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利用抑肌素基因 5 端調節區建立篩選調節抑肌素基因表現之
中草藥分子生物分析平台

Establishment of molecular biological assay system for
screening myostatin inhibitor in Chinese herbs

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Author

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ABSTRACT

Myostatin, also known as growth and differentiation factor 8 (GDF-8), is one of the major regulators of skeletal muscle development. The phenotypes of myostatin-mutant cattle and mice have clearly established myostatin as a negative regulator of skeletal muscle growth. It has been found that the growth rate of myoblasts cultured in serum from double-muscling cattle fetuses was higher than myoblasts in normal-muscling cattle fetuses serum. Deletion and inhibition of myostatin in animals lead mainly to increased muscle mass and reduced fat mass. It is important to find an efficient way to regulate myostatin activity to improve animal growth. Myostatin 5'-regulatory region was used to construct the reporter plasmid. L8 rat myoblasts were transfected with reporter plasmids. After selection, transfected L8 cells were used to screen the CMH (Chinese medicine herbs) with myostatin regulators. Twenty seven CMHs were examined by using this molecular biological screening assay. Four CMHs were identified containing the myostatin regulators. It is concluded this molecular screening system is useful to identify CMH containing myostatin regulator.

LITERATURE REVIEW

Myostatin

Introduction of myostatin

Myostatin, also known as growth and differentiation factor 8 (GDF-8), is one of the major regulators of skeletal muscle development (Beyer *et al.*, 2013). It is a member of the transforming growth factor- β (TGF- β) superfamily. Myostatin plays a negative regulator in skeletal muscle growth. Overexpression of myostatin induces skeletal muscle atrophy and the mutations in the myostatin-encoding region are response for muscle hypertrophy and hyperplasia (Swatland and Kieffe, 1974). The phenotypes of myostatin-mutant cattle and mice have clearly established myostatin as a negative regulator of skeletal muscle growth. It has been found that the growth rate of myoblasts cultured in serum from double-muscle fetuses was higher than myoblasts in normal-muscle serum (Gerrard and Judge, 1993). Furthermore, addition of myostatin to culture medium inhibits the proliferation of myoblasts (Thomas *et al.*, 2000; Rios *et al.*, 2001; Taylor *et al.*, 2001). Moreover, the degree of inhibition is dose dependent and reversible (Thomas *et al.*, 2000).

Activation of the myostatin and effect of myostatin on muscle development

Myostatin protein is predominantly synthesized in skeletal muscle as a 375-amino acid protein. It is proteolytically processed and then mature C-terminal myostatin is secreted. Pro-myostatin dimerizes by a disulfide bond near the carboxy(C)-terminus and is systematically cleaved at the RSRR (Arg-Ser-Arg-Arg) site by a calcium-dependent serine proteinase, which was known as furin convertase. (Molloy *et al.*, 1994, 1999; Lee and McPherron, 2001). This proteolysis generates the N-terminal propeptide and a C-terminal receptor-binding domain (McFarlane *et*

al., 2005). The latent myostatin complex forms as the propeptide noncovalently binds the C-terminal region, and this prevents myostatin from binding to the target receptor (McFarlane *et al.*, 2005). Members of the bone morphogenetic protein-1/tolloid (BMP-1/TLD) family are metalloproteinases that can cleave the propeptide and release mature myostatin (Figure 1) (Wolfman *et al.*, 2003). Mature myostatin binds to type IIB receptor and mainly transduces the signal via the Smad pathway (

Figure 2). Among mammals, myostatin gene contain 3 exons and the myostatin gene among different species are highly homologous, indicating a conserved function (Table 1) (Lee and McPherron, 2001; Shen, 2010). Myostatin appears to down regulate both prenatal and postnatal myogenesis (Deveaux *et al.*, 2001; Pirruccello-Straub *et al.*, 2018). In the previous research shows that myostatin down regulates the muscle fiber number by reducing the myoblast proliferation and differentiation during early myogenesis (Deveaux *et al.*, 2001). Myostatin appears to be a target for compensatory muscle growth in farm animals, and for muscle wasting and cachexia in illness (Pirruccello-Straub *et al.*, 2018).

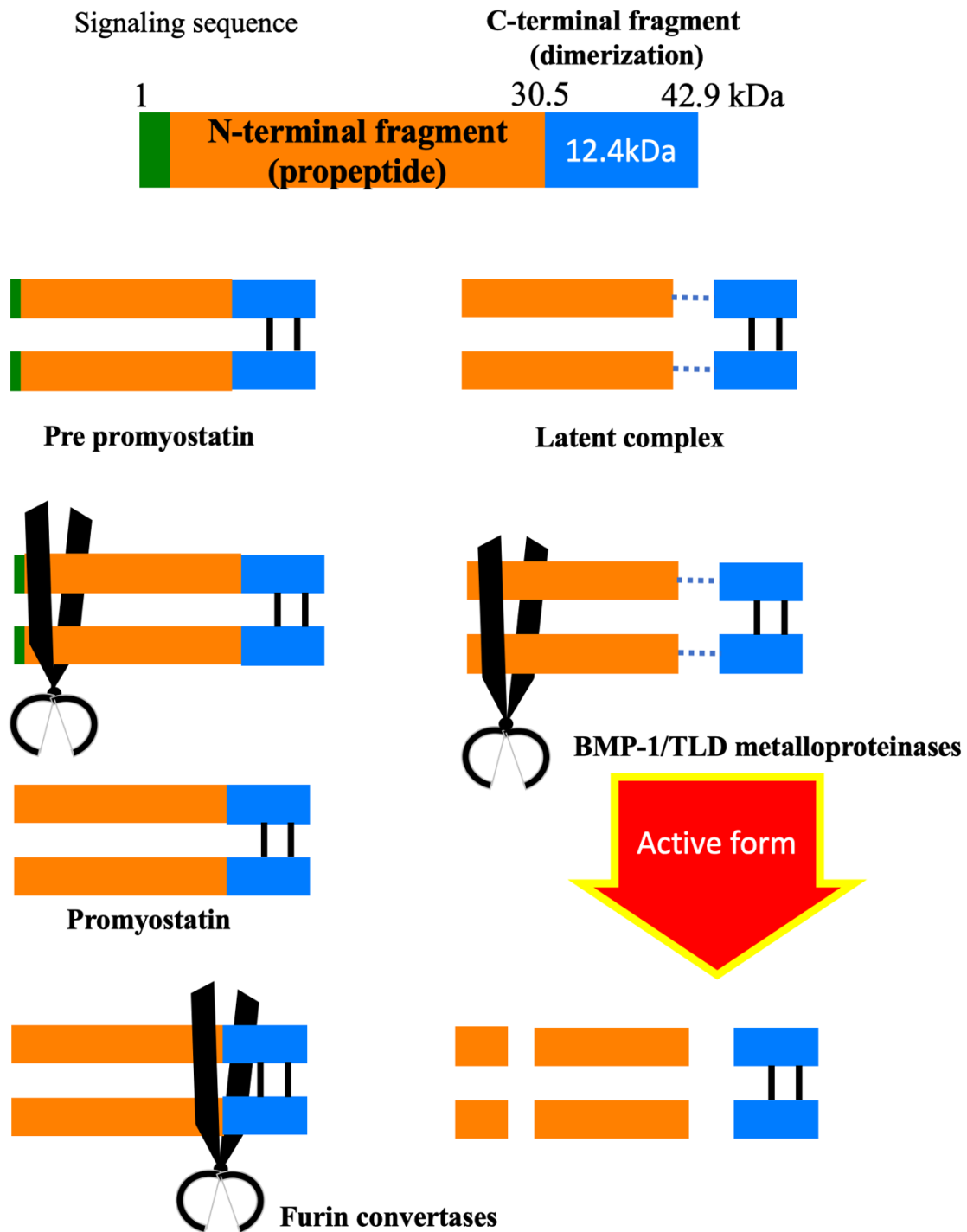


Figure 1. The activation of myostatin by furin convertase and BMP-1/TLD metalloproteinase

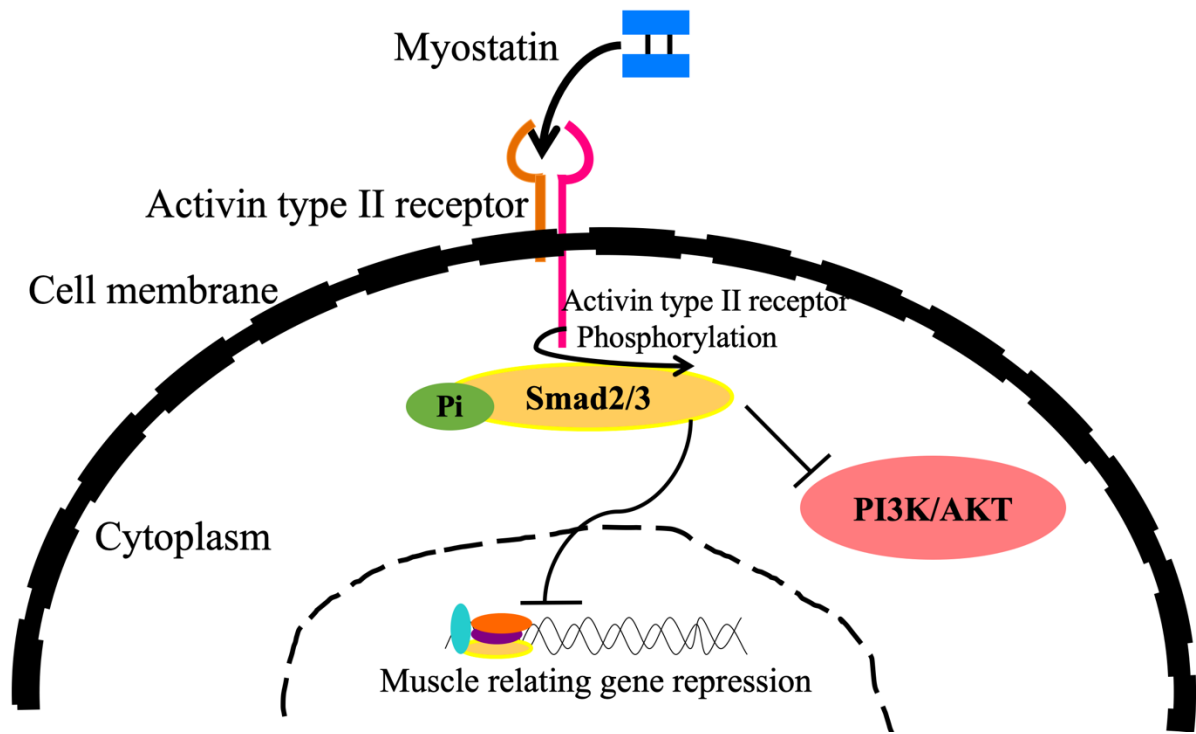


Figure 2. The signal transduction pathway of myostatin

Table 1. Homology matrix of myostatin amino acid sequences in 13 species (Shen, 2010)

Human (<i>Homo sapiens</i> _NP_005250.1)	100%													
Chimpanzee (<i>Pan troglodytes</i> _NP_001073388.1)	99.7%	100%												
Monkey (<i>Macaca mulatta</i> _NP_001073588.1)	99.7%	100%	100%											
Pig (<i>Sus scrofa</i> _O18831.1)	98%	98%	98%	100%										
Dog (<i>Canis lupus</i> _NP_001002959.1)	96%	96%	96%	97%	100%									
Cattle (<i>Bos taurus</i> _NP_001001525.1)	94%	94%	94%	96%	93%	100%								
Sheep (<i>Ovis aries</i> _O18830.1)	94%	94%	94%	93%	93%	93%	100%							
Goat (<i>Capra hircus</i> _Q6T5B8.2)	94%	94%	94%	95%	93%	93%	100%	100%						
Mouse (<i>Mus musculus</i> _NP_034964.1)	96%	97%	97%	96%	94%	93%	92%	92%	100%					
Rat (<i>Rattus norvegicus</i> _NP_034964.1)	95%	95%	95%	95%	93%	91%	91%	91%	98%	100%				
Chicken (<i>Gallus gallus</i> _NP_001001461.1)	92%	92%	92%	91%	90%	88%	88%	88%	91%	90%	100%			
Goose (<i>Anser anser</i> _AAL35276.1)	91%	91%	91%	90%	89%	87%	87%	87%	90%	89%	97%	100%		
Duck (<i>Anas platyrhynchos</i> _Q8UWE0.1)	92%	92%	92%	91%	90%	88%	88%	88%	91%	90%	98%	98%	100%	

Regulation of myostatin gene expression

To regulate the level of specific gene expression, there are various methods. Some strategies are demonstrated the regulation of myostatin gene expression. 5' regulatory region single nucleotide polymorphism (SNP), and SNP in myostatin gene affected myostatin gene expression. In addition, the amount of active protein, antagonist and antibody of myostatin are also can be used to regulate the myostatin activity.

SNP analysis of myostatin gene showed that myostatin 435GG/447AA genotype in both Duroc x Landrace (DL) and Duroc x Yorkshire x Landrace (DYL) pigs had significantly lower lean weight, lean percentage, anterior-end meat and posterior-end meat, while they had significantly thicker backfat either measured by ultrasound or carcass measurement (Jiang *et al.*, 2002; Tu *et al.*, 2013). The haploptype-1 (A435-A447-A879) of myostatin promoter up-regulated two-fold of the reporter gene expression in cell culture system (Bongiorni *et al.*, 2014). The result from Liu and coworkers (Liu *et al.*, 2010) results showed that G/A435-A/A447-T/A879 genotype resulted in significantly higher body weight on Day 21 (BW21), and significantly lower average daily weight gain from Day 21 to Day 70 (ADG2) in Duroc pigs (Liu *et al.*, 2010). The result from Shen (2010) showed 5' regulatory region was highly conserved in bird and in mammal (Figure 3).

The inhibitor of myostatin or myostatin receptor antibody can be considered for therapy of skeletal muscle dystrophy or atrophy. Injection of myostatin antigen in mice induced production of maternal myostatin antibody. The offspring, both male and female, contained higher crude protein and lower crude fat compared with control group (Liang *et al.*, 2007). Moreover, the JA16 Ab (myostatin monoclonal antibody)-treated mice gained approximately 10%

weight than the control and untreated mice (Whittemore *et al.*, 2003). In mdx mouse, model of Duchenne muscular dystrophy (DMD), the decreased serum creatine kinase activity and increased extensor digitorum longus (EDL) mass were found in myostatin antibody-treated group (Bogdanovich *et al.*, 2002). In addition, the antibody specifically against myostatin receptor, activin type II receptor (ActR II), increased the animal muscle growth. The interaction of ActR II and its antibody prevented myostatin from binding to ActR II, and interrupted myostatin intracellular signal transduction pathway (Lach-Trifilieff *et al.*, 2005). In poultry, the animals from direct injection of myostatin antibody into yolk had significantly heavier carcass, leg muscles and breast muscles than that in the control group (Kim *et al.*, 2007).

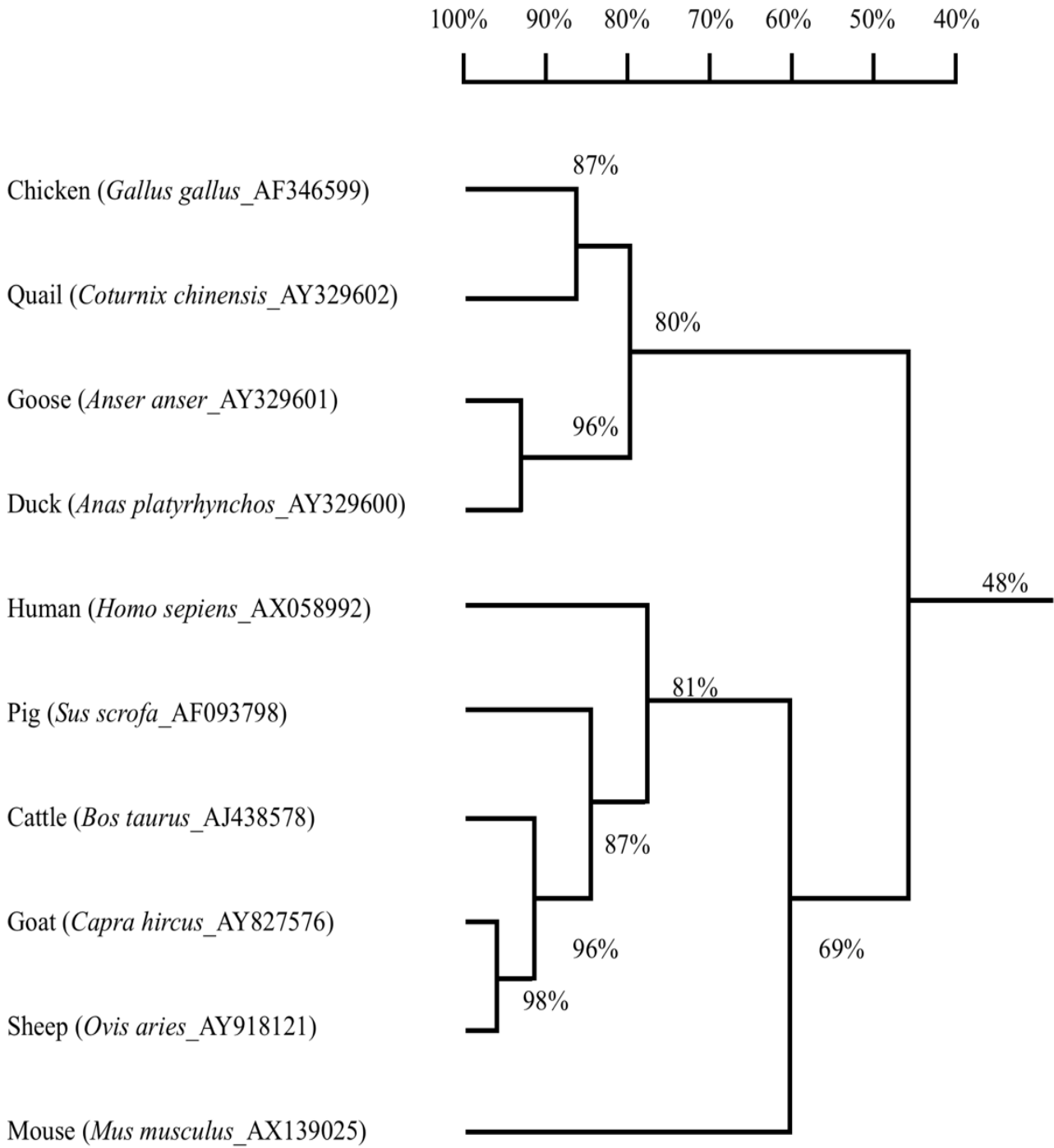


Figure 3. The 5' regulatory region sequence of myostatin homologous tree in 10 species (Shen, 2010)

Effect of myostatin on myoblast proliferation and differentiation

Myoblast fusion is a basic step within the muscle differentiation in most organisms. To achieve the complex of the system, the high expression level of *Pax* and *Mp20* are critical during myoblast fusion. In contrast to *Pax* and *Mp20*, low expression level of *mspo* increases myoblast fusion (Bataillé *et al.*, 2010). Typically, these cells migrate long distance prior to fuse. In addition, the first fusion that occurs throughout embryogenesis, needs a smORF-encoding microprotein, Minion (microprotein inducer of fusion). Lacking Minion, myoblasts are still aligned but fail to form syncytial cell (Millay *et al.*, 2013; Zhang *et al.*, 2017). Myoblast fusion involves cell migration, adhesion and transduction. The regeneration and/or growth of muscle involves proliferation and differentiation of muscle satellite cells (Wei *et al.*, 2015).

Myoblast proliferation

Myostatin affects the rate of myoblast proliferation by regulating cell cycle progression. In mouse myoblast C₂C₁₂ cell culture, the overexpressing endogenous myostatin inhibited myoblast proliferation through cell-cycle arrest in the G₁ and G₂ phases. The cells prevent to transit to S phase where DNA synthesis occurs and M phase where cell division occurs, respectively (Dominique *et al.*, 2006). The cell cycle arrest of G₁ phase is mediated through up regulation of one of the cyclin-dependent kinase inhibitors, p21 (Manceau *et al.*, 2008). In addition, myostatin also trend to reduce levels of cyclin dependent kinase-2 (Cdk2). The combination of the increased p21, and the decreased Cdk2 level resulted in a marked reduction of Cdk2 activity. Rb, a substrate of Cdk2, accumulated in an active hypophosphorylated state resulting in inhibition of cell cycle progression at G₁ phase (Thomas *et al.*, 2000). Rios and coworkers (Rios *et*

al., 2002) also reported an increase in the expression of p21 in the presence of ectopically overexpressed myostatin.

Myoblast differentiation

In mouse C₂C₁₂ myoblasts, myostatin expression increases during differentiation (Rios *et al.*, 2001). In addition, myostatin has also been shown to inhibit the myoblast differentiation of (Rios *et al.*, 2002). To inhibit the myoblast differentiation, myostatin inhibit the expression of the MRFs, MyoD, myogenin Myf5 (Langley *et al.*, 2002) as well as the late myogenic differentiation marker, myosin heavy chain (MHC) (Rios *et al.*, 2002; Wei *et al.*, 2015). Interestingly, the expression of p21, in contrast to proliferating myoblasts, is also down-regulated by myostatin during differentiation (Langley *et al.*, 2002). The mechanism by which myostatin regulates MyoD has been partially elucidated. Langley and coworkers (Langley *et al.*, 2002) showed that the presence of myostatin increased the phosphorylation of Smad 3, and increased the association of MyoD and Smad 3, which is known to repress MyoD activity (Liu *et al.*, 2001). In addition, in embryo, myostatin signaling promotes the initiation of the terminal differentiation by activating MyoD expression in myoblasts (Manceau *et al.*, 2008). It has also been shown that cells made quiescent by myostatin are not the same as reserve cells. During myoblast differentiation, a subset of myoblasts, termed reserve cells, remain quiescent and undifferentiated but retain the capacity to proliferate and differentiate. Reserve cells are characterized by a high expression of p130 and Myf5 and low MyoD expression. Myostatin-induced quiescent cells, however, do not express high protein level of p130 or Myf5 in myoblast differentiation, although they have decreased MyoD and MRF expression (Langley *et al.*, 2002).

Characteristic of myostatin in muscle development

Deletion and inhibition of myostatin in animals lead mainly to increased muscle mass and reduced fat mass. In myostatin-null mice and double-muscling cattle breeds without myostatin present, there is an increase of muscle fiber number and hyperplasia of muscle cells were observed (Holmes and Ashmore, 1972; Gerrard and Judge, 1993; McPherron *et al.*, 1996). Meanwhile, the results in mouse C₂C₁₂ myoblasts had demonstrated that myostatin control muscle mass by inhibiting cell proliferation and protein synthesis. Myostatin shows negatively regulate satellite cell activation (Taylor *et al.*, 2001; Zhang *et al.*, 2015). Forkhead box O (FoxO) is a key element of myostatin expression. The report showed that inhibition of the FoxO pathway increased in skeletal muscles during aged animal muscle atrophy (Reed *et al.*, 2012).

Licorice

Introduction of licorice

Glycyrrhiza uralensis, also known as licorice, has been used in food additive and medicinal herb in Asia for thousand years. It is perennial herb native from Mediterranean region, central to southern Russia and Turkey to Iran (Dastagir *et al.*, 2016). Raw licorice roots mainly from the *Glycyrrhiza* species, such as *G. glabra*, *G. uralensis* and *G. inflata*, are usually used as medicinal material (Simmler *et al.*, 2015). Licorice is multifunctional herb, and its special taste made it have been researched since 1940s. The pharmaceutical effect of licorice was first reported in 1950s and many scientists investigated licorice components analysis and its bioactivity in the same year. In 1950s, succus licorice was the major licorice used in daily life. Licorice was usually utilized for gastric ulcer therapy, but the effective molecule nor the medicinal mechanism was unknown. In traditional Chinese pharmacy, licorice was considered as drug to rejuvenate physical condition for fighting fatigue. In Egypt, Greece and Rome, licorices have been used since ancient times, and even now licorice is still used in the daily life (Dastagir *et al.*, 2016).

Bioactivity of licorice

Most reports showed that licorices have anti-germ activity (Arifullah *et al.*, 2013; Ren *et al.*, 2015). One of the components in licorice, liquiritigenin, has estradiol (E₂) activity which can activate estrogen receptor-alpha and beta (ER α and β) in MCF-7 and T47D breast cancer cells (Lecomte *et al.*, 2017). Licorice has been used in hepatitis therapy for a long time. Chalcones, one of the licorice components, were relatively more active than other compounds for protecting liver (Kuang *et al.*, 2017). The active compounds of licorice ethanol extract were increased to the PPAR-gamma ligand-binding activity which increased myoblast

and liver cell glucose sensitivity (Kuroda *et al.*, 2010). Some components of licorice have ability to induce hepatic carcinoma apoptosis through ROS dependent pathway, such as LicA (Choi *et al.*, 2014). In vivo study showed that glabridin in *G. glabra*, served as a dietary supplement, down regulated the sensitivity of oxidation in low-density lipoprotein (LDL), therefore, increased the possibility of the atherosclerotic lesion space (Fuhrman and Aviram, 2001; Grassi *et al.*, 2010). These results may be associated with the absorption and binding of glabridin to the low-density lipoprotein and protection of the low-density lipoprotein from oxidization (Fuhrman *et al.*, 1997; Messier *et al.*, 2011).

The methanol extract of *G. glabra rhizomes*, at a dose of 150 mg/kg, has anti-arthritic activity in male rats mediated by inhibition of leukocyte migration, autoantigen production, and exhibition of anti-proteinase activity (Choudhary *et al.*, 2015). Ahn and coworkers (2013) declared that *G. glabra* effectively inhibits the adipogenesis of 3T3-L1 cells. Glabridin has a therapeutic effect on metabolic disorders, such as diabetes and hyperglycemia, by modulating glucose metabolism through AMPK (AMP activated protein kinase) in skeletal muscle cells (Sawada *et al.*, 2014). Yoshioka and coworkers (2018) demonstrated that licorice extract activates mTOR, p70S6K, and regulation of phosphorylation of FoxO3a. These factors were involved in protein synthesis, thus help to prevent type 2 diabetes mellitus-induced muscle atrophy.

Component Analysis of Licorice

Licorice contains many polar and nonpolar molecules. Licorice root contains chemical compounds such as triterpene saponins, flavonoids, coumarins, and other phenolics (Hosseinzadeh *et al.*, 2015). In licorice, flavonoids contain diverse structures (Figure 4), which includes flavanones, chalcones, flavones, and

isoflavones (Zhang and Ye, 2009) (Table 2). Some of the active constituents of *Glycyrrhiza spp.* that belong to these flavonoid structures (Hosseinzadeh *et al.*, 2015). Chalcones compose with open-chain flavonoid which has two aromatic rings connected by three-carbon α , β unsaturated ketone (Figure 4) (Franceschelli *et al.*, 2017). Mahapatra *et al.* (2015) showed that chalcones had efficiency effect on diabetes management and therapy.

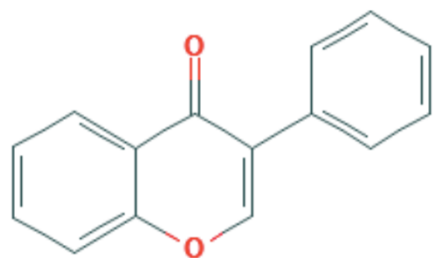
Liquiritin, isoliquiritin and isoliquirigenin inhibit the growth of non-small cell lung cancer cells. The chemical constituents of licorice are extensively investigated by completely different authors (Hayashi *et al.*, 2016; Siracusa *et al.*, 2011). Although, few studies were done on the nutritious composition of *G. glabra*. Nutritionally, licorice may be a source of proteins, amino acids, polysaccharides and straight forward sugars, minerals (such as calcium, phosphorus, sodium, potassium, iron, magnesium, silicon, selenium, manganese, zinc, and copper), pectins, resins, starches, sterols, gums, oestrogens, tannins, phytosterols (sitosterol and stigmasterol), coumarins, vitamins (B1, B2, B3, B5, E, and C), and glycosides. An oversized variety of biological compounds have additionally been isolated, such as triterpenes, saponins (responsible for the sweet taste), and flavonoids (Rizzato *et al.*, 2017; Wang *et al.*, 2015). Recently, one of licorice main component, glabridin, was found to inhibit dexamethasone- induced protein degradation in C₂C₁₂ myotubes and the muscle atrophy in mice. Glabridin reduced dexamethasone-induced expression of MuRF1 and Cbl-b through the inhibition of glucocorticoid receptor activation and acted as an antagonist (Yoshioka *et al.*, 2019). Kinoshita *et al.* (2017) showed licorice flavonoid oil (LFO) in elderly diet significantly improved their body trunk muscle mass and body fat mass. Therefore, licorice may be used as the diet additives for improving animal muscle mass. In this thesis, the molecular screening system was established to screen the myostatin regulator in CMH, licorice was chosen to

investigate the effect of licorice ethanol extract on the proliferation and differentiation in L8 myoblasts.

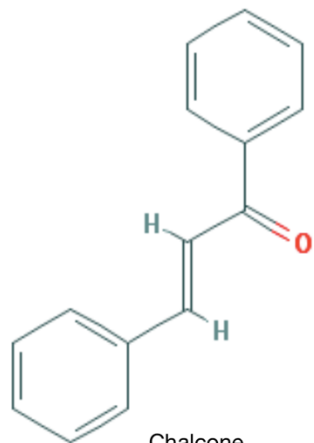


Table 2. The flavonoid compounds in licorice

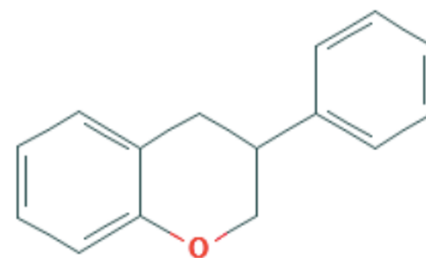
Structure	Compounds
Chalcones	isoliquiritigenin (Simmler <i>et al.</i> , 2014) isoliquiritigenin apioside (Simmler <i>et al.</i> , 2014) isoliquiritin (Simmler <i>et al.</i> , 2014) neoisoliquiritin (Zhang and Ye, 2009) licochalcone A and B (Zhang and Ye, 2009)
Flavanones	liquiritigenin (Xu <i>et al.</i> , 2013) liquiritin (Xu <i>et al.</i> , 2013) liquiritin apioside (Xu <i>et al.</i> , 2013)
Flavones (flavonols)	licoflavonol (Qiao <i>et al.</i> , 2014; Zhang and Ye, 2009) isolicoflavonol (Qiao <i>et al.</i> , 2014; Zhang and Ye, 2009)
Isoflavones	licoisoflavone (Zhang and Ye, 2009; Qiao <i>et al.</i> , 2014) licoricone (Zhang and Ye, 2009; Qiao <i>et al.</i> , 2014) gancaonin (Zhang and Ye, 2009; Qiao <i>et al.</i> , 2014) angustone (Zhang and Ye, 2009; Qiao <i>et al.</i> , 2014) isoangustone (Zhang and Ye, 2009; Qiao <i>et al.</i> , 2014) glabrone (Zhang and Ye, 2009)
Isoflavans	glabridin (Zhang and Ye, 2009) hispaglabridin A and B (Zhang and Ye, 2009)



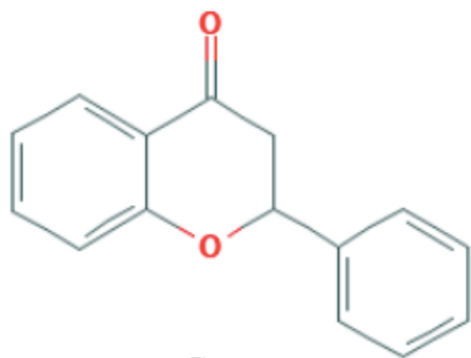
Isoflavone



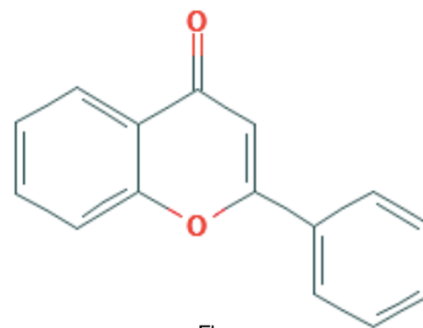
Chalcone



Isoflavan



Flavanone



Flavone

Figure 4. The chemical structures of major flavonoids in licorice.

MATERIALS AND METHODS

Plasmid construction

The primers for the construction of the plasmid were purchased from Mission Biotechnology. The primer sequences of mouse myostatin 5' regulatory region was designed according to *Mus musculus* AX139025. The primer sequences are listed below:

Forward mMSTN-PL-F-BglII-5': 5'-ATA AGA TCT CCT TTT TAA GTC
CTA AGT CAC ACG G-3'

Reverse mMSTN-PL-R-BamHI-3': 5'-ATA GGA TCC CCA GGG AGT
CCT GTA TAC TG-3'

Template DNA is isolated from C₂C₁₂ myoblast. The PCR reaction mixture contains 500 ng genomic DNA, 0.4 μM forward primer, 0.4 μM reverse primer, 0.3 mM dNTPs, 4 μL 5X KAPA HiFi Fidelity Buffer, KAPA 0.5U HiFi HotStart DNA Polymerase (Kapa Biosystems, Roche, Basel, Switzerland). The reaction conditions are listed in Table 3. The restriction enzyme cutting site were added on the primers which use in PCR. After PCR, the product was purified by phenol chloroform extraction method. Restriction enzyme BglII 10 U/μL (A/GATCT) and BamHI (G/GATCC) with bovine serum albumin (BSA) 10 mg/mL (Promega, Madison, Wisconsin, United State). One μg pMetLuc2 reporter plasmid and 1 μg PCR product were digested with *Bg*III and *Bam*HI. And then, the 100 ng plasmid and 100ng PCR fragments were ligated by 0.5 U T4 DNA ligase (Promega, Madison, Wisconsin, United State) (Figure 1). The plasmid was confirmed by sequencing (Mission

Biotech, Taipei, Taiwan), and then the insert sequence was aligned with mouse 5' regulatory region sequence (*Mus musculus* AX139025).



Table 3. The PCR condition for amplification of mouse myostatin 5'-regulatory region

	Temperature (°C)	Time	Number of cycles
Denaturation	95°C	1 min	1
Denaturation	94°C	1 min	
Annealing	65°C	1 min	30
Extension	72°C	1 min	
Extension	72°C	5 min	1
	4°C	overnight	

Transfection of pMet Luc2 mMSTN into L8 cells

L8 cells were transfected with pMetLuc2 mMSTN by using Lipofectamine 2000 (Invitrogen, California, United States) accordance to manufacturer's recommendations.

L8 cells were cultured in DMEM with 10% FBS (Fetal bovine serum; Hyclone, GE Healthcare, Illinois, United States). Lipofectamine® reagent (9 µL) was diluted in 150 µL Opti-MEM® medium. Four µg plasmid DNA was added in 150 µL of Opti-MEM, and mixed gently and incubated for 5 minutes at room temperature. The diluted Lipofectamine 2000 and the diluted DNA were mixed and incubated for 30 minutes. DNA-Lipofectamine 2000 mixture (250 µL) was added into petri dish containing cells and medium, then mixed gently by rocking the dish back and forth. The cells were incubated 24 hrs. The medium was replaced by fresh medium containing 200 µg/mL G418. Consequently, the single cell colony is selected for future use.

Chinese medicinal herb extraction

Twenty seven CMHs were tested in this study. The CMHs were grinded to powder with grinder at 25000-30000 rpm. Herb powders (0.5 g) were added into 10 mL ethanol and stirred for 10 hrs. The mixture was filtered with No.1 filter paper, and then put the mixture in hood to evaporate the alcohol. Gel-like product was collected and recorded the weight. The final product was dissolved in alcohol to final concentration of 40 mg/mL.

Chinese medical herb screening

The pMetLuc2 mMSTN transfected cells were cultured in DMEM with 10% FBS, and 1% Penicillin/streptomycin (10000 Unit/mL; Gibco). The L8 cells were plated in 96 well plate (Corning Inc., New York, United States). The final concentration of each well is 12000 cells/well. After the cells seeded for 24 h, cells were then treated with CMH ethanol extract. After 24 hrs treatment, the supernatant were collected for Luciferase activity assay. The cells were further performed MTT assay.

Protein extraction

The culture medium was removed and cells were washed three times with iced-cold PBS, pH 7.3. Then the cells were scraped with rubber policeman and collected in 500 μ L homogenization solution [20 mM Tris (pH7.5), 0.25 mM Sucrose, 1 mM EGTA, 5 mM EDTA, 1 mM phenylmethane sulfonyl fluoride (PMSF), 25 μ g/mL leupeptin, 50 mM 2-mercaptoethanol (2-Me), 1%NP-40]. After sonication, the lysate was centrifuged at 14000 xg for 20 min at 4°C. Cytosolic proteins were measured by the Bradford method using bovine serum albumin as standards (Bradford, 1976).

Western blot analysis

Lysate from the L8 myoblast was mixed with 6X sample buffer containing 62.5 mM Tris, 2% (w/v) sodium dodecyl sulfate, 10% (v/v) glycerol, 5% (v/v) 2-mercaptoethanol, and 0.0125% (w/v) bromophenol blue. The mixture was incubated at 100°C for 5 min and subjected to sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE). The separated proteins in the

gels were transferred onto the 0.45 μm polyvinylidene fluoride membrane. The membrane was incubated with a blocking solution consisting of 3% gelatin in TTBS (TBS with 0.05% tween20) for 1 hr at room temperature and hybridized with primary antibodies (myostatin 1:1000, GTX32624, GeneTex, California, United States; GAPDH 1:2000, MAB347, Chemicon, Burlington, Masschusettes, United States) for 1 hr at room temperature. And then, the membrane was incubated with horseradish peroxidase-conjugated secondary antibody for 1 hr at room temperature. Target bands were visualized using Bio-Rad Clarity Western ECL Substrate and detected by illuminance capturer (Vilber FUSION SL4 Spectra, France). The density of the specific band was determined using ImageJ image analysis software (National Institutes of Health, Bethesda, Maryland, United States).

Cell viability assay

MTT (5 mg/mL; 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide), Sigma-Aldrich, St. Louis, Missouri, United States) was prepared in Phosphate Buffered Saline (PBS buffer, 137 mM NaCl, 2.7 mM KCl, 4.3 mM Na_2HPO_4 , 1.47 mM KH_2PO_4 ; pH 7.4.) and filtered to sterilize and remove insoluble residue. Ten μL filtered MTT solution (yellow) was added to CMH treated cells in 96-well plate (Corning, New York, United States) and incubated at 37°C for 4 h. One hundred μL solubilization solution (10% SDS in 0.01M HCl) was added to each well and incubated for 24 hrs, and then used spectrostar nano microplate reader (BMG Labtech, Ortenberg, Germany) to read the plate with wavelength 595 nm (Denizot and Lang, 1986).

Luciferase assay

Luciferase substrate kit was obtained from Clontech (TaKaRa, Japan). To prepare the 10x substrate stock solution, lyophilized secreted luciferase substrate was added substrate buffer, and then diluted the 10x substrate stock solution 1:10 in reaction buffer. Fifteen μL culture media from each sample was transferred into the 96-well plate, and then add 2.5 μL of 1x substrate to each sample. The luciferase activities were read by luminometer (Mierbeta TriLux, Perkin Elmer, Waltham, Massachusetts, United States) after mixing well.

HPLC (high performance liquid chromatography)

The dried root of *Glycyrrhiza uralensis Fisch* was obtained from a traditional pharmaceutical store in Taichung, Taiwan.

Alcohol dissolved samples were removed large particles by centrifuging at 13000 rpm for 10 min before loading. The Reverse phase HPLC separation is accomplished by using an isocratic elution with flow rate of 4.0 mL/min. Column temperature was 30°C. The mobile phase consists of water with ethanol (5% to 100% ethanol in 30 min). The injection volume was 200 μL of each sample. The total run time was 50 minutes for each injection. Column 5C18 AR-II K54606 (10 mm x 250 mm) (Cosmosil, Nacalai Tesque Inc., Japan) was used to separate the licorice crude extract and licorice fraction 3 (G3). Column 5C18 AR-II K54524 (4.6 mm x 250 mm) (Cosmosil, Nacalai Tesque Inc., Japan) was used to separate licorice fraction 3-3 (G3-3).

Statistical analysis

All experiments were carried out three to five times in triplicates. SAS was utilized to analyze data. In western blot and luciferase quantification, the data

were analyzed by proc ANOVA post-hoc Tukey's Honest Significant Difference test ($P<0.05$). The others were analyzed by proc GLM post-hoc Dunnett's test ($P<0.05$).



RESULTS

Construction of pMetLuc2mMSTNPL

Mouse myostatin 5'-regulatory region was amplified PCR and then inserted into pMetLuc2 plasmid (Figure 6). The myostatin 5'-regulatory region was confirmed by sequencing. The result showed the sequence was identical to the sequence in GenBank (*mus musculus*, AX139025) (Figure 5). The plasmid is referred to pMetLuc2mMSTNPL. The expression of reporter gene, luciferase, in this plasmid is under the control of myostatin 5'-regulatory region.

Screening CMH for inhibition of luciferase activity by molecular biological screening platform

Plasmid pMetLuc2mMSTNPL was used to transfect L8 myoblasts. After G418 selection, the morphology of transfected cell did not change when compare to the origin cells. The transfected L8 cells were used to screen the CMH for the inhibition of luciferase activity. Total 27 CMH were used to screen for the inhibition of luciferase activity (Figure 7). Amount 27 CMHs, licorice, perilla, houttuynia cordata, and jiaogulan inhibited the activity of luciferase in transfected cells. In addition, the inhibition of luciferase activity is negative correlated with cell growth. Licorice ethanol extract decreased luciferase activity at 200 $\mu\text{g/mL}$ and 400 $\mu\text{g/mL}$ in transfected L8 cells. In the meantime, it increased L8 cell growth (Figure 8A). However, high concentration of licorice ethanol extract showed cytotoxicity in transfected cells. The perilla ethanol extract showed similar effect on the inhibition of luciferase activity and cell growth (Figure 8B). Perilla extract decreased luciferase activity at 2000 $\mu\text{g/mL}$ and 4000 $\mu\text{g/mL}$ and increase cell growth at 500 $\mu\text{g/mL}$, 1000 $\mu\text{g/mL}$ and 2000 $\mu\text{g/mL}$. *Houttuynia cordata*

ethanol extract increased transfected L8 cell growth at 1000 $\mu\text{g}/\text{mL}$, 2000 $\mu\text{g}/\text{mL}$ and 4000 $\mu\text{g}/\text{mL}$, and it decreased luciferase activity at 4000 $\mu\text{g}/\text{mL}$ and 8000 $\mu\text{g}/\text{mL}$ (Figure 9A). Jiaogulan ethanol extract increased L8 cell growth at 2000 $\mu\text{g}/\text{mL}$ and 4000 $\mu\text{g}/\text{mL}$, and it decreased luciferase activity at 2000 $\mu\text{g}/\text{mL}$ and 4000 $\mu\text{g}/\text{mL}$ (Figure 9B).

Positive correlation of luciferase activity with myostatin expression level in transfected L8 cells

To confirm myostatin protein quantity is positively correlated with luciferase activity in transfected L8 cells, Western blot analysis was performed. The transfected L8 cells were treated with licorice extract A (20 $\mu\text{g}/\text{mL}$) and B (10 $\mu\text{g}/\text{mL}$) for 24 h. The results showed that licorice A significantly decreased myostatin protein level ($p < 0.01$) (Figure10). In addition, the luciferase activity was inhibited by Licorice A (Figure11). The results demonstrate that the luciferase activity is positive correlated with myostatin protein level. It indicated that the luciferase expression through the pMetLuc2mMSTNPL plasmid in L8 cell can be the indicator of the myostatin protein level.

AY204900_Mus_ 1 CCTTTT-AAGTCCTAAGTCACACGGAGTTCTATGTCCTCAAAATGTTGCT 49
insert_seq. 1 CCTTTTAAAGTCCTAAGTCACACGGAGTTCTATGTCCTCAAAATGTTGCT 50
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insert_seq. 51 CAGCCTCTACCCTGTCTACCTGGATGATTTTCTCTCTCAAACCTGAGAGTC 100
AY204900_Mus_ 100 TGTAACCTATTAAGCATTAAGTACACACACACCCCTGACCCCAGCGGGCTC 149
insert_seq. 101 TGTAACCTATTAAGCATTAAGTACACACACACCCCTGACCCCAGCGGGCTC 150
AY204900_Mus_ 150 CATTCTCCATTCTCCCCTGTGCTTAAAAGAAGCTGCCCTGGAGTTTCAGT 199
insert_seq. 151 CATTCTCCATTCTCCCCTGTGCTTAAAAGAAGCTGCCCTGGAGTTTCAGT 200
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Figure 5. (continued)

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insert_seq.	651	AGTTGAGAACTAAAGATAAGAGAAGCTAAGTACTGTGCCGTCTTTGTCA	700
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insert_seq.	701	TCGACTTAGAAGAGGCCAAAATTGAGATTTGAACTCAGGTTTATTTGACTC	750
AY204900_Mus_	750	TTCAGTCTCAGCTCACAATGGCAGTACAGTCTAAAAAAAAAAAAAAAAATCAC	799
insert_seq.	751	TTCAGTCTCAGCTCACAATGGCAGTACAGTCTAAAAAAAAAAAAAAAAATCAC	800
AY204900_Mus_	800	AGGATCAATTTCCCTCTGAGGTATATAGCAGCATGTGTAATGATAATTATG	849
insert_seq.	801	AGGATCAATTTCCCTCTGAGGTATATAGCAGCATGTGTAATGATAATTATG	850
AY204900_Mus_	850	ACATCGAAAAGAATTCTATGCAGAAAATGAATTTCCAGACAAATCTGA	899
insert_seq.	851	ACATCGAAAAGAATTCTATGCAGAAAATGAATTTCCAGACAAATCTGA	900
AY204900_Mus_	900	CTTTATAGGCCTGCTCTAATATTGTCTTGTATAAAGAGGGCCAGATCACC	949
insert_seq.	901	CTTTATAGGCCTGCTCTAATATTGTCTTGTATAAAGAGGGCCAGATCACC	950
AY204900_Mus_	950	TCAGGGTGTCTGCTTTGTGTCTGGTTTTCCCTTCATCTTTAATGGTGGGCA	999
insert_seq.	951	TCAGGGTGTCTGCTTTGTGTCTGGTTTTCCCTTCATCTTTAATGGTGGGCA	1000
AY204900_Mus_	1000	AATCTAGTACATTATGGAAGCCCACTTTTTTTTTCCCTCAAGAGATATAGA	1049
insert_seq.	1001	AATCTAGTACATTATGGAAGCCCACTTTTTTTTTCCCTCAAGAGATATAGA	1050
AY204900_Mus_	1050	TGCCTCTTAAAAATTTGATGAAAATGCATTAACCTTTCAAGCTACTGAGC	1099
insert_seq.	1051	TGCCTCTTAAAAATTTGATGAAAATGCATTAACCTTTCAAGCTACTGAGC	1100
AY204900_Mus_	1100	TGCATTTTAGTTCACTGAGGCAGTAAATTGGGTGTATACTGTACAGGAAT	1149
insert_seq.	1101	TGCATTTTAGTTCACTGAGGCAGTAAATTGGGTGTATACTGTACAGGAAT	1150
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insert_seq.	1151	GGTGGTGACCTAAAAATAAATATTTGATACAAGCCACCATAGTCTCTTGG	1200
AY204900_Mus_	1200	GGTGTGTGTAAGGGGAGTAATGAATTAATAATCTAAAGACTCCTCAGCTT	1249
insert_seq.	1201	GGTGTGTGTAAGGGGAGTAATGAATTAATAATCTAAAGACTCCTCAGCTT	1250

Figure 5. (continued)

AY204900_Mus_	1250	CCCAAACAGGAGGAGGAACTCTGTGGCCTGGAAGCGTCCTCTGTCCCTGC	1299
insert_seq.	1251	CCCAAACAGGAGGAGGAACTCTGTGGCCTGGAAGCGTCCTCTGTCCCTGC	1300
AY204900_Mus_	1300	TGCTGTGTTTGTTCAGCTCTTTAAGAGTTCACCCCATTCGATCTTGTGGC	1349
insert_seq.	1301	TGCTGTGTTTGTTCAGCTCTTTAAGAGTTCACCCCATTCGATCTTGTGGC	1350
AY204900_Mus_	1350	TCCTAAAGCCAAGGGTGAAAGTTTGATCCTTGCAGAGGCCACTTAAATTC	1399
insert_seq.	1351	TCCTAAAGCCAAGGGTGAAAGTTTGATCCTTGCAGAGGCCACTTAAATTC	1400
AY204900_Mus_	1400	AGAGAACAAAAAGCACCATTCTCTGCCCTAGACTCTAGCCAGATCCCTG	1449
insert_seq.	1401	AGAGAACAAAAAGCACCATTCTCTGCCCTAGACTCTAGCCAGATCCCTG	1450
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AY204900_Mus_	1500	CCTGTGACAGTATGGGAACGCAACAAAGGACACCCCTCTACATGCGACTT	1549
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AY204900_Mus_	1550	GCTCTTTGTGTGCTCACGGGACCTGACATCATTCACAGAGAACACAGATT	1599
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AY204900_Mus_	1600	GCACTTTACTGTGTCAGCCCTGGAAGTCTGAGTCAAACGAAATAATGCTCC	1649
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AY204900_Mus_	1700	GCCTCCTCGCTGACAGGATTCTGTTGGCAATCAAAAAAAAAAAAAAAAAAAAA	1749
insert_seq.	1701	GCCTCCTCGCTGACAGGATTCTGTTGGCAATCAAAAAAAAAAAAAAAAAAAAA	1750
AY204900_Mus_	1750	AAAAGCAACACTCAGTCTTTAGTCTGTATCTCTGTAATAGAAAATAGCAA	1799
insert_seq.	1751	AAAAGCAACACTCAGTCTTTAGTCTGTATCTCTGTAATAGAAAATAGCAA	1800
AY204900_Mus_	1800	TACTTATAAGCTGAAATCAAGCACAGGTTTTATGTTAGTCAAAGCCATTA	1849
insert_seq.	1801	TACTTATAAGCTGAAATCAAGCACAGGTTTTATGTTAGTCAAAGCCATTA	1850

Figure 5. (continued)

AY204900_Mus_	1850	AGCTATCAAAAAGTAAACCCATGTACACAGAAACGTCCCAGGACTGGTTTG	1899
insert_seq.	1851	AGCTATCAAAAAGTAAACCCATGTACACAGAAACGTCCCAGGACTGGTTTG	1900
AY204900_Mus_	1900	TAATATGTCCTGACAAATAAGCCATGAAAACAAGCTCCTCAAATTACTGA	1949
insert_seq.	1901	TAATATGTCCTGACAAATAAGCCATGAAAACAAGCTCCTCAAATTACTGA	1950
AY204900_Mus_	1950	TGCAACTTTTTTAGCAGGGTCACAAACTCAGCTTTCTTTAAATTAAGTCAG	1999
insert_seq.	1951	TGCAACTTTTTTAGCAGGGTCACAAACTCAGCTTTCTTTAAATTAAGTCAG	2000
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insert_seq.	2001	CTCTTCCTAGTTTTTACTTCTCTAATTACCCAGCACTTAACGCATATTTT	2050
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insert_seq.	2301	CTCGACTGTAACAAAATACTGCTTGGTGACTTGTGACAGACAGGGTTTTA	2350
AY204900_Mus_	2350	ACCTCTGACAGCGAGATTCATTGTGGAGCAGGAGCCAATCATAGATCCTG	2399
insert_seq.	2351	ACCTCTGACAGCGAGATTCATTGTGGAGCAGGAGCCAATCATAGATCCTG	2400
AY204900_Mus_	2400	ACGACACTTGTCTCCTCTAAGTTGGAAATATAAAAAGCCACTTGAATACA	2449
insert_seq.	2401	ACGACACTTGTCTCCTCTAAGTTGGAAATATAAAAAGCCACTTGAATACA	2450
AY204900_Mus_	2450	GTATACAGGACTCCCTGG	2467
insert_seq.	2451	GTATACAGGACTCCCTGG	2468

Figure 5. The sequence of plasmid insert aligned with NCBI Mouse myostatin 5'-regulatory region (*Mus musculus*, AX139025). Labeled the location of E-Box, CAAT Box and TATA Box.

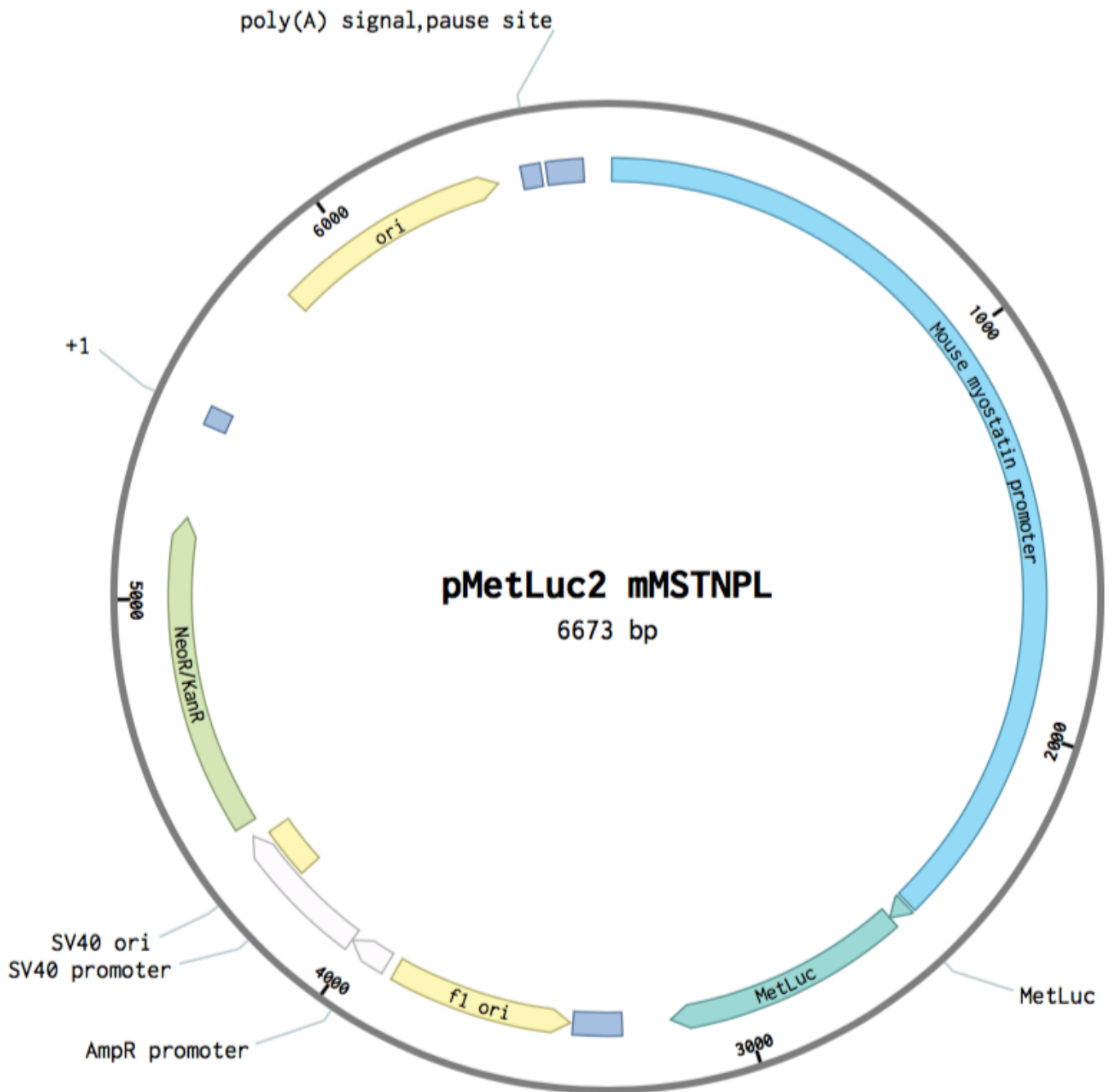


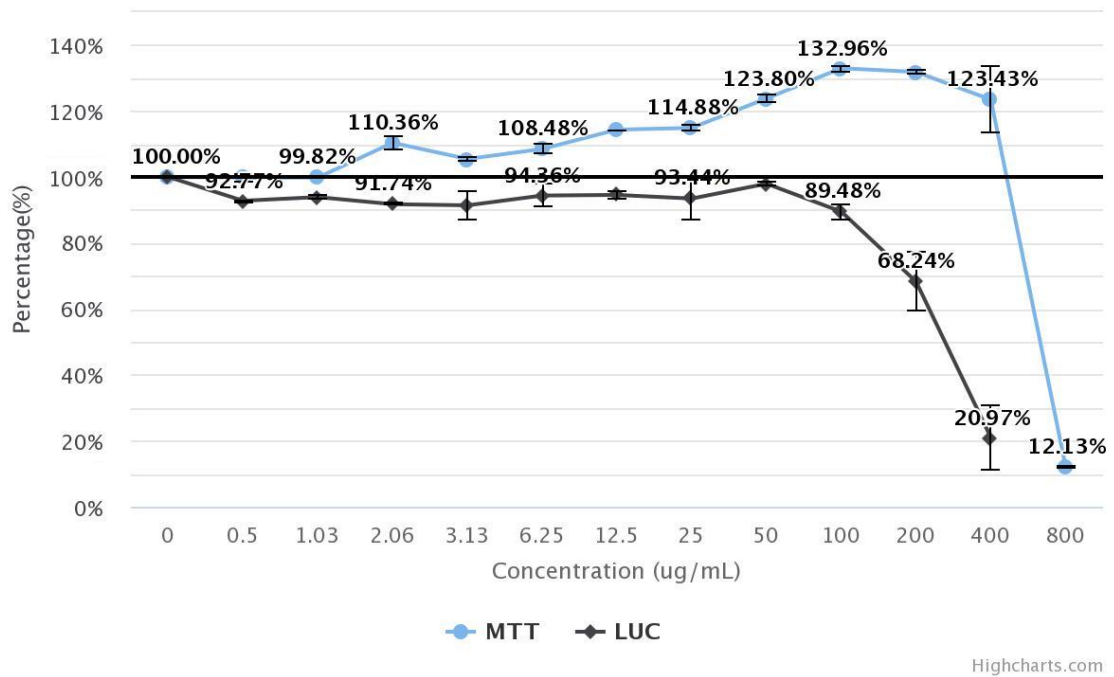
Figure 6. The plasmid map of pMetLuc2mMSTNPL. This plasmid was inserted a 2468 bp mouse myostatin 5'-regulatory region.

黃芩	白朮	甘草	艾葉	麥芽	杜仲	黃柏
Skullcap	Bai zhu	Licorice	Artemisia indica.	Malt	Eucommia ulmoides	Huang bai
小金英	牛蒡	黃耆	紫蘇	黨蔘	天麻	大蒜
Ixeris chinensis	greater burdock	Astragalus	Perilla	Codonopsis pilosula	Gastrodia elata	Allium sativum
玉竹	肉桂	麝香草	迷迭香	小茴香	肉豆蔻	秦皮
Solomon's Seal	Chinese cinnamon	Thyme	Rosemary	Fennel Bulbs	Nutmeg	Qinpi
洋香菜	魚腥草	陳皮	絞股藍	茯苓	川七	
Parsley	Houttuynia cordata	Chenpi	Jiaogulan	Wolfiporia extensa	Anredera cordifolia	

Figure 7. The Chinese name and common name of 27 CMHs. Black square: Herb showed inhibition of luciferase activity.

(A)

甘草(Licorice)



(B)

Perilla (紫蘇)

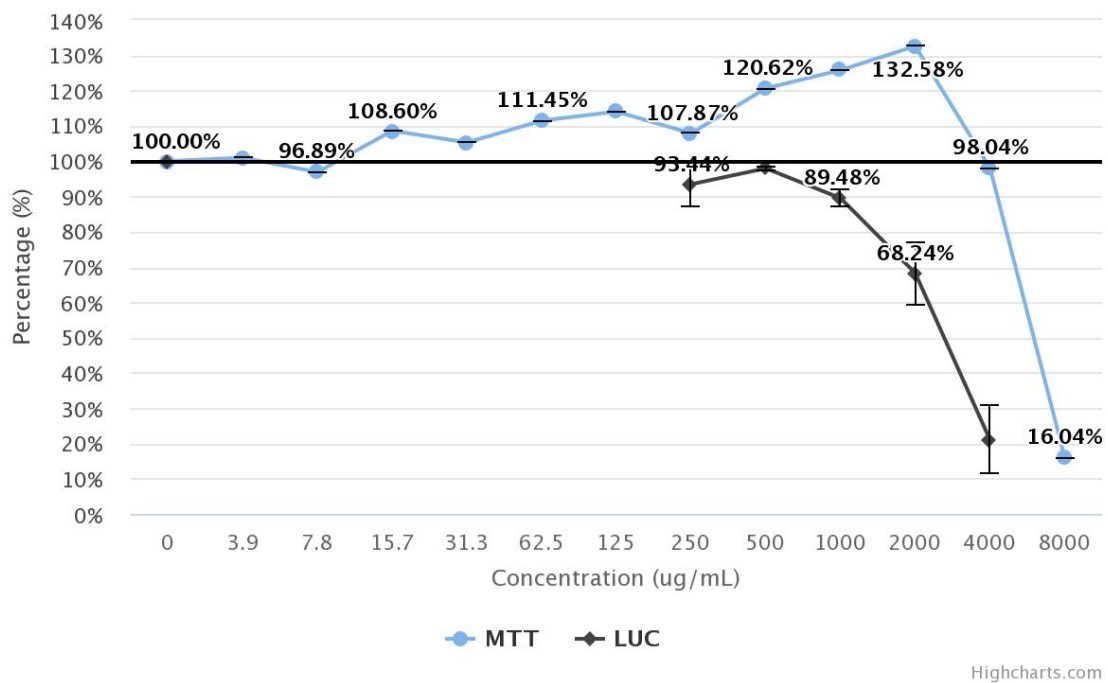
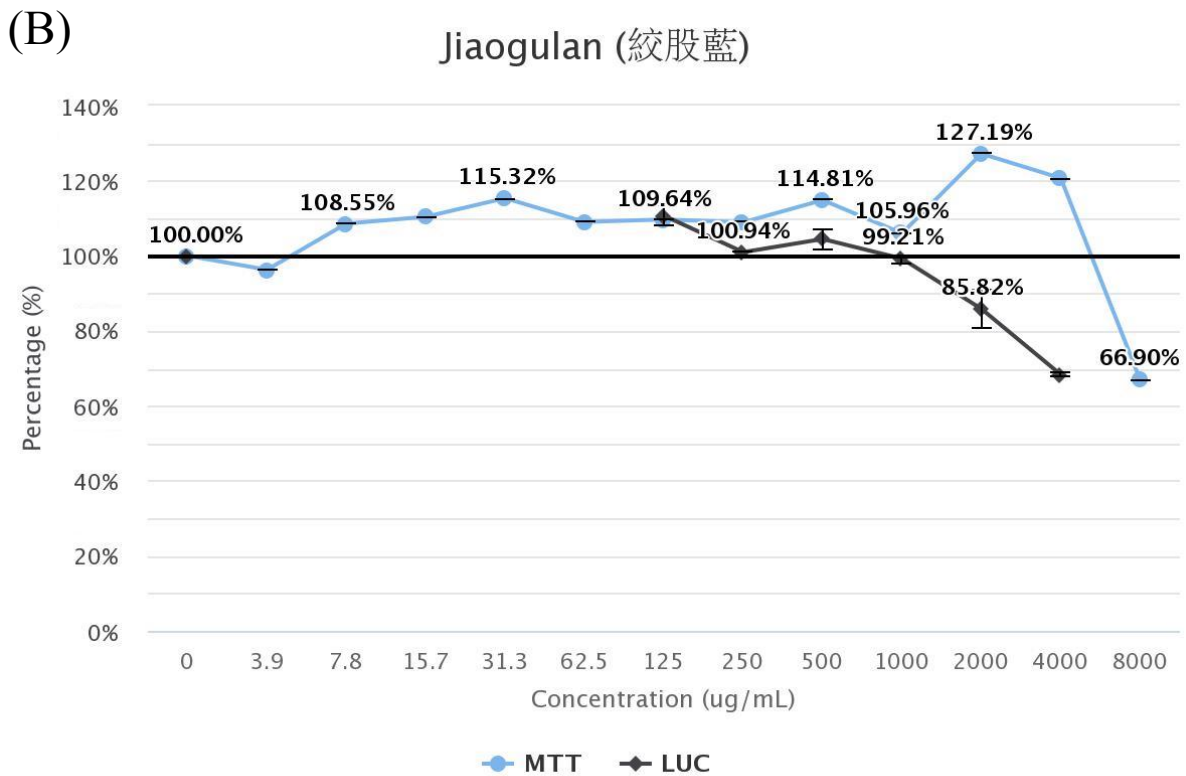
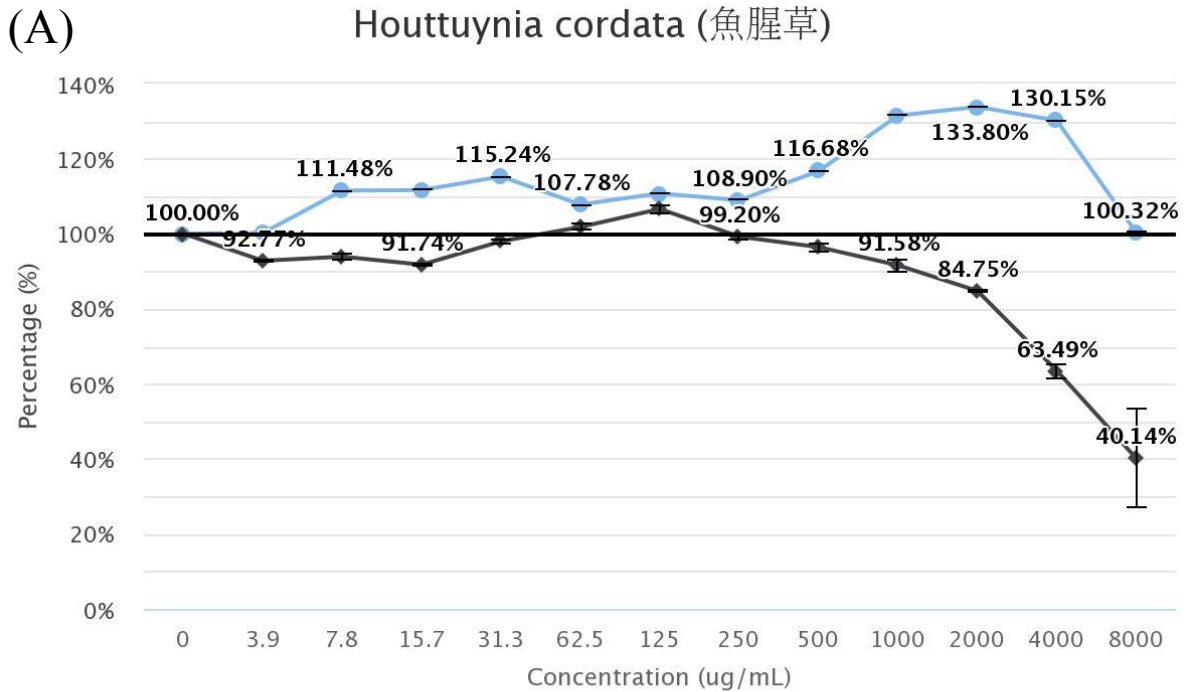


Figure 8. Effect of licorice ethanol extract (A) and perilla ethanol extract (B) on luciferase activity and cell growth in transfected L8 cells.



Highcharts.com

Figure 9. Effect of *Houttuynia cordata* ethanol extract (A) and jiaogulan ethanol extract (B) on luciferase activity and cell growth in transfected L8 cells.

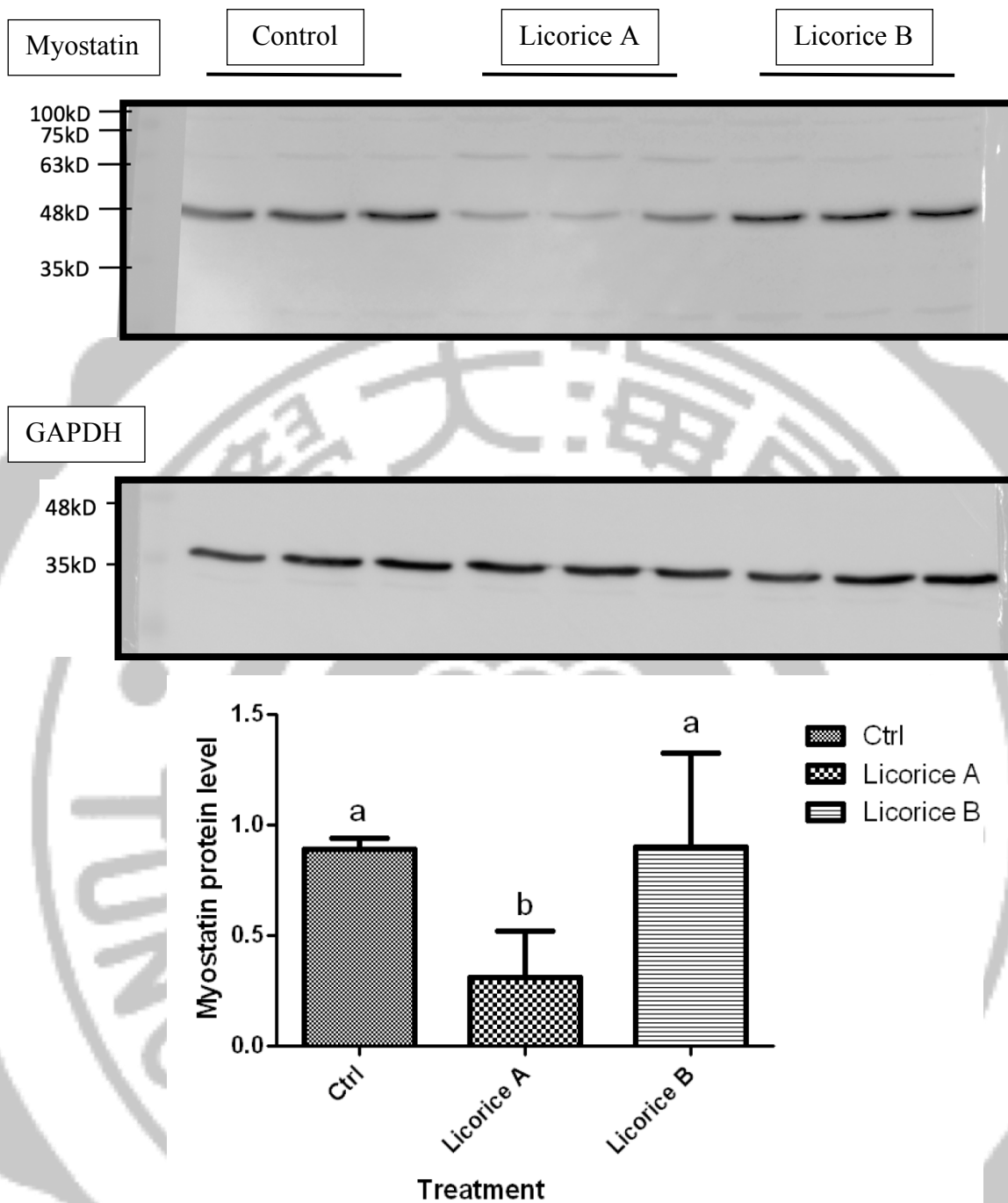


Figure 10. Effect of licorice ethanol extract on myostatin protein levels in transfected L8 cells. The cells were treated with different concentration of licorice ethanol extract (Licorice A: 20 $\mu\text{g}/\text{mL}$ and Licorice B: 10 $\mu\text{g}/\text{mL}$), and GAPDH served as internal control.

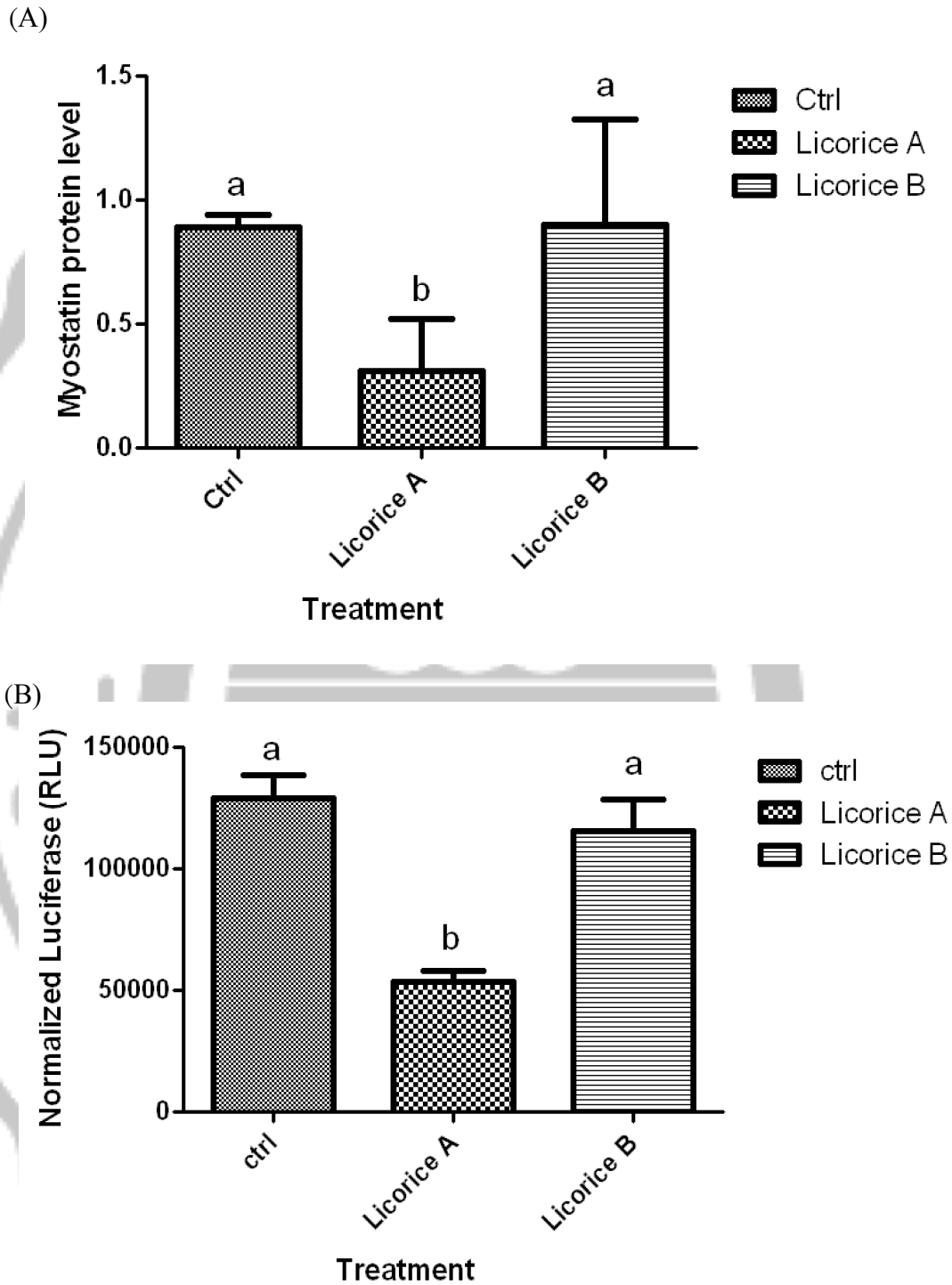


Figure 11. Effect of licorice ethanol extract on myostatin concentrations and luciferase activities. A: myostatin protein level from figure 10; B: luciferase activity

The isolation of active fraction from licorice ethanol extract by HPLC

To identify the active ingredient from licorice, high performance liquid chromatography (HPLC) was used. During the HPLC separation, 2.4 mL licorice ethanol extract was applied to polar solid phase and eluted in five fractions based on retention time, G1 (0-10 min), G2 (10-20 min), G3 (20-30 min), G4 (30-40 min), G5 (40-50 min) (Figure 12B). The ethanol gradient was from 5% to 100% in 30 min, and then 100% ethanol for another 15 min. The fractions were dried by vacuum centrifugation, then dissolved in ethanol (40 mg/mL). Each fraction was determined the inhibition of luciferase activity by the method described previously. Amount the 5 fractions, licorice fraction 3 (G3) significantly increased transfected L8 cell growth, and it decreased luciferase activity (Figure 13-15). Pure glabridin, which is the main compound of licorice ethanol extract, also applied to the screening platform. Glabridin increased L8 cell growth, in addition, luciferase activity was decreased by glabridin.

To identify the active ingredient in G3, G3 was further applied into polar solid phase and eluted in five fractions based on retention time, G3-1 (0-10 min), G3-2 (10-20 min), G3-3 (20-30 min), G3-4 (30-40 min), G3-5 (40-50 min) (Figure 16B.). The ethanol gradient for G3 was 45% for 5 min, 45% to 100% for 40 min, and then kept 100% for 5 min. The fractions were dried by vacuum centrifugation, then dissolved in ethanol (40 mg/mL). Each fraction was determined the inhibition of luciferase activity by the method described previously. Licorice fraction G3-2 and G3-3 increased transfected L8 cell growth, in the same time, they inhibited luciferase activity in transfected L8 cell (Figure 17). The glabridin was applied into the same column with the same elution condition. The result showed that the retention time for glabridin is before 20 min (Figure 16A). This result imply that glabridin may contribute the effect of G3-2

on cell growth and luciferase activity, while G3-3 contains other unknown component(s) affected the cell growth and luciferase activity.



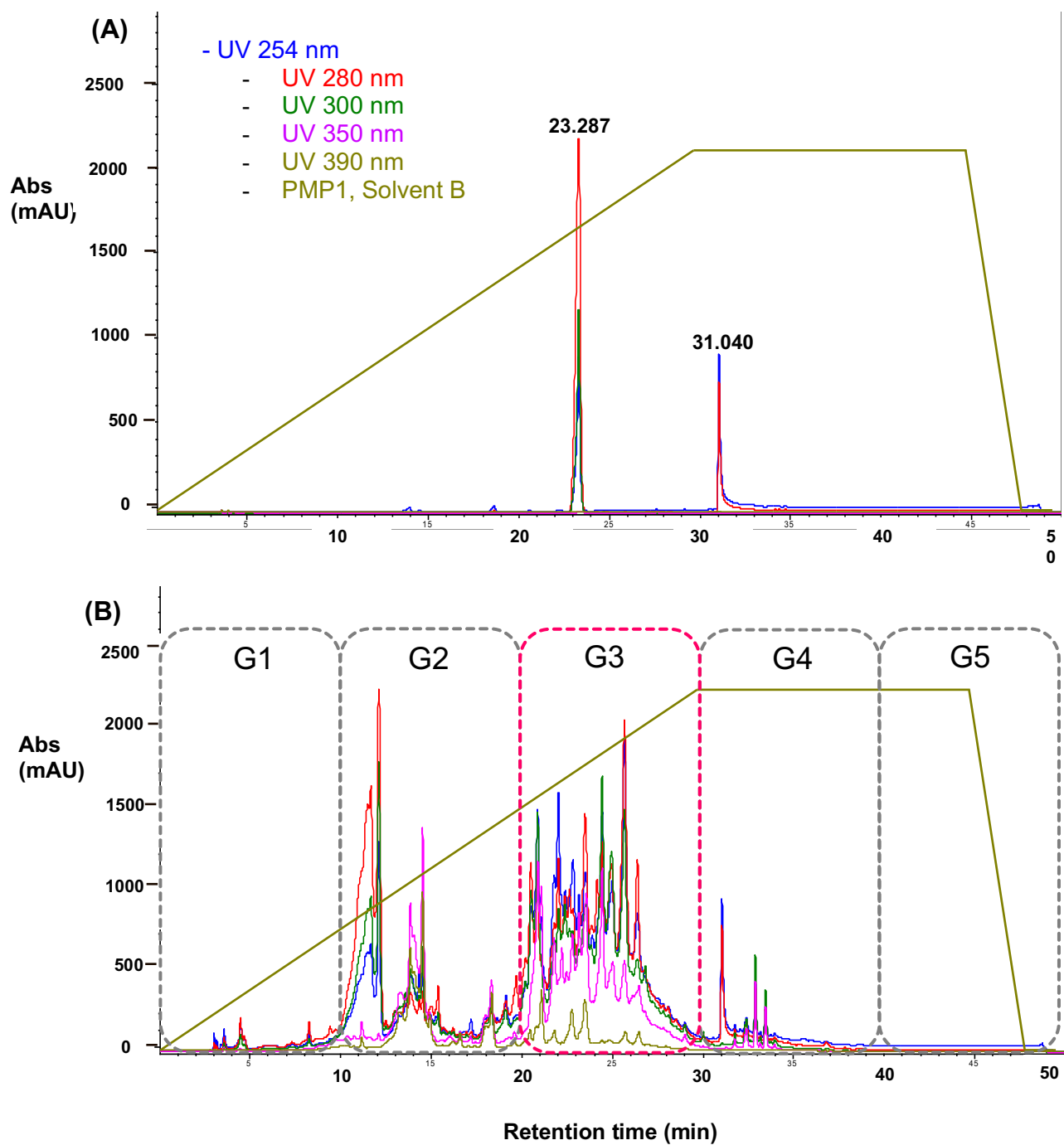


Figure 12. HPLC profile of glabridin (A) and licorice ethanol extract (B). The elution was separated in five fraction (10 min for each).

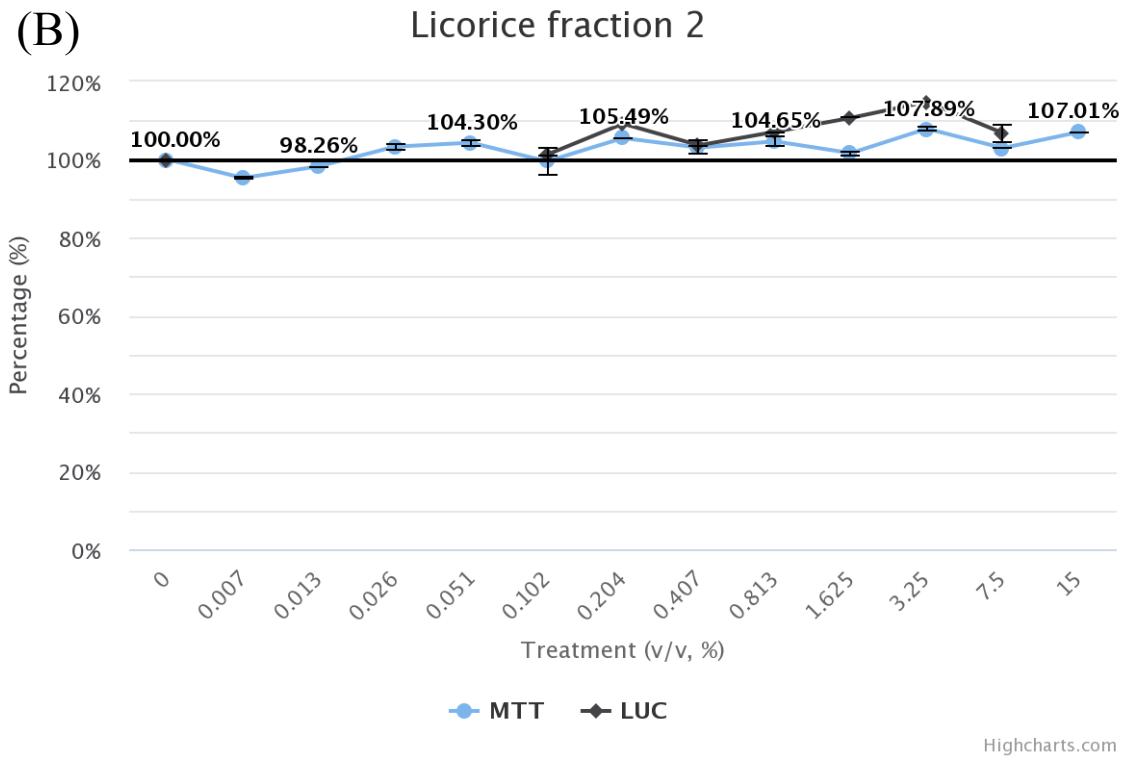
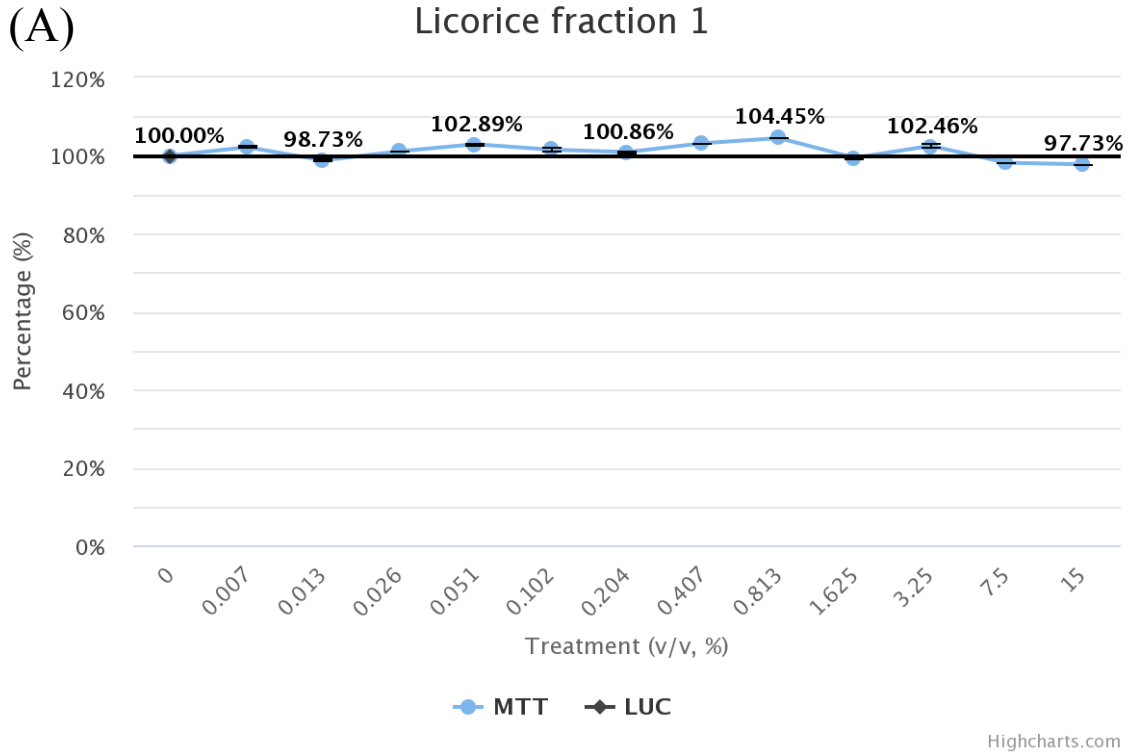


Figure 13. Effect of licorice fraction 1 and fraction 2 by HPLC on cell growth and luciferase activity in transfected L8 cells.

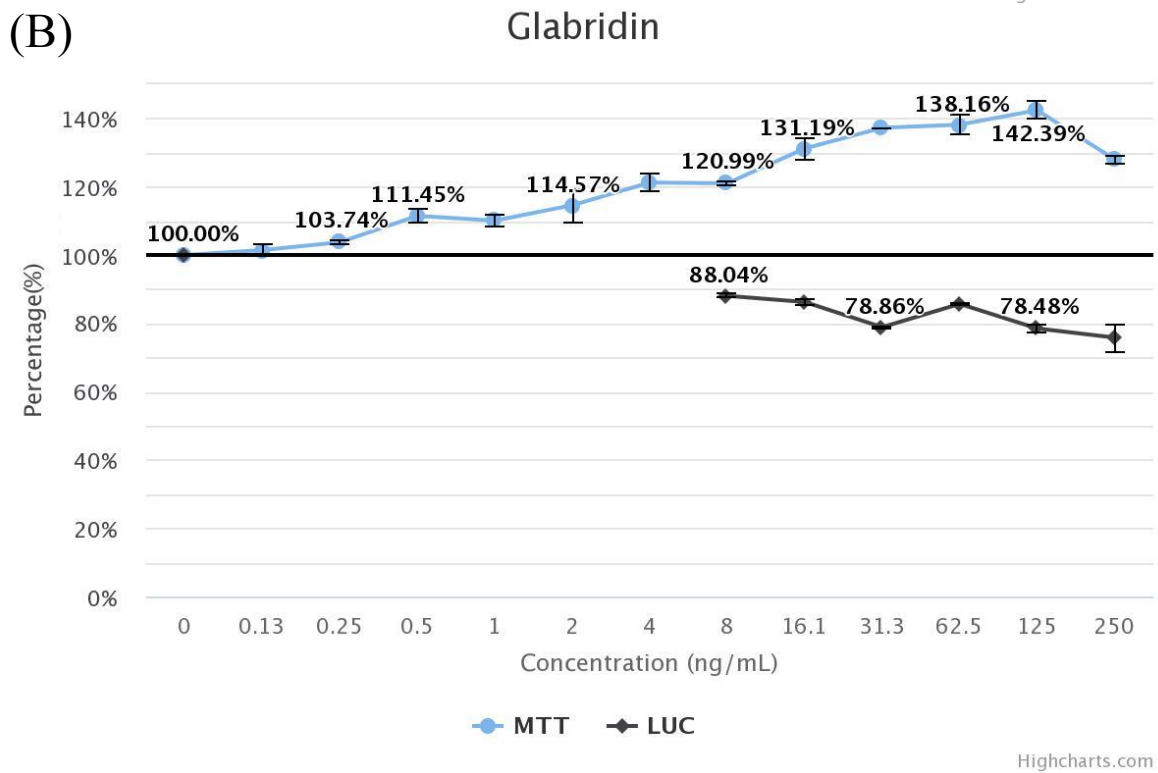
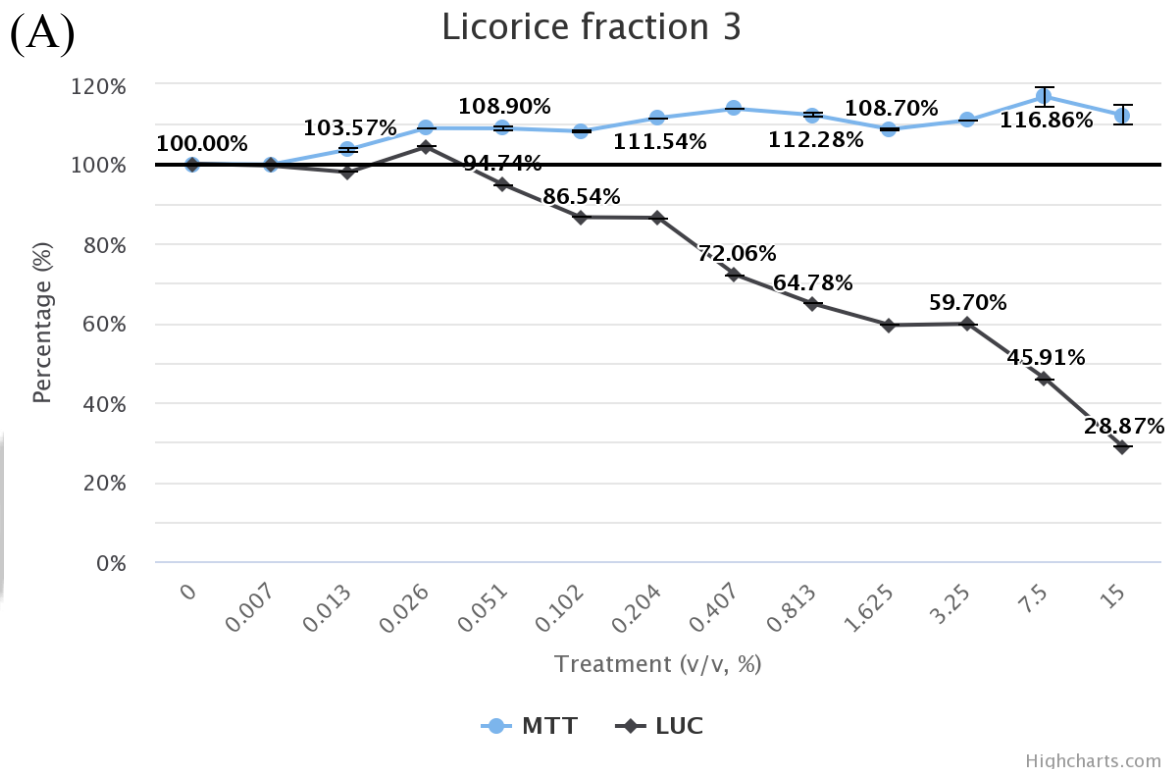


Figure 14. Effect of licorice fraction 3 and glabridin by HPLC on cell growth and luciferase activity in transfected L8 cells.

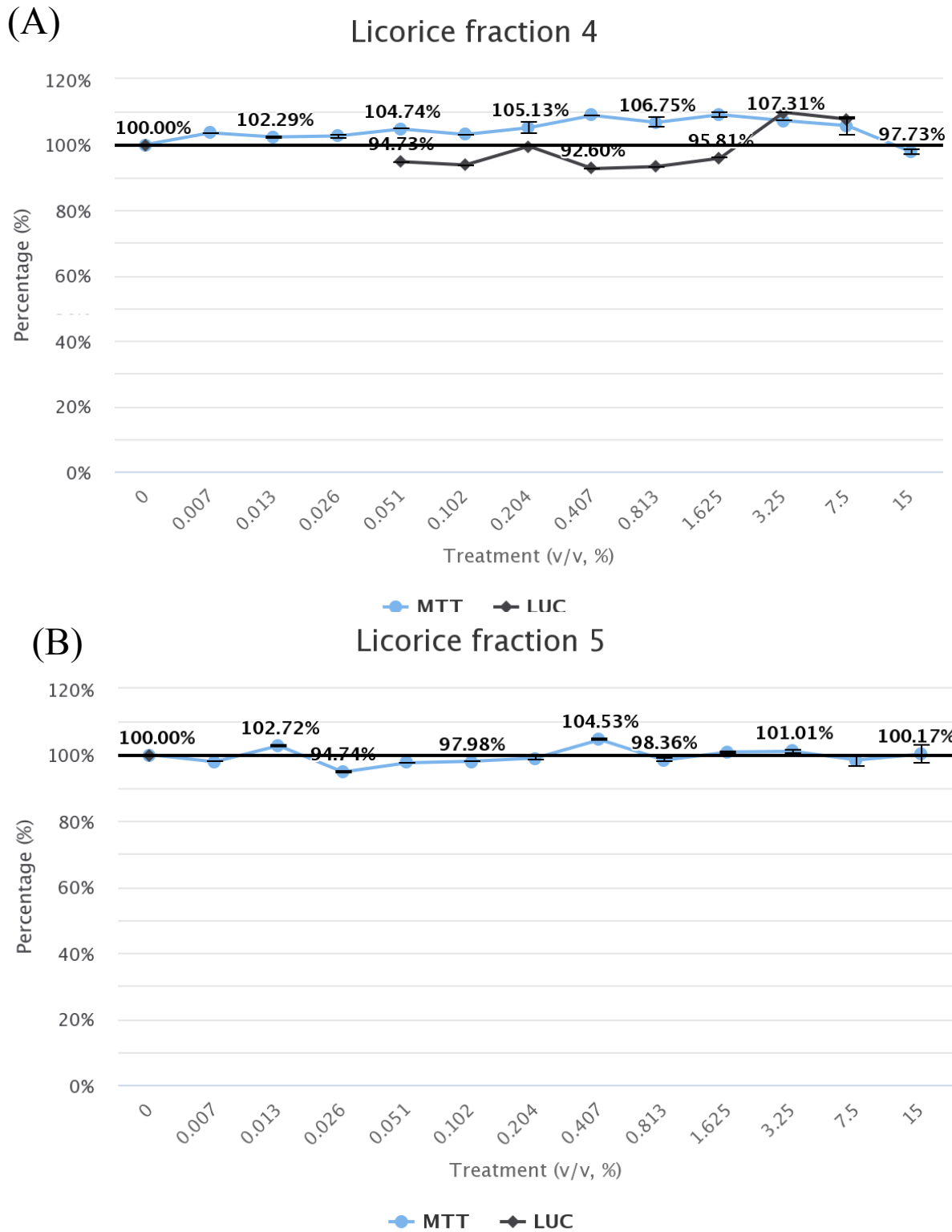


Figure 15. Effect of licorice fraction 4 and fraction 5 by HPLC on cell growth and luciferase activity in transfected L8 cells.

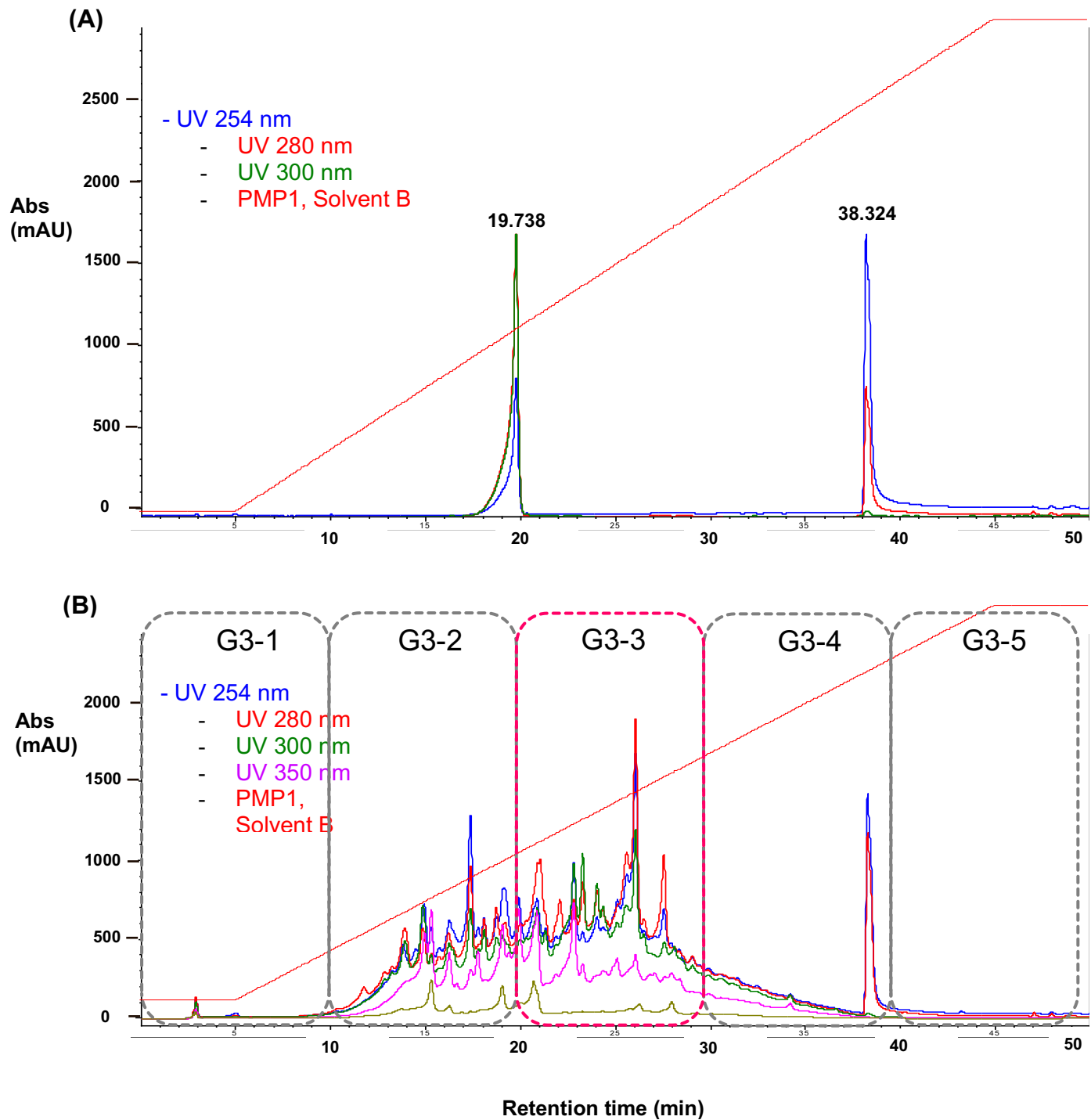
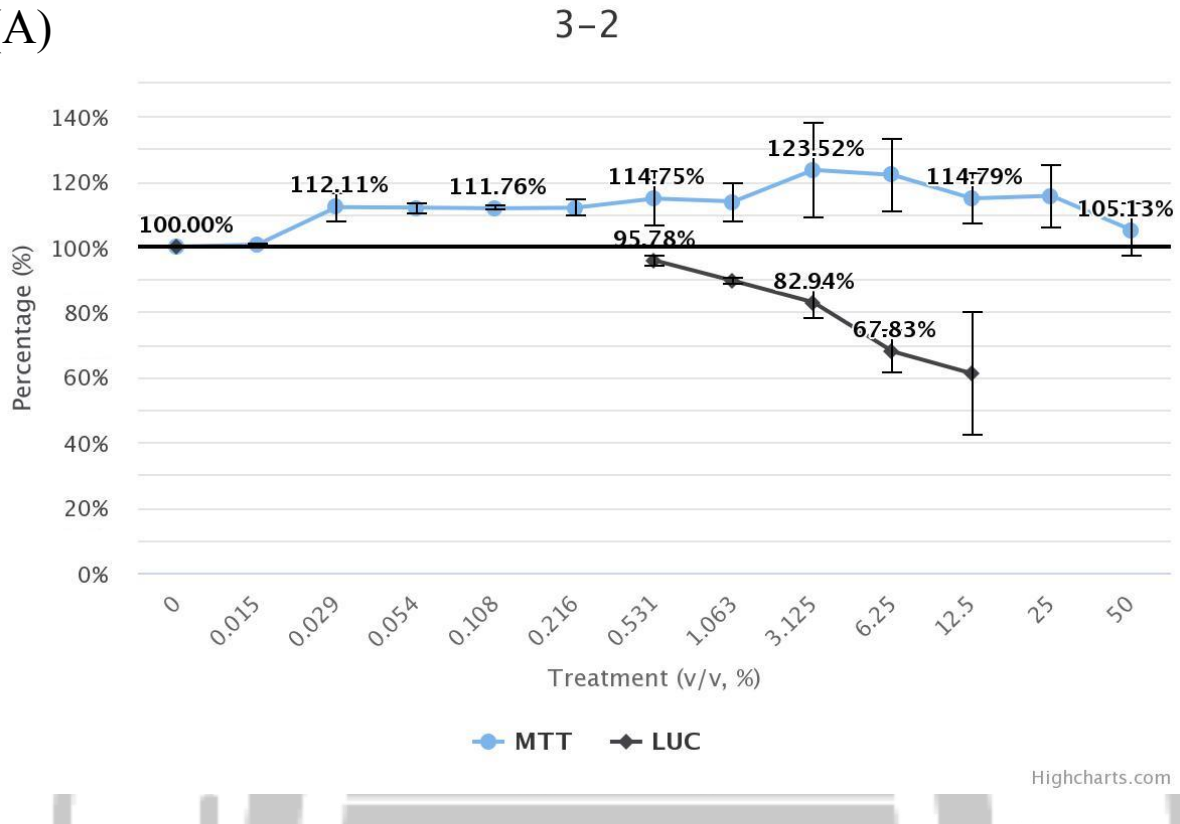


Figure 16. HPLC elution profile of glabridin (A) and licorice fraction G3 (B).

The elution was separated in five fractions (10 min for each).

(A)



(B)

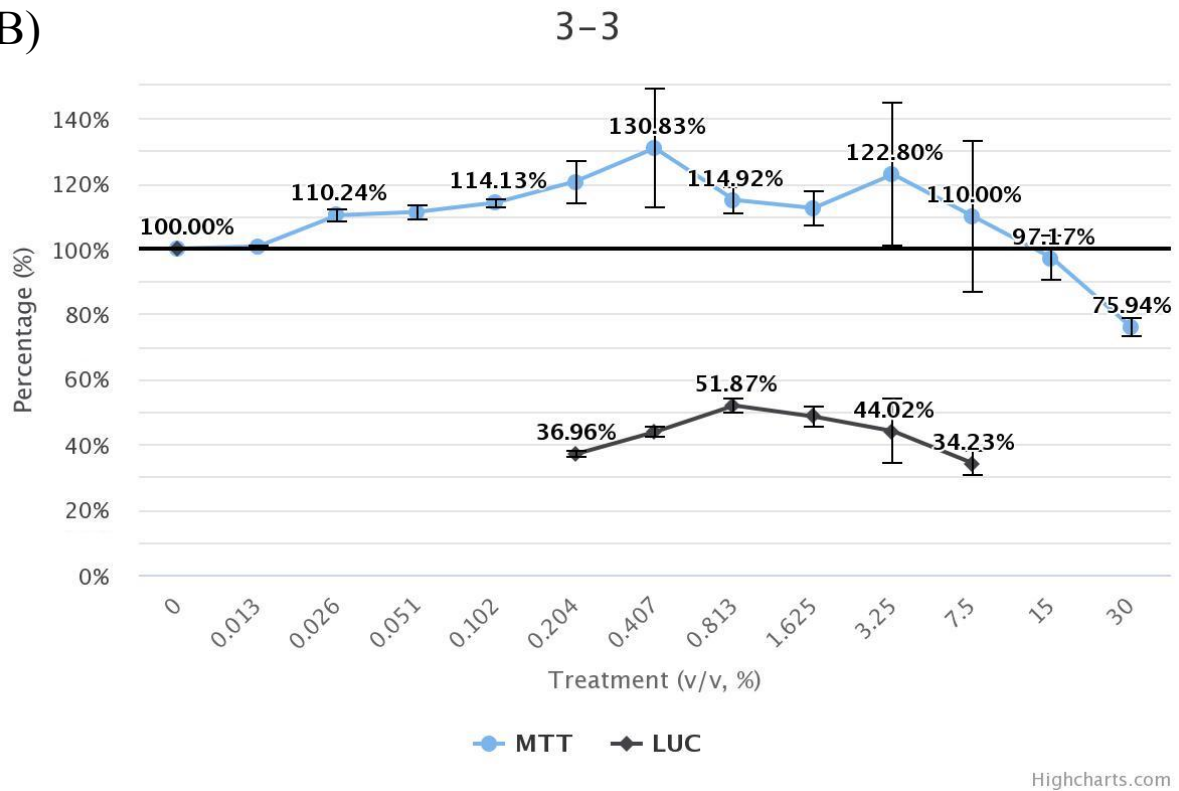


Figure 17. Effect of licorice fractions G3-2 and G3-3 on cell growth and luciferase activity in L8 cells.

DISCUSSION

Myostatin, is one of the major regulators of skeletal muscle development. It is a member of the transforming growth factor- β (TGF- β) superfamily. Myostatin plays a negative regulator in skeletal muscle growth. Overexpression of myostatin induces skeletal muscle atrophy and the mutations in the myostatin-encoding region are response for muscle hypertrophy and hyperplasia (Swatland and Kieffe, 1974). Passive vaccination and monoclonal antibody were used to increase animal growth and induced muscle hypertrophy (Bogdanovich *et al.*, 2002; Liang *et al.*, 2007). However, these methods were too time consuming and labor intensive. CMHs were used as daily health care or therapy in Asia for long time. Recently, more and more CMHs were added in animal feed to enhance animal growth. Unfortunately, usage of CMHs in animal growth does not have systemic analysis, so, it is important to establish a simple platform to screen the effective CMHs before using in animal feed.

Myostatin is one of major factor involved in muscle growth. Effective CMHs on regulation of myostatin gene expression are the candidate for animal growth enhancer. To establish a myostatin gene expression regulator screen system, mouse myostatin 5' regulatory region was inserted into pMetLuc2 plasmid (Figure 6). The plasmids were transfected onto L8 myoblasts and selected by G418. The expression of reporter gene, luciferase, in L8 myoblasts is under the control of myostatin 5'-regulatory region.

Twenty seven CMHs were screened by L8-myostatin molecular scanning system. Amount 27 CMHs, licorice, perilla, houttuynia cordata, and jiaogulan inhibited the activity of luciferase in transfected cells. In addition, the inhibition of luciferase activity is correlated with the enhancement of cell growth. These

results implied that these 4 CMHs promoted the L8 myoblast cell growth mediated by the inhibition of myostatin gene expression (Figure 8-9).

To confirm myostatin protein concentration is positive correlated with luciferase activity in transfected L8 cells, Western blot analysis was used. The transfected L8 cells were treated with licorice extract A (20 $\mu\text{g/mL}$) and B (10 $\mu\text{g/mL}$) for 24 h. The results showed that licorice A significantly decreased myostatin protein level. In addition, the luciferase activity was inhibited by Licorice A (Figure 10). The results imply that the luciferase activity is positive correlated with myostatin protein level (Figure 11). These results demonstrated that the luciferase expression through the pMetLuc2mMSTNPL plasmid in L8 cell can be the indicator of the myostatin protein level.

Licorice flavonoid oil (LFO) diet supplement significantly improved elderly body trunk muscle mass and body fat mass (Kinoshita *et al.*, 2017). Recently, one of licorice major ingredient, glabridin, was found to inhibit dexamethasone-induced protein degradation to prevent C₂C₁₂ myotubes and the muscle atrophy in mice (Yoshioka *et al.*, 2019). Glabridin is antagonist of dexamethasone. It binds to glucocorticoid receptor to block the action of glucocorticoid. In this study, licorice ethanol extract inhibited the L8 myoblasts cell proliferation mediated by the inhibition of myostatin gene expression (Figure 8A). However, glabridin enhanced the L8 myoblast proliferation, but did not decrease the myostatin gene expression (Figure 14B). To identify the active ingredients in licorice, HPLC was used to perform the separation of licorice components. Enhancement of cell proliferation/inhibition of myostatin gene expression activities were identified in Fraction 3 (G3) (Figure 14A). In the chromatogram profile, G3 and glabridin are overlapped (Figure 12). To confirm the G3 active compound is not glabridin, G3 was applied to HPLC again. Licorice

fraction G3-2 and G3-3 increased transfected L8 cell growth, and inhibited luciferase activity in transfected L8 cell (Figure 17). The glabridin was applied into the same column with the same elution condition. The result showed that the retention time for glabridin is before 20 min (Figure 16A). This result implies that glabridin may contribute the effect of G3-2 on cell growth, while G3-3 contains other unknown component(s) affected the myoblast cell growth and luciferase activity.

In the future, identification of active ingredient in G3-3 of licorice will be carry on by LC-MS. Animal study will also be conducted. In addition, more CMHs will be screened by this molecular screening system to find more nature animal growth enhancer in CMHs.

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