, 1994; Iinuma *et al.*, 1996a, 1996b, 1996c; Ito *et al.*, 1997; Kosela *et al.*, 1999; 2000; Dechathai *et al.*, 2005; 2006; 2008). Some possess potent biological activities, for example, the five xanthones isolated from the

barks of *G. dulcis* have shown inhibitory effects on *Plasmodium falciparum* (Likhitwitayawuid *et al.*, 1998), the three prenylated paraxanthonoids from its fruits are cytotoxic against cancer cell lines (Soemiati *et al.*, 2004), and several phenolic compounds isolated from the seeds, fruits and flowers also act as anti-oxidants and anti-bacterial agents (Deachathai *et al.*, 2005; 2006; 2008; Hutadilok-Towatana *et al.*, 2007). Lately, anti-atherogenesis (Decha-Dier *et al.*, 2008), anti-restenosis (Pinkaew *et al.*, 2009) and anti-tumor angiogenesis (Pang *et al.*, 2009) effects of a biflavoniod, morelloflavone, isolated from *G. dulcis* leaves have been demonstrated.



Figure 12 Leaves, fruit, and flowers of Garcinia dulcis.

2.4 Morelloflavone

Morelloflavone (IUPAC Name: 8-[(2S,3R)-5,7-dihydroxy-2-(4hydroxyphenyl)-4-oxo-2, 3-dihydrochromen-3-yl]-2-(3,4-dihydroxyphenyl)-5, 7dihydroxychromen-4-one) represents a biflavonoid consisting of a flavanone, naringenin covalently linked to a flavone, luteolin (Verbeek *et al.*, 2004) (Fig. 13). This compound, also known as fukugetin (Konoshima *et al.*, 1969), is abundant in plants of *Garcinia* spp. It was firstly found in *Garcinia morella* (Karanjgaokar *et al.*, 1967) and is a common constituent among *Garcinia* spp., for example, *G. volkensii* (Herbin *et al.*, 1970), *G. talboti* (Joshi *et al.*, 1970), *G. livingstonei* (Pelter *et al.*, 1971; Mbwambo *et al.*, 2006), *G. terpnophylla and G. echinocarpa* (Bandaranayake *et al.*,

1975), G. multiflora (Chen et al., 1975; Lin et al., 1997), G. dulcis (Ansari et al., 1976; Deachathai et al., 2005; 2006; 2008), G. indica (Cotterill et al., 1977), G. densivenia (Waterman and Crichton, 1980), G. xanthochymus (Baslas and Pradeep, 1979, 1981), G. quadrifaria (Waterman and Hussain, 1982), G. spicata (Gunatilaka et al., 1984), G. nervosa (Babu et al., 1988), G. atroviridis (Permana et al., 2003), G. scortechinii (Sukpondma et al., 2005), G. cowa (Shen et al., 2006), G. subelliptica (Masuda et al., 2005; Terashima et al., 2008), and G. cymosa (Elfita et al., 2009). The study of this naturally occurring compound is interesting from its pharmacological point of view. Morelloflavone has been found to be an inhibitor of secretory phospholipase A₂ (Gil et al., 1997) and exerts potent anti-inflammatory effects in mice (Gil et al., 1997; Castardo et al., 2008). It also possesses anti-HIV activity by inhibiting both HIV-1 reverse transcriptase (HIV-1 RT) in vitro and HIV-1 (strain LAV-1) in human lymphocytes (Lin et al., 1997). In 2002, Li and co-workers demonstrated that morelloflavone inhibited fatty acid synthase, which became a potential antifungal target (Li et al., 2002). This compound isolated from the leaves of Rheedia gardneriana as fuguketin, also shows antibacterial activity (Verdi et al., 2003). Its strong inhibition on tyrosinase, the major enzyme responsible for skin melanization, has been found (Masuda et al., 2005). Among the anti-plasmodial compounds isolated from Endodesmia calophylloides, morelloflavone has been found to be active against Plasmodium falciparum W2 strain which is resistant to chloroquine and other drugs (Ngouamegne et al., 2008). Moreover, antioxidant activities of morelloflavone have been observed in various experimental systems (Sanz et al., 1994; Gil et al., 1997; Deachathai et al., 2005; Hutadilok-Towatana et al., 2007). Decha-Dier and co-workers have discovered anti-atherogenic effects of morelloflavone in hypercholesterolemic-induced rabbits (Decha-Dier et al., 2008). This biflavonoid also prevents restenosis by blocking injury-induced neointimal hyperplasia via the inhibition of vascular smooth muscle cell migration (Pinkaew et al., 2008). Pang and co-workers have recently shown that morelloflavone exerts antitumor angiogenesis by targeting the activation of Rho-GTPases and ERK signaling pathways (Pang et al., 2009).

Although morelloflavone reduces blood cholesterol levels and thus prevents progression of atherosclerosis in high fat-diet induced hypercholesterolemic animals (Decha-Dier *et al.*, 2008), the mechanism underlying such ability of this compound is still unknown. It was postulated that morelloflavone might exert its activity *via* inhibition of cholesterol *de novo* synthesis, and therefore, the effect of morelloflavone on HMG-CoA reductase, the key regulating enzyme of cholesterol pathway in mammals, was explored in this study.

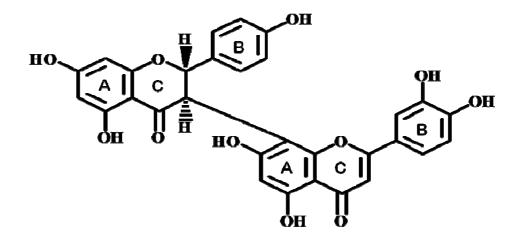


Figure 13 Structure of morelloflavone (MW = 556) consisting of naringenin (left) and luteolin (right) subunits.

CHAPTER 3

METERIALS AND METHODS

3.1 Materials

3.1.1 Morelloflavone

Morelloflavone (94% pure) was obtained from Dr. Decha Pinkaew. It was isolated from *Garcinia dulcis* leaves as previously described (Pinkaew *et al.*, 2009).

3.1.2 Recombinant plasmids and competent cells

The recombinant plasmids pET 17b-CD-HMG and *Escherichia coli* BL21 competent cells were kindly provided by Dr. Sriwan Wongwisansri of Bioassay Laboratory, National Center for Genetic Engineering and Biotechnology (BIOTEC), Thailand.

3.1.3 Animals

Male Wistar rats at about 8 weeks of age and weighing 230-270 g, were obtained from Animal House Facility Unit, Faculty of Science, Prince of Songkla University.

3.1.4 Instruments

| Name | Model | Manufacturer |
|-----------------------------------|---------|--------------------|
| Analytical balance | BP110S | Sartorius, Germany |
| Autoclave | HA-300M | Hirayama, Japan |
| Bench-top refrigerated centrifuge | 5804R | Eppendorf, Germany |

| Name | Model | Manufacturer |
|--|---------------------------|----------------------|
| Fraction collector | 2110 | Bio-Rad, USA |
| Heat block | D1100 | Labnet, USA |
| Microtube pump | MP-3 | Eyela, Japan |
| PCR machine | Mastercycler | Eppendorf, Germany |
| Polyacrylamide gel electrophoresis unit | AE-6450 | ATTO, Japan |
| Submarine type electrophoresis unit | Mupid Ex | Takara Bio, USA |
| Superspeed refrigerated centrifuge | J-30I | Beckman, USA |
| Ultracentrifuge | XL Series | Beckman, USA |
| Ultrasonicator | MSE ultrasonic power unit | MSE, England |
| UV-visible spectrophotometer | 8453 | Hewlett-Packard, USA |

3.1.5 Chemicals All chemicals used were of analytical grade.

| Name | Manufacturer |
|---------------------------|--------------------|
| Acrylamide | Merck, Germany |
| Agarose | Sigma-Aldrich, USA |
| Ammonium persulfate | Merck, Germany |
| Ampicillin, sodium salt | Sigma-Aldrich, USA |
| Blue Sepharose | Sigma-Aldrich, USA |
| Bovine serum albumin | Sigma-Aldrich, USA |
| Bromophenol blue | Fluka, Switzerland |
| Cholesterol, 95% pure | Sigma-Aldrich, USA |
| Dimethyl sulfoxide (DMSO) | Fluka, Switzerland |

| Name | Manufacturer |
|---|-------------------------|
| Dipotassium hydrogenphosphate | Fluka, Switzerland |
| Dithiotreitol (DTT) | Sigma-Aldrich, USA |
| Ethidium bromide | Sigma-Aldrich, USA |
| Ethylenediamine tetraacetic acid (EDTA) | Fluka, Switzerland |
| Folin-Ciocalteau's phenol reagent | Carlo Erba, Spain |
| Genistein from Glycine max (soybean), 98% pure | Sigma-Aldrich, USA |
| Glycerol | Analar, England |
| Glycine | Sigma-Aldrich, USA |
| D, L-3 hydroxy-3-methylglutarylcoenzyme A | Sigma-Aldrich, USA |
| (HMG-CoA) Isopropyl β-D-thiogalactopyranoside | Sigma-Aldrich, USA |
| LB media | Sigma-Aldrich, USA |
| Luteolin from synthesis, ≥98% pure | Sigma-Aldrich, USA |
| Naringenin from synthesis, ~95% pure | Sigma-Aldrich, USA |
| β -nicotinamide adenine dinucleotide phosphate, | Sigma-Aldrich, USA |
| reduced form (NADPH) Olive oil, extra virgin (Sabroso [®]) | Ihoros De Ibarra, Spain |
| Pravastatin, sodium salt | Sigma-Aldrich, USA |
| Potassium chloride (KCl) | Carlo Erba, Spain |
| Potassium dihydrogen phosphate | Merck, Germany |
| SOC medium | Sigma-Aldrich, USA |
| Sodium chloride (NaCl) | Merck, Germany |
| Sucrose | Carlo Erba, Spain |
| Tetramethylethylenediamine (TEMED) | Sigma-Aldrich, USA |
| Tris (hydroxymethyl) aminomethane | Vivantis, Italy |

3.2 Methods

Part 1: In vitro study

3.2.1 Preparation of HMG-CoA reductase

HMG-CoA reductase was expressed in *E. coli* according to Frimpong and co-workers (Frimpong *et al.*, 1993). The recombinant plasmid pET 17b-CD-HMG which contain HMG-CoA reductase gene was transformed to *E. coli* BL21 competent cell by following the on-line protocol of Sosnick Lab, University of Chicago, USA (http://sosnick.uchicago.edu/competent_cells.html). Firstly, 5 μ l of the plasmid solution was added to 10 μ l of the competent cells, and the resulting mixture was then put in an ice-bath for 10 min. The cell mixture was incubated at 42 °C for 90 sec, and placed back on ice for 2 min to reduce damage to the *E. coli* cells. Subsequently, 10 μ l of SOC medium and 1 ml of LB broth (without antibiotic) were added, and the mixture was further incubated at 37 °C by swirling the reaction tube every 15 min for 1 h. The resulting culture (200 μ l) was spread on LB agar (3 ml/plate) containing 100 μ g/ml ampicillin as a selective marker and further incubated at 37 °C.

After 12-16 h of culture, a freshly transformed colony was picked and inoculated into 10 ml LB broth containing 100 µg/ml of ampicillin and incubated at 37 °C overnight. An inoculum was diluted in 200 ml LB broth containing 100 µg/ml of ampicillin and cultured until its absorbance at 600 nm reached 0.8-1.0. Expression of the house mouse (*Mus musculus*) HMG-CoA reductase catalytic domain was induced by 0.5 mM isopropyl β -D-thiogalactopyranoside. Culturing was continued for another 5.5 h. The 200 ml culture was divided into four 50 ml aliquots. Each aliquot was centrifuged (10,000×g, 15 min at 4 °C), and the pellet was kept at -80 °C until used. The cell pellet was resuspended in 2 ml of 20 mM phosphate buffer, pH 7.3 containing 50 mM NaCl, 10% (v/v) glycerol, 100 mM sucrose, and 10 mM DTT on ice. This suspension was then used to prepare cell-free extract for the enzyme purification. To prepare cell-free extract, cells were lysed by sonication at 23% amplitude for 3 min (pulse on 6 sec, off 9 sec). Cell debris was separated by centrifugation (20,442×g, 10 min at 4 °C) and the supernatant subsequently filtered through a 0.2 μ m polysulfone syringe filter (Acrodisc[®], GelmanSciences, USA). The supernatant was subjected immediately to a Blue Sepharose column (1×5 cm) pre-equilibrated with 20 mM phosphate buffer, pH 7.3 containing 50 mM NaCl, 10% (v/v) glycerol, 100 mM sucrose, and 10 mM DTT. Elution was carried out using a linear gradient of 0 to 2 M NaCl in the same buffer at a flow rate of 0.25 ml/min, 4°C. Each fraction (1 ml) was monitored for protein at OD 280 nm and HMG-CoA reductase activity at OD 340 nm. The active fractions were then analyzed by SDS-polyacrylamide gel electrophoresis.

3.2.2 Kinetic studies of HMG-CoA reductase

The HMG-CoA-dependent oxidation of NADPH was monitored at 340 nm and 37 °C, with some modifications from the procedure developed in the Bioassay Laboratory, BIOTEC, Thailand (S. Wongwisansri, personal communication). The standard assay mixture contained 200 μ M D, L-HMG-CoA, 200 μ M NADPH, 100 mM NaCl, 1.0 mM EDTA, 10 mM DTT, and 100 mM phosphate buffer, pH 6.8 and a final volume of 150 μ l. The reaction mixture containing the enzyme (0.76 μ g protein) and all components, except NADPH, was pre-incubated at 37 °C for 6 min. The reaction was started when NADPH was added and its initial velocity was measured as the decreasing rate of absorbance at 340 nm (\triangle OD₃₄₀/min). The rate of reaction in the unit of \triangle OD₃₄₀/min was then translated into the enzyme specific activity from following calculation:

 $OD_{340} = \epsilon cl$

Where ϵ is millimolar absorptivity or millimolar extinction coefficient of NADPH which is equal to 6.22 mM⁻¹ cm⁻¹

c is the concentration of NADPH in mM

l is the optical path length in cm

Enzyme activity or reaction rate (V) was measured as the reduction of OD_{340} (ΔOD_{340}) over time investigated,

Therefore

$$V = \Delta c = OD_{340} / min / 6.22 mM^{-1} cm^{-1} \times cm^{1}$$
$$V = OD_{340} / min \times 0.1608 mM / min$$

V OD3407 **HIII** × 0.10

For 150 μ l reaction

 $V = OD_{340} / min \times 0.02412 \ \mu mol/min$

Specific activity (V_{spec}) was calculated as reaction rate per milligram protein used, therefore

 $V_{spec} = [OD_{340} / min \times 0.02412 \ \mu mol/min] / \mu g \text{ protein used}$

Enzyme kinetic parameters (k_m and V_{max}), were evaluated using the non-linear regression method based on Michaelis-Menten equation and the type of inhibition was identified using Lineweaver-Burk plot.

To investigate its inhibitory effect on enzyme activity, the test compound dissolved in DMSO at various concentrations, was added into the reaction mixture. For the HMG-CoA inhibition study, the NADPH concentration was fixed at 200 μ M, and, HMG-CoA was used in the range of 25-150 μ M. With respect to NADPH, HMG-CoA was fixed at 200 μ M, and NADPH was 50-100 μ M. The type of inhibition was identified using Lineweaver-Burk plot.

Kinetic parameters were evaluated using the non-linear regression method based on Michaelis-Menten equation:

$$V_i = \frac{V_{max} \times S}{K_m(1 + I/K_i) + S}$$

Where V_i is the initial velocity in the absence and presence of inhibitor; S and I, are concentrations of the substrate and the inhibitor, respectively; V_{max} , the maximum velocity; K_m , the Michaelis-Menten constant; and K_i is the inhibition constant defined as [E][I]/[EI]; [E] = enzyme concentration, [I] = inhibitor concentration, [EI] = enzyme-inhibitor complex concentration.

3.2.3 Protein assay

Samples were determined for their protein concentrations by the Lowry method (Lowry *et al.*, 1951) using bovine serum albumin as a standard.

3.2.4 SDS-polyacrylamide gel electrophoresis

The samples were electrophoresed in a slab gel, composed of 3% stacking gel and 12% separating gel according to the method of Laemmli (Laemmli, 1970). Each sample and a standard protein marker solution (Sigma-Aldrich, USA) containing phosphorylase b (94 kDa), albumin (67 kDa), ovalbumin (43 kDa), carbonic anhydrase (30 kDa), trypsin inhibitor (20.1 kDa), and α -lactalbumin (14 kDa), were mixed with a sample buffer (0.2 M Tris-HCl, pH 6.8 with 4% SDS, 40% (v/v) glycerol, 8 mM EDTA, 0.4% DTT, and 0.4% bromophenol blue as the tracking dye) in a ratio of sample: buffer = 3: 1 (v/v). They were then boiled for 5 min and loaded on the gel. Electrophoresis was carried out in the electrode buffer (25 mM Tris-HCl, 0.192 M glycine, 0.1% SDS, pH 8.3) at a constant current of 100 Volts for 1 h 45 min. After electrophoresis, protein bands were stained with Coomassie Blue. Migration distances of the samples, standard markers and bromophenol blue were measured and calculated for their relative mobility (R_f) by the following equation:

Relative mobility $(R_f) =$ <u>Migration distance of protein</u>

Migration distance of bromophenol blue

A standard curve plotted between logarithms of the standard molecular weights and their $R_{\rm f}$ values was used to estimate the molecular weight of protein sample.

Part 2: In vivo study

3.2.5 Animal treatments

All procedures concerning animal treatments and experimentation in this study were reviewed and approved by the Institutional Committee for Ethical Use of Experimental Animals at Prince of Songkla University (approval no. 22/51). Male Wistar rats were randomly divided into 4 groups (n = 5). Group 1 and Group 4 were fed pelletized commercial chow (normal) diet (C.P. Mice Feed[®], Charoen Phokphand, Thailand) and accessed to drinking-water *ad libitum*, whereas Group 2 and Group 3 received normal diet supplemented with 1% (w/w) cholesterol to induce hypercholesterolemia.

Each rat was housed separately in an identical stainless steel cage $(28\times30\times40 \text{ cm})$ and maintained in an air-conditioned room at 22 ± 3 °C, 50-60% relative humidity, and 12 h-light/dark cycle. Blood was taken twice a month for plasma cholesterol analysis. Before blood sampling, they were overnight-fasted. Heparinized blood was obtained from ocular bed puncture under ether anesthesia. It was centrifuged at 1,700×g for 15 min to separate plasma.

When the rats in Group 2 and Group 3 had their plasma cholesterol levels higher than 100 mg/dl, the experiments were started. For 6 weeks of study, animals in Group 1 were given normal diet and intragastrically administered with olive oil (5 ml) daily; Group 2: normal diet plus 5 ml cholesterol suspension in olive oil (1 g% body weight/day); Group 3: normal diet plus 5 ml suspension containing cholesterol (1 g% body weight/day) and morelloflavone (0.01 g% body weight/day); Group 4: normal diet plus 5 ml morelloflavone suspension in olive oil (0.01 g% body weight/day).

All rats were observed daily for any abnormalities of activities and behaviors. They were weighed individually once a week throughout the experimental period. Food and water intakes were also measured daily. At the end of experiments, the animals were fasted overnight. Blood was collected from the abdominal aorta under ether anesthesia and then prepared for plasma. Their livers were removed immediately after cervical dislocation and stored at -80 °C for mRNA analysis and hepatic HMG-CoA reductase assays.

3.2.6 Blood lipids measurement

Plasma HDL-C, triglyceride, and total cholesterol concentrations were determined by the colorimetric methods using an autoanalyzer (Cobas[®] Mira, Roche, Switzerland).

3.2.7 Preparation of liver microsome

The frozen liver was excised and immediately placed in an ice-cold 50 mM potassium phosphate buffer, pH 7.0 containing 0.2 M sucrose and 2 mM DTT (2 ml/g liver). The homogenization was performed in three strokes with a motor-driven Teflon pestle in a Potter-Elvehjem type glass homogenizer. The homogenate was

centrifuged for 10 min at $15,000 \times g$, 4 °C and the supernatant solution was centrifuged at $100,000 \times g$ for 75 min, and 4 °C. The precipitate was re-suspended in the same buffer with 50 mM EDTA and then centrifuged at $100,000 \times g$ for 60 min, and 4 °C to obtain microsomal pellet (Kleinsek *et al.*, 1977). It was kept frozen at -20 °C until use.

After thawing in an ice-bath, the microsomes were manually homogenized with a Teflon pestle and a glass homogenizer, in a solubilization buffer consisting of 0.1 M sucrose, 2 mM DTT, 50 mM KCl, and 30 mM EDTA in 50 mM potassium phosphate buffer, pH 7.0 (5 ml/g liver) (Heller and Gould, 1974). The homogenates were quantified for their protein contents and further used for HMG-CoA reductase assays.

3.2.8 Hepatic HMG-CoA reductase assay

The assay mixture contained 200 μ M D, L-HMG-CoA, 200 μ M NADPH, 100 mM NaCl, 1.0 mM EDTA, 10 mM DTT, and 100 mM phosphate buffer, pH 6.8 at a final volume of 150 μ l. The homogenate of liver microsomes in 100 mM phosphate buffer, pH 6.8 containing 100 mM NaCl, 1.0 mM EDTA, and 10 mM DTT was pre-incubated at 37 °C for 2 min. The reaction was then started by addition of D, L-HMG-CoA, and followed by NADPH. The absorbance at 340 nm was continuously measured for 3 min. The initial velocity of each sample was used to calculate the specific activity of the enzyme.

3.2.9 Determination of hepatic mRNA expression of HMG-CoA reductase and LDL receptor

Total mRNA was isolated from rat liver tissues based on the protocol of GenluteTM Direct mRNA Miniprep kit (Sigma-Aldrich, USA). Reverse transcription was carried out by using oneStep RT-PCR kit (Quigen, USA) and operated by a thermal cycler PCR. The appropriate primers of the target gene were designed corresponding to the positions of 104-123 and 2327-2346 of Wistar rat HMG-CoA reductase cDNA (Ihara-Watanabe *et al.*, 1999) as follows:

sense primer:5'-ACAATGTTGTCAAGACTTTT-3'antisense primer:5'-CCTCCTATGCTACCAGCCCAT-3'

The primers for LDL receptor gene were constructed according to Chen and Cheng (2005):

sense primer:5'-CAGCTCTGTGTGAACCT-3'antisense primer:5'-TTCTTCAGGTTGGGGATCA-3'

| Step | Duration | Temperature | |
|------------------------|----------|-------------|--|
| Reverse transcription | 30 min | 50 °C | |
| Initial PCR activation | 15 min | 95 °C | |
| Three step cycling | | | |
| - Denaturation | 1 min | 94 °C | |
| - Annealing | 1 min | 68 °C | |
| - Extension | 1 min | 72 °C | |
| Number of cycler | 40 | | |
| Final extension | 10 min | 72 °C | |

The program of the thermal cycler was set up as follows:

The PCR products were electrophoresed on a 2% agarose gel at 100 Volts for 45 min. The gel was stained by ethidium bromide. DNA bands were detected under UV light illumination and their densities were quantified by a Gel Document System using Labworks 4.6 Image Acquisition and Analysis Software (Bio Imagine System, UVP, USA).

3.2.10 Statistical analysis

All data were expressed as mean \pm SEM. They were analyzed for statistical significance by one-way analysis of variance using SPSS 11.5 (SPSS Institute, USA) and individual comparisons were obtained by LSD's multiple-range test. Significance was judged at p < 0.05.

CHAPTER 4

RESULTS AND DISCUSSION

Part I: In vitro study

4.1 Preparation of the catalytic portion of mouse HMG-CoA reductase

The DNA portion harboring a catalytic domain of mouse HMG-CoA reductase was successfully introduced into *E. coli* cells *via* the plasmid vector. After its expression was induced, the harvested protein was applied to Blue Sepharose affinity column. Upon eluting with 0-2.0 M NaCl gradient, enzyme-containing fractions were obtained at 0.40-0.64 M NaCl as in Figure 14. They were then subjected to SDS-polyacrylamide gel electrophoresis (SDS-PAGE), and the results are shown in Figure 15. Each fraction appeared as one major protein band on SDS-PAGE with an estimated molecular weight of 50 kDa, the same as reported previously (Wongwisansri and Kitikara, 2004). These enzyme fractions were then pooled, concentrated, and assayed for protein and the enzyme activity. Each preparation batch showed not less than 26% of protein yield and was purified about 4.5 folds from the starting material.

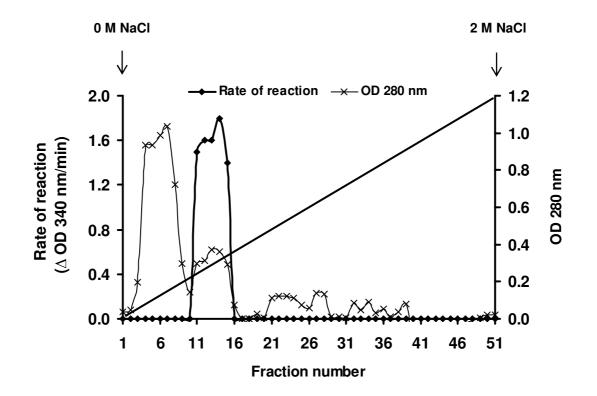


Figure 14 Chromatogram of Blue Sepharose column (1×5 cm). The bound proteins were eluted with a linear gradient of 0-2.0 M NaCl in 20 mM phosphate buffer, pH 7.3 containing 50 mM NaCl, 10% (v/v) glycerol, 100 mM sucrose, and 10 mM DTT at a flow rate of 0.25 ml/min. Each fraction (1 ml) was collected and measured for protein and the enzyme activity as described in Chapter 3.

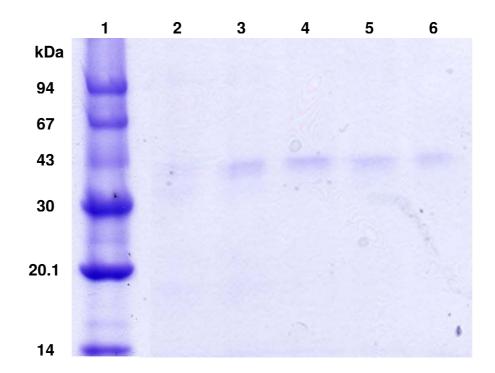


Figure 15 SDS-PAGE of HMG-CoA reductase fractions on 12 % polyacrylamide slab gel. The protein bands were stained with Coomassie Blue. Lane 1; low molecular weight markers: phosphorylase b (94 kDa), albumin (67 kDa), ovalbumin (43 kDa), carbonic anhydrase (30 kDa), trypsin inhibitor (20.1 kDa), and α-lactalbumin (14 kDa). Lane 2-6; HMG-CoA reductase fractions eluted from Blue Sepharose column containing protein of 21, 25, 32, 24, and 25 µg, respectively.

4.2 Enzyme kinetics and characterization

The HMG-CoA reductase preparation was tested for its activity. The enzyme was catalytically active yielding a high rate of oxidation and HMG-CoA concentration dependent as shown in Figure 16.

Its kinetic parameters, Michaelis-Menten constant (K_m) and maximum rate of reaction (V_{max}), were evaluated from the Lineweaver-Burk plots (Figure 16) using activity assay conditions as described in Chapter 3.

When the standard enzyme reaction was performed, V_{max} value was calculated as 5 ± 0.26 µmole NADPH/min/mg protein. Under the same conditions, K_m values for the two substrates, D, L-HMG-CoA and NADPH, were 74 ± 0.06 and 133 ± 0.21 µM, respectively, similar to those previously obtained (S. Wongwisansri, personnel communication).

The recombinant HMG-CoA reductase was also assayed in the presence of test substance at various concentrations. The Lineweaver-Burk plots (Figure 17 and Figure 18) were employed to predict the type of inhibition. Morelloflavone (A) competed with HMG-CoA, whereas it was noncompetitive towards NADPH, the second substrate of the enzyme. The inhibition constants (K_i) of this compound with regard to HMG-CoA and NADPH were 80.87 \pm 0.06 μ M and 103 \pm 0.07 μ M, respectively. From these results, it was concluded that morelloflavone occupied a portion of the binding site of HMG-CoA, thus blocking access of this substrate to the active site, whereas the nicotinamide binding pocket of NADPH was unoccupied by the inhibitor molecule.

In order to investigate which of morelloflavone subunits would participate in such binding, naringenin (B) and luteolin (C) were examined for their enzyme inhibitory activity. As evident from the results in Figure 17, both aglycone isoflavones were inhibitory to HMG-CoA reductase in the same manner as morelloflavone. They competed with HMG-CoA at the K_i values of $83.59 \pm 4.37 \,\mu$ M and $83.58 \pm 0.94 \,\mu$ M, respectively, comparable to morelloflavone but rather weaker than genistein (D), a HMG-CoA reductase inhibitor from soy (Sung *et al.*, 2004) (K_i = $49 \pm 1.28 \,\mu$ M). The more hydrophobic characteristics may allow genistein to bind more strongly to the enzyme than naringenin and luteolin (Breton *et al.*, 1975). These isoflavones were also noncompetitive with respect to NADPH (Figure 18). Their K_i values were: narigenin $182 \pm 67 \mu$ M, luteolin $188 \pm 14 \mu$ M, and genistein $104 \pm 0.64 \mu$ M. These findings thus suggest that morelloflavone molecule would function by binding with a hydrophobic pocket of the enzyme active site *via* either subunit. In this regard, a common structural feature of isoflavone binding to the enzyme is most likely "ring B". This cyclic structure is favorable for the enzyme binding based on the fact that it resembles a pyran ring of both HMG and statin as shown in Figure 19. It has been reported, however, that in addition to bonds formed by the HMG-like moiety, compactin and simvastatin exhibit binding *via* a decalin ring structure (Istvan, 2003), and this might the same case for "ring A and ring C" of isoflavones.

In this study, inhibition constant values of pravastatin (E) for HMG-CoA and NADPH were $0.6 \pm 0.02 \mu$ M. and $23.43 \pm 0.93 \mu$ M, respectively (Figure 17 and Figure 18). Although morelloflavone had relatively high K_i values compared with statins which usually inhibit the enzyme in the nanomolar range (Istvan, 2002), this compound appeared to be a novel naturally occurring HMG-CoA reductase inhibitor. However, structural mechanism for morelloflavone inhibition of HMG-CoA reductase needed to be further elucidated. Through x-ray crystallographic studies, statins have been found to basically use their common HMG-like moiety to occupy the active site of the enzyme, with the hydrophobic groups of the statins being positioned in a shallow groove formed by rearrangement of the C-terminal residues of the enzyme catalytic domain and thus sterically preventing the substrate from binding (Istvan, 2003).

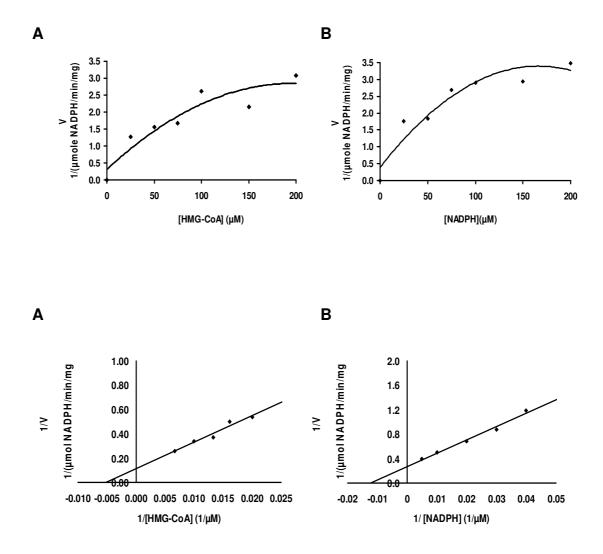


Figure 16 Michaelis-Menten plots (upper panel) and Lineweaver-Burk plots (lower panel) of HMG-CoA reductase for the two substrates, HMG-CoA (A), and NADPH (B). All analyses were conducted in six replicates.

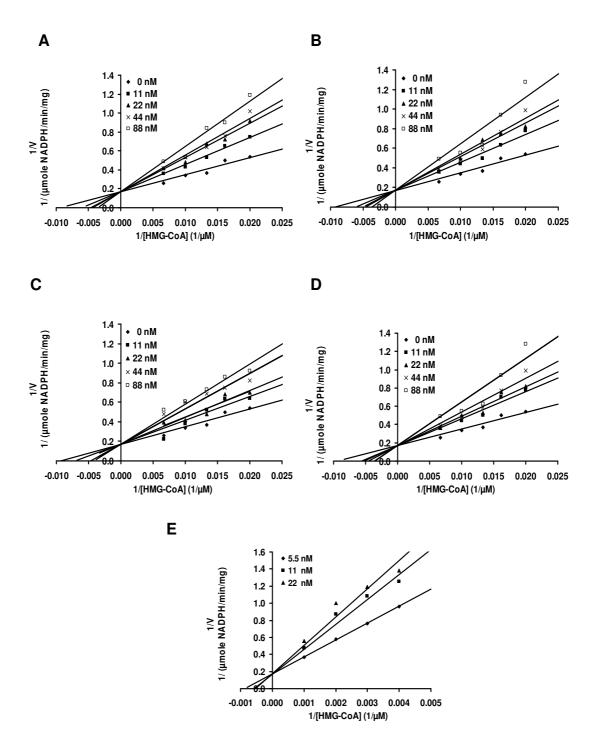


Figure 17 Lineweaver-Burk plots for the inhibition of HMG-CoA reductase by morelloflavone (A), naringenin (B), luteolin (C), genistein (D), and pravastatin (E), with respect to HMG-CoA. All analyses were conducted in six replicates.

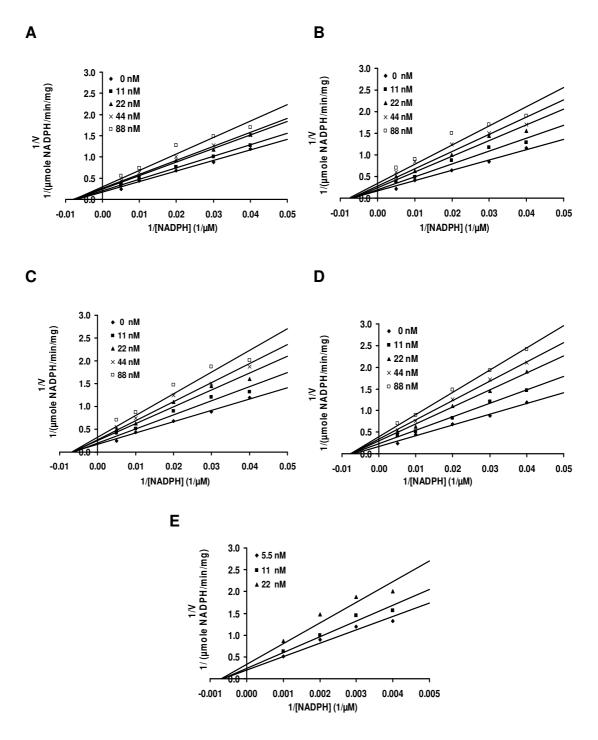


Figure 18 Lineweaver-Burk plots for the inhibition of HMG-CoA reductase by morelloflavone (A), naringenin (B), luteolin (C), genistein (D), and pravastatin (E), with respect to NADPH. All analyses were conducted in six replicates.

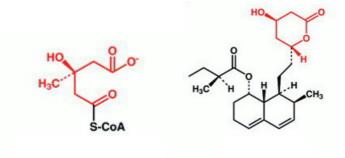


Figure 19 Cyclic structures of HMG-CoA (left) and statin drug (right) with their pyran rings as designated in red (www.nature.com/statins).

Part II: In vivo study

4.3 Effects on animal growth and food intake

Throughout the experimental period, daily food intake among the control rats (Group 1) and those receiving 0.01% (w/w) morelloflavone (Group 4) were not altered whereas both untreated (Group 2) and morelloflavone-treated hypercholesterolemic animals (Group 3) consumed lesser amounts of food until the end of study (Appendix I). This apparent effect has been reported previously in high-fat diet rats (Bahramikia and Yazdanparast, 2008). Surprisingly, their body weight levels were unchanged throughout the entire period (Appendix II). As a result, average food intake per body weight ratios of Group 2 and Group 3 were significantly lower than the rest (Figure 20).

To examine whether the treatments had any effects on the vital organ responsible for detoxification, all rat livers were excised, weighed and grossexamined at the termination. Their relative liver weight to final body weight ratios are also shown in Figure 21. A high-cholesterol containing diet has been reported to induce an over-weight liver from hepatic lipids accumulation (Seifalian *et al.*, 1999; Jayasooriya *et al.*, 2000). The same evidence was also obtained in this study (Appendix III). Therefore, those significantly greater ratios of liver weight to body weight in hypercholesterolemic groups as compared to the normal controls were attributed to increased liver weights among these animals. Due to hepatic enlargement, both cholesterol groups did not lose weight despite lower food intake.

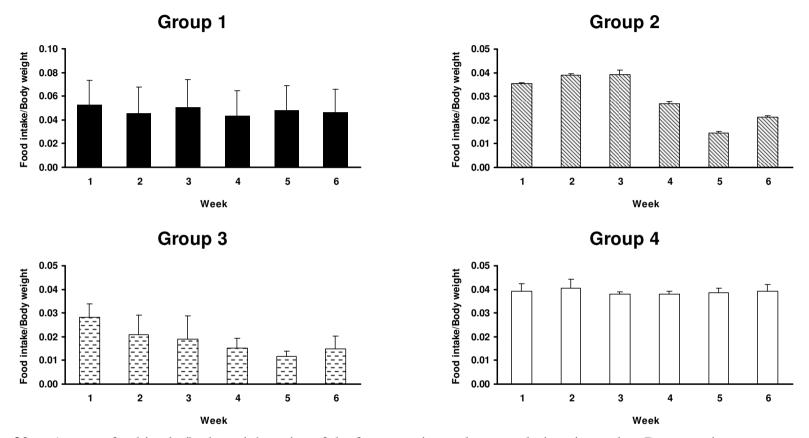


Figure 20 Average food intake/body weight ratios of the four experimental groups during six weeks. Data are given as mean ± SEM (n = 4). Group 1 = normal, Group 2 = hypercholesterolemic, Group 3 = hypercholesterolemic + morelloflavone, Group 4 = normal + morelloflavone.

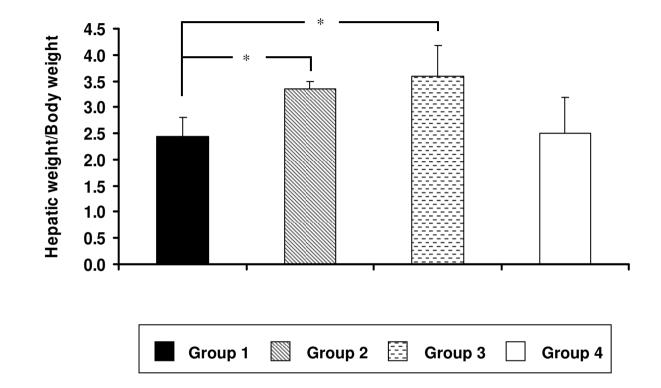


Figure 21 Rat hepatic weight/body weight ratios at the final week. Data are given as mean \pm SEM (n = 4), * p < 0.05. Group 1 = normal, Group 2 = hypercholesterolemic, Group 3 = hypercholesterolemic + morelloflavone, Group 4 = normal + morelloflavone.

4.4 Effects on blood lipid profiles

Oral administration of 0.01% (w/w) morelloflavone for 4 months has previously shown its anti-hyperlipidemic effects in the rabbits receiving highcholesterol diet (Decha-Dier et al., 2008). In the present study, however, the serum total cholesterol levels in hypercholesterolemic rats were not significantly improved after receiving the same dose of this compound continuously for 6 weeks as shown in Figure 22A. These controversial results are still unknown, yet most likely reflect differences in lipoprotein metabolism among animal species (Tall, 1993). Highcholesterol fed rats seem to respond to the HMG-CoA reductase inhibitors variably. Statins do not contribute to the plasma cholesterol-lowering action in this animal model (Endo et al., 1979; Lee et al., 2001) as they do in hamsters and rabbits (Ma et al., 1986), whereas anti-HMG-CoA reductase flavonoids such as hesperitin (Lee et al., 1999), quercetin (Bok et al., 1999), naringenin (Shin et al., 1999), and eriocitrin (Miyake *et al.*, 2006), have shown their hypocholesterolemic effects in these rats. In normocholesterolemic rats (Group 4), however, morelloflavone caused some but not significantly increased total cholesterol levels as compared to those of Group 1 (Figure 22A). Meanwhile, this compound seemed to stimulate the good cholesterol "HDL-C" in these animals. In contrast to the other groups, their plasma HDL-C levels did not decline with time but became slightly increased (Figure 22B). The reduction of this plasma lipoprotein fraction normally occurs in aging (Walter, 2009) and is speculated to be another risk factor for developing atherosclerosis (Nofer et al., 2002). From the above results, it was therefore postulated that such alteration of circulating HDL-C in Group 4-rats would partially account for their increased plasma total cholesterol levels, although the true reason is still unknown. It should be noted, however, that flavonoids have been demonstrated to increase HDL-C in various hypercholesterolemic subjects including rats (Kim et al., 2003). This compound produced no influence on blood triglyceride level in any rat group as shown in Figure 22C.

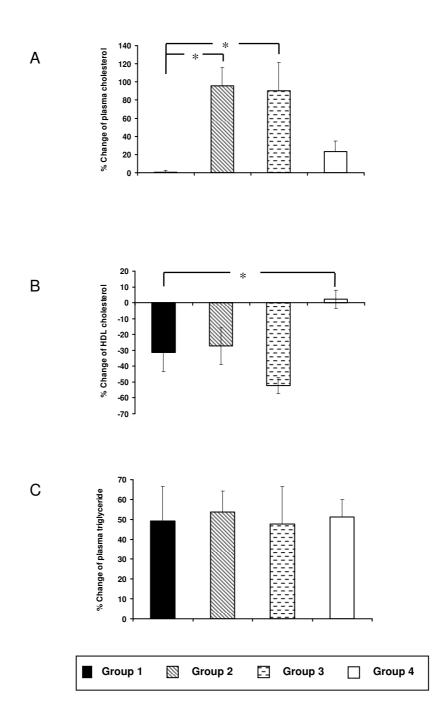


Figure 22Relative changes of plasma cholesterol (A), HDL-C (B), and
triglyceride (C) among the experimental rats. Means were normalized
with 0 week value in each group. Data are given as mean \pm SEM (n =
4), * p < 0.05. Group 1 = normal, Group 2 = hypercholesterolemic,
Group 3 = hypercholesterolemic + morelloflavone, Group 4 = normal
+ morelloflavone.

4.5 Effects on HMG-CoA reductase and LDL receptor mRNA expressions

To investigate whether morelloflavone would affect HMG Co-A reductase in the experimental rats, both the activity and the mRNA expression of this enzyme were measured in their livers. Among the four treatment groups, levels of hepatic HMG-CoA reductase mRNA (A) and its activity (B) correlated well (Figure 23). As anticipated, dietary cholesterol in rat livers of group 2 and 3 significantly inhibited the enzyme (Figure 23A). In mammalian cells, the intracellular concentration of cholesterol is maintained by a balance between endogenous synthesis and exogenous supply. This balance is achieved through feedback regulation on the biosynthesis pathway, and also on the LDL receptor synthesis (Brown and Goldstein, 1990). Intake of high-cholesterol diet increases the intracellular sterol pool, which alternatively suppresses the expression of HMG-CoA reductase mRNA via sterol regulatory elements (SREs) (White et al., 1997). In the presence of HMG-CoA reductase inhibitor, however, the *de novo* cholesterol biosynthetic pathway is depressed resulting in a decreasing pool. As a compensatory mechanism, HMG-CoA reductase gene is up-regulated leading to an elevation of the mRNA expression (Bergstrom et al., 1998). The phenomena whereby HMG-CoA reductase inhibitors trigger an over-expression of HMG-CoA reductase gene transcription are welldocumented (Bergstrom et al., 1998; Pitman et al., 1998). In the current study, such in vivo effect of morelloflavone was also evidenced (Figure 23 B). It was probable that either the compound itself or its metabolite would limit the availability of newly synthesized cholesterol, thereby promoting a hepatic expression of HMG-CoA reductase mRNA with a simultaneous increase in the enzyme activity in groups 3 and 4 (Figure 23A).

LDL receptor which plays an important role in blood cholesterol clearance is also modulated by cholesterol mainly from exogenous source (Brown and Goldstein, 1990). Its gene is down-regulated by high-cholesterol diet through the regulatory process mediated by negative effectors of transcription (SREBPs) (White *et al.*, 1997). Such LDL lowering effect of dietary cholesterol also occurred in this study (Figure 23C). When cellular cholesterol content becomes lower, SREBPs undergo proteolytic cleavage and the amino-terminal domain then activates LDL receptor gene expression (Horton *et al.*, 2002). Although, there is a suggestion that

HMG-CoA reductase and LDL receptor may be regulated independently in the liver (Spady *et al.*, 1985), a coordinate regulation of the expression of the genes coding for both proteins was demonstrated in this study (Figure 23). In accompany with HMG-CoA reductase gene (A), LDL receptor gene (B) was enhanced in the rat livers after morelloflavone treatment (Figure 23C). Accordingly, an increased cellular uptake of this lipoprotein *via* the receptor might explain those higher HDL levels in Figure 22.

Based on the above results, morelloflavone supplementation seemed to improve the lipid metabolism in animal fed high-cholesterol diet through hepatic responses to HMG-CoA reductase inhibition. However, more extensive investigation in an appropriate animal model to confirm the effects of this compound on the plasma and hepatic lipids level, apolipoproteins level, hepatic cholesterol-regulating enzymes and sterol excretion is required to establish its mechanism of action.

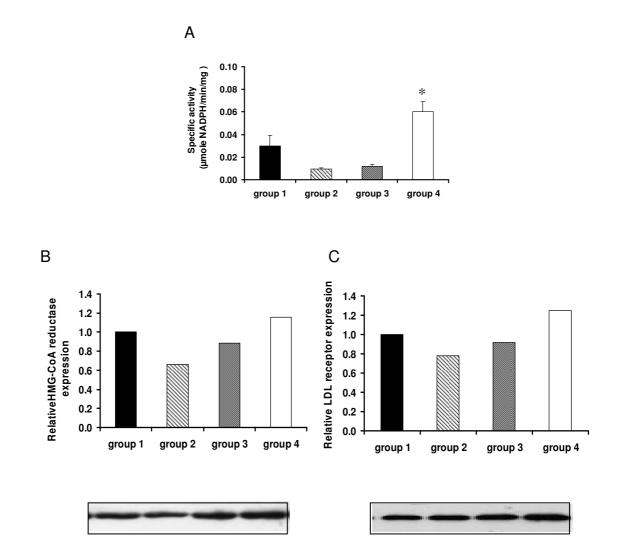


Figure 23 Hepatic HMG-CoA reductase activity (A), mRNA expressions of HMG-CoA reductase (B), and LDL receptor (C) of the experimental rats. The values of the enzyme activity are given as mean \pm SEM, (n = 3), * p < 0.05. Representative results of liver HMG-CoA reductase and LDL receptor mRNA levels are shown in each column. Group 1 =Group normal, Group 2 hypercholesterolemic, 3 = = hypercholesterolemic + morelloflavone, Group 4 = normal + morelloflavone.

CHAPTER 5

CONCLUSIONS

House mouse catalytic portion of HMG-CoA reductase was successfully produced from *E. coli* strain BL21 by using the recombinant plasmid pET 17b-CD-HMG. The partially purified enzyme containing a major protein of 50 kDa was obtained. Morelloflavone isolated from *Garcinia dulcis* leaves showed its inhibitory effects on this enzyme *in vitro* by competing with HMG-CoA, whereas it was noncompetitive towards NADPH. The same compound and both isoflavone subunits, naringenin and luteolin, were found equally effective at the enzyme inhibition. Based on these results, it was postulated that morelloflavone could exert the HMG-CoA reductase inhibitory activity through binding of the enzyme active site with either of its subunits.

The animal study further revealed that morelloflavone caused a significant increase of circulating HDL particles in normal rats, and also up-regulated hepatic HMG-CoA reductase and LDL receptor mRNA expressions in both normal and hypercholesterolemic animals in response to the enzyme inhibition.

The findings from this study thus suggested that morelloflavone derived from *G. dulcis* could be considered as an effective HMG-CoA reductase inhibitor that might be developed into a new hypocholesterolemic agent in the future. However, further studies are necessary to confirm the above effects of this compound and to establish its mechanism of action.

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APPENDIX

| Appendix I: Average daily food intake (g) of the four experimental groups during 6- |
|---|
| week period |

| | Group 1 | Group 2 | Group 3 | Group 4 |
|--------|-------------|------------|------------|------------|
| Week 1 | 27.86±11.14 | 17.00±0.42 | 13.06±2.02 | 18.03±0.59 |
| Week 2 | 24.54±12.09 | 18.18±0.35 | 10.38±3.07 | 19.29±1.23 |
| Week 3 | 26.75±12.47 | 17.96±0.81 | 9.03±2.89 | 17.77±0.92 |
| Week 4 | 23.18±11.05 | 12.04±0.52 | 7.03±1.13* | 17.34±0.49 |
| Week 5 | 25.32±11.31 | 6.57±0.30* | 5.09±1.15* | 17.57±1.37 |
| Week 6 | 24.32±10.73 | 9.32±0.49* | 6.80±2.21* | 18.00±1.39 |

Data are given as mean \pm SEM (n = 4), * p < 0.05. Group 1 = normal, Group 2 = hypercholesterolemic, Group 3 = hypercholesterolemic + morelloflavone, Group 4 = normal + morelloflavone.

Appendix II: Average body weight (g) of the four experimental groups during 6week period

| | Group 1 | Group 2 | Group 3 | Group 4 |
|--------|--------------|--------------|--------------|--------------|
| Week 1 | 524.75±12.49 | 480.25±10.59 | 448.80±30.18 | 454.80±20.56 |
| Week 2 | 504.50±27.40 | 465.25±7.42 | 439.60±32.49 | 461.40±20.65 |
| Week 3 | 496.50±27.24 | 456.75±10.09 | 433.80±31.72 | 458.60±21.53 |
| Week 4 | 495.50±28.69 | 445.50±12.22 | 443.80±30.14 | 450.80±20.65 |
| Week 5 | 497.75±26.68 | 448.75±10.57 | 439.80±35.70 | 465.80±22.02 |
| Week 6 | 498.75±25.77 | 440.50±10.05 | 429.00±44.18 | 477.80±29.83 |

Data are given as mean \pm SEM (n = 4), Group 1 = normal, Group 2 = hypercholesterolemic, Group 3 = hypercholesterolemic + morelloflavone, Group 4 = normal + morelloflavone.

Appendix III: Average hepatic weight (g) of the four experimental groups at the end of study period

| Group | Weight (g) |
|---------|-------------|
| Group 1 | 12.29±1.51 |
| Group 2 | 14.92±1.52* |
| Group 3 | 14.72±0.28* |
| Group 4 | 11.30±1.03 |

Data are given as mean \pm SEM (n = 4), * p < 0.05. Group 1 = normal, Group 2 = hypercholesterolemic, Group 3 = hypercholesterolemic + morelloflavone, Group 4 = normal + morelloflavone.

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