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Evidence-based research on traditional Japanese medicine, Kampo, in treatment of gastrointestinal cancer in JapanJianjun Gao^{1,2}, Peipei Song¹, Fanghua Qi¹, Norihiro Kokudo¹, Xianjun Qu², Wei Tang^{1,2,*}¹ Hepato-Biliary-Pancreatic Surgery Division, Department of Surgery, Graduate School of Medicine, The University of Tokyo, Tokyo, Japan;² Shandong University China-Japan Cooperation Center for Drug Discovery & Screening, Ji'nan, Shandong, China.

ABSTRACT: Gastrointestinal cancer is a great threat to human health in Japan. Conventional anti-cancer therapies including surgery, radiation, and chemotherapy are the main strategies and play important roles in curing this disease or extending the life of patients with these cancers. On the other hand, patients undergo great suffering induced by these treatments. Kampo, the Japanese traditional medicine, has been used in clinics to reduce side effects and to improve the quality of life of gastrointestinal cancer patients in Japan. In order to testify to the efficacy and safety of these Kampo medicines and to clarify the underlying mechanisms, a number of clinical and basic studies were implemented in the past several decades. These studies suggested the benefits of Kampo medicine as an adjuvant to conventional anti-cancer therapies in treating gastrointestinal cancer. Since the safety and efficacy as well as quality control of traditional medicine have long been focused worldwide, the development course of Kampo medicine may provide reference to other countries in the world.

Keywords: Kampo, traditional medicine, gastrointestinal cancer, side effects, quality of life

1. Introduction

Traditional medicine, defined in contrast to Western medicine, is the sum total of knowledge, skills and practices based on theories, beliefs and experiences indigenous to different cultures that are used to maintain health, as well as to prevent, diagnose, improve or

treat physical and mental illnesses (1). In the world, traditional medicine is generally available, affordable, and commonly used in large parts of Asia, Africa, and Latin America (2). In some of the African countries such as Uganda, Tanzania and Ethiopia, 60-90% of the population depends on traditional medicine for primary health care at the present time (3). Moreover, the use of traditional medicine as complementary/alternative medicine (CAM) has been reported in 70-90% of populations in developed countries such as Canada, France, Germany and Italy (3). In Asia, the use of traditional medicine in Japan (Kampo) has a history of thousands of years and plays an important role to insure human health. Indeed, it is reality that traditional medicine can treat various chronic conditions (4). However, evidence of safety and efficacy as well as quality control of traditional medicine have long been focused and highlighted by policy-makers, medical professionals and/or the public in many countries in the world (5-9).

Cancer has been the leading cause of death in Japan since 1981 (10). In 2010, a total of 353,000 people died of this disease, accounting for one in every three deaths, according to the statistics published by the Japan Ministry of Health, Labor and Welfare. Among causes of cancer deaths in 2009, stomach, colon, rectum, liver, gallbladder and bile ducts, and pancreas cancers lead to 169,932 deaths which make up 49.4% of all deaths induced by malignant tumors (11). Thus gastrointestinal tumors are extremely serious cancer types and the most common causes of cancer-related death in Japan at the present stage. Due to early detection by mass screening and application of modern cancer treatment strategies including surgery, radiation, and chemotherapy, the 5-year survival rates of patients with gastrointestinal cancers have shown steady improvement since 1962 according to the hospital-based cancer registry of Japan National Cancer Center Hospital (12). On the other hand, those conventional therapies inflict great suffering and require stoic endurance on the part of the patients. In this context, Kampo was introduced into gastrointestinal tumorthrapy, aiming to deal with problems such as the side effects of radiotherapy and chemotherapy and various types of general malaise (13-15).

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2. Research on Kampo medicine in treatment of gastrointestinal cancer

Kampo medicine plays an important role in cancer prevention in high-risk groups, enhancement of tumor immunity, improvement of general condition after operations, and attenuation of adverse reactions to chemotherapy and radiation. Although several herbs were supposed to have direct antitumor activity, Kampo medicine mainly acts as a kind of adjuvant to conventional antitumor treatments in clinical practice, contributing to the maintenance of a good quality of life for cancer patients (16). In this regard, Kampo formulas such as Juzentaihoto, Ninjin'yoeito, Hochuekkito, Shosaikoto, Daikenchuto, and Hangeshashinto are often prescribed to patients with gastrointestinal cancers in Japan (Table 1) (17-21).

2.1. Clinical studies on Kampo medicine used for care of gastrointestinal cancer

The approval of the currently used Kampo for application in clinics is without rigorous clinical trials from phases I to III but simply based on that Kampo medicine had passed the test of a thousand years of historical experience in Japan (22). Although the efficacy of Kampo is believed by many prescribers and patients, objective evaluation of those Kampo medicines to attain scientific evidence

is trend-driven. In this regard, a number of clinical trials were carried out to testify to the efficacy and safety of Kampo in Japan. In order to disseminate evidence-based medicine in Kampo products, the Japan Society for Oriental Medicine established the Special Committee for Evidence-based Medicine (EBM) in 2001, aiming to present evidence from "good" studies of Kampo products such as randomized controlled trials (RCTs) published from 1986 when the quality of Kampo formulations reached the present levels. According to their studies, based on the criteria that *i*) studies that employed Kampo formulations are approved by Japan government as a research object, *ii*) studies that used rational methods for experimental design including randomized controlled trials (RCTs), quasi-randomized controlled trials (quasi-RCTs), crossover trials, and meta-analyses, it was demonstrated that there are a total of 359 randomized controlled clinical trials and 1 meta-analysis on Kampo products from the year 1986 (including 1986) to June 2010 (23). Based on that evidence, fifty nine Kampo medicines were recommended and included in Japanese Clinical Practice Guidelines by March 31, 2011 (24).

The efficacy of Kampo medicine in preventing cancer occurrence, enhancing the immune system, and/or reducing the side effects of conventional therapies has been testified through EBM based clinical trials (Table 2). The combination of operations with chemotherapy is the most common treatment strategy

Table 1. Kampo medicines that are currently used for treatment of gastrointestinal cancers in Japan

Kampo	Formulation*	Composition of crude drugs**	Approved indications
Juzentaihoto	Granules	Astragalus Root, Cinnamon Bark, Rehmannia Root, Peony Root, Cnidium Rhizome, Atractylodes Lancea Rhizome, Japanese Angelica Root, Ginseng, Poria Sclerotium, Glycyrrhiza	Relief of declined constitution after recovery from disease, fatigue and malaise, anorexia, perspiration during sleep, cold limbs, and anemia.
Ninjin'yoeito	Granules	Rehmannia Root, Japanese Angelica Root, Atractylodes Rhizome, Poria Sclerotium, Ginseng, Cinnamon Bark, Polygala Root, Peony Root, Citrus Unshiu Peel, Astragalus Root, Glycyrrhiza, Schisandra Fruit	Relief of declined constitution after recovery from disease, fatigue and malaise, anorexia, perspiration during sleep, cold limbs, and anemia.
Hochuekkito	Granules	Astragalus Root, Atractylodes Lancea Rhizome, Ginseng, Japanese Angelica Root, Bupleurum Root, Jujube, Citrus Unshiu Peel, Glycyrrhiza, Cimicifuga Rhizome, Ginger	Indicated for the symptoms/conditions of patients having delicate constitution, reduced digestive functions, and severe fatigability of limbs.
Shosaikoto	Granules	Bupleurum Root, Pinellia Tuber, Scutellaria Root, Jujube, Ginseng, Glycyrrhiza, Ginger	Relief of the following symptoms of those patients with moderately strong constitution, right upper abdominal tenderness accompanied by fullness and discomfort, coated tongue, oral cavity discomfort, anorexia, and/or those with slight fever and nausea; Improvement of liver dysfunction due to chronic hepatitis.
Daikenchuto	Granules	Processed Ginger, Ginseng, Zanthoxylum Fruit	Relief of abdominal cold feeling and pain accompanied by abdominal flatulence.
Hangeshashinto	Granules	Pinellia Tuber, Scutellaria Root, Processed Ginger, Glycyrrhiza, Jujube, Ginseng, Coptis Rhizome	Relief of the symptoms of those patients with blocked feeling in the stomach pit and occasional nausea, vomiting, anorexia, borborygmus, and a tendency to loose stools or diarrhea.

* Granules are a mixture of extract of crude drugs indicated and certain medicinal accessory materials.

** All the crude drugs comply with the Japanese Pharmacopoeia.

Table 2. Clinical trials of Kampo in treatment of gastrointestinal cancer in Japan

Kampo Formula	Tumor	Study Design	Study Purpose	Conclusion	Ref.
Juzentaihoto	Gastric cancer	RCT-envelope 33 patients	Improvement effect on host-immunity in patients undergoing postoperative adjuvant chemotherapy (tegafur/uracil (UFT) 300 mg/day).	Juzentaihoto is useful in gastric cancer patients undergoing postoperative adjuvant UFT.	27
Juzentaihoto	Gastric cancer	RCT 94 patients	Efficacy of Juzentaihoto combined with oral 5-fluorouracil (5-FU) as postoperative adjuvant chemotherapy in patients with surgically treated gastric cancer.	Combination of Juzentaihoto with oral 5-FU is effective for patients with surgically treated stage III or IV gastric cancer.	28
Juzentaihoto	Colorectal cancer	RCT-envelope 44 patients	To elucidate the mechanism by which Juzentaihoto reduces the adverse reaction to treatment with 5-FU (hepatopathy) by determining the distribution of 5-FU in tissues of patients with colorectal cancer receiving slow-release tegafur preoperatively.	Administration of Juzentaihoto in patients receiving slow-release tegafur capsules increases 5-FU concentration in tumor tissues but decreases 5-FU concentration in normal tissues, enhancing the tumor selectivity of tegafur. This effect may be partly due to the modulation by Juzentaihoto of thymidine phosphorylase activity in tissues and of cytochrome P-450.	36
Juzentaihoto	Colorectal cancer	RCT 168 patients	To evaluate the clinical efficacy of Juzentaihoto for the prevention of postoperative recurrence of colorectal cancer.	Juzentaihoto may have a metastasis-suppressive effect, but since these are interim reports, the follow-up is still ongoing.	37 38
Juzentaihoto	Hepatocellular carcinoma	RCT 20 patients	To evaluate the effect of Juzentaihoto for reducing adverse effects of spongel + lipiodol + phosphatidyl choline + cisplatin treatment in transarterial embolization (TAE) for hepatocellular carcinoma.	Juzentaihoto significantly suppresses nausea/vomiting after TAE with spongel + lipiodol + phosphatidyl choline + cisplatin for hepatocellular carcinoma.	34
Juzentaihoto	Esophageal, gastric, or colorectal cancer	RCT-envelope 174 patients	Effect on the cell-mediated immunity of postoperative patients with esophageal, gastric, or colorectal cancer.	Juzentaihoto postoperatively administered for treatment of esophageal, gastric, or colorectal cancer may act as a biological response modifier (BRM).	39
Juzentaihoto	Gastric and colorectal cancer	RCT-envelope 284 patients	To evaluate the efficacy of Juzentaihoto for reducing adverse effects and improving quality of life (QOL) in postoperative patients undergoing chemotherapy (UFT 4 capsules/day) for gastric, colorectal, or breast cancer (curative resection/non-curative resection).	Juzentaihoto reduces the number of adverse drug reactions and improves QOL in postoperative patients on chemotherapy (UFT 4 capsules/day) for gastric, colorectal, or breast cancer.	29 30
Ninjin'yoeito	Gastric cancer	RCT-envelope 46 patients	Efficacy for reducing adverse effects and improving performance status in patients undergoing postoperative adjuvant chemotherapy (fluoropyrimidine anticancer drug).	Ninjin'yoeito tends to suppress the decreases in RBC count and platelet count but not the decrease in WBC count and does not improve performance status in patients undergoing postoperative adjuvant fluoropyrimidine-based chemotherapy for gastric cancer.	31
Ninjin'yoeito	Colorectal cancer	RCT-envelope 23 patients	Immunostimulation and improvement of nutritional status in postoperative patients with colorectal cancer.	Ninjin'yoeito significantly promotes improvement of lymphocyte count and PHA-stimulated lymphocyte proliferation in postoperative patients with colorectal cancer, suggesting its role as a possible biological response modifier.	40
Shosaikoto	Colorectal cancer	RCT 20 patients	Immunostimulation and suppression of liver metastasis in postoperative patients with colorectal cancer.	Saiko agents increased PHA-stimulated lymphocyte proliferation and NK cell activity, evaluated by CD57 and CD16, suggesting its immunostimulating effect.	41
Shosaikoto	Liver cancer	quasi-RCT 95 patients	Preventive effect on the progression of cirrhosis to liver cancer.	While not significant, the Shosaikoto treatment tends to lower the incidence of liver cancer and AFP.	33
Hochuekkito	Large intestine carcinoma	RCT-envelope 20 patients	To evaluate the efficacy of 1-week preoperative treatment with Hochuekkito for improving pre- and post-operative nutritional status and immune function in patients scheduled to undergo laparotomy for large intestine carcinoma.	Preoperative treatment with Hochuekkito may be useful for early recovery from surgery for large intestine carcinoma.	42

Continued

Hochuekkito	Gastric/colon cancer	RCT 51 patients	To evaluate the efficacy of preoperative administration of Hochuekkito for postoperative systemic inflammatory response syndrome (SIRS) in gastric/colon cancer.	Preoperative administration of Hochuekkito significantly suppresses the postoperative inflammatory response to surgical wounding.	25
Hochuekkito	Gastric or colorectal cancer	RCT 48 patients	To evaluate whether preoperative administration of Hochuekkito relieves surgical stress in patients with gastric or colorectal cancer.	Preoperative administration of Hochuekkito extract granules reduces the response to surgical stress and may be helpful for accelerating postoperative recovery.	26
Daikenchuto	Large intestine carcinoma	RCT 30 patients	To evaluate the reduction in the number of days to postoperative flatulence and the anti-inflammatory efficacy of Daikenchuto in patients who underwent laparotomy for large intestine carcinoma.	Daikenchuto is useful in promoting flatulence and inhibiting inflammation after surgery for large intestine carcinoma.	35
Daikenchuto	Colorectal cancer	RCT 175 patients	To evaluate the effects of Daikenchuto on intestinal obstruction following colorectal cancer surgery.	Daikenchuto extract fine granules do not prevent ileus following colorectal cancer surgery, but do result in the reduction of postoperative abdominal pain and irregular bowel movements.	43
Daikenchuto	Liver carcinoma	RCT 20 patients	To evaluate the anti-inflammatory efficacy of Daikenchuto in postoperative patients with liver carcinoma.	Daikenchuto may be useful in inhibiting early postoperative inflammation after surgery for liver carcinoma.	35
Daikenchuto	Gastric and colorectal cancer	RCT-envelope 20 patients	To evaluate the efficacy of Hangeshashinto for delayed diarrhea induced by irinotecan in patients with metastatic gastric and colorectal cancer.	Hangeshashinto is a useful supportive therapy from the viewpoint of QOL in patients treated for advanced gastric and colorectal cancer with S-1/irinotecan combination therapy.	32

Abbreviations: RCT-envelope, randomized controlled trial using sealed envelopes for allocation; RCT, randomized controlled trial; quasi-RCT, quasi-randomized controlled trial.

for gastric cancer. It was demonstrated that preoperative administration of the Kampo medicine Hochuekkito could be helpful in suppressing the postoperative inflammatory response to surgical wounding and accelerating postoperative recovery (25,26). In addition, Juzentaihoto, Ninjin'yoeito, and Hangeshashinto were exhibited to have abilities in improving host-immunity and reducing side effects in gastric cancer patients undergoing postoperative adjuvant chemotherapy including tegafur-uracil (UFT), 5-FU, fluoropyrimidine, and irinotecan treatments (27-32). The application of Kampo medicine in treatment of liver cancer were also studied (33-35). For example, Juzentaihoto was found to suppresses nausea/vomiting after transarterial embolization (TAE) with spongel + lipiodol + phosphatidyl choline + cisplatin for hepatocellular carcinoma (34). In addition, Daikenchuto may be useful in inhibiting early postoperative inflammation after surgery for liver carcinoma (35). The efficacy of Kampo medicine in treating large intestine cancer and colorectal cancer were extensively investigated. A series of Kampo products including Juzentaihoto, Ninjin'yoeito, Shosaikoto, Hochuekkito, Daikenchuto, and Hangeshashinto were proved to be effective in stimulating immune system, improving nutritional status, acting as biological response modifier, enhancing the tumor selectivity of chemotherapy drugs, or improving the quality of life of cancer patients (Table 2).

These clinical studies suggested the benefits of Kampo medicine in combination with conventional anti-cancer strategies in treatment of gastrointestinal cancers.

2.2. Basic studies on the mechanisms underlying the effects of Kampo medicine

Mechanisms underlying the effects of those Kampo medicines have been extensively investigated. Studies indicated that the immunity regulation effects of **Juzentaihoto** and **Hochuekkito** were related with enhancement of humoral immunity, cell-mediated immunity, natural killer (NK) activity, macrophage activity, and production of immunity related cytokines. It was found that oral administration of Juzentaihoto inhibited liver metastasis of colon cancer cells implanted in mice. However, this inhibitory effect was not observed in mice pretreated with 2-chloroadenosine which had an effect to inactivate macrophages. Hochuekkito enhanced the *in vitro* cytotoxic activity of activated T lymphocytes from healthy volunteers and cancer patients (44). In a mouse cancer metastasis model prepared by implanting mouse colon cancer cells (Colon 26-L5), removal of NK cells led to the disappearance of the inhibitory effects of Hochuekkito on cancer metastasis (45). In addition, oral administration of Hochuekkito to mice enhanced the anti-tumor cytotoxic activity of peritoneal exudate

cells (46) and inhibited restraint of a stress-induced decrease of serum interleukin (IL)-12 in tumor bearing mice (47). These studies indicated that Juzentaihoto and Hochuekkito may exhibit indirect antitumor activity through enhancing the organism's antineoplastic immune response.

Shosaikoto was shown to lower the incidence of liver cancer in clinical studies (33). Basic studies indicated that Shosaikoto has ability to prevent hepatopathy. In hepatocytes isolated from rats, Shosaikoto inhibited antibody-dependent cellular cytotoxicity (ADCC) responses and cell impairment induced by the culture supernatant of activated macrophages (48). In addition, Shosaikoto could inhibit liver fibrosis, which may decrease the risk of liver cancer. In rat liver stellate cells, Shosaikoto inhibited proliferation and transformation to myofibroblast-like cells, and suppressed type I/III procollagen mRNA expression (49). In another study, a diet containing Shosaikoto was given to a rat fibrosis model induced by dimethylnitrosamine or porcine serum. It inhibited an increase in the malondialdehyde level *in vivo*. Furthermore, it inhibited oxidative stress in rat liver stellate cells and hepatocytes *in vitro* (50). These studies illustrated the direct protective effects of Shosaikoto on liver cells. Also, Shosaikoto was found to exhibit an immune stimulatory effect. Oral administration of Shosaikoto to mice and rats activated macrophages (48). Other studies demonstrated that Shosaikoto showed a regulatory effect on production of cytokines. For example, oral administration of Shosaikoto increased the production of IL-1, IL-2, IL-1 β , IL-6, IL-10, GM-CSF, TNF- α , and IFN- γ and decreased the excessive production of IL-4 and IL-5 in peripheral blood mononuclear cells derived from healthy individuals or chronic hepatitis patients (48,51-53). The activities of lymphokine-activated killer (LAK) cells and NK cells in human peripheral blood mononuclear cells were also enhanced after oral administration of Shosaikoto (48). All these experimental studies elucidated the mechanisms underlying the anti-tumor effects of Shosaikoto.

Daikenchuto showed benefits to patients with ileus following colorectal cancer surgery in clinical studies. The pharmacological effects of Daikenchuto are displayed in two aspects (54). First, oral administration of Daikenchuto promoted gastrointestinal motility through enhancing the contraction of stomach/duodenum in the resting phase in healthy subjects. Second, oral administration of Daikenchuto increased the plasma motilin, gastrin, plasma vasoactive intestinal polypeptide (VIP), calcitonin gene related peptide (CGRP), substance P, and serotonin levels in healthy subjects or patients with paralytic ileus after extensive hysterectomy with pelvic lymphadenectomy (55-57). Mechanisms behind the pharmacological actions of Daikenchuto were illustrated in animal studies.

Daikenchuto improved chlorpromazine-induced decrease of small intestinal and distal colonic transit in mice. It was found that atropine and a cholecystokinin receptor antagonist, lorglumide, could antagonize the activities of Daikenchuto (58). In isolated rabbit jejunum, Daikenchuto also enhanced spontaneous contractions and reversed the decrease of spontaneous contractions inhibited by atropine. In the longitudinal muscle of isolated guinea pig ileum, Daikenchuto induced contractions and acetylcholine (ACh) release. The contraction induction effect could be inhibited by pretreatment or concurrent use of atropine, tetrodotoxin, noradrenaline, substance P receptor antagonist, or one 5-HT₄ receptor inhibitor ICS205930, but not by pre-exposure to hexamethonium (59,60). These experimental studies on laboratory animals suggested that the contractile response induced by Daikenchuto is partially mediated by ACh released from cholinergic nerve endings and that 5-HT₄, substance P, and cholecystokinin receptors would be involved in this effect of Daikenchuto.

Hangeshashinto was found to be useful in relieving delayed diarrhea induced by irinotecan which is usually employed to treat metastatic gastric and colorectal cancer (32). Studies indicated that Hangeshashinto exhibits pharmacological effects *via* actions including protection of the gastric mucosa, anti-inflammatory actions, and enhancement of large intestinal water absorption. Oral administration of Hangeshashinto to rats inhibited taurocholic acid-induced decreases in the gastric mucosa levels of phospholipids, a reduction of gastric mucosal potential differences, and gastric mucosa back diffusion of H⁺ (61). In addition, Hangeshashinto inhibited an ethanol-induced decrease in the volume of mucus in the superficial gastric mucosa and the deep mucosa of the gastric body (61). These results demonstrated the protective effects of Hangeshashinto on gastric mucosa. In regard to its anti-inflammatory actions, oral pretreatment with Hangeshashinto in rats inhibited irinotecan induced production of prostaglandin E₂ in the large intestinal mucosa (62). In terms of enhancement of large intestinal water absorption, oral administration of Hangeshashinto to rats enhanced water absorption in the large intestine (63). In another study, oral pre-treatment of this drug to rats inhibited the irinotecan induced reduction of water absorption in the large intestine (62). Thus, these studies suggested the efficacy of Hangeshashinto in reducing side effects induced by the chemotherapy drug irinotecan.

3. Quality control of Kampo medicine

Japan is currently the only developed country where traditional medicine, *i.e.* Kampo, is widely believed by the public, officially recognized by the Government, and mostly (148 of all 210 formulae) covered by the

national health insurance system. Thus far, much progress has been made in scientifically evaluating Kampo medicine. These achievements may be ascribed to, if not all, the standardization of Kampo products. In order to improve quality control of Kampo medicine, an Advisory Committee for Kampo Drugs which is in close association with the Ministry of Health, Labor and Welfare was established in 1982. Since the implementation of Good Manufacturing Practice Law in 1986, equal standards were applied to all pharmaceutical drugs including Western drugs and Kampo drugs. Moreover, guidelines for ethical extract products in oriental medicine formulations were developed in 1985 (64). One characteristic of Kampo is that every formulation has consisted of fixed combinations of herbs in standardized proportions, which is different from traditional Chinese medicine with modifying formulas. Although each Kampo product may be produced by various manufacturers, it is composed of exactly the same ingredients under the Ministry's standardization methodology. Moreover, the herbs used in the formulas must have the required levels of at least two marker components in order for the formula to be approved as a medicament (65). Because of that, the quality of crude drugs receives a high degree of attention. Since the majority of crude drugs are imported from different countries, they were subjected to rigorous checks such as quantification of the possible active ingredients to guarantee consistency in quality. The herbs also undergo tests for any possible contaminants of agricultural pesticides or environmental pollutants, especially heavy metals. Microbial tests for bacteria, yeast and mold are also carried out. These measures ensure the quality of Kampo products from the very beginning. It is the high and consistent quality of Kampo products that have prepared the groundwork for clinical and basic studies on the therapeutic value of Kampo medicine.

4. Prospects and conclusions

The evaluation of the efficacy and safety of traditional medicine including Kampo is not without challenges. The clinical application of Kampo medicine has long been based on its own diagnosis theory called 'Sho' in Japanese which refers to a particular pathological status of a patient and is patterned according to the patient's constitution and symptoms. The term 'Sho' is different from the disease entities understood in terms of modern Western medicine. However, the currently used evaluation model employed in clinical studies are nearly based on the disease terms defined by Western medicine. Considering the characteristics of Kampo medicine, *i.e.* the uncertainty of active ingredients, the combination of diagnosis theories of Kampo medicine and Western medicine would be more rational in patients selected and grouped in clinical trials in

the future. Despite all of this, in terms of scientific evaluation and quality control of traditional medicine, some meaningful conclusions on Kampo medicine are put forward as an exploration for reducing the side effects and improving the quality of gastrointestinal cancer patients subjected to conventional anti-cancer therapies. These achievements may provide reference for the development and dissemination of other traditional medicines in the world.

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Brief Report

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Synthesis of solasodine glycoside derivatives and evaluation of their cytotoxic effects on human cancer cellsChangzhi Cui¹, Xuesen Wen¹, Min Cui¹, Jian Gao¹, Bin Sun^{1,2}, Hongxiang Lou^{1,*}¹ School of Pharmaceutical Sciences, Shandong University, Ji'nan, Shandong, China;² National Glycoengineering Research Center, Shandong University, Ji'nan, Shandong, China.

ABSTRACT: Solasodine glycosides, such as solamargine, have been proved to be very important anti-cancer agents. In order to discover more potent cytotoxic agents and explore the preliminary structure activity relationship, a new series of solasodine glycosides 2-9 were synthesized *via* a transglycosylation strategy, and their cytotoxic activity against a panel of human cancer cell lines (MCF-7, KB, K562, and PC3 cells) were evaluated by MTT assays. The results indicated that compounds 2, 8, and 9 with the substitute moiety of rhamnose, 2-hydroxyethoxymethyl, and 1,3-dihydroxypropan-2-yloxy-methyl, respectively, exhibited quite strong anticancer activity. The underlying mechanism tests demonstrated that these compounds could induce apoptosis detected by DAPI staining, and Annexin V and propidium iodide binding. Cell cycle analysis indicated that the cancer cells were predominantly arrested at the G2/M phase when exposure to these compounds was examined by flow cytometry. These compounds may serve as lead candidates in the development of novel chemotherapeutics for cancer treatment.

Keywords: Saponins, solasodine, cytotoxicity, apoptosis

1. Introduction

Natural derived products which belong to the traditional medicinal system are affluent resources for new drug development. Steroidal glycosides are a series of natural products and display a wide spectrum of structural diversity with versatile biological activities, such as anti-inflammatory, antibacterial, antiparasitic, antifungal, and anticancer activities (1-5). Both the

sugar residue and aglycone in the structure play very important roles in their biological performance (6-8).

Solasodine, a solanum type of steroid alkaloid with a C27 cholestane skeleton, was first discovered from the fruit of the devil's apple in free and conjugated forms (9). Solasodine conjugates displayed uneventful anticancer activity, but the mixture of the solasodine glycosides (BEC: 33% solasonine, 33% solamargine, and 34% of their corresponding di- and monoglycosides) discovered from the Devil's Apple plant (9-15) have been clinically applied as an antineoplastic agent.

Solasodine glycosides exhibited obvious anticancer activities when compared with its aglycone (9-14). Meanwhile, the cytotoxic activities of solasodine glycosides were generally dependent on the presence of sugars, in particular rhamnose (9-21), and were mediated by the rhamnose-binding receptors that occur on the membrane of cancer cells (21).

In order to discover more potent cytotoxic agents for cancer therapy with the simplified structures which were facilitated for industrial preparation, six monosaccharides including rhamnose as well as two open-loop saccharide analogues were introduced to solasodine and compounds 2-9 were then prepared accordingly (Scheme 1).

2. Materials and Methods**2.1. Chemicals**

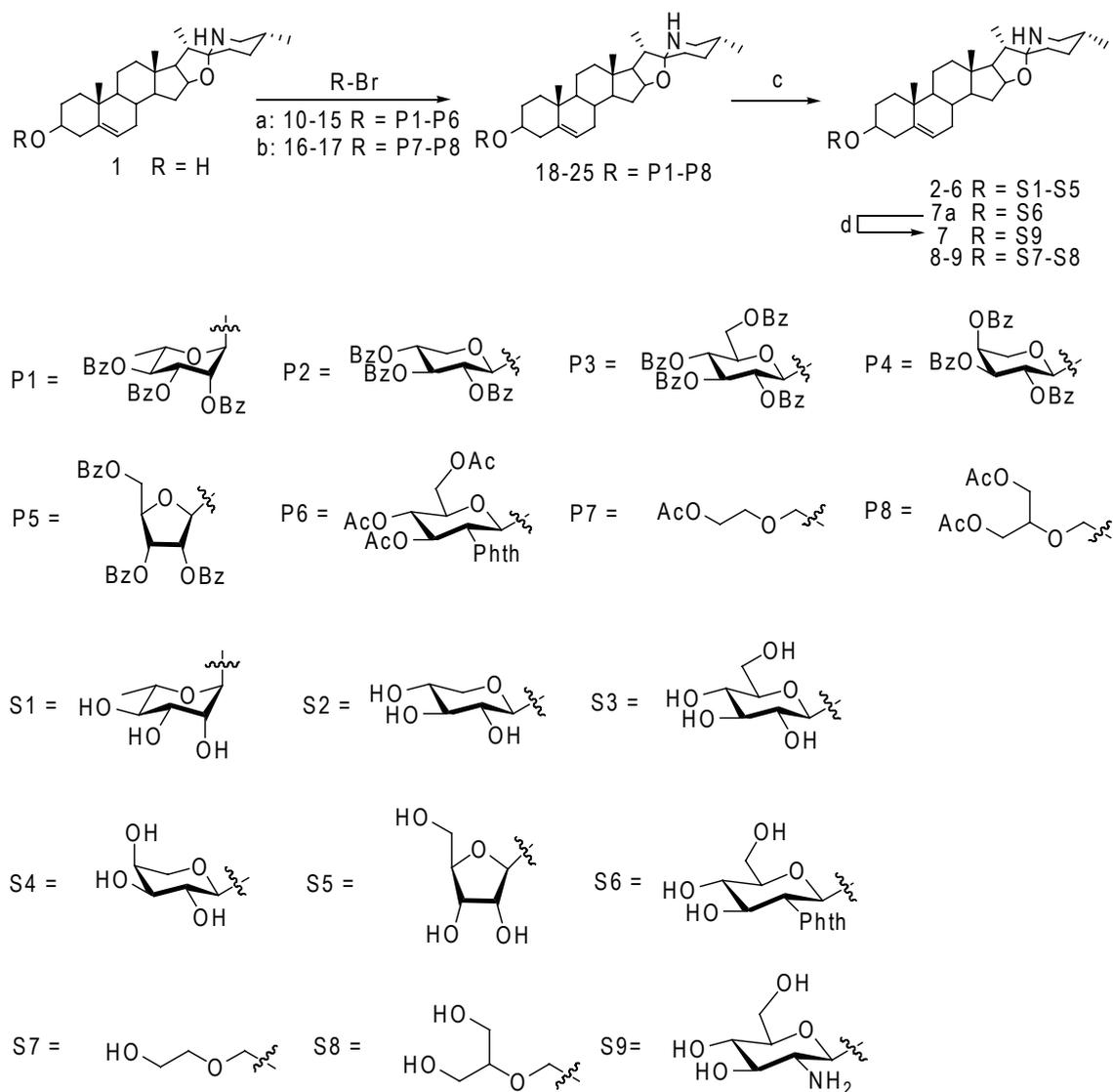
Solasodine **1** (Scheme 1) was synthesized using five straightforward sequential reactions as reported previously (22). Six glycosyl bromide **10-15** (Scheme 1) was then employed to assemble a small library of solasodine glycosides using classic Koenigs-Knorr glycosylation. In addition, two open-loop saccharide analogues were transferred to the solasodine using 2-(1*H*-benzotriazole-1-yl)-1,1,3,3-tetramethyluronium tetrafluoroborate (DIEA) as a catalyst.

As shown in Scheme 1, glycosyl donors **10-14** were prepared from the corresponding benzoylated monosaccharide using a solution of 45% HBr/HOAc in CH₂Cl₂. Compound **15** was easily prepared from

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Scheme 1. Reagents and conditions. (a) AgOTf, 4 Å molecular sieve, -20°C , solasodine, 12 h; (b) DIEA, CH_2Cl_2 , room temperature, 24 h; (c) $\text{CH}_3\text{OH}/\text{CH}_3\text{ONa}$, 4 h, 98%; (d) CH_3NH_2 , 4 h, 97%.

commercial glucosamine hydrochloride (23). Compound **16** was synthesized from 1,3-dioxolane using AcBr, and **17** was prepared from 1,3-dioxolan-4-yl-methanol (24). And then each donor of **10-15** was condensed with solasodine **1**, respectively, with catalysis from silver trifluoromethanesulfonate (AgOTf) to provide protected **18-23**. Under the promotion of DIEA, coupling of open-loop donor **16** or **17** with solasodine **1** afforded the intermediates **24** and **25**, respectively. At last, MeOH/MeONa or CH_3NH_2 was used to hydrolyze the tosyl, acetyl and phthalic anhydride to provide target compounds **2-9**. The β -bond linkage between glycosyl and aglycone in compounds **2-7** was confirmed. Specific information on synthesis of targeted compounds **2-9** is indicated in the Appendix.

2.2. Cell lines and cell culture

Human squamous cell carcinoma (KB), breast

adenocarcinoma (MCF-7), myelogenous leukemia (K562), prostate cancer (PC3), human umbilical vein endothelial (ECV304), and human hepatocyte cells (HL7702) were maintained in RPMI 1640 medium (HyClone, Thermo Fisher Scientific Inc., Waltham, MA, USA). The medium was supplemented with 10% fetal bovine serum (FBS) (Gibco, Invitrogen, Carlsbad, CA, USA), 100 $\mu\text{g}/\text{mL}$ penicillin and 100 $\mu\text{g}/\text{mL}$ streptomycin. Cells were cultured in a humidified atmosphere of 5% CO_2 at 37°C .

2.3. MTT assay

Cells were seeded into 96-well culture plates at a density of $4-5 \times 10^3$ cells per well and cultured for 24 h. Thereafter the cells were treated with various concentrations of tested compounds and incubated for 24-48 h. Cell viability was assessed by MTT assay. The cells were incubated with 5 mg/mL MTT solution at 37°C for 4 h, and the resulting crystals were dissolved

in dimethyl sulfoxide (DMSO). The optical density (OD) was measured using a plate microreader (Bio-Rad 680, Bio-Rad Co., Hercules, USA). Cell viability was calculated as follows:

$$\text{Cell viability (\%)} = 100 \times (\text{OD}_{\text{sample}} - \text{OD}_{\text{blank}}) / (\text{OD}_{\text{control}} - \text{OD}_{\text{blank}})$$

2.4. Cell morphology examination

To assess the effect of compounds **2**, **8**, and **9** on apoptosis, MCF-7 cells were seeded in 24-well plates. After treatment with compounds **2**, **8** and **9** for 24 h cells were washed twice with PBS, and were observed under the microscope. Images were processed with Adobe Photoshop 7.0.

2.5. 4',6-Diamino-2-phenylindole (DAPI) staining

MCF-7 cells were seeded onto 12-mm round, glass cover slips in 24-well plates. After 24 h treatment with compounds **2**, **8**, and **9**, cells were stained with 4 µg/mL DAPI for 10 min at room temperature. The cover slips containing cells were then mounted on microscope slides using mounting medium and analyzed with fluorescence microscopy. Fluorescence images were processed using AutoQuant X 2.1 software from Cybernetics, Inc. (Bethesda, MD, USA).

2.6. Apoptosis analysis

Apoptosis was evaluated with Annexin V/propidium iodide (PI) staining using an FITC Annexin V Apoptosis Detection Kit purchased from BD Biosciences (San Jose, CA, USA). MCF-7 cells were seeded at a density of 1×10^5 /mL into 6-well plates. After 24 h incubation with compounds **2**, **8**, and **9**, the cells were washed twice with ice-cold phosphate buffered saline (PBS). Each cell sample was transferred to individual tubes and centrifuged at $200 \times g$ for 5 min. The supernatant was removed, and the cells were resuspended in 400 µL of annexin V-FITC binding buffer and incubated at room temperature in the dark for 15 min with 5 µL annexin V-FITC and PI 10 µL (50 µg/mL) was added for another 5 min. Apoptosis was measured using flow cytometry (Becton, Dickinson and Company, Franklin Lakes, NJ, USA) and WinMDI 2.9 analysis software.

2.7. Cell-cycle analysis

Cell cycle was analyzed using flow cytometry of PI-stained cells. MCF-7 cells were cultured in the presence of different concentrations of compounds **2**, **8**, and **9**. After designated time intervals, cells were fixed in 70% ethanol overnight at 4°C and washed once with PBS. Then, the cells were incubated with 1 U/mL of RNase A (DNase free) for 30 min at 37°C and 10 µg/mL of

PI for 1 h at room temperature in the dark. Cell cycle distribution was examined by flow cytometry, and data was analyzed using the Modfit program (Becton, Dickinson and Company).

2.8. Statistical analysis

All experiments were performed at least three times. Statistical analysis was performed with an analysis of variance (ANOVA) followed by the Turkey's *t*-test. $p < 0.05$ were considered statistically significant.

3. Results and Discussion

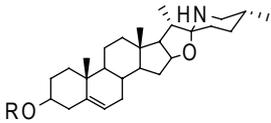
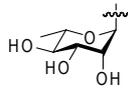
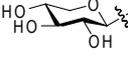
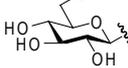
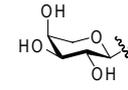
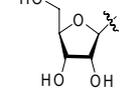
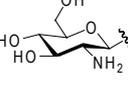
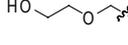
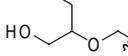
3.1. Cytotoxic effects of synthesized compounds on human cancer cells

The anticancer effect of synthesized saponins **2-9** as well as solasodine **1** against human KB, K562, MCF-7, PC3, EVC304, and HL7702 cells were preliminarily evaluated by MTT assay (25). As shown in Table 1, compounds **2**, **3**, **8**, and **9** were found to possess relatively potent cytotoxic activity against PC3 cells, with IC₅₀ values ranging from 7.2 to 18.4 µM. In K562 cells, compounds **2** and **9** with IC₅₀ values of 18.8 and 17.0 µM showed obvious superiority to other synthesized compounds in suppressing cell proliferation. Similarly, compounds **2**, **8**, and **9** exhibited excellent proliferation inhibitory activity against MCF-7 cells, with IC₅₀ values of 12.9 and 17 µM, respectively. On the other hand, compounds **2**, **8**, and **9** showed weak cytotoxicity in human normal cell lines ECV304 and HL7702 (Table 1). These results indicated that the saponins with rhamnose (*e.g.* compound **2**) or open-loop saccharide analogues (*e.g.* compounds **8** and **9**) exhibited more potent anticancer activity than the other derivatives synthesized in the current study.

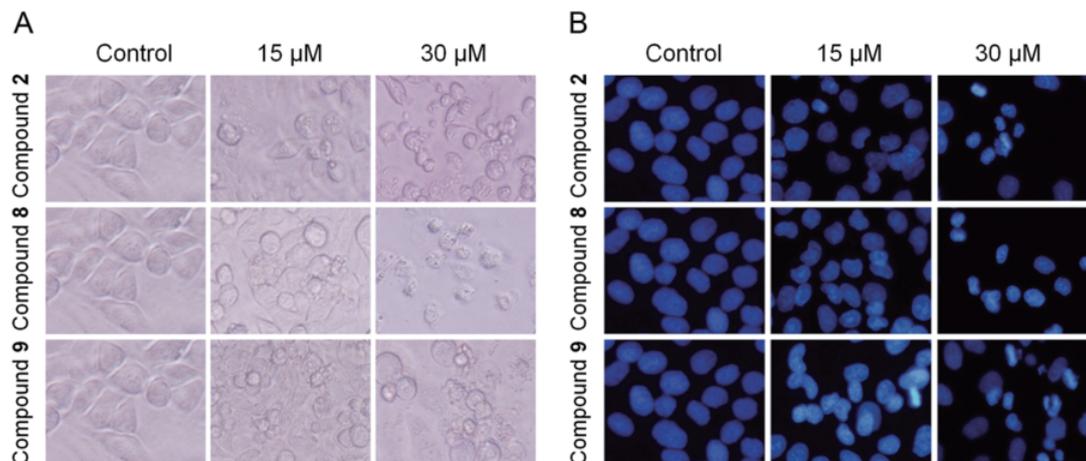
3.2. Compounds **2**, **8**, and **9** induced cell morphologic alteration in MCF-7 cells

To examine the effect of compounds **2**, **8**, and **9** on cell morphology during cell death, MCF-7 cells were treated with various concentrations (0, 15, 30 µM) of these compounds for 24 h, respectively. As shown in Figure 1A, the morphological character of cells changed greatly. Cell shrinkage and membrane integrity loss appeared, which are typical features of apoptosis (26,27). Apoptotic cells were also defined on the basis of characteristic nuclear morphology including condensation of chromatin producing crescent shapes around the periphery of the nucleus and apoptotic bodies (28,29). As shown in Figure 1B, the non-treated cells had rounded and intact nuclei with diffuse DAPI staining, while the cells treated with compounds **2**, **8**, and **9** had nuclei that were smaller and brighter

Table 1. The cytotoxicity of compounds 1-9 on human cell lines

Compound	R	IC ₅₀ (μM)					
		KB	K562	MCF-7	PC3	ECV304	HL7702
Solamargine (Ref. 30)		7.8	8.0	8.2	5.9	ND	ND
1	H-	ND	ND	ND	13.6	ND	ND
2		29.1	18.8	14.2	18.4	34.0	36.0
3		ND	51.0	53.0	16.5	ND	ND
4		ND	43.0	77.0	27.9	ND	ND
5		ND	44.0	64.0	23.1	ND	ND
6		ND	ND	ND	66.9	ND	ND
7		ND	37.0	98.0	21.9	ND	ND
8		28.9	112.0	12.9	17.4	> 40	> 40
9		29.4	17.0	26.0	7.2	> 40	> 40

ND: not detected.

**Figure 1. Morphological changes of MCF-7 cells after treatment with compounds 2, 8, and 9, respectively, for 24 h. Cell morphology was examined directly under light microscopy (A) or detected *via* fluorescence microscopy after the cells were stained with DAPI (B).**

stained with condensed chromatin forming crescent-shaped profiles around the periphery of the nucleus or separate globular structures (apoptotic bodies). The result implied that compounds **2**, **8**, and **9** induced cell apoptosis in MCF-7 cells.

3.3. Quantification of apoptotic cells solicited by compounds **2**, **8**, and **9** in MCF-7 cells

To further quantify the extent of compounds **2**, **8**, **9**-induced apoptosis, MCF-7 cells double stained with annexin V and PI were analyzed by flow cytometry. The analysis results revealed that the proportion of cells stained with annexin V increased with compounds **2**, **8**, and **9** treatment (Figure 2). The percentages of apoptotic cells increased from 0.88%, 0.07%, and 0.07% to 8.58%, 9.16%, and 10.56% when cells were exposed to 15 μ M compounds **2**, **8**, and **9**, respectively, for 24 h. The proportion of apoptotic cells went up to 30.36%, 36.71%, and 34.18% when cells were treated with 30 μ M of the above agents.

3.4. Compounds **2**, **8**, and **9** induced cell cycle arrest at G₂/M phase in MCF-7 cells

To determine whether compounds **2**, **8**, and **9** exerted cell growth inhibition activity *via* the induction of cell cycle arrest in addition to apoptosis, we next focused on the effect of compounds **2**, **8**, and **9** on the cell cycle of MCF-7 cells. We observed that compounds **2**, **8**, and **9** treatment led to cells accumulating in the G₂/M phase compared with non-treated cells (Figure 3). The ratio of cells in the G₂/M phase following the treatment with 15 μ M compounds **2**, **8**, and **9** for 24 h increased from 6.60%, 5.79%, and 4.35% to 11.18%, 11.41%, and 7.71%, respectively. The values rose to 14.33%, 17.26%, and 17.84%, respectively, when cells were incubated with 30 μ M of the above compounds for 24 h.

In summary, solasodine **1** together with its eight glycoside derivatives **2-9** were synthesized *via* a Koenigs-Knorr strategy. The preliminary structure-activity relationship (SAR) analysis indicated that

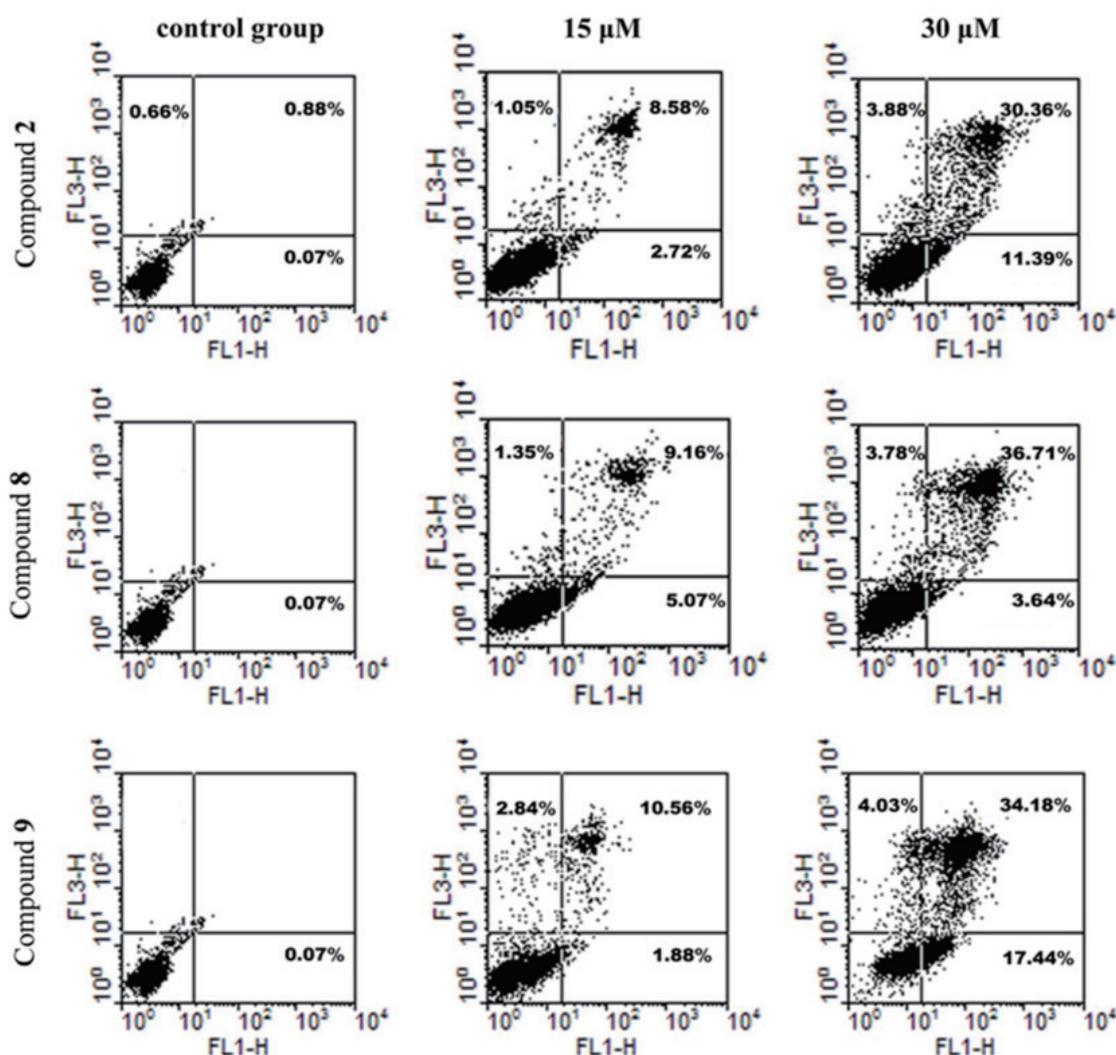


Figure 2. Flow cytometry analysis of cell apoptosis. MCF-7 cells were treated with compounds **2**, **8**, and **9**, respectively, for 24 h and then stained with Annexin-V/PI.

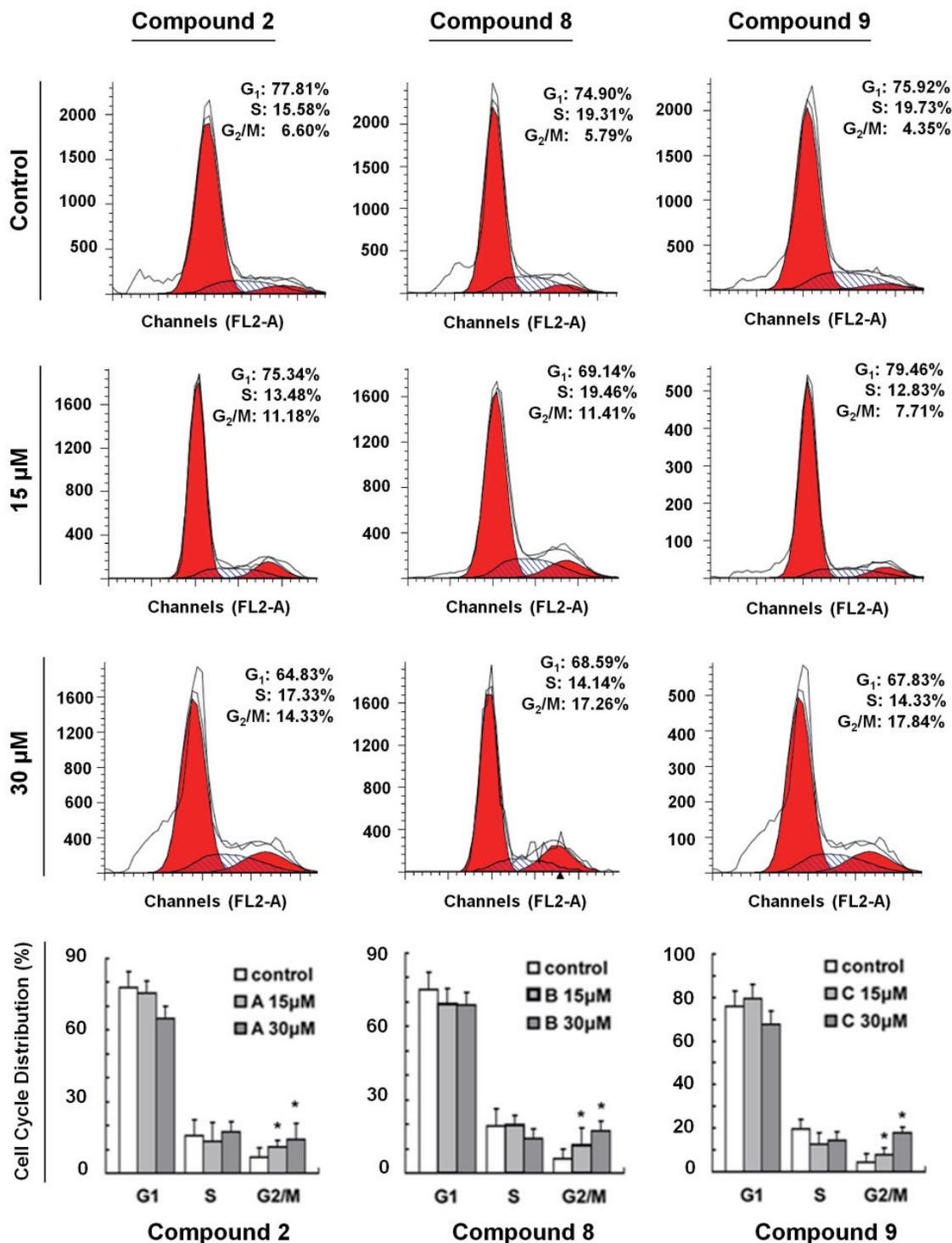


Figure 3. DNA content analysis. MCF-7 cells were treated with compounds 2, 8, and 9, respectively, for 24 h and then stained with PI. The fluorescence of PI-stained cells was analyzed using flow cytometry. Cell cycle distribution was analyzed based on the DNA content. * $p < 0.05$ vs. control.

the rhamnose substituent was essential and the chain substituents in compounds 8 and 9 were also very important for its anticancer activity. The other sugar substituted derivatives exhibited less anticancer activity. These results were consistent with the previous report (21). Furthermore, it was demonstrated that the chain and rhamnose substituents exhibited similar cytotoxic potency. Our underlying work suggested that compounds

2, 8, and 9 have the ability to kill cancer cells *via* a form of apoptosis-like cell death. Moreover, compounds 2, 8, and 9-treated cells were arrested predominantly at the G₂/M phase. This study demonstrated that compounds 2, 8, and 9 may merit further investigation as potential therapeutic leads for the development of novel anticancer drugs. Much more work concerning the biochemical mechanism is ongoing.

Acknowledgements

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Appendix

Flash column chromatography was performed on silica gel (200-300 mesh, Qindao Ocean Chemical Co., Qingdao, Shandong, China). Analytical thin-layer chromatography (TLC) was carried out on precoated

Silica Gel 60 F₂₅₄ plates (E. Merck, Darmstadt, Germany) with detection by fluorescence and/or by charring with 30% (v/v) H₂SO₄ in EtOH. Dichloromethane was distilled from CaH₂. The chemicals and materials were purchased from Alfa-Aesar (Ward Hill, MA, USA) and were used as received. NMR spectra were recorded on a Bruker Avance DRX 600 spectrometer (Bruker BioSpin, Rheinstetten, Germany) at 600 (¹H) and 150 (¹³C) MHz, respectively. Elucidations of chemical structures were based on ¹H, ¹³C, ¹H-¹H COSY, HMBC, and HMQC NMR experiments. Signals are reported as follows: s (singlet), d (doublet), t (triplet), q (quintet), m (multiplet), and coupling constants are reported in Hz (hertz). Melting points were determined in an X-6 melting-point apparatus and are uncorrected. Optical rotations were measured on a Perkin-Elmer 241 MC polarimeter (Waltham, MA, USA). Mass spectral data (HR-ESI-MS and HR-FAB-MS) were obtained on LTQ-Orbitrap XL (Thermo Fisher Scientific Inc.) and JEOL JMS-DX-303HF (JEOL, Tokyo, Japan) in the positive ion mode, respectively.

Synthesis of compound **1** (Ref. 28)

Yield 18%, white solid; ¹H-NMR (600 MHz, CDCl₃): 5.37 (br. s, H-C(6)); 4.31 (m, Ha-C(16)); 3.54 (m, Ha-C(3)); 2.68 (m, CH₂-(26)); 1.19 (s, Me-(19)); 1.10 (s, Me-(21)); 0.97 (s, Me-(27)); 0.84 (s, Me-(18)). ESI-MS: 414.3 ([M + H]⁺).

Synthesis of compounds **10-14**

HBr/HOAc (1.0 mmol) was added to a solution of each of the protected sugars (0.5 mmol) in 4 mL CH₂Cl₂ at 0°C. The reaction mixture was stirred at 0°C for 12 h, the progress of the reaction was monitored by TLC. The solvent was removed under reduced pressure to afford a residue. Purification of the residue on a silica gel column with a 6:1 solution of petroleum ether-EtOAc gave **10-14** (yield 40-50%, oil).

Synthesis of compound *D*-glucopyranosyl bromide (**15**)

Triethylamine (1.7 mL, 11.6 mmol) was added to the suspension of *D*-galactosamine hydrochloride (1 g, 4.65 mmol) in DMF (30 mL). After 20 min stirring at room temperature, phthalic anhydride (688 mg, 4.65 mmol) was added to this solution, and the mixture was stirred for 2 h at 50°C under N₂ atmosphere. The reaction mixture was allowed to stand until its temperature dropped to room temperature, excess amounts of triethylamine (5 mL) and acetic anhydride (5 mL) were added. After a 2-day stirring at room temperature under a N₂ atmosphere, the reaction mixture was concentrated in vacuo, mixed with water (150 mL), extracted three times with petroleum ether-EtOAc 1:1, and dried over Na₂SO₄. The organic layer was concentrated and separated by silicagel column chromatography (petroleum ether-

EtOAc 1:1) to give the protected *D*-galactosamine (1.82 g, 82%). The resulting protected *D*-galactosamine (1.395 g, 2.92 mmol) was mixed with acetic anhydride (0.69 mL, 7.31 mmol), and treated with HBr-AcOH (30%, 16 mL, excess) with ice-cooling. After stirring for an hour at room temperature under a N₂ atmosphere, the mixture was concentrated in vacuo. The reaction mixture was mixed with saturated NaHCO₃, extracted three times with CH₂Cl₂, dried over Na₂SO₄, and concentrated to give **15** (1.41 g, 97%, Crude).

Synthesis of compounds **16 and 17**

AcBr (5.05 mmol) was added to a solution of each of the acetyl protected rings (5 mmol) in 6 mL anhydrous CH₂Cl₂ at room temperature. The mixture was stirred at the same temperature for 12 h. The solvent was removed under reduced pressure to afford **16 and 17** (yield 70-80%, oil).

Synthesis of compounds **18-23**

AgOTf (33 mg, 0.13 mmol) was added to a stirred mixture of brominated sugar (1.01 mmol) and solasodine (0.414 g, 1.0 mmol) in anhydrous CH₂Cl₂ (8 mL) with a 4 Å molecular sieve at -20°C under a N₂ atmosphere. The reaction mixture was stirred under this condition for 90 min. The solution was filtered and the solvent was evaporated under reduced pressure. The residue was subjected to silica column chromatography with a 40:1 solution of CH₂Cl₂-MeOH to afford **18-23** as white solids (yield 20-70%).

Synthesis of compounds **24 and 25**

Ethyl-diisopropylamine (DIEA) (3 mL) was added to a solution of solasodine (0.414 g, 1 mmol) in 5 mL CH₂Cl₂ at room temperature. After stirring for 15 min, compounds **16 and 17** (1.01 mmol) were added to the mixture and stirred for 12 h. The mixture was concentrated and purified by silica column chromatography with a 40:1 solution of CH₂Cl₂-MeOH to afford **24 and 25** (yield 70-80%).

Synthesis of compounds **2-6, 7a, 8, and 9**

Compounds **18-25** were dissolved in a solution of MeOH-CH₂Cl₂ (2:1, 10 mL). To the solution was added 2 g/100 mL MeONa/MeOH until a pH 9-10 was attained. The mixture was stirred at room temperature for 4 h, neutralized with Amberlite IR-120 (H⁺), filtered and concentrated. The residue was added to a silica gel column with a 45:1 solution of CH₂Cl₂-MeOH to afford compounds **2-6, 7a, 8, and 9** (yield 95-100%).

Solasodine 3-O-L-rhamnoside (2). ¹H-NMR (DMSO, 600 MHz) δ: 0.76 (3H, s, Me-18), 0.89 (3H, s, Me-27), 0.97 (3H, s, Me-21), 1.10 (3H, s, Me-19),

2.64 (2H, m, H-26), 3.54 (1H, s, H-3), 4.54 (1H, s, H-16), 4.71 (1H, d, $J = 19.2$ Hz, H-1'), 5.33 (1H, s, H-6); $^{13}\text{C-NMR}$ (DMSO, 600 MHz) δ : 140.7; 121.6; 98.6; 75.8; 71.4; 71.1; 69.0; 56.1; 49.8; 41.9; 40.5; 39.4; 38.5; 36.9; 32.2; 32.0; 31.2; 29.7; 20.7; 19.5; 18.6; 18.4. ESI-MS: 559.8 ($[\text{M} + \text{H}]^+$).

Solasodine 3-O-D-xylosidase (3). $^1\text{H-NMR}$ (DMSO, 600 MHz) δ : 0.76 (3H, s, Me-18), 0.89 (3H, s, Me-27), 0.97 (3H, s, Me-21), 1.10 (3H, s, Me-19), 2.67 (2H, m, H-26), 3.63 (1H, s, H-3), 4.54 (1H, s, H-16), 4.94 (1H, d, $J = 17.4$ Hz, H-1'), 5.33 (1H, s, H-6); $^{13}\text{C-NMR}$ (DMSO, 600 MHz) δ : 140.93; 121.4; 105.2; 102.2; 97.3; 83.2; 77.7; 73.7; 70.0; 61.6; 56.4; 49.8; 46.2; 41.5; 40.5; 38.7; 37.2; 36.8; 32.2; 31.9; 31.4; 29.8; 28.0; 20.7; 19.5; 18.7; 16.2; 15.0. ESI-MS: 545.7 ($[\text{M} + \text{H}]^+$).

Solasodine 3-O-D-galactosidase (4). $^1\text{H-NMR}$ (DMSO, 600 MHz) δ : 0.74 (3H, s, Me-18), 0.86 (3H, d, $J = 7.2$ Hz, Me-27), 0.97 (3H, s, Me-21), 1.05 (3H, t, $J = 7.2$ Hz, Me-19), 2.37 (2H, m, H-26), 3.61 (1H, s, H-3), 4.17 (1H, s, H-16), 4.18 (1H, d, $J = 17.4$ Hz, H-1'), 5.32 (1H, m, H-6); $^{13}\text{C-NMR}$ (DMSO, 600 MHz) δ : 141.0; 121.6; 101.8; 78.3; 77.2; 75.5; 74.0; 71.0; 68.6; 62.8; 60.9; 56.5; 47.6; 41.3; 38.8; 36.8; 29.8; 19.9; 19.6; 16.6; 15.8. ESI-MS: 575.7 ($[\text{M} + \text{H}]^+$).

Solasodine 3-O-L-arabinoside (5). $^1\text{H-NMR}$ (CD_3OD , 600 MHz) δ : 0.83 (3H, s, Me-18), 0.87 (3H, d, $J = 6.0$ Hz, Me-27), 0.99 (3H, d, $J = 7.2$ Hz, Me-21), 1.04 (3H, s, Me-19), 2.64 (2H, m, H-26), 3.58 (1H, m, H-3), 3.70 (2H, m, H-2'), 3.96 (2H, m, H-1'), 4.32 (1H, s, H-16), 5.37 (1H, s, H-6); $^{13}\text{C-NMR}$ (CD_3OD , 600 MHz) δ : 140.0; 122.0; 100.7; 98.5; 78.5; 72.7; 71.6; 67.6; 65.2; 62.7; 56.5; 50.1; 47.5; 41.3; 40.5; 39.8; 38.7; 37.2; 36.9; 34.0; 32.1; 31.4; 30.1; 29.6; 20.9; 19.4; 16.4; 15.2. ESI-MS: 545.7 ($[\text{M} + \text{H}]^+$).

Solasodine 3-O-D-riboside (6). $^1\text{H-NMR}$ (DMSO, 600 MHz) δ : 0.75 (3H, s, Me-18), 0.88 (3H, d, $J = 6.6$ Hz, Me-27), 0.96 (3H, s, Me-21), 1.07 (3H, d, $J = 6.0$ Hz, Me-19), 2.69 (2H, s, H-26), 3.67 (1H, s, H-3), 4.53 (1H, m, H-16), 4.85 (1H, t, $J = 6.0$ Hz, H-1'), 5.33 (1H, d, $J = 4.8$ Hz, H-6); $^{13}\text{C-NMR}$ (DMSO, 600 MHz) δ : 140.4; 121.7; 98.7; 97.9; 76.3; 70.9; 68.5; 63.7; 61.4;

55.8; 54.9; 49.3; 48.5; 45.2; 41.0; 40.3; 38.7; 38.2; 36.8; 31.7; 31.4; 31.0; 30.1; 20.3; 19.1; 18.3; 17.3; 15.8. ESI-MS: 545.7 ($[\text{M} + \text{H}]^+$).

Solasodine 3-O-(2'-hydroxyethoxy)methyl ether (8). $^1\text{H-NMR}$ (CD_3OD , 600 MHz) δ : 0.84 (3H, s, Me-18), 0.87 (3H, d, $J = 6.0$ Hz, Me-27), 0.98 (3H, d, $J = 4.8$ Hz, Me-21), 1.05 (3H, s, Me-19), 2.66 (2H, m, H-26), 3.50 (1H, m, H-3), 3.75 (4H, m, H-2',3'), 4.32 (1H, s, H-16), 4.81 (2H, m, H-1'), 5.38 (1H, s, H-6); $^{13}\text{C-NMR}$ (CD_3OD , 600 MHz) δ : 140.5; 121.7; 98.3; 94.2; 71.0; 62.3; 56.5; 50.1; 41.2; 40.5; 40.0; 39.4; 37.1; 36.9; 32.2; 31.4; 28.8; 20.8; 19.4; 16.5; 15.4. ESI-MS: 487.7 ($[\text{M} + \text{H}]^+$).

Solasodine 3-O-(1',3'-dihydroxypropan-2'-yloxy)methyl ether (9). $^1\text{H-NMR}$ (DMSO, 600 MHz) δ : 0.75 (3H, s, Me-18), 0.85 (3H, m, Me-27), 0.97 (3H, s, Me-21), 1.05 (3H, t, $J = 7.2$ Hz, Me-19), 3.09 (1H, m, H-2'), 3.24 (2H, m, H-26), 3.61 (1H, m, H-3), 4.38 (1H, t, $J = 4.8$ Hz, H-16), 5.32 (1H, d, $J = 4.8$ Hz, H-6); $^{13}\text{C-NMR}$ (DMSO, 600 MHz) δ : 140.9; 120.6; 95.8; 94.5; 79.4; 76.7; 61.6; 60.7; 59.1; 58.7; 56.4; 55.5; 54.9; 41.4; 40.9; 37.2; 36.7; 32.2; 31.4; 29.2; 20.7; 19.5; 16.4; 15.2. ESI-MS: 487.7 ($[\text{M} + \text{H}]^+$).

Synthesis of compound 7

$\text{CH}_3\text{NH}_2 \cdot \text{H}_2\text{O}$ (1 mL) was added to a solution of compound **7a** (1 mmol) in 3 mL anhydrous MeOH at room temperature. The mixture was stirred at this temperature for 4 h. The mixture was concentrated and purified by silica column chromatography with a 40:1 solution of CH_2Cl_2 -MeOH to afford **7** (yield: 90%).

Solasodine 3-O-glucosaminidase (7). $^1\text{H-NMR}$ (DMSO, 600 MHz) δ : 0.75 (3H, s, Me-18), 0.78 (3H, d, $J = 6.6$ Hz, Me-27), 0.88 (3H, d, $J = 6.6$ Hz, Me-21), 0.98 (3H, s, Me-19), 2.67 (2H, m, H-26), 3.66 (1H, d, $J = 10.8$ Hz, H-3), 4.56 (1H, s, H-16), 5.21 (1H, s, H-1'), 5.34 (1H, d, $J = 4.2$ Hz, H-6); $^{13}\text{C-NMR}$ (DMSO, 600 MHz) δ : 140.7; 121.7; 98.0; 77.7; 73.9; 70.5; 65.5; 62.7; 61.2; 59.0; 56.8; 50.0; 41.4; 38.3; 37.2; 36.8; 34.2; 32.2; 31.5; 29.5; 20.8; 19.9; 19.6; 16.6; 15.7. ESI-MS: 574.8 ($[\text{M} + \text{H}]^+$).

Identification and evaluation of agents isolated from traditionally used herbs against *Ophiophagus hannah* venom

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ABSTRACT: The aim of this study was firstly to identify active molecules in herbs, that are traditionally used for the treatment of snake bite, such as *Curcuma antinaia*, *Curcuma contravenenum*, *Andrographis paniculata*, and *Tanacetum parthenium*; secondly to test similar structurally related molecules and finally to prepare and evaluate an efficient formulation against *Ophiophagus hannah* venom intoxication. Three labdane based compounds, including labdane dialdehyde, labdane lactone, and labdane trialdehyde and two lactones including 14-deoxy-11,12-didehydroandrographolide and parthenolide were isolated by column chromatography and characterised. Using the isolated rat phrenic nerve-hemidiaphragm preparation, the antagonistic effect of crude extracts, isolated compounds and prepared formulations were measured *in vitro* on the inhibition of the neuromuscular transmission. Inhibition on muscle contraction, produced by the 5 µg/mL venom, was reversed by test agents in organ bath preparations. A labdane trialdehyde, isolated from *C. contravenenum*, was identified as the best antagonising agent in the low micromolar range. Tests on formulations of the most potent *C. contravenenum* extract showed, that the suppository with witepsol H15 was an effective medicine against *O. hannah* venom. This study elucidated the active compounds, accounting for the antivenin activity of traditionally used herbs and suggested the most suitable formulation, which may help to develop potent medicines for the treatment of snake bite in the future.

Keywords: Antivenin activity, rat nerve-hemidiaphragm, labdane dialdehyde, labdane trialdehyde, parthenolide, desoxy-andrographolide

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1. Introduction

The actual incidence and the severity of snake poisoning are currently highly undervalued (1). The importance of snake bites is considered a major occupational disease causing both disabilities and mortalities. This disease is causing devastation to individuals, who are involved in agricultural work in the tropical regions worldwide. Inaccurate epidemiological data resulted in the underestimation of this international problem, which requires both high attention and sincere efforts to alleviate its burden (2). Based on these facts, the World Health Organization (WHO) is calling for new, proven and affordable treatments. Traditional medicine or herbal medicine has long been used for the treatment of snake bite worldwide for its affirmed effectiveness, easy availability, and fine economic affordability. However, the active ingredients contained in these herbs and most effective formulations are still needed to be elucidated.

We have previously reported a labdane dialdehyde structure (Figure 1), which was isolated from a novel *Curcuma zedoaroides* species and exhibited well antivenin activity (3). Species in the genus *Curcuma* including *Curcuma antinaia* and *Curcuma contravenenum* were in regular use in Thailand against cobra intoxication, but they are very hard to find on local markets in Isarn today. Besides of those herbs, another two medical plants including *Andrographis paniculata* and *Tanacetum parthenium* are widely applied for the treatment of snake bite in India and China. *Andrographis paniculata* is in use in India as a snake venom antidote.

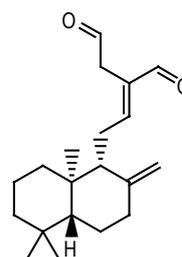


Figure 1. Chemical structure of labdane dialdehyde.

The leaves of *A. paniculata*, locally known as Nilavembu, were grinded into a paste and applied topically at the site of the snake's bite (4). *Tanacetum parthenium*, a member of the Asteraceae family, which is also known as the Compositae family, represents a herb well known for its medicinal properties. Since ancient times this herb was used by the Greeks and the Egyptians as well as the early Europeans for the treatment of a number of illnesses, such as headaches, stomach ache, menstrual pain, joint pain, fever. The Chinese used this herb due to its healing properties against insects and snake bites (5). In order to further elucidate the ingredients, that account for antivenin activity of these herbs and develop formulations, that are effective against snake venom, we prepared and tested the *in vitro* antivenin activity of the crude extracts, purified compounds and certain formulations against *Ophiophagus hannah* venom in the present study.

2. Materials and Methods

2.1. Materials

Lyophilized *O. hannah* venom was obtained from the Queen Suavabha Memorial Institute, Bangkok, Thailand. The venom was dissolved in normal saline, aliquoted and kept at -20°C as stock solution. Herbs including *C. zedoaroides*, *C. antinaia*, *C. contravenenum*, *A. paniculata*, and *T. parthenium* were purchased from the King Cobra Village, Khon Kaen Province, Thailand. The chemicals and solvents were purchased from Aldrich (Gillingham, UK) and Lancaster Synthesis (Lancaster, UK). Mass spectra were obtained by Atmospheric Pressure Chemical Ionisation (APCI), using a Hewlett-Packard 5989b quadrupole instrument (Vienna, Austria). Both proton and carbon NMR spectra were obtained on a Bruker AC 250 instrument (Follanden, Switzerland), calibrated with the solvent reference peak. Infra-red spectra were plotted from KBr discs on a Mattson 300 FTIR spectrophotometer (Coventry, UK).

Laboratory and HPLC grade dichloromethane, petroleum ether 60-80°C, ethyl acetate, chloroform-d and methanol were purchased from Fisher-Scientific (Waltham, MA, USA). The Soxhlet extraction system was the Quickfit (C5/23) model with (24/29) joint from BÜCHI, Switzerland. The heating mantle used was the Heidolph EKT 3001 from Sigma-Aldrich, UK. The rotary evaporator used was the BÜCHI Rotavapor Model R-144, Switzerland. $^1\text{H-NMR}$ and $^{13}\text{C-NMR}$ were recorded in Bruker advance 400 in chloroform-d. Column chromatography was carried out on silica gel and thin-layer chromatography (TLC) on TLC-silica gel 60 F₂₅₄.

2.2. Preparation of crude extracts and isolation of the tested compounds

The rhizomes of *C. zedoaroides*, *C. antinaia*, and *C. contravenenum* (30 g of the dried, powdered rhizomes)

were extracted with dichloromethane using a Soxhlet extraction apparatus, respectively. Once the Soxhlet extractions were completed the solvent was removed under reduced pressure from the round bottom flasks using a rotary evaporator. The extract content of *C. zedoaroides*, *C. antinaia*, and *C. contravenenum* were gathered and the entry code **Kae1**, **Mia3**, and **Rat7** was assigned. Column chromatography of the crude extract was performed using petroleum ether:ethyl acetate (60:40) as solvent system and the compounds were detected with UV light, permanganate, and 2,4-dinitrophenylhydrazine (DNPH). The **Kae1** extract finally provided labdane dialdehyde; **Mia3** extract finally provided labdane dialdehyde and labdane lactone; **Rat7** extract finally provided labdane dialdehyde, labdane lactone, and labdane trialdehyde. The spectra data of these three compounds are shown in the Appendix.

Fifty grams of dried and ground leaves of *A. paniculata* were extracted with 500 mL of methanol using a Soxhlet extractor. The solvent was evaporated off solvent using rotary evaporation. TLC analysis was performed with a dichloromethane:methanol (95:5) solvent mixture, and both andrographolide and 14-deoxy-11,12-didehydroandrographolide, were purified by column chromatography. The spectra data of 14-deoxy-11,12-didehydroandrographolide are shown in the Appendix.

Fifty grams of grinded plant material *T. parthenium* was extracted with 500 mL of chloroform using a Soxhlet extractor. The extraction was monitored by TLC with ethyl acetate:petroleum ether 60-80°C (70:30) and the TLC plate was developed with vanillin reagent and heated at 400°C for 1 min. Two point five gram of crude extract was dissolved in a minimum volume of ethyl acetate in the heat and 1/3 of the volume of petrol ether was added and cooled on ice. The yellowish sample was recrystallised from ethyl acetate:petroleum ether (70:30) to give parthenolide, 130 mg, as an off white powder. The spectra data of this compound are shown in the Appendix.

2.3. Preparation of *C. contravenenum* formulations

Ethanol solution (entry code: OHRat7): For the preparation of the ethanolic solution, 0.5 g of the concentrated extract of *C. contravenenum* was dissolved in 5 mL of ethanol and divided into two portions of 2.5 mL each. To one of the samples **OHRat7** 3 drops of trifluoroacetic acid were added. Both samples were incubated at 36°C for a period of 1 week, then tested.

Water for injection (entry code: AqRat7): For the preparation of a solution for the injection, 2.5 equivalents of an aqueous solution of 10% $\text{Na}_2\text{S}_2\text{O}_5$ to 0.2 g of the extract in 0.1 mL DMSO were mixed until homogenous and water was added until a final volume of 4 mL.

Suppositories (entry code: Rat7S): Suppositories (3 g) containing 30% active ingredients were prepared by the fusion method as previously described (6). Sixty six grams of *C. contravenenum* rhizomes, giving 1.98 g crude extract, were extracted with 300 mL of acetone as described. The

Table 1. Antivenin activity of crude extracts from *Curcuma* species

Entry Code	Source	Yield	Composition	Muscle contraction response
Kae1	<i>C. zedoaroides</i>	7.0%	Labdane dialdehyde, 79%	32.1% (50 µg/mL); 63.1% (100 µg/mL)
Mia3	<i>C. antinaia</i>	2.6%	Labdane dialdehyde, 45% Labdane lactone, 39%	26.4% (50 µg/mL); 56.5% (100 µg/mL)
Rat7	<i>C. contravenenum</i>	3.0%	Labdane dialdehyde, 32% Labdane lactone, 37% Labdane trialdehyde, 8%	53.5% (50 µg/mL); 82.6% (100 µg/mL)

Measurement time is 30 min; full contraction 100%, venom 0% at 5 µg/mL.

solvent was evaporated off to about a third of the volume and 6 g witepsol H15, a synthetic fat was added and the complete solvent was then evaporated off in vacuum. The homogeneous melt was poured into the suppository moulds and after cooling the suppositories were obtained. The melting point of the prepared suppositories was detected at 32°C.

2.4. *In vitro* antivenin activity assay

Sprague Dawley rats (200-250 g) were obtained from the Animal House, Faculty of Medicine, Khon Kaen University. The treatment procedures, according to current UK legislation, were approved by the bioethics committee, Faculty of Medicine, Khon Kaen University (HO 2434-76). Animals had free access to fresh water and food pellets. They were exposed to automated 12 h light cycles.

Rat phrenic nerve-hemidiaphragms were prepared according to the staff of the Department of Pharmacology, University of Edinburgh (1970) and the contractile responses were studied. The entire nerve-muscle preparation was submerged in 50 mL Krebs's solution with carbogen and the temperature was maintained at 37°C. The phrenic nerve was stimulated with a rectangular-wave pulse of 0.5 msec/0.5 Hz through a bipolar platinum electrode, using a Grass Model S-48 stimulator. Muscle contraction was recorded with a force transducer and Grass Polygraph recorder. The indicated doses of crude extract (**Kae1**, **Mia3**, and **Rat7**), test solutions of the isolated compounds in DMSO, and prepared formulations (**OHRat7**, **AqRat7**, and **Rat7S**) were added, respectively, to a circulating water bath at 37°C for 30 min and then venom was added and 30 min later the muscle contraction was accessed. Two phrenic nerve-hemidiaphragm preparations were obtained from one animal. Control, DMSO, was set to 100%, 5 µg/mL venom to 0%. The % test response (muscle contraction) was measured at 30 min in presence of the test compounds and the venom. In the control group, 5 µg/mL venom alone gradually and completely inhibited the indirectly-evoked twitches within 30 min (0% response). In presence of the antidote the contraction remained at a certain percentage of the full contraction. This effect was used here to screen the 3 *Curcuma* plant extracts, the isolated compounds and the formulations of the best plant.

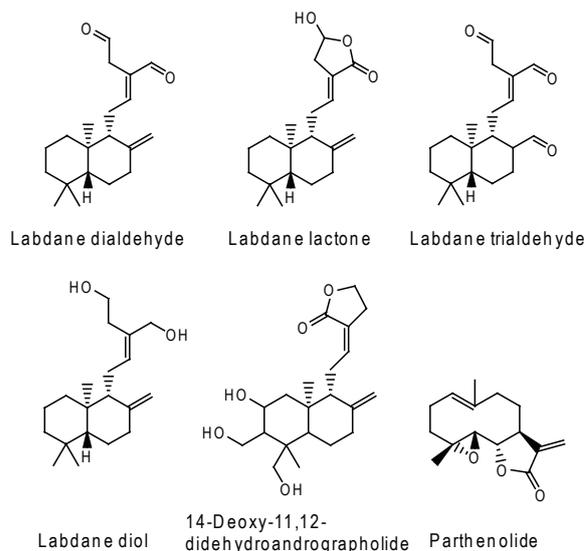


Figure 2. Chemical structures of the isolated compounds.

3. Results and Discussion

3.1. *In vitro* antivenin activity of crude extracts from the genus *Curcuma*

The protecting effects of crude extracts **Kae1**, **Mia3**, and **Rat7** from *C. zedoaroides*, *C. antinaia*, and *C. contravenenum* against *O. hannah* venom on the neuromuscular transmission of the rat phrenic nerve-hemidiaphragm are shown in Table 1. The **Kae1** extract was obtained in the highest yield (7%) and at a 100 µg/mL dose the response was still above 60%. The best protection was determined for **Rat7** extract (yield 3.0%), giving for the 100 µg/mL organ bath concentration a more than 80% of the original contraction of the diaphragm. **Mia3** extract was obtained in the lowest yield (2.6%) and demonstrated the lowest antivenom activity. The muscle contract response was determined as 56.5% when exposed to 100 µg/mL of this extract.

3.2. Analysis of the ingredients in the crude extracts and evaluation of their antivenin activity

The crude extracts from the five medicinal herbs, used in the present study, were further purified and generated the test compounds (Figure 2). The antivenin activities of

these agents were examined and the results are shown in Table 2. The crude extract **Kae1** from *C. zedoaroides* is a very good source of the labdane dialdehyde, which was obtained from this species in 79% yield and excellent purity, by simple column chromatography. A 10 µg/mL concentration of this compound showed 83% protection against the snake venom.

The crude extract **Mia3** from *C. antinaia* provided in addition to labdane dialdehyde, a second labdane derivative, identified as labdane lactone (Figure 2), which showed a bioactivity about 60%. Its formation can be understood if we assume that the labdane dialdehyde is oxidised to the corresponding carboxylic acid and the second aldehyde is able to form the hydroxyl-lactone structure of labdane lactone. The proportions of labdane dialdehyde and labdane lactone in **Mia3** were detected as 45% and 39%, respectively. Labdane lactone was previously isolated from members of the Zingiberaceae family, which are traditionally used as a medicine against inflammatory diseases. This anti-inflammatory agent regulates NF-κB-regulated cellular responses in particular it inhibited NF-κB activation, suppressed phosphorylation, p65 nuclear translocation and reporter gene transcription (11).

From *C. contravenenum* extract **Rat7**, in addition to labdane dialdehyde and labdane lactone, a third molecule labdane trialdehyde (Figure 2) was finally isolated. The percentages of labdane dialdehyde, labdane lactone, and labdane trialdehyde in this extract were determined as 32%, 37%, and 8%, respectively. Labdane trialdehyde

maintained nearly a full diaphragm contraction and with 99.5% protection it is considered the best venom antidote, reported to date. Comparing the activities of these three molecules, the labdane trialdehyde is the most potent antivenin agent, followed by labdane dialdehyde and labdane lactone. Through analyzing the relationship between activities and structures of the above isolated compounds, the succindialdehyde structure may be essential and if the dialdehyde moiety was reduced to a diol the activity was lost (labdane diol, Figure 2). The labdane trialdehyde was previously isolated from myoga extracts on the search for inhibitors of human platelet aggregation and human 5-lipoxygenase. This compound was found to be a potent inhibitor of human platelet aggregation and human 5-lipoxygenase (7). The percentage, in which it is present in this curcuma species, is with 8% a good new source of this agent and therefore, it can be evaluated for further therapeutic applications.

Guided by previous enzyme-linked immunosorbent assay (ELISA) study (8) *T. parthenium* (Feverfew) and *A. paniculata* were also investigated in addition to the *Curcuma* plant. From *A. paniculata*, andrographolide and 14-deoxy-11,12-didehydroandrographolide were isolated and tested. The desoxy-derivative displayed a weak inhibition of 32% at the 10 µM test concentration while andrographolide was found inactive as antidote. As 14-deoxy-11,12-didehydroandrographolide is only present in less than 0.01% in the traditional formulation, known as Nilavembu, it cannot be recommended. The concentration and resulting of this the active amount of 14-deoxy-11,12-didehydroandrographolide is simply too low to have any medicinal effect. *T. parthenium* is a herb well known for its medicinal properties. The Chinese used this herb due to its healing properties against insects and snake bites. Overall parthenolide isolated from this herb had reasonable antivenom activity of about 54% and it can easily be isolated in good quantities by crystallisation. It is the only European herb, which is readily available in a commercial formulation as a tablet from e.g. simply supplements (9).

3.3. In vitro antivenin activity of *C. contravenenum* formulations

C. contravenenum is the easiest plant in terms of plant production and has the biggest rhizome based on the total weight of the entire plant. It gave a 3.0% yield of the crude extract (**Rat7**) which exhibited most potent antivenin activity. For these reasons, three formulations of **Rat7** extract were prepared and tested on their antivenin activity (Table 3). The ethanolic extract of **Rat7** was

Table 2. Antivenin activity of isolated compounds

Compound	Muscle contraction response
Labdane dialdehyde	
1 µM	66.1%
10 µM	82.6%
Labdane diol	
1 µM	0
10 µM	2.1%
Labdane lactone	
1 µM	45.6%
10 µM	61.9%
Labdane trialdehyde	
1 µM	83.5%
10 µM	99.5%
14-Deoxy-11,12-didehydroandrographolide	
1 µM	12.2%
10 µM	32.4%
Parthenolide	
1 µM	29.4%
10 µM	54.0%

Measurement time is 30 min; full contraction 100%, venom 0% at 5 µg/mL.

Table 3. Antivenin activity of crude extracts from *Curcuma* species

Entry Code	Formulation	Muscle contraction response
OHRat7	10% in ethanol	1.1% (500 µg/mL); 0.2% (1,000 µg/mL)
AqRat7	Hydrogensulfite adduct; 5% in water	5.1% (1,000 µg/mL); 6.3% (5,000 µg/mL)
Rat7S	Suppository 30%	46.1% (200 µg/mL); 77.4% (400 µg/mL)

Measurement time is 30 min; full contraction 100%, venom 0% at 5 µg/mL.

found inactive, and therefore formulations in Thai whisky cannot be recommended. Chemically, the aldehyde functionality has been converted into an ethoxy semi-acetale. Thus, a protected, chemically unreactive structure was formed without any bioactivity. The formulation as hydrogen sulfite adduct was supposed to work as an injection in water, but again with the loss of the chemical reactivity the bioactivity was also lost, at least in the *in vitro* (*ex vivo*) experiment. The formulation of the extract **Rat7** as a suppository worked well; *in vitro* 77% response remained for the 400 µg/mL dose, compared with the original extract of **Rat7** at a 100 µg/mL dose, which gave 83% venom protection. Based on a good *in vitro-in vivo* correlation, we obtained from a previous study, 1-2 suppositories (3 g) should translate into a working anti-venom medication.

Three curcuma species, *i.e.* *C. zedoaroides*, *C. antinaia*, and *C. contravenenum*, and two other traditional medicines *A. paniculata*, and *T. parthenium* were confirmed active against *O. hannah* venom in our *ex-vivo* assay. In addition to the labdane dialdehyde, which was discovered in *C. zedoaroides* in our previous study, labdane lactone and labdane trialdehyde, isolated from curcuma species *C. antinaia* and *C. contravenenum* were found effective now against the venom. Labdane trialdehyde is the best anti-neurotoxic agent known to date. However, the isolated labdane trialdehyde is unstable. It is supposed, that the natural plant formulation is stabilising labdane trialdehyde with the labdane dialdehyde. Feverfew extract at high doses may be used for snake venom intoxication as a common European alternative. It should be noted, that the only efficient formulation is a suppository, in addition to the preparation of the freshly grinded root of the fresh rhizome. These results provided evidences about the usefulness of some traditional medicines as antidotes and gave clues on the drug development in the future. Further studies are ongoing to replace the *in vitro* antivenin assay, used in this study, by an *in vitro* method, in which chicken intestine is used and not laboratory animal tissue.

Acknowledgements

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Appendix

Labdane dialdehyde. 2-[2-(5,5,8a-Trimethyl-2-methylene-decahydro-naphthalen-1-yl)-ethylidene]-succinaldehyde. The ¹H-NMR and ¹³C-NMR data are identical with Lattmann *et al.* (10).

¹H-NMR (CDCl₃, 300 MHz): δ: 9.7 + 9.5 (1H, 1H, CHO), 6.75 (1H, t, C12H), 4.90 + 4.40 (1H, 1H, s, C17H), 3.40 (2H, d, C14H), 2.40 (3H, m, C11H, C6H), 2.0-1.0 (10H, m), 0.9 + 0.8 + 0.7 (9H, s, 3×Me); ¹³C-NMR: δ: 196.38, 192.60 CHO, C15 + C16; 146.96, 133.78 C8, C13, 159.05 C12, 106.33 C17, 55.20, 54.18 C6, C7, 32.22, 39.34 C1, C5, 34.15, 23.83, 14.23 Me, 40.54, 38.30, 38.15, 36.78, 23.55, 23.52, 18.23 C14, C11, C9, C10, C2, C3, C4.

Labdane lactone. 5-Hydroxy-3-[2-(5,5,8a-trimethyl-2-methylene-decahydro-naphthalen-1-yl)-ethylidene]-dihydro-furan-2-one. The $^1\text{H-NMR}$ and $^{13}\text{C-NMR}$ data are identical with Kunnumakkara *et al.* (11).

$^1\text{H-NMR}$ (CDCl_3 , 300 MHz): δ : 6.69 (br s, 1H), 5.94 (br s, 1H), 4.81 (s, 1H), 4.36 (br s, 1H), 2.99 (m, 1H), 2.71 (m, 1H), 2.40-2.36 (m, 2H), 2.20 (m, 1H), 2.05-1.08 (m, 12H), 0.89 (s, 3H), 0.82 (s, 3H), 0.71 (s, 3H); $^{13}\text{C-NMR}$ (75 MHz): δ : 170.65, 148.01, 143.45, 124.48, 107.52, 96.46, 56.02, 55.21, 41.90, 39.94, 39.12, 37.68, 33.57, 33.45, 25.58, 23.99, 21.62, 19.21, 14.23.

Labdane trialdehyde. 2-[2-(2-Formyl-5,5,8a-trimethyl-decahydro-naphthalen-1-yl)-ethylidene]-succinaldehyde. The $^1\text{H-NMR}$ and $^{13}\text{C-NMR}$ data are identical with Suebsasana *et al.* (7).

$^1\text{H-NMR}$ (CDCl_3 , 300 MHz): δ : 9.96 (s, 1H, H-17), 9.62 (s, 1H, H-15), 9.50 (s, 1H, H-16), 7.10 (dd, 1H, $J = 6.0$, 8.8, H-12), 3.54 (d, 1H, $J = 17$, H-14a), 3.48 (d, 1H, $J = 17$, H-14b), 2.85 (m, 1H, $J = 8.8, 12, 15.2$, H-11a), 2.65 (m, 1H, $J = 4, 6, 15.2$, H-11b), 2.50 (m, 1H, H-8), 2.35 (m, 1H, H-7b), 1.92 (m, 1H, H-5), 1.75 (m, 1H, H-1b), 1.60 (m, 2H, H-6), 1.42 (m, 1H, H-3b), 1.40 (m, 2H, H-2), 1.40 (m, 1H, H-7a), 1.22 (m, 1H, H-3a), 1.15 (m, 1H, H-1a), 1.05 (m, 1H, H-9), 0.88 (s, 2H, H-18), 0.82 (s, 3H, H-19), 0.80 (s, 3H, H-20); $^{13}\text{C-NMR}$ (75 MHz): δ : 205.37 (C-17), 198.84 (C-15), 194.80 (C-16), 158.72 (C-12), 138.35 (C-13), 56.87 (C-9), 55.03 (C-5), 49.23 (C-8), 43.16 (C-3), 40.29 (C-14), 39.88 (C-10), 39.88 (C-1), 34.25 (C-4), 34.25 (C-18), 25.65 (C-7), 27.14 (C-11), 22.31 (C-19), 20.11 (C-6), 19.82 (C-2), 16.16 (C-20).

14-Deoxy-11,12-didehydro-andrographolide. The $^1\text{H-NMR}$ and $^{13}\text{C-NMR}$ data are identical with Suebsasana *et al.* (7).

$^1\text{H-NMR}$ (CDCl_3 , 300 MHz): δ : 7.43 (1H, t, $J = 1.76$ Hz, H-14), 6.85 (1H, dd, $J = 10.1$ and 15.8 Hz, H-12), 6.15 (1H, d, $J = 15.8$ Hz, H-11), 4.86 (2H, d, $J = 1.3$ Hz, H-15), 4.75 (1H, d, $J = 1.8$ Hz, 17a), 4.49 (1H, d, $J = 1.8$ Hz, 17b), 4.12 (1H, d, $J = 11.0$ Hz, H-19a), 3.39 (1H, t, $J = 5.3$ Hz, H-3), 3.38 (1H, d, $J = 11.4$ Hz, H-19b), 1.22 (3H, s, H-18), 0.83 (3H, s, H-20); $^{13}\text{C-NMR}$: δ : 172.2 (C=O, C-16), 148.0 (C, C-8), 142.8 (CH, C-12), 136.0 (CH, C-11), 129.2 (C, C-13), 121.1 (CH, C-14), 109.2 (CH₂, C-17), 80.8 (CH, C-3), 69.5 (CH₂, C-15), 64.2 (CH₂, C-19), 61.7 (CH, C-9), 54.7 (CH, C-5), 43.0 (C, C-4), 38.5 (C, C-10), 38.2 (CH₂, C-1), 36.6 (CH₂, C-7), 28.1 (CH₂, C-2), 22.9 (CH₂, C-6), 22.6 (CH₃, C-18), 15.9 (CH₃, C-20).

Parthenolide. The $^1\text{H-NMR}$ and $^{13}\text{C-NMR}$ data are identical with Tiunan *et al.* (12).

$^1\text{H-NMR}$ (CDCl_3 , 300 MHz): δ : 6.34 (d, $J = 3.6$ Hz, H-13 α), 5.62 (d, $J = 3.0$ Hz, H-13 β), 5.21 (dd, $J = 2.7$, 12.0 Hz, H-1), 3.86 (t, $J = 8.4$ Hz, H-6), 2.79 (d, $J = 9.0$ Hz, H-5), 2.74 to 2.82 (m, H-7), 2.32 to 2.44 (m, H-9 β), 2.32 to 2.49 (m, H-2 β), 2.11 to 2.21 (m, H-2 α , H-3 β , H-8 α , H-9 α), 1.72 (s, H-14), 1.70 to 1.77 (m, H-8 β), 1.31 (s, H-15), 1.20 to 1.28 (m, H-3 α). $^{13}\text{C-NMR}$ (CDCl_3 , 75.5 MHz): δ : 169.3 (C-12), 139.2 (C-11), 134.6 (C-10), 125.3 (C-1), 121.3 (C-13), 82.4 (C-6), 66.4 (C-5), 61.5 (C-4), 47.7 (C-7), 41.2 (C-9), 36.3 (C-3), 30.6 (C-8), 24.1 (C-2), 17.3 (C-15), 16.9 (C-14).

***LKB1*, *TP16*, *EGFR*, and *KRAS* somatic mutations in lung adenocarcinomas from a Chiba Prefecture, Japan cohort**

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ABSTRACT: The discovery of somatic mutations in cancer-related genes has been applied to understand the genetic basis of cancer. Here we report somatic mutations of two tumor suppressor genes: *LKB1* (exons 1, 4, and 8) and *TP16* (*CDKN2A*) (exons 1 and 2); and two oncogenes: epidermal growth factor receptor *EGFR* (exons 18-21) and *KRAS* (exon 2) in 97 lung adenocarcinoma tissues in a cohort from the Kujukuri coast area of Chiba, Japan. In the *LKB1* gene, only F354L substitutions were observed in 14 of the 97 tissue samples (14.4%). In the *TP16* gene, only two deletions were observed in contrast to previous reports. On the other hand, the *EGFR* gene was highly mutated (38.1%) and mainly L858R substitutions occurred (23.7%) as well as insertions and deletions. In the *KRAS* gene, 10 substitutions at codon 12 were observed (10.3%). Co-occurrence of *EGFR* and *KRAS* somatic mutations was identified in one patient, those of *EGFR* and *LKB1* were in three patients, and those of *KRAS* and *LKB1* were in four patients. The lower rates of *LKB1*, *TP16*, and *KRAS* somatic mutations in lung adenocarcinomas are characteristic of the Kujukuri cohort as compared to Caucasians.

Keywords: Chromatograms, tumor suppressor genes, oncogenes, mTOR pathway, RAS pathway, co-occurrence of mutations

1. Introduction

The *LKB1* tumor suppressor gene, which encodes serine-threonine kinase 11 (STK11), has been shown to regulate cell cycle progression, apoptosis, and cell polarity (1). Previous reports have suggested a *LKB1* somatic mutation rate as high as 30% in non-small cell lung carcinoma (NSCLC) tumors derived from Caucasian patients (2,3),

while being infrequent in NSCLC Asian patients (3%) (4). Another review (5) reported that *LKB1* somatic mutations were observed in 20.18% of lung adenocarcinoma by analyzing the Catalog of Somatic Mutations in Cancer (COSMIC) (6), a large-scale database curated by the Wellcome Trust Sanger Institute. The *TP16* gene (*CDKN2A*, cyclin-dependent kinase inhibitor 2A) is also a tumor suppressor gene and has been reported to show a high rate of deletions as a major type of somatic mutation in cancer, consisting of 24.57% of lung adenocarcinomas in the COSMIC database (5). In contrast to these tumor suppressor genes, *EGFR* and *KRAS* oncogenes play a central role in tumorigenesis. The tyrosine kinase activity of *EGFR* phosphorylates tyrosine residues of target proteins, including *KRAS*, to initiate multiple signaling pathways resulting in cell proliferation, migration, metastasis, resistance to apoptosis, and angiogenesis (7,8). Small-molecule inhibitors of *EGFR* tyrosine kinase activity (TKIs) such as gefitinib and erlotinib provide a good concept for anticancer drugs. However, clinical trials have revealed significant variability in the TKI response depending on *EGFR* and *KRAS* gene mutations (9). Interestingly, somatic *EGFR* and *KRAS* gene mutations, which are upstream and downstream of the RAS pathway, respectively, have been reported to be mutually exclusive. Thus, concurrent *EGFR* and *KRAS* mutations have been very rare (5).

Lung adenocarcinoma is the most common form of lung cancer, which has an average 5-year survival rate of 15%, mainly because of late-stage detection and paucity of late-stage treatment options (10,11). To determine the genetic basis of lung adenocarcinoma in a cohort from the Kujukuri coast area in Chiba prefecture, Japan, we analyzed exons 1, 4, and 8 of *LKB1*, exons 1 and 2 of *TP16* which includes almost the entire sequence of the protein coding region, exons 18-21 of *EGFR*, and exon 1 of *KRAS*, according to the histogram of somatic mutations in *LKB1*, *TP16*, *EGFR* and *KRAS* from the COSMIC database (6).

2. Materials and Methods

2.1. Patients

Primary tumor samples from patients with lung adeno-

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carcinoma were obtained from 97 randomly selected patients (54 males and 43 females) with lung cancer who were diagnosed and had undergone surgical resection at Asahi General Hospital, Asahi-Shi, located in the Kujukuri coast area, Chiba prefecture, Japan from September 2004 to March 2009. This study was conducted according to the principles expressed in the Declaration of Helsinki. Asahi General Hospital obtained written informed consent from all subjects. The Asahi General Hospital Institutional Review Board approved use of human tissue in this study according to the Ethical Guidelines of the Ministry of Health, Labour, and Welfare of Japan. The data were analyzed anonymously. All patients were pathologically diagnosed as having primary lung adenocarcinoma and subclassified into histological subtypes by examining more than 10 histological slides. The age distribution was: 5 patients under 50-years old; 12 patients 50- to 59-years-old; 28 patients 60- to 69-years-old; 39 patients 70- to 79-years-old; and 13 patients 80 or over (average = 69.11, S.D. = 9.94).

2.2. DNA extraction and sequencing analysis

Genomic DNA extraction was performed using a simple 96-well plate based DNA extraction. A few milligrams of the frozen specimen was digested with Proteinase K at 55°C for 2 h in 100 µL of extraction buffer (10 mM Tris-HCl pH 7.5, 5 mM EDTA, 0.5% Tween 20) followed by heat inactivation at 95°C for 10 min. The preparation was centrifuged at 3000 × g for 15 min, and 2-5 µL of the supernatant was used as a PCR template.

Mutations were detected using PCR-based direct sequencing of three *LKB1* exons (exons 1, 4, 8), two *TP16* exons (exons 1, 2), four *EGFR* exons (exons 18-21), and exon 1 of the *KRAS* gene. The primers for PCR amplification and sequencing analysis are shown in Table 1.

PCR amplification was done in 50 µL reactions containing 0.2 µM of each primer pair and 0.25 units of Pfu DNA polymerase (Celestar-Lexico Sciences, Chiba, Japan).

LKB1 exons were amplified by initial denaturation at 94°C for 5 min followed by 40 cycles of 10 sec at 98°C, 30 sec at 68°C, and a 1 min final extension at 72°C. *TP16* exons were amplified by initial denaturation at 94°C for 5 min followed by 40 cycles of 10 sec at 98°C, 30 sec at 65°C, 30 sec at 72°C, and a 1 min final extension at 72°C. *TP16* exon 2 amplification was performed with 5% of DMSO. *EGFR* exons were amplified by initial denaturation at 94°C for 1 min followed by 35 cycles of 5 sec at 98°C, 30 sec at 65°C, 1 min at 72°C, and a final extension at 72°C for 1 min. The exon 1 region of *KRAS* was amplified by initial denaturation at 94°C for 1 min followed by 40 cycles at 98°C for 5 sec, 50°C for 30 sec, and 72°C for 30 sec, and a final extension at 72°C for 5 min.

The PCR products were purified using a GFX96 PCR purification kit (GE Healthcare Life Sciences, Little Chalfont, UK) and sequenced directly using a 3730xl DNA Analyzer (Applied Biosystems, Foster City, CA, USA) with 1 µM of each sequencing primer.

Several samples that were difficult to determine the mutation status by the direct sequencing method were cloned into pUC118 before sequencing.

3. Results

Somatic mutations in *LKB1*, *TP16*, *EGFR*, and *KRAS* genes observed in the cohort from the Kujukuri coast area of Chiba prefecture are summarized in Table 2. The sequence chromatograms are shown in Figures 1, 2, and 3 for single nucleotide substitutions in *LKB1*, *EGFR*, and *KRAS*; two *TP16* deletions; and *EGFR* deletions and insertions, respectively.

Table 1. PCR primer sequences and sequencing primer sequences for exons 1, 4 and 8 of *LKB1*, exons 1 and 2 of *TP16*, exons 18-21 of *EGFR*, and exon 1 of *KRAS*

Gene	Exon	PCR primer sequences	Sequencing primer sequences
<i>LKB1</i>	exon 1	F 5'-TGGAGAAGGGAAGTCGGAACACAAGG-3' R 5'-GCCAGACGGGTCCAGCTCAG-3'	F 5'-GGGAAGTCGGAACACAAGGAAGGAC-3' R 5'-GAACCATCAGCACCGTGACTGGC-3'
	exon 4	F 5'-CCAGCTGGGCCTGTGGTGTTC-3' R 5'-CTGGTCCGGCAGGTGTCGTC-3'	F 5'-CCCCTGTGAGGGGCAGGGAG-3' R 5'-AACGGGTGCACTGCCTGTGG-3'
	exon 8	F 5'-AGAGGACATGGCTGAGCTTCTGTGG-3' R 5'-GGGACGTGGGATTGGCCACC-3'	F 5'-CAGAGGAGCTGGGTGGAAAACTGG-3' R 5'-TGCAGACAGGCAGGCACCCTG-3'
<i>TP16</i>	exon 1	F 5'-AGGGTCGGAGGGGGCTCTTC-3' R 5'-CTGATTCCAATTCCCCTGCAAACCTTCG-3'	F 5'-AAGAGGAGGGGCTGGCTGGTC-3' R 5'-CTCCAGAGTCGCCCGCCATC-3'
	exon 2	F 5'-GGGCTTGTGTGGGGTCTGC-3' R 5'-GGCGCTGAGCTGAGGCAAG-3'	F 5'-TGCGGTGAGGGGGCTTAC-3' R 5'-TCCCGGGCTGAACTTCTGTGC-3'
<i>EGFR</i>	exon 18	F 5'-TGTCTTCCAAATGAGCTGGCAAGTG-3' R 5'-GGAGTTCCCAAACACTCAGTGAACAAAAG-3'	F 5'-GTGTCCTGGCACCCAAGC-3' R 5'-CCCCACAGACCATGAGAG-3'
	exon 19	F 5'-GGGTGCATCGCTGGTAACATCC-3' R 5'-GATGTGGAGATGAGCAGGGTCTAG-3'	F 5'-GCAGCATGTGGCACCATCTC-3' R 5'-CAGAGCAGCTGCCAGACATG-3'
	exon 20	F 5'-CTCAAGATCGATTCATGCGTCTTCAC-3' R 5'-CACACACATATCCCCATGGCAAACCTC-3'	F 5'-GCATTCATGCGTCTTACCTG-3' R 5'-CATATCCCCATGGCAAACCTTTC-3'
	exon 21	F 5'-GTCAGCAGCGGGTTACATCTTCTTC-3' R 5'-CTCACCCAGAATGTCTGGAGAGC-3'	F 5'-CAGCCATAAGTCTCGACGTG-3' R 5'-TCCTCCCCTGCATGTGTTAAAC-3'
	<i>KRAS</i>	exon 1	F 5'-AGGTACTGGTGGAGTATTGATAGTG-3' R 5'-CATACTCCCAAGGAAAGTAAAGTCC-3'

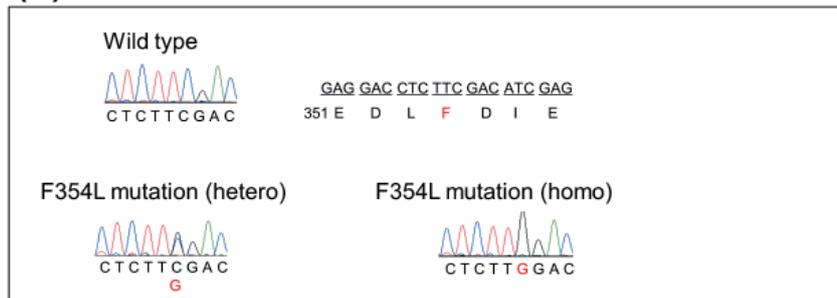
F, forward. R, reverse.

Table 2. Somatic mutations in exons 1, 4 and 8 of *LKB1*, exons 1 and 2 of *TPI6*, exons 18-21 of *EGFR*, and exon 1 of *KRAS* observed in the Kujukuri cohort

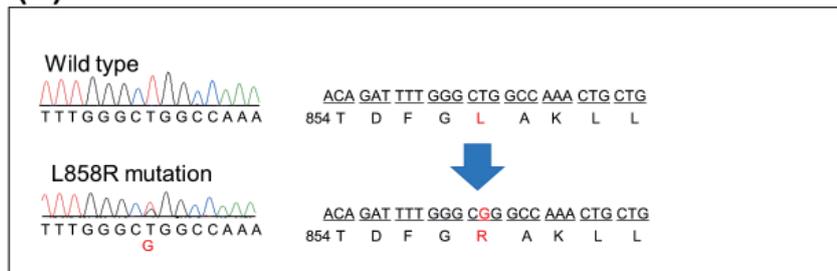
Gene/exon	Somatic mutations	Number	Male	Female
<i>LKB1</i> (exons 1, 4, 8)	ttc/ttg (F354L)	14	9	5
	<i>TPI6</i> (exons 1, 2)			
<i>TPI6</i> (exons 1, 2)	del549c	1	1	0
	del666-673	1	1	0
<i>EGFR</i> (exons 18-21)	ctg/cgg (L858R)	23	11	12
	D770-N771 ins SVD	1	0	1
	cgc/tgc (R776C), ctg/cgg (L858R)	1	0	1
	746ELREA deletion	7	6	1
	746ELREA deletion, ctg/cgg (L858R)	1	1	0
	ggc/gcc (G719A)	1	0	1
	747LRE deletion, A750P, ctg/cgg (L858R)	1	0	1
	P772-H773 ins YNP, H773Y	1	0	1
	gtg/ctg (V834L), ctg/cgg (L858R)	1	0	1
	<i>KRAS</i> (codon 1)	ggt/gtt (G12V)	1	0
ggt/tgt (G12C)		4	2	2
ggt/gat (G12D)		3	2	1
ggt/gat (G12A)		2	2	0

The two right columns show the numbers related to gender.

(A) *LKB1* F354L mutation



(B) *EGFR* L858R mutation



(C) *KRAS* codon 12 mutation

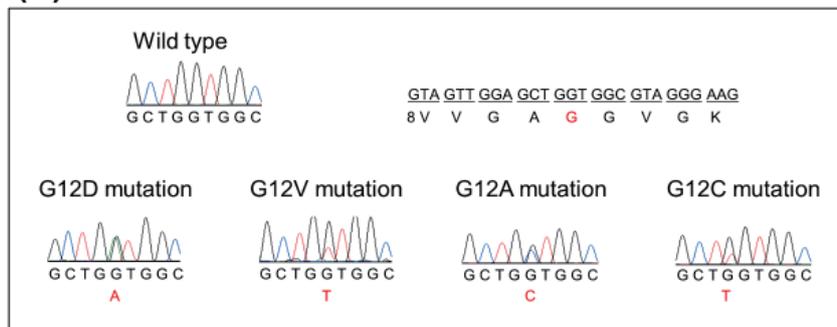


Figure 1. Wave charts of analyzed nucleotide sequences and corresponding amino acid sequences. (A) *LKB1* wild type nucleotide sequence and base substitutions (hetero and homo) at F354L. (B) *EGFR* wild type nucleotide sequence and base substitution at F858R. (C) *KRAS* wild type nucleotide sequence and base substitutions at codon 12.

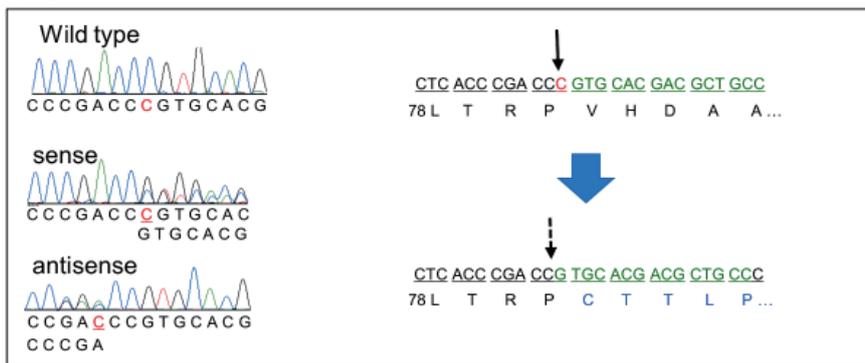
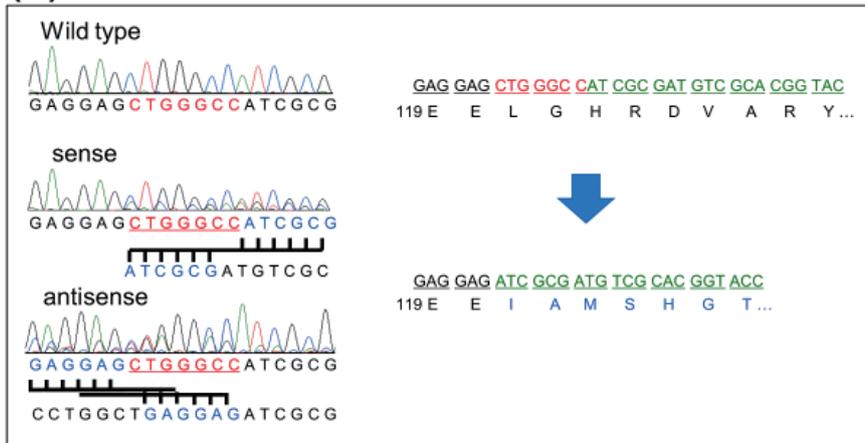
(A) 243c deletion mutation**(B) 361-367 deletion mutation**

Figure 2. Wave charts of nucleotide sequences and corresponding amino acid sequences of wild type and two deletions observed in the *TP16* gene. (A) 549C deletion mutation. (B) 666-673 deletion mutation.

3.1. *LKB1* and *TP16* mutations

We analyzed the sequence of exons 1, 4, and 8 of the *LKB1* gene and the sequence of the *TP16* gene (exons 1 and 2). Of the 97 lung adenocarcinoma tissue samples, only 14 samples (14.4%) were observed to contain F354L substitutions in exon 8 of the *LKB1* gene, of which two samples showed biallelic mutations. Figure 1A shows the three types of *LKB1* sequences observed in chromatograms: wild type, hetero F354L mutation, and biallelic (homo) F354L mutation. On the other hand, only 2 of the 97 tissue samples (2.1%) showed deletions in exon 2 of the *TP16* gene (cytosine deletion at 549 changed the amino acid sequence after 82Val; and the deletion from 666 to 673 changed the amino acid sequence after 121Leu; the numbers of nucleotides and residues are from NM_000077.4 and NP_000068.1 in the NCBI database), as shown in Figures 2A and 2B.

3.2. *EGFR* and *KRAS* mutations

The major somatic mutations of the *EGFR* gene observed in the cohort were L858R substitutions at

exon 21, as shown in the Figure 1B chromatogram. There were 23 samples of the 97 tissue samples (23.7%) that showed only this substitution and four samples that showed other types of mutations in addition to this substitution (Table 2). Another major mutation was the 746-ELREA-750 deletion at exon 19 that was observed in 7 samples (7.2%). The sequencing result chromatogram showed this deletion in Figure 3A. Figures 3B and 3C showed the sequencing results of insertion mutations. Figure 3B showed the insertion of three amino acids YNP between 772Pro and 773His in exon 20, and the histidine residue was substituted by Tyr. Figure 3C showed the insertion of three amino acids SVD between 770Asp and 771Asn in exon 20. The other mutations observed are summarized in Table 2. In total, 37 of 97 samples (38.1%) displayed somatic mutations in the *EGFR* gene.

The somatic mutations observed in the *KRAS* gene were only substitutions at codon 12 of amino acid residue Gly to Asp (3 samples), to Val (1 sample), to Ala (2 samples), and to Cys (4 samples). A total of 10 samples showed somatic mutations in *KRAS* (10.3%). Figure 1C shows the sequences of the somatic mutations at *KRAS* codon 12 on chromatograms.

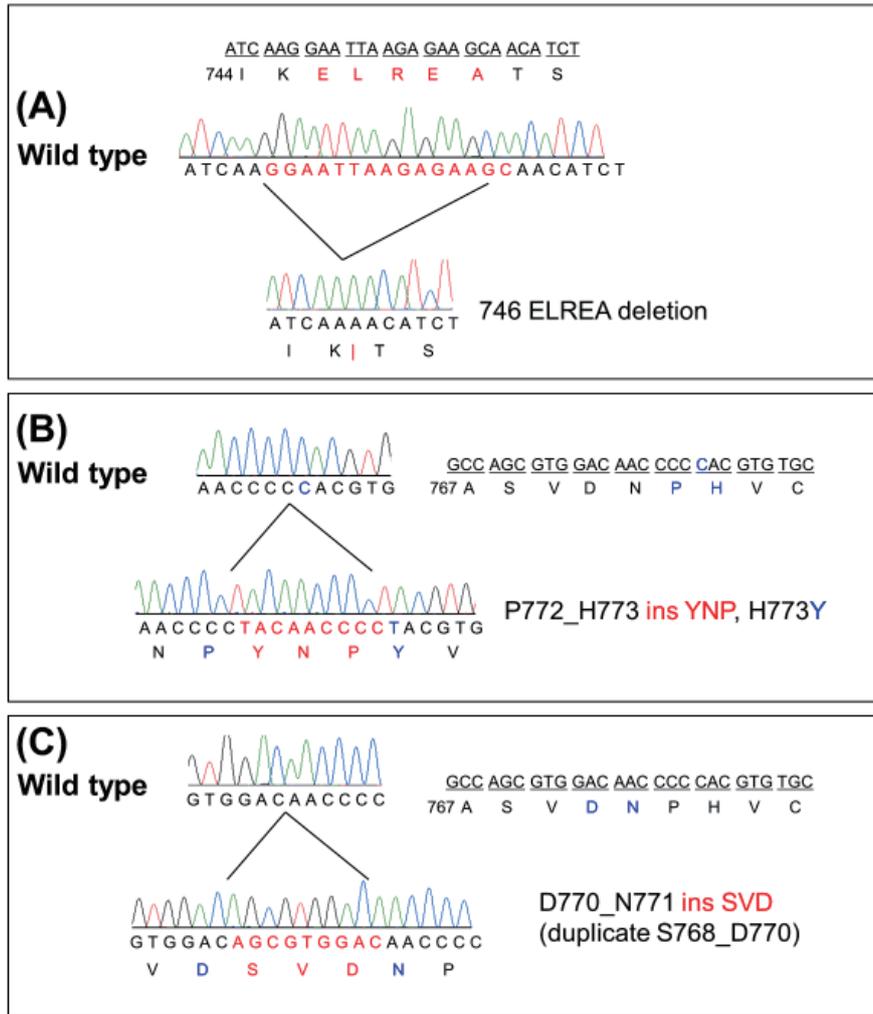


Figure 3. Wave charts of nucleotide sequences and corresponding amino acid sequences of wild type, deletions, and insertions observed in the *EGFR* gene. (A) 746ELREA deletion. (B) P772_H773 insertion YNP with H773Y. (C) D770_N771 insertion SVD (duplication S768_D770).

3.3. Concurrency of mutations

The concurrent *EGFR* and *KRAS* mutations have been reported to be very rare (5), meaning the occurrence of those mutations were mutually exclusive in lung adenocarcinoma. We observed that only one of the 97 tissue samples harbored the concurrent *EGFR* and *KRAS* mutations. In contrast, co-occurrence of *EGFR* and *LKB1* mutations were observed in three samples, while co-occurrence of *KRAS* and *LKB1* mutations were observed in four samples.

4. Discussion

We analyzed somatic mutations in tumor suppressor genes *LKB1* and *TP16*, and in oncogenes *EGFR* and *KRAS*, using 97 lung adenocarcinoma tissue samples in a cohort from the Kujukuri coast area of Chiba prefecture, Japan. Compared with the review paper that analyzed the COSMIC database (5), we showed the rate of *TP16* somatic mutations in the Kujukuri cohort

to be very small (2.1%), while the COSMIC database showed a 24.57% mutation rate. The *LKB1* mutation rate was 14.4% from our observations, while that of the COSMIC database was 20.18%. In particular, only F354L substitutions in the *LKB1* gene were observed in the Kujukuri cohort.

As for *EGFR*, the somatic mutation rate was 38.1% from our observations, which is a little bit higher than the rate from the COSMIC database (35.85%). *KRAS* also showed a lower somatic mutation rate in the Kujukuri cohort (10.3%) than that from the COSMIC database (23.02%). The lower rates of somatic mutations for *LKB1*, *TP16*, and *KRAS* genes in lung adenocarcinomas are characteristic of the Kujukuri cohort.

EGFR mutations have been more commonly found in tumors from patients who never smoked cigarettes (12), while *KRAS* mutations have been present in those with significant tobacco exposure (9,13). Unfortunately, we could not follow the history of smoking or non-smoking patients from clinical records of the cohort,

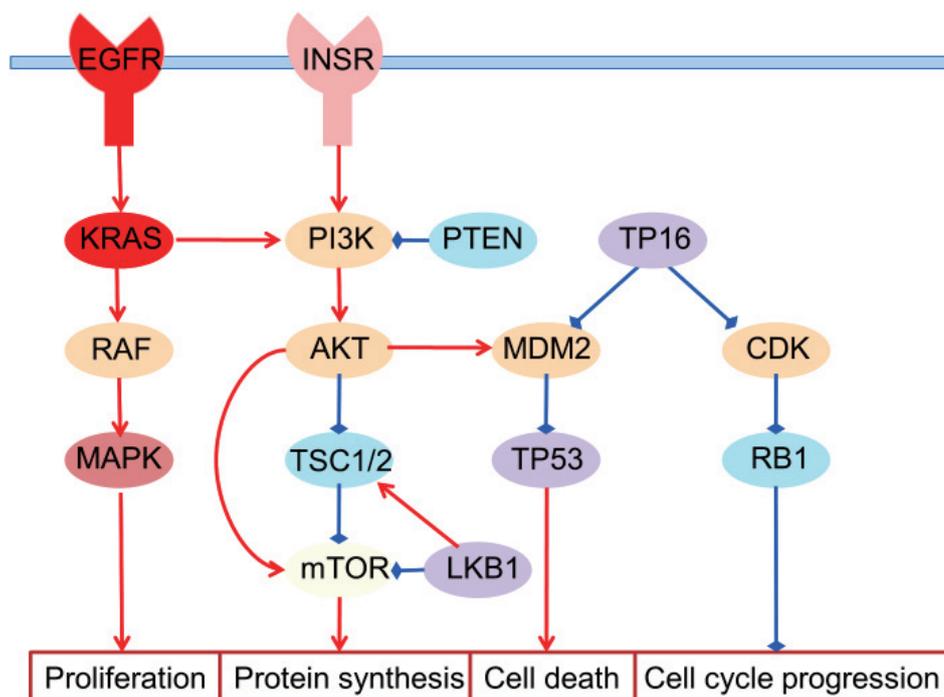


Figure 4. Pathways relating to *LKB1*, *TP16*, *EGFR*, and *KRAS*. The pathways were summarized from Kyoto Encyclopedia of Genes and Genomes (KEGG). Red lines represent activation and blue lines represent inhibition.

except for the 11 patients who were never-smokers. Among the 11 patients, we found four *EGFR* mutations (36.3% = 4/11) that included three L858R substitutions and one insertion, and a single *KRAS* mutation (9.1% = 1/11). The percentages of *EGFR* and *KRAS* mutations in the 11 never-smoking patients were not significantly different from those in the remaining 86 patients (*EGFR*: 38.3% = 33/86, and *KRAS*: 10.5% = 9/86), suggesting that *EGFR* and *KRAS* mutations in the cohort are not associated with tobacco use. Contrary to previous reports describing the association between *EGFR* mutations and non-smoking history (12), recent papers have reported no association with smoking or non-smoking history in Asian patients (14-16).

Gender also has been reported to be unconnected with *EGFR* mutations in Asian patients (14,15). We also found no association between gender and any type of *EGFR* mutations in the cohort. However, focusing on the 746ELREA deletions, 6 out of 7 patients with this type of deletion were male patients. Moreover, we investigated the distribution of mutations in the four genes across different clinicopathological subgroups based on tumor size, clinical stage, and histological subtypes including acinar, papillary, and bronchioloalveolar carcinoma patterns. However, no significant association with clinicopathological subgroups was found (data not shown), which is consistent with a previous report (17).

One interesting topic in analyzing genetic somatic mutations is the co-occurrence of mutations in two

genes. Co-occurrence of *EGFR* and *KRAS* mutations, which lie upstream and downstream of the RAS pathway respectively, has been reported to be mutually exclusive (5). In the review paper that analyzed the COSMIC database (5), the co-occurrence of *TP16* and *KRAS* mutations was reported in pancreatic cancer, lung large cell cancer, colon cancer, kidney tumors, and lung adenocarcinoma. However, in the present study, co-occurrence of *TP16* and *KRAS* mutations was not observed because of the lower mutation rate of *TP16* in the Kujukuri cohort. On the other hand, as to *LKB1* we co-sequenced *LKB1* and *KRAS* in the 97 lung adenocarcinomas tissue samples, and found four co-occurring mutations in these genes. So, mutations in these genes may not be mutually exclusive and therefore these genes may not be located on the same pathway. Co-occurrence of *LKB1* and *EGFR* mutations was also observed in three samples. Taken together, *LKB1* might act separately from the RAS pathway to suppress tumorigenesis. According to the known pathways summarized from the Kyoto Encyclopedia of Genes and Genomes (KEGG) (Figure 4) related to the *LKB1*, *TP16*, *EGFR*, and *KRAS* genes pathways, *LKB1* attenuates protein synthesis by inhibiting the mTOR pathway (18), that is separated from the RAS pathway.

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Inhibition of morphine tolerance is mediated by painful stimuli *via* central mechanisms

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ABSTRACT: Tolerance to morphine analgesia following repeated administration disturbs the continuation of opioid therapy for severe pain. Emerging evidence suggests that the development of morphine tolerance may be antagonized by painful stimuli. To clarify the detailed mechanisms of these phenomena, we examined the effects of several pain stimuli on morphine-induced tolerance. Subcutaneous (*s.c.*) injection of morphine (10 mg/kg) produced an analgesic effect, which was evaluated by tail-pinch test. Morphine-induced analgesia was diminished by repeated administration of morphine (10 mg/kg, *s.c.*) once a day for 5 days, demonstrating the development of tolerance. Morphine analgesic tolerance was suppressed by nerve injury-induced neuropathic pain and formalin- or carrageenan-induced inflammatory pain. Tolerance to serum corticosterone elevation by morphine (10 mg/kg), which was evaluated by fluorometric assay, was also suppressed by formalin-induced inflammatory pain. Moreover, morphine analgesia induced by intracerebroventricular (10 nmol) or intrathecal (5 nmol) injection was diminished by repeated administration of morphine *s.c.*, and this was also suppressed by carrageenan-induced inflammatory pain. These results suggest that morphine tolerance is inhibited by several pain stimuli, including neuropathic and inflammatory pain, through central mechanisms.

Keywords: Opioid, analgesia, neuropathic pain, inflammatory pain

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1. Introduction

Opioid ligands including morphine are potent analgesics that are widely used for the treatment of severe pain, *e.g.*, cancer progression (1,2). In addition to other serious adverse effects such as respiratory depression and constipation, tolerance and dependence may also disturb the continuation of opioid therapy (3). Therefore, an understanding of the molecular mechanisms underlying opioid tolerance and dependence is required. Several hypotheses concerning opioid tolerance have been published. For instance, alterations of neural networks and glial activities after chronic exposure to opioids were revealed (4-6). These phenomena might be responsible for the opioid tolerance and dependence.

In contrast, emerging evidence suggests that the development of opioid tolerance may be antagonized by painful stimuli (7,8). Experimental inflammatory pain or surgical pain suppressed the morphine tolerance (9,10). Despite several lines of evidence, the effects of different types of pain on opioid tolerance have not been yet clarified. Moreover, detailed mechanisms underlying the inhibition of opioid tolerance by painful stimuli are unclear. Therefore, we examined the effects of several pain stimuli (*i.e.*, neuropathic and inflammatory pain) on morphine-induced tolerance, and investigated whether the central nervous system (CNS) is a key region relating to these phenomena.

2. Materials and Methods

2.1. Animals

Male ICR mice (15-25 g) were obtained from Japan SLC, Inc. (Osaka, Japan), and they were housed in an air-conditioned (23-24°C, 60-70% relative humidity) vivarium with a 12 h dark/light cycle (light on from 8:00 to 20:00). Water and food were available *ad libitum*. All experimental procedures were approved by the Animal Research Committee of Wakayama Medical University and complied with the ethical guidelines of the International Association for the Study of Pain (11).

2.2. Morphine administration

Morphine hydrochloride (Takeda, Osaka, Japan) was dissolved in physiological saline, and administered to mice by subcutaneous (*s.c.*, 0.1 mL/10 g), intracerebroventricular (*i.c.v.*, 5 μ L), or intrathecal (*i.t.*, 5 μ L) injection. The *i.c.v.* injections were administered into the left lateral ventricle, and the *i.t.* injections were administered into the region between spinal L5 and L6 segments as described in a previous report (12,13). For the *i.c.v.* and *i.t.* injections, Hamilton microsyringes fitted with 32-gauge or 26-gauge needles were used, respectively.

2.3. Tail-pinch test

The analgesic effects of morphine were evaluated by the tail-pinch test as described in our previous report (14). Briefly, the mouse tail root was pressed with a 6-mm-wide flattened clip, which was adjusted to about 500 g pressure. Nociceptive responses were indicated by the latency required for response to the pressure by biting the clip. To avoid tissue damage, a cut-off time of 15 sec was set. The clip was applied every 15 min following morphine administration over a 120 min period. The percentage of maximum possible effect (%MPE) was calculated using the following formula: %MPE = 100 \times (measured latency – baseline latency)/(15 – baseline latency). The magnitude of analgesia was evaluated by the area under the curve (AUC) calculated from the time course of %MPE.

2.4. Estimation of serum corticosterone (SCS) level

Because the level of SCS due to circadian rhythm was most stable before noon each day, the blood was collected for SCS determination at 10:00-11:30. Mice were killed by decapitation and trunk blood was collected 1 h after morphine injection. The serum was separated by centrifugation and SCS level was measured by fluorometric assay according to the method of Zenkar and Bernstein (15).

2.5. Pain models

Three pain models were used in this study. For the neuropathic pain model, mice were anesthetized with sodium pentobarbital (70 mg/kg, *i.p.*) and the sciatic nerves of both hind limbs were exposed through a small incision. Then, 1/3 of the nerve thickness was tightly ligated with a silk suture (16,17). As a sham control, the incision site was closed without ligation. To induce acute inflammatory pain, mice were given 2% formalin or 1% λ -carrageenan into the intraplantar (*i.pl.*, 20 μ L) surface of both hind paws (18,19). Formalin and carrageenan were dissolved in phosphate buffered saline (PBS, vehicle control). The *i.pl.* injection was performed using a Hamilton microsyringe fitted with a 30-gauge needle.

2.6. Statistical analysis

Data are presented as the mean \pm S.E.M. Statistical analysis was performed using a one-way analysis of variance (ANOVA) followed by Tukey multiple comparisons test, or two-way ANOVA followed by Bonferroni multiple comparisons test. Significance was established at $p < 0.05$.

3. Results

3.1. Evaluation of morphine analgesia and its tolerance

Subcutaneous injection of morphine (10 mg/kg) prolonged the withdrawal latency at 15 min after administration. The prolonged latency lasted for 75 min and was restored within 120 min, indicating the analgesic effect of morphine. Saline injected as a control had no effect on the pain threshold (Figure 1A). By repeated administration of morphine (10 mg/kg) once a day for 5 days, morphine-induced analgesia was gradually diminished, demonstrating the development of analgesic tolerance as shown in AUC (Figure 1B).

3.2. Inhibition of morphine analgesic tolerance by nerve ligation-induced neuropathic pain

To examine whether an analgesic tolerance to morphine is affected by neuropathic pain, mice were given partial sciatic nerve ligation (PSL) 1 day before the first morphine administration. Morphine (10 mg/kg) was administered once a day for 5 days, and morphine-induced analgesia was evaluated on days 1, 3 and 5. On day 1, the time-course of morphine analgesia in PSL-operated mice was similar to that in sham-operated mice (Figure 2A), indicating that morphine analgesia was not affected by neuropathic pain. In contrast, on day 5, the analgesic

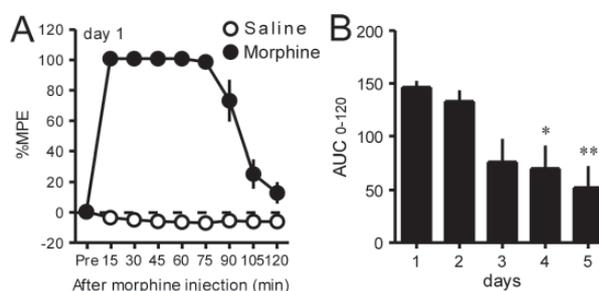


Figure 1. Evaluation of morphine analgesia and tolerance. (A) Time course of morphine analgesia on day 1. Morphine (10 mg/kg) or saline was subcutaneously (*s.c.*) administered, and the analgesic effects during a 120 min period after injection were evaluated by tail-pinch test. Analgesic effects are shown as time course of %MPE. (B) The development of tolerance to morphine analgesia, estimated by AUC. Morphine (10 mg/kg, *s.c.*) was repeatedly administered once a day for 5 days, and analgesic effect on each day is shown as AUC calculated from time course of %MPE. Data are presented as the mean \pm S.E.M. of 6-8 mice. %MPE: the percentage of maximum possible effect; AUC: area under the curve. * $p < 0.05$, ** $p < 0.01$ vs. day 1.

effect in PSL-operated mice was greater than that in sham-operated mice (Figures 2B and 2C), indicating that the development of tolerance to morphine analgesia was significantly suppressed by PSL-induced neuropathic pain.

3.3. Inhibition of morphine analgesic tolerance by formalin- or carrageenan-induced inflammatory pain

We examined the effects of inflammatory pain on the development of analgesic tolerance to morphine. First, 2% formalin was *i.pl.* injected into mice 2 h before

the first morphine administration. Repeated morphine administration and the evaluation of morphine tolerance were performed as shown in Figure 2. On day 1, there was no difference between morphine (10 mg/kg)-induced analgesia in formalin-treated mice and that in PBS-treated control mice, indicating that morphine analgesia was not affected by inflammatory pain (Figure 3A). In contrast, on day 5, the analgesic effect in FOR-treated mice was greater than that in PBS-treated mice (Figure 3B). As shown in AUC, the development of morphine tolerance was significantly suppressed by formalin-induced

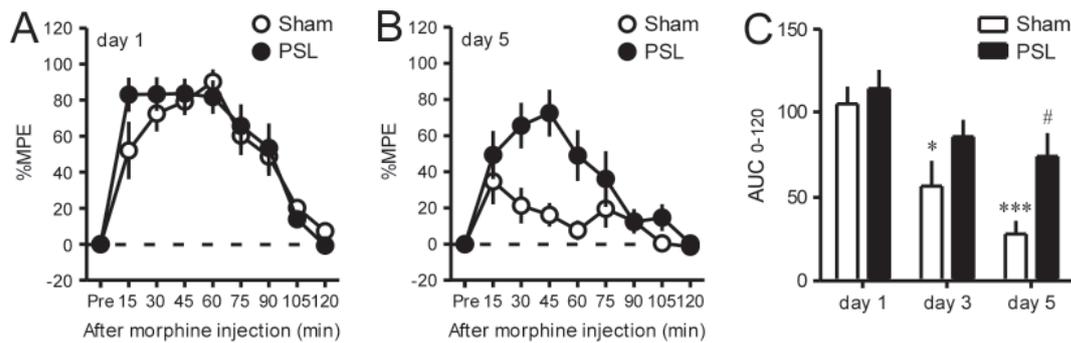


Figure 2. Inhibition of morphine analgesic tolerance by nerve ligation-induced neuropathic pain. Partial sciatic nerve ligation (PSL) was performed 1 day before morphine administration. As sham control, nerve was exposed. Morphine (10 mg/kg, *s.c.*) was administered once a day for 5 days, and morphine analgesia was evaluated for 120 min on days 1, 3, and 5. Analgesic effects of morphine on days 1 (A) and 5 (B) are shown as time course of %MPE and AUC (C). Data are presented as the mean \pm S.E.M. of 8-9 mice. %MPE: the percentage of maximum possible effect; AUC: area under the curve. * $p < 0.05$, *** $p < 0.001$ vs. day 1. # $p < 0.05$ vs. sham.

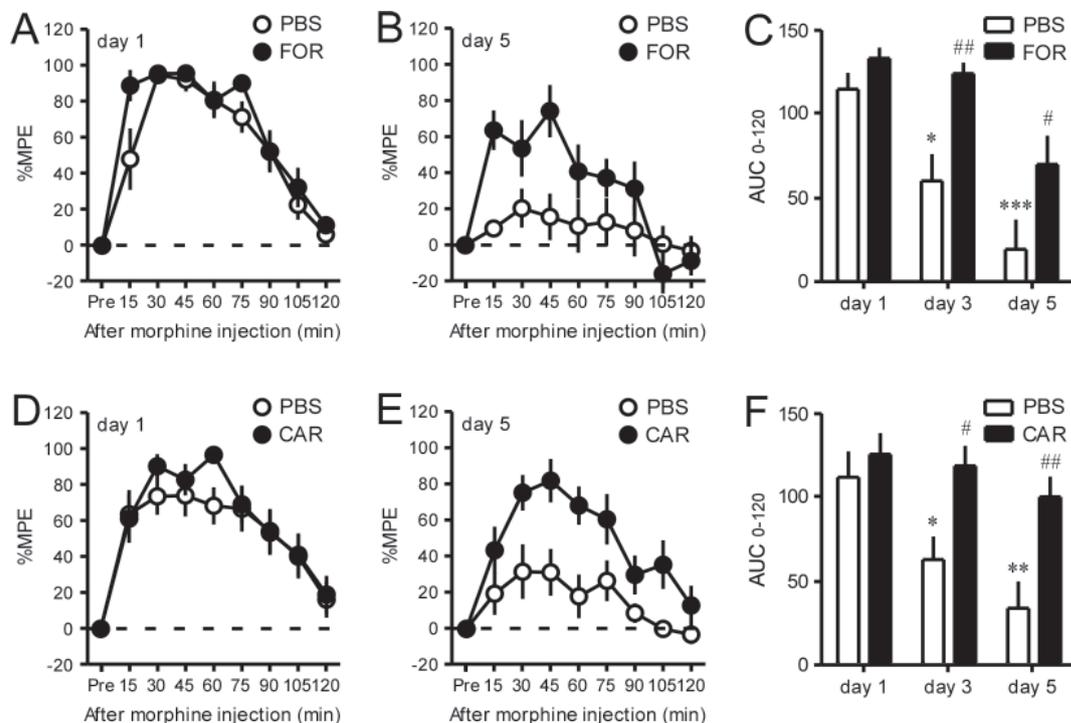


Figure 3. Inhibition of morphine analgesic tolerance by formalin- or carrageenan-induced inflammatory pain. 2% formalin (FOR; A-C) or 1% carrageenan (CAR; D-F) was *i.pl.* injected 2 h or 3 h before morphine administration on day 1, respectively. As vehicle control, PBS was injected. Morphine (10 mg/kg, *s.c.*) was administered once a day for 5 days, and morphine-induced analgesia was evaluated for 120 min on days 1, 3, and 5. Analgesic effects of morphine in FOR- or CAR-treated group on days 1 (A, D) and 5 (B, E) are shown as time course of %MPE (A, B, D, E), and AUC (C, F). Data are presented as the mean \pm S.E.M. of 8-9 mice. %MPE: the percentage of maximum possible effect; AUC: area under the curve. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ vs. day 1. # $p < 0.05$, ## $p < 0.01$ vs. PBS.

inflammatory pain on days 3 and 5 (Figure 3C). Second, as another inflammatory pain model, 1% carrageenan was *i.pl.* injected into mice 3 h before the first morphine administration. On day 1, there was no difference between morphine (10 mg/kg)-induced analgesia in carrageenan-treated mice and that in PBS-treated control mice, suggesting that carrageenan injection also had no effect on morphine-induced analgesia on day 1 (Figure 3D). On day 3 and 5, the analgesic effect in carrageenan-treated mice was greater than that in PBS-treated mice (Figures 3E and 3F).

3.4. Effects of formalin-induced inflammatory pain on tolerance to SCS elevation by morphine

Morphine produces not only analgesia but also SCS elevation. To confirm the inhibition of morphine analgesic tolerance by inflammatory pain, we examined the effect of formalin-induced inflammatory pain on tolerance to SCS elevation by repeated morphine administration. Morphine (10 mg/kg) was administered once a day for 5 days, and the SCS level was evaluated 1 h after morphine injection (10 mg/kg) on day 6. Basal SCS level was 6.8 ± 1.3 $\mu\text{g/dL}$ ($n = 5$) in naive mice. After repeated saline injection for 5 days, SCS was markedly elevated by morphine. The morphine-induced SCS elevation was significantly suppressed after repeated morphine administration for 5 days, while tolerance was developed following *i.pl.* administration of PBS. The repeated morphine-induced suppression of SCS elevation was reversed by formalin *i.pl.*, indicating the inhibition of tolerance by formalin-induced inflammatory pain (Figure 4).

3.5. Inhibition of tolerance to intracerebroventricular or intrathecal morphine analgesia by carrageenan-induced inflammatory pain

Generally, it is believed that morphine develops analgesic

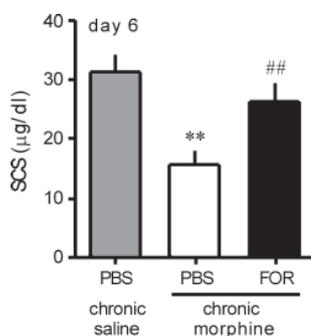


Figure 4. Inhibitory effect of intraplantar formalin on tolerance to morphine-induced corticosterone elevation. 2% formalin (FOR) was *i.pl.* injected 2 h before morphine administration on day 1. As vehicle control, PBS was injected. Morphine (10 mg/kg, *s.c.*) or saline was administered once a day for 5 days, and SCS elevation at 1 h after morphine (10 mg/kg, *s.c.*) administration was evaluated on day 6. Data are presented as the mean \pm S.E.M. of 5-6 mice. ** $p < 0.01$ vs. chronic saline. ## $p < 0.01$ vs. PBS.

tolerance through a central mechanism, although morphine acts on both the peripheral and central nervous systems. To investigate the key region, *i.e.*, supraspinal or spinal sites, of the inhibitory effects of painful stimuli on tolerance to morphine analgesia, we examined the effects of carrageenan-induced inflammatory pain on tolerance to *i.c.v.* or *i.t.* morphine. The analgesic tolerance to morphine was developed by the administration of morphine (10 mg/kg, *s.c.*) once a day for 3 days (on days 2, 3, and 4), and morphine *i.c.v.* or *i.t.*-induced analgesia was evaluated on days 1 and 5. On day 1, there was no effect of carrageenan *i.pl.* injection on the time course of morphine *i.c.v.* (10 nmol)-induced analgesia (Figure 5A). In PBS *i.pl.*-treated mice, the analgesic effect of *i.c.v.* morphine on day 5 was significantly diminished in comparison with that on day 1 by repeated morphine *s.c.* administration, which indicated the development of analgesic tolerance to *i.c.v.* morphine. However, on day 5, the analgesic effect of morphine *i.c.v.* in carrageenan *i.pl.*-treated mice was greater than that in PBS-treated mice (Figures 5B and 5C). On day 1, there was no effect of carrageenan *i.pl.* injection on morphine *i.t.* (5 nmol)-induced analgesia (Figure 5D). However, on day 5, the analgesic effect of morphine *i.t.* in carrageenan *i.pl.*-treated mice was greater than that in PBS-treated mice (Figures 5E and 5F). These results suggest that the development of tolerance to *i.c.v.* or *i.t.* morphine analgesia was significantly inhibited by carrageenan-induced inflammatory pain on day 5.

4. Discussion

It is generally considered that opioid tolerance may be antagonized under painful conditions in clinic (1,8). Here, we demonstrated that morphine-induced analgesia was gradually decreased by repeated morphine administration in mice. This indicated the development of tolerance to morphine analgesia, which is consistent with previous reports (20,21). Therefore, we examined the effect of neuropathic and inflammatory pain on the development of tolerance to morphine analgesia in mice. PSL-induced neuropathic pain (*i.e.*, allodynia and thermal hyperalgesia) is usually observed on the next day of PSL, and lasts for at least 14 days (22). After *i.pl.* injection of formalin or carrageenan, hyperalgesia and edema were observed, which are typical prognostics of inflammation, and these phenomena lasted for 7-14 days (18,19). On day 1, morphine analgesia evaluated by the time-course and AUC was not affected by both PSL-induced neuropathic pain and formalin- or carrageenan-induced inflammatory pain. We found that tolerance to morphine analgesia was inhibited by not only PSL-induced neuropathic pain, but also formalin- or carrageenan-induced inflammatory pain.

To confirm the inhibitory effects of pain stimuli on morphine tolerance, we focused on SCS elevation, which is one of characteristic effects of morphine. Normally, acute administration of morphine increases

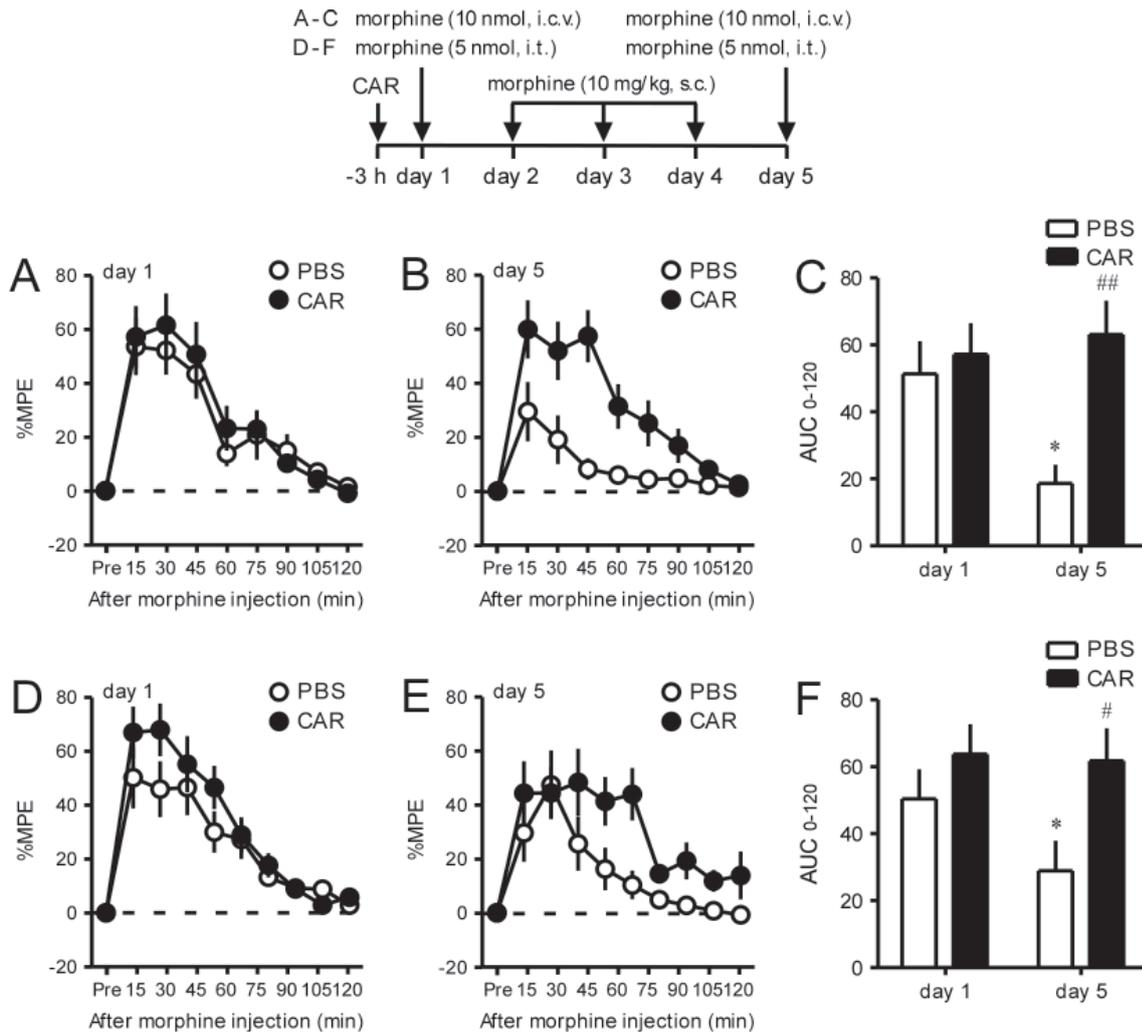


Figure 5. Inhibition of intracerebroventricular or intrathecal morphine tolerance by carrageenan-induced inflammatory pain. 1% carrageenan (CAR) was *i.pl.* injected 3 h before morphine administration on day 1. As vehicle control, PBS was injected. Morphine was administered *i.c.v.* (10 nmol, A-C) or *i.t.* (5 nmol, D-F) on days 1 and 5. For the development of tolerance, morphine (10 mg/kg) was *s.c.* administered on days 2, 3, and 4. Morphine-induced analgesia was evaluated for 120 min on days 1 and 5. Experimental schedules are indicated above. Analgesic effects of *i.c.v.* or *i.t.* morphine on days 1 (A, D) and 5 (B, E) are shown as time course of %MPE, respectively, and AUC (C, F) was calculated from time course of %MPE. Data are presented as the mean \pm S.E.M. of 11-13 mice. *i.c.v.*: intracerebroventricular; *i.t.*: intrathecal; %MPE: the percentage of maximum possible effect; AUC: area under the curve. * $p < 0.05$ vs. day 1. # $p < 0.05$, ## $p < 0.01$ vs. PBS.

the SCS level in a dose-dependent manner. This effect is also diminished by repeated administration of morphine, indicating the development of tolerance (23). The tolerance to SCS elevation was also suppressed by formalin-induced inflammatory pain, strongly suggesting that painful stimuli might affect the mechanism of tolerance through not only an analgesic effect, but also an endocrine effect. It is well known that morphine acts on the hypothalamus to release corticotropin-releasing factor, which then releases adrenocorticotrophic hormone (ACTH) from the pituitary. The released ACTH works on the adrenal cortex and results in the elevation of SCS (24). These observations indicate that tolerance to SCS elevation by morphine may be mediated by a supraspinal site, and are consistent with analgesic tolerance.

Several lines of evidence indicate the potential molecular mechanisms of opioid tolerance. It is well

known that opioid receptors are 7-transmembrane G protein-coupled receptors that are classified in three types, μ , δ , and κ (25). All of the opioid receptors are associated with an intracellular G α i protein, and are thought to produce analgesic effects through the activation of G α i-related pathways (26). Notably, the μ -opioid receptor largely contributes to opioid analgesia, and desensitization of the μ -opioid receptor may be responsible for opioid tolerance (27). According to the cAMP hypothesis reported previously (28), opioid tolerance develops due to receptor internalization and subsequent desensitization at the single cell level (29). The relative activity versus endocytosis hypothesis was also suggested as a mechanism to explain the receptor internalization-dependent desensitization (3,28). Furthermore, the relationship between the activation of protein kinase C (PKC) and receptor desensitization was reported. Indeed, blockade of the

PKC pathway inhibited the development of morphine tolerance (30). Recently, growing evidence suggests that opioid tolerance is mediated by anti-opioidergic neural networks in the CNS, *i.e.*, the midbrain (6). Namely, anti-opioidergic neural networks in the CNS were enhanced by chronic morphine treatment, and cholecystokinin, neuropeptide, and glutamate, which are candidates of typical anti-opioid mediators, prevented the opioid effects (31-33). For example, concomitant treatment with MK801 (NMDA glutamatergic receptor antagonist) suppressed the development of tolerance to morphine analgesia. Furthermore, chronic morphine exposure activates microglia and astrocytes in the CNS (5). These cells express μ -opioid receptor and produce several inflammatory mediators following morphine treatment (34). For instance, the critical roles of interleukin (IL)-1 β and IL-6 in morphine tolerance were identified (21,35). Taken together, anti-opioidergic neural networks and inflammation in the CNS largely participate in the opioid tolerance.

It was reported that analgesic effects were produced through not only the CNS but also the peripheral nervous system, *e.g.* primary afferents (36). To clarify a key region underlying inhibition of morphine tolerance by painful stimuli, we evaluated the effect of carrageenan-induced inflammatory pain on the analgesia and its tolerance by *i.c.v.*- or *i.t.*- administered morphine. In these experiments, analgesic tests after morphine *i.c.v.* or *i.t.* administration were performed on days 1, and 5. To avoid excess tissue damage by the manipulation of *i.c.v.* and *i.t.*, morphine was systemically (*s.c.*) administered on days 2, 3, and 4 for the development of tolerance. Tolerances to both *i.c.v.*- and *i.t.*-administered morphine were inhibited by carrageenan-induced inflammatory pain. These results indicate that interaction between opioid actions and pain stimuli in the CNS (midbrain and spinal cord) is crucial for the inhibition of opioid tolerance mediated by painful stimuli.

Although we could not determine the exact mechanisms underlying the inhibition of morphine tolerance by painful stimuli in this study, a possible mechanism is suggested. Previous reports showed that the inhibition of morphine tolerance and dependence by painful stimuli was partially mediated by activation of the endogenous κ opioid system (10,37). Therefore, this mechanism our observations in the midbrain and spinal cord presented here. However, further research into the mechanisms is still required.

In conclusion, morphine analgesic tolerance is inhibited by several pain stimuli, including neuropathic and inflammatory pain, through central mechanisms. The mechanistic details remain to be established, but these findings suggest that the clinical use of opioids should not be limited by its adverse tolerance, and we hope that these findings may lead to the development of more efficacious therapeutics for patients suffering from severe pain.

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Comparison of antioxidant activity of compounds isolated from guava leaves and a stability study of the most active compound

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ABSTRACT: In the present study, quercetin (QT), morin (MR), and quercetin-3-O-glucopyranoside (QG) isolated from guava leaves were comparatively tested for antioxidant activity using DPPH, ABTS, and FRAP methods. QT was the most active among them. The free radical scavenging activity of QT was approximately four times higher than MR and two times higher than QG. The reducing power of QT was eight times higher than MR and two times higher than QG. A mixture of QT with MR or QG showed interesting combination effect. The synergistic antioxidant activity was obtained when QT was mixed with MR whereas the antagonistic effect was found when mixed with QG. The stability study of QT in liquid preparations indicated that the decomposition reaction rate of QT could be explained by a kinetic model assuming a first-order chemical reaction. The aqueous solution of QT was rapidly decomposed with $t_{1/2}$ of approximately five days whereas QT entrapped in chitosan nanoparticles was five times longer. It was concluded that QT was the most active antioxidant from guava leaves. Entrapment of QT in chitosan nanoparticles could significantly enhance its stability.

Keywords: Guava, quercetin, antioxidant, combination effect, stability, chitosan nanoparticles

1. Introduction

Guava (*Psidium guajava* L.), a plant of the Myrtaceae family, is found widespread in hot climate countries including Thailand. Traditional healers of different countries use its leaves to treat diarrhea, one of the leading causes of mortality and morbidity especially in developing countries (1). The scientific evaluation of guava with respect to diarrhea has been well studied (2-5). The extract of guava leaves has been reported

to have narcotic-like activity (6), antinociceptive (7), antimutagenic (8), anticough (9), and central nervous system (CNS) depressant activity (10). Our previous study on the screening for biological actions of several plants commonly grown in Thailand reported that fruits and leaves of many edible plants possess high antioxidant activity (11). Among these, guava leaf extract has shown the highest antioxidant capacity (12). It was reported that the antioxidant activity of guava leaves is higher than its fruit (13). A phytochemical study of guava leaves has demonstrated many compounds including lipids, carbohydrates, proteins, vitamins, essential oils, tannins, saponins, flavonoids, sterols and triterpenes (14,15). Those polyphenolic compounds are believed to be the major principles that provide the plant with high antioxidant properties (16-19). However, a literature survey revealed that no data has been made to compare the potency of the active compounds as well as the interaction effects among them. The present study aimed to compare the antioxidant capacity of the active compounds isolated from guava leaves. Moreover, the interaction of the most active compound in combination with others as well as its stability was also our interest.

2. Materials and Methods

2.1. Plant materials

Mature leaves of guava (*P. guajava*) were collected during January-February from the botanical garden of Chiang Mai University, Thailand. The leaves were washed with clean water then dried in a hot air oven at 50°C for 48 h. After drying, they were pulverized into fine powder. The guava leaf powder obtained which passed a No. 60 sieve was collected for further study.

2.2. Chemicals

Potassium persulfate, ferrous sulfate ($\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$), ferric chloride ($\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$), 2-diphenyl-1-picrylhydrazyl (DPPH), and 2,2'-azinobis-(3-ethylbenzothiazoline-6-sulfonic acid) diammonium salt (ABTS) were purchased from Sigma-Aldrich (St. Louis, MO, USA). Trolox was from Aldrich Chemical Company (Steinheim, Germany) and 2,4,6-tri (2-pyridyl)-S-triazine (TPTZ) was

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from Fluka Chemicals (Buchs, Switzerland). Column chromatography were carried out on Cosmosil 75 μm C₁₈-OPN (Nacalai Tesque, Kyoto, Japan), Sephadex LH-20 (GE Healthcare Biosciences AB, Uppsala, Sweden), Silica gel 60 (230-400 mesh) (Merck, Darmstadt, Germany), Toyopearl HW-40C (TOSOH, Tokyo, Japan) and MCI-gel (Mitsubishi chemical corporation, Tokyo, Japan). Pre-coated RP-18 F₂₅₄ (0.25 mm) and organic solvents (AR grade) were from Merck (Darmstadt, Germany). Water was deionized and purified using a Milli-Q system (Millipore, Bedford, MA, USA). Sodium alginate was from Sigma-Aldrich (St. Louis, MO, USA), chitosan (95% DD, MW 22 kDa) was a gift from Seafresh Industrial PCL, Thailand. Other chemicals were of the highest available grade and used as obtained.

2.3. Compound extraction

Three active compounds, quercetin (QT), morin (MR), and quercetin-3-*O*-glucopyranoside (QG), were isolated according to the description in our previous report (20). Briefly, the methanol (MeOH) extract from the dried powder of guava leaves was subjected to column chromatography (Cosmosil C18-OPN, 75 μm , 6.0 cm (i.d.) \times 40 cm) and eluted stepwise with MeOH-aqueous mixtures of different polarity (5 to 100% aqueous MeOH). Fractions were collected and subjected to chemical analysis using thin-layer chromatography (TLC) and antioxidant activity was evaluated by the DPPH method. The most active fractions were further isolated by column chromatography using Sephadex LH-20 and silica gel 60 columns to obtain QT, Toyopearl HW-40C and Cosmosil 75 μm C18-OPN columns to obtain MR, and Cosmosil 75 μm C18-OPN and MCI-gel columns to yield QG.

2.4. DPPH assay

The DPPH assay is considered to be a good *in vitro* model widely used to assess free radical scavenging efficacy in a relatively short time. In the present study, this method was used to screen the antioxidant power of the extracted fractions and the isolated bioactive pure compounds of guava leaves. In its free radical form, DPPH has an absorbance at 515 nm which disappears upon reduction by an antioxidant compound to become a stable diamagnetic molecule with a result of a color change from purple to yellow (21). The test could be taken as an indication of the hydrogen donating ability of the tested compounds. The assay was done using the method described previously by Gamez *et al.* (22) with some modification. Briefly, a sample in MeOH (100 μL) was added to a solution of 200 μM DPPH radicals in 100 μL . The reaction mixture was left to stand for 30 min at room temperature in the dark. The scavenging activity of samples was estimated by measuring the absorption of the mixture at 515 nm, which reflected to

the amount of DPPH radicals remaining in the solution. The scavenging activity was expressed as the IC₅₀, the concentration of samples required for scavenging 50% of DPPH radicals in the solution.

2.5. ABTS assay

The ABTS assay is an excellent method used for determining the antioxidant activity of a broad diversity of substances, such as hydrogen-donating antioxidants or scavengers of aqueous phase radicals and of chain-breaking antioxidants or scavengers of lipid peroxy radicals (23). In the present study, it was used for confirmation on the free radical scavenging potential of the pure and combined compounds isolated from guava leaves. The assay was done in the same manner as previously described by Okonogi *et al.* (11). Briefly, the pre-formed radical monocation of ABTS was generated by reacting ABTS solution (7 mM) with 2.45 mM potassium persulfate (K₂S₂O₈). The mixture was allowed to stand for 15 h in the dark at room temperature. The solution was diluted with ethanol to obtain an absorbance of 0.7 ± 0.2 units at 750 nm. The test samples were separately dissolved in ethanol to yield an appropriate concentration. An aliquot of 20 μL of an ethanolic test solution of each sample was added to 180 μL of ABTS free radical cation solution. The absorbance, monitored for 5 min, was measured spectrophotometrically at 750 nm using a microtitre plate reader. The free radical scavenging activity of each sample was expressed as trolox equivalent antioxidant capacity (TEAC), which was obtained by comparing the absorbance change at 750 nm in a reaction mixture containing a test material with that containing trolox. This index is defined as the millimolar concentration of a trolox solution whose antioxidant capacity is equivalent to 1.0 mg of the test sample (24).

2.6. Ferric reducing ability of plasma (FRAP) method

This method was used to confirm antioxidant activity by mechanism of reducing action of the pure and combined compounds isolated from guava leaves. The reducing power of the test samples was determined by using a FRAP assay described by Benzie & Strain (25) with some modification. Briefly, the FRAP reagent containing 2.5 mL of 10 mM TPTZ solution in 40 mM HCl plus 2.5 mL of 20 mM FeCl₃ and 25 mL of 0.3 M acetate buffer, pH 3.6, was freshly prepared. The test samples were dissolved in ethanol to an optimal concentration. An aliquot of 20 μL test solution was mixed with 180 μL of FRAP reagent. The absorption of the reaction mixture was measured at 595 nm by a microtitre plate reader. Ethanolic solutions of known Fe (II) concentration, in the range of 50-500 μM (FeSO₄), were used as a calibration curve. The reducing power was expressed as equivalent concentration (EC). This parameter was defined as the concentration of the test

sample having a ferric reducing ability equivalent to that of 1 mM FeSO₄.

2.7. Interaction effect on antioxidant activity

The mixture of QT and MR or QT and QG at a weight ratio of 1:1 was subjected to antioxidant test using ABTS and FRAP assays. The experimental antioxidant activity (*E*) of the mixture expressed as TEAC or EC values respectively was determined as described in Sections 2.5 and 2.6, respectively. The theoretical antioxidant activity (*T*) is the sum of the experimental activity of each compound fraction, calculated using individual antioxidant activity in the following equation: $T = [(X_A \times E_A) + (X_B \times E_B)]$, where X_A and X_B represent the mole fraction of compound A and compound B, respectively, and E_A and E_B represent the antioxidant activity of 100% A and B, respectively (26).

2.8. Preparation and characterization of QT loaded chitosan nanoparticles

QT loaded chitosan nanoparticles were prepared according to the ionic gelation principle described previously by Sarmiento *et al.* (27) with some modification. Chitosan at 0.5 mg/mL in 1% (v/v) acetic aqueous solution was prepared. A portion of 1 mL of 0.1 mg/mL QT in water-propylene glycol mixture was mixed with 10 mL of 0.03% (w/v) sodium alginate solution. Under magnetic stirring at room temperature, 0.3 mL of 0.18 mM calcium chloride solution was added followed by 1 mL of chitosan solution. Opalescent colloidal dispersion was formed spontaneously. The hydrodynamic diameter and zeta potential of the nanoparticles were measured by dynamic light scattering (DLS) using a Malvern system (Zetasizer, version 5.00, Malvern Instruments Ltd., Malvern, UK) consisting of computerized auto-titrate and DLS software. Measurements were taken at an angle of 173°. For entrapment efficiency determination, the dispersion was subjected to ultra centrifugation at 20,000 g, 4°C for 30 min. The separated nanoparticles obtained were dissolved in DMSO prior to analysis by HPLC using a C-18 column (150 mm × 4.6 mm, 5 μm; Agilent, USA) with a mobile phase of acetonitrile/0.1% (w/v) phosphoric acid solution (30:70, v/v) and detected by UV detector at 373 nm.

2.9. Stability testing

The aqueous solution of QT (0.1 mg/mL) and colloidal dispersion of QT loaded nanoparticles of equivalent QT concentration were kept in light protected and non-protected containers for 4 weeks. The amount of QT remaining in each sample during the storage time was determined periodically by HPLC as described in Section 2.8.

3. Results and Discussion

3.1. Antioxidant activity of the isolated compounds

The antioxidant activity can be evaluated by different methods depending on the mechanism of antioxidant action. In this study, three compounds, *i.e.* QT, MR, and QG, structures as shown in Figure 1, were isolated from guava leaves and subsequently subjected to the comparative antioxidant test using DPPH and ABTS methods for the free radical scavenging mechanism and the FRAP method for reducing action. The results as shown in Table 1 indicated that QT was the most active antioxidant with an IC₅₀ value of 1.20 ± 0.02 μg/mL. The results from ABTS confirmed that QT possessed excellent free radical scavenging activity with a TEAC value of 57.54 ± 0.07 mM/mg. It was noticed from the IC₅₀ and the TEAC values that the free radical scavenging potential of QT was approximately four times higher than MR and two times higher than QG. The principle of the FRAP method is based on the reduction of a ferric-tripyridyltriazine complex to its ferrous colored form in the presence of antioxidants. The reducing power property indicates that the antioxidant compounds are electron donors and can reduce the oxidized intermediates of the lipid peroxidation process, so that they can act as primary and secondary antioxidants (28). The results indicated that QT, MR, and QG possessed reducing activity but not at the same level. These results clearly indicated

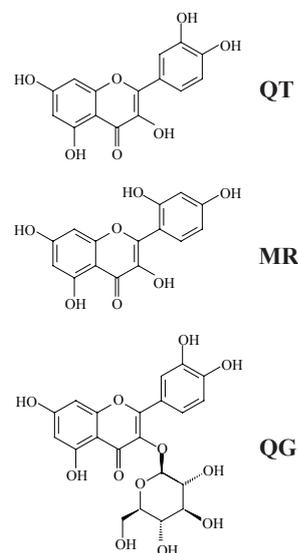


Figure 1. Chemical structure of quercetin (QT), morin (MR), and quercetin-3-O-glucopyranoside (QG).

Table 1. Antioxidant activity of the isolated compounds

Isolated compound	IC ₅₀ (μg/mL)	TEAC (mM/mg)	EC (mM/mg)
QT	1.20 ± 0.02	57.54 ± 0.07	72.69 ± 1.06
MR	5.41 ± 0.20	14.59 ± 0.62	8.56 ± 0.33
QG	3.58 ± 0.05	32.35 ± 0.12	42.52 ± 1.08

that QT was the most potent reducing substance with an EC value of 72.69 ± 1.06 mM/mg. This value was eight and two times higher than MR and QG, respectively. From this point of view, it was confirmed that QT was the most active antioxidant compound among them. According to its obviously high EC value, QT was considered to be a good electron donor and could terminate oxidation chain reactions by reducing the oxidized intermediates into a stable form.

3.2. Effect of compound combination on antioxidant power

As QT showed the strongest antioxidant activity among the three active compounds isolated from guava leaves, this compound was combined with each of the other two in order to examine interaction effects. The calculated theoretical antioxidant activities (T) of the mixtures were compared with the experimental activity (E). If E is greater than T ($E/T > 1$), then a positive interaction happened among the compounds, thus displaying synergism. On the other hand, if E is less than T ($E/T < 1$) a negative interaction happened, thus displaying antagonism. An additive effect was obtained when no reaction occurred and the ratio of E/T showed a value of 1. The results of QT in combination with MR or QG are shown in Table 2. It was found that the combination of QT with MR yielded a significant synergistic antioxidant effect, whereas the combination with QG gave an antagonistic antioxidant action. Synergistic effects of complex mixtures are thought to be important in plant defenses against a harmful environment. Plants usually present defenses as a set of compounds, not as individual ones, and it is thought that the minor constituents may act as synergists, enhancing the effect of the major constituents through a variety of mechanisms (29). In the present study, our results demonstrated that the antioxidant activity of QT as a major active component was synergized by a minor component, MR.

3.3. Stability study

This experiment was done in order to examine the effect of light and aging on the chemical stability of QT extracted from guava leaves. The results revealed that

Table 2. Antioxidant activity of the combined isolated compounds

Antioxidant activity	T	E	E/T	Interaction effect
TEAC (mM/mg)				
QT + MR	36.06	57.72	1.60	Synergism
QT + QG	44.95	35.91	0.79	Antagonism
EC (mM/mg)				
QT + MR	40.62	67.54	1.66	Synergism
QT + QG	60.79	40.84	0.67	Antagonism

Mixture of 1:1 weight ratio.

QT decreased rapidly during the first week of storage and further gradually decomposed according to the exponential degradation profiles demonstrated in Figure 2. A high reduction within one week of approximately 70% of QT was found in the samples without light protection as shown in Figure 2A and 30% in the light protection samples as demonstrated in Figure 2B. The result suggested that QT degradation was enhanced by photolysis. This result of light-induced decomposition of QT in our study was in agreement with the report of Smith *et al.* (30). The results provided valuable information about the degradation aspect of QT. This revealed the relevance to the real conditions of QT encountered in pharmaceutical or cosmetic products that is limited. To overcome this drawback, in the present investigation, QT was encapsulated in nanoparticles of chitosan. It was found that QT loaded chitosan nanoparticles were formed successfully by the gelation method. The hydrodynamic diameter and zeta potential of the nanoparticles obtained were found to be 310.1 ± 7.1 nm and -35.9 ± 0.5 mV, respectively. This confirmed that QT loaded particles obtained were in the nanometer size range. The resulting zeta potential value, from the viewpoint of the DLVO theory (31,32), indicated the high stability of the QT loaded nanoparticle colloidal system obtained. The entrapment efficiency was found to be $96.26 \pm 0.1\%$ indicating that QT was quantitatively encapsulated in the nanoparticles. The degradation profile of QT in the nanoparticles was demonstrated to be an exponential curve as shown in Figure 3. It was noted that the degradation of QT entrapped in the nanoparticles was also enhanced by light. However, the amount of QT entrapped in the nanoparticles when compared with that of non-entrapped samples, was found to be significantly higher at the same time of storage. This result indicated that the degradation of QT in the nanoparticles was obviously reduced.

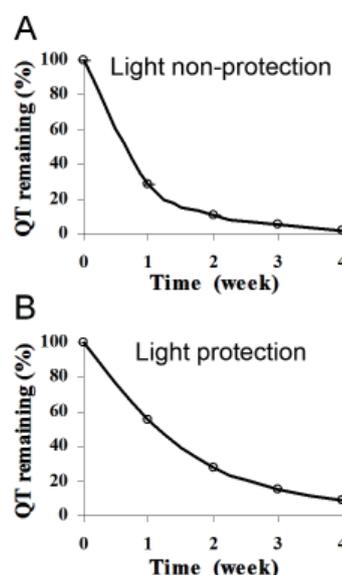


Figure 2. Effect of aging on QT solution kept in light non-protected (A) and light protected (B) containers.

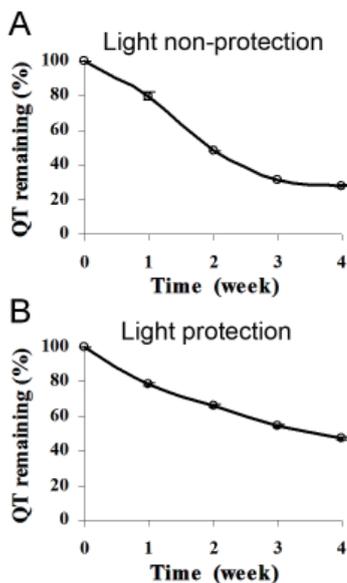


Figure 3. Effect of aging on QT loaded chitosan nanoparticles kept in light non-protected (A) and light protected (B) containers.

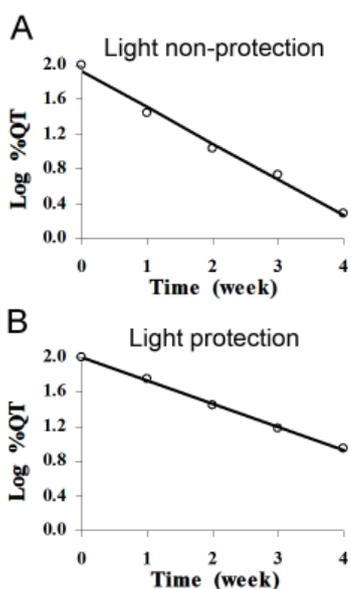


Figure 4. Plot of the logarithm concentration of QT remaining in light non-protected (A) and light protected (B) containers versus time of storage.

The relation of logarithm concentration of the remaining QT and the storage time showed a linear regression of $y = 1.93 - 0.42x$ ($r^2 = 0.991$) and $y = 1.99 - 0.27x$ ($r^2 = 0.998$) for the light non-protected and protected samples, respectively as shown in Figure 4. This result suggested that the decomposition of QT could be explained by a kinetic model assuming a first-order chemical reaction in the following equation: $dC/C = -kdt$ where C is the remaining amount of QT at time t , k is the degradation rate constant, and t is the time point of storage. Similarly with the non-entrapped QT, the linear relationship, of $y = 2.00 - 0.15x$ ($r^2 = 0.967$) and $y = 1.99 - 0.08x$ ($r^2 = 0.993$), was obtained from the light non-protected and protected

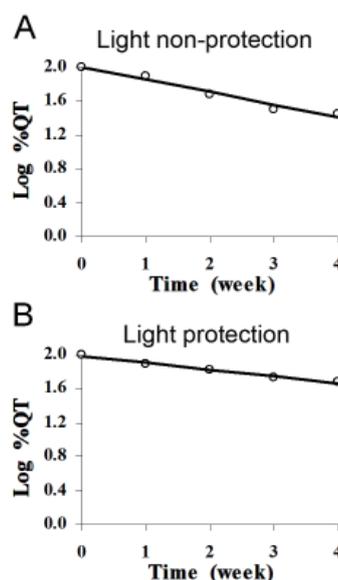


Figure 5. Plot of the logarithm concentration of QT loaded chitosan nanoparticles remaining in light non-protected (A) and light protected (B) containers versus time of storage.

Table 3. Experimental first order kinetic parameters and correlation coefficient of the degradation of QT in different samples

QT samples	r^2	$k_{(OBS)}$ (week ⁻¹)	$t_{1/2}$ (days)
NE-QT-L	0.991	0.961	5.049
NE-QT-NL	0.998	0.612	9.704
NN-QT-L	0.967	0.347	13.966
NN-QT-NL	0.993	0.187	25.995

NE-QT-L: non-entrapped QT solution in light non-protected container; NE-QT-NL: non-entrapped QT solution in light protected container; NN-QT-L: QT entrapped in nanoparticles dispersion in light non-protected container; NN-QT-NL: QT entrapped in nanoparticles dispersion in light protected container.

nanoparticles, respectively as shown in Figure 5. This indicated that the degradation kinetics of QT entrapped in the nanoparticles also followed a first-order kinetic reaction. The degradation rate of QT from each sample could be simply observed from the slope of the straight lines, the higher the slope the higher the decomposition rate. The experimental kinetic parameters of QT degradation are provided in Table 3. The observed first order kinetic rate constant, $k_{(OBS)}$, was obtained from the slope value multiplied by 2.303. The half-life, $t_{1/2}$, of QT from each sample was calculated from the $k_{(OBS)}$. It was found that QT entrapped in chitosan nanoparticles and protected from light was the most stable. The $t_{1/2}$ of QT from this sample was approximately five times longer than that of the non-entrapped sample kept in the light non-protected container.

4. Conclusion

QT, MR, and QG were isolated from guava leaves. Among these three compounds, QT was found to have the most active antioxidant activity through free radical scavenging and reducing mechanisms. The antioxidant activity of QT was synergized by MR and antagonized by QG. The

stability study of QT in liquid preparations revealed that QT was unstable. The degradation of QT was enhanced by light and through a first-order kinetic reaction. Entrapment of QT in chitosan nanoparticles could significantly improve the stability of QT.

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was used because an accurate concentration of the DCCA solution is easy to prepare due to the fact that it is a solid compound. As shown in Figure 1, DCCA can be released as OCl^- in aqueous solution (21). Until now, DCCA has been used for the production of OCl^- , which has in turn been utilized for chemiluminescent detection of sulfide ion in spring water (21). To evaluate the scavenging effect of ROS, aminophenyl fluorescein (APF) (22) was employed in this study. It has been reported that APF can emit fluorescence originating from fluorescein in the presence of OCl^- (22).

Taking these reagents together, we constructed a flow-injection analysis (FIA) system (Figure 2) in order to develop a reliable, simple, and rapid whole assay operation method. APF was mixed manually with each test compound and the mixed solution was injected into the FIA system. The injected solution was reacted in-line with OCl^- produced from DCCA. The fluorescence intensity of fluorescein generated from non-fluorescent APF would be significantly attenuated if the test compound had a scavenging effect on the OCl^- . Thus, the change of the fluorescence intensity can be used to estimate the anti-oxidant activity of the test compounds. In the present study, several compounds known as anti-oxidants were assayed by the proposed FIA system and the feasibility of the FIA system was verified.

2. Materials and Methods

2.1. Materials

Aminophenyl fluorescein (APF) was purchased from Sekisui Medical Co., Ltd. (Tokyo, Japan). Dichloroisocyanuric acid, sodium salt (DCCA) was purchased from Sigma-Aldrich (St. Louis, MO, USA). *N,N*-Dimethylformamide (DMF) was purchased from Nacalai Tesque, Inc. (Kyoto, Japan). Sodium hypochlorite containing 6% active chlorine was

purchased from Koso Chemical (Tokyo, Japan). Water used throughout this investigation was processed by a Milli-Q Water Purification System (Millipore Corporation, Milford, MA, USA).

2.2. Sample preparations

A commercially available 5.0 mM APF solution (in DMF) was diluted with additional DMF to create a 10 μM APF stock solution. The APF stock solution was stable for several months in a refrigerator. Each sample solution was prepared by mixing a test compound in 500 μL of 0.1 M sodium acetate (in H_2O) with 500 μL of 10 μM APF. The blank solution was a mixture of 500 μL of 0.1 M sodium acetate and the same volume of 10 μM APF. Twenty microliters of either a sample or the blank solution were injected manually into the FIA system through a Rheodyne 7125 injector equipped with a 20 μL sample loop.

2.3. FIA system

A schematic diagram of the FIA system constructed in this experiment is shown in Figure 2. The two elution pumps were Shimadzu LC-10AS (Shimadzu Corp., Kyoto, Japan). All elution tubing and connectors used were of PEEK™ materials. Fluorescence intensity was monitored using a Shimadzu RF-535 fluorescence detector and the excitation and emission wavelengths were set at a constant 490 and 515 nm, respectively. The mobile phase solutions A and B were 0.1 M sodium acetate and 50 μM DCCA in H_2O , respectively. The flow rate for both solutions was 2.0 mL/min.

2.4. Preparation of test compounds

The test compounds dithiothreitol, reduced glutathione, and 3-methyl-1-phenyl-5-pyrazolone (MPP) were all purchased from Nacalai Tesque, Inc.

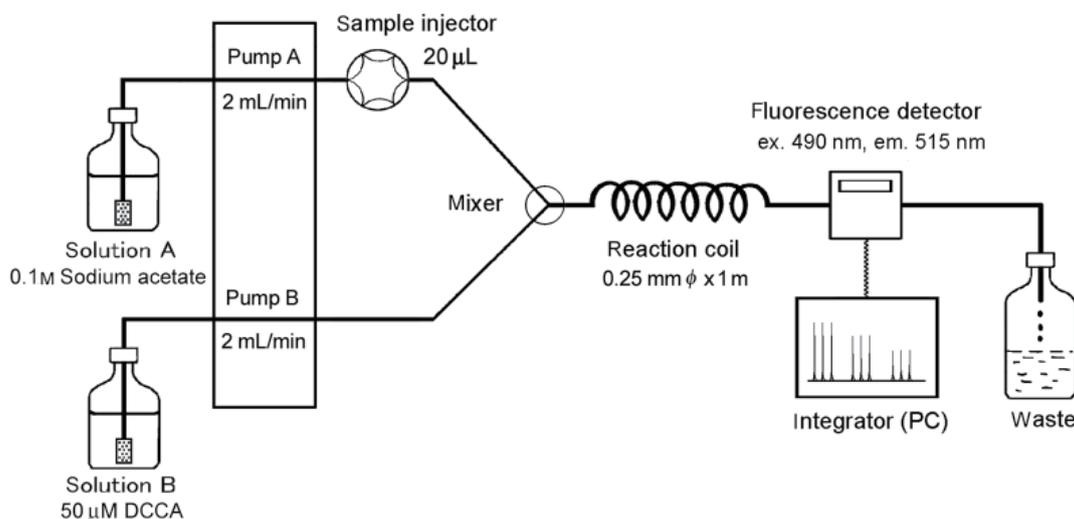


Figure 2. Block diagram of the proposed FIA system.

Quercetin dihydrate was purchased from Wako Pure Chemical Industries, Ltd. (Osaka, Japan). Melatonin, quinolinic acid, and kynurenine were purchased from Sigma-Aldrich (St. Louis, MO, USA). These compounds were dissolved in H₂O to prepare a 1.0 mM stock solution. In the case of the lipophilic compounds such as the pyrazolone derivatives, a water-soluble organic solvent such as DMF was used. The stock solution was diluted with 0.1 M sodium acetate in H₂O to prepare a 0.05-20 μM solution, and 500 μL of the aliquot was used for the evaluation of anti-oxidant activity by the FIA system described above.

2.5. Evaluation of anti-oxidant activity

In the absence of a test compound, the fluorescence peak height obtained in the FIA chart was designated as the blank value (100%) ($n = 5$). Subsequently, a dose-response curve for each test compound was drawn by plotting the fluorescence peak height at each concentration of test compound against the concentration of the test compound ($n = 5$). The concentration of the test compound indicating 50% of the blank value was regarded as the EC₅₀ of the compound according to previous papers (23,24). Thus, in the present study, the EC₅₀ value (μM) was evaluated as the anti-oxidant activity of the test compound.

3. Results and Discussion

3.1. Source of hypochlorite anion

Among several kinds of ROS such as HO·, O₂⁻, ¹O₂, H₂O₂, ONOO⁻, HOCl, and ROO·, in the present study, we chose HOCl as the ROS for the present FIA assay. As a source of OCl⁻, we first investigated the use of sodium hypochlorite (NaOCl); however, the relative standard deviation (RSD) value of the sample was too large, and therefore stable precise data could not be obtained (data not shown). For this reason, it was speculated that the poor precision may have arisen from the physical properties of NaOCl, because NaOCl is a liquid compound. Commercially available NaOCl is a 5-7% solution, and therefore the concentration is not exact. Thus, DCCA was chosen as the source of OCl⁻, because although DCCA is a solid compound, it can release OCl⁻ in aqueous solution (Figure 1).

3.2. Effect of pH and kinds of base

In the present study, APF was used for the ROS indicator. APF fluoresces, which originates from fluorescein in the presence of a hydroxyl radical or HOCl, although APF itself shows no fluorescence (22). The responsiveness of APF to HOCl was three times higher than it was to a hydroxyl radical, and therefore a subtle scavenging effect of OCl⁻ by a test compound could be detected using APF. APF can react with OCl⁻ to emit intense fluorescence at 515 nm in a basic solution. A pH range for the carrier solution of 8.0-11.0 was therefore recommended. For preparing the carrier solution at pH 8.0-11.0, several bases were examined as to whether sufficient data could be obtained. Table 1 lists the fluorescence peak height and the RSD of each base examined in the present study. As listed in Table 1, the use of 0.1 M sodium acetate (CH₃CO₂Na) in H₂O gave the highest peak height and the smallest RSD values. Therefore, 0.1 M CH₃CO₂Na was used as the carrier solution.

3.3. Effect of concentrations of APF and DCCA

Figures 3A and 3B show the change of fluorescence intensity (peak height) as a function of the concentrations of DCCA and APF, respectively. The observed fluorescence peak height increased with increasing concentration of both DCCA and APF. In the preliminary experiment, a decrease of fluorescence peak height was not dependent on the concentration of the test compound at DCCA concentrations above 50 μM in carrier solution B (Figure 2) (data not shown); therefore, the DCCA concentration was set at 50 μM. The obtained peak height was 7.0×10^5 at 5.0 μM APF; therefore, the concentration of 5.0 μM APF was sufficient for the evaluation of EC₅₀ values for each test compound. Reproducibility of the FIA system was estimated using relative standard deviations of peak height for each measured point, RSD < 4.5% for DCCA (Figure 3A) and RSD < 4.4% for APF (Figure 3B).

3.4. Effect of flow rate

Figure 4 shows a representative FIA chart of reduced glutathione obtained at a flow rate of 2.0 mL/min. The observed peak heights apparently decreased with increasing concentration of reduced glutathione, indicating that the anti-oxidant activity of reduced glutathione could also be evaluated by the present FIA

Table 1. Effects of carrier solution on the fluorescence peak height and RSD (%)

Carrier solution	Mean value of peak height	RSD (%)
0.01 M Sodium phosphate buffer in H ₂ O (pH 7.4)	3,200	5.3-19.7
0.1 M Sodium acetate in H ₂ O (pH 8.5)	531,689	0.2-2.7
0.1 M Sodium bicarbonate in H ₂ O (pH 8.7)	40,700	1.5-8.7
0.1 M Sodium carbonate in H ₂ O (pH 10.5)	7,250	0.7-5.2

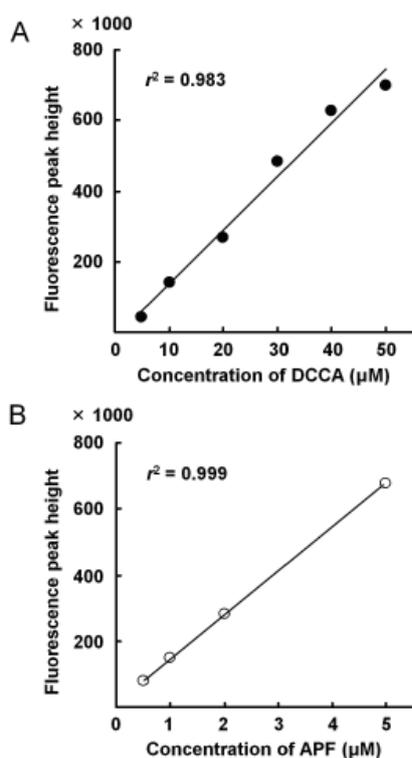


Figure 3. Fluorescence intensity of fluorescein produced from APF as a function of concentration by the present FIA system. Concentration of DCCA (A) and APF (B).

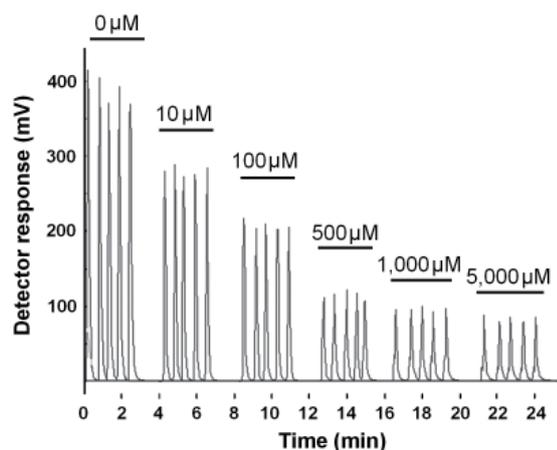


Figure 4. Representative FIA chart of 0, 10, 100, 500, 1,000, and 5,000 µM reduced glutathione, obtained at a flow rate of 2.0 mL/min. Numerals in this figure represent the concentrations of reduced glutathione.

system. When the flow rate was slow (0.5 mL/min), the peak top was divided. Although the reason why the peak shape deteriorated is not yet understood, a clear peak shape would be required for an accurate evaluation. In addition, a higher flow rate can contribute to a reduced analysis time. Thus, considering the peak shape and total analysis time, a flow rate of 2.0 mL/min was chosen in this study.

3.5. Anti-oxidant activity of test compounds

Figure 5 shows a representative dose-response curve

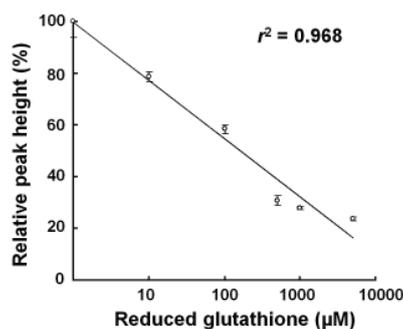


Figure 5. Dose-response curve of reduced glutathione evaluated by the present FIA system.

Table 2. EC₅₀ values of test compounds, obtained by the proposed method

Test compounds	EC ₅₀ (µM)
Ascorbic acid	307.2
Reduced glutathione	162.6
Quercetin dihydrate	1,024
Dithiothreitol	211.1
Edaravone (3-methyl-1-phenyl-5-pyrazolone)	203.1
Quinolinic acid	1,723
Kynurenine	934.2
Melatonin	1,517

of reduced glutathione evaluated by the present FIA system. At concentrations of reduced glutathione above 1,000 µM, the plotted point was excluded from the linear regression line; therefore, only the points within the linear regression line were adopted to determine the EC₅₀ value. Similarly, several compounds were tested for anti-oxidant activity using the present FIA system.

In the present study, the test compounds other than reduced glutathione were ascorbic acid, quercetin dihydrate, dithiothreitol, edaravone, quinolinic acid, kynurenine, and melatonin. The EC₅₀ values for each compound were calculated and are listed in Table 2. Among them, reduced glutathione exhibited the smallest EC₅₀ value. To our knowledge, thus far there have been no reports revealing that the anti-oxidant activity against OCl⁻ of reduced glutathione was more effective than that of ascorbic acid, which is a well-known antioxidant. In the case of scavenging OCl⁻ in an *in vitro* experiment, our studies indicated that reduced glutathione might be more effective than ascorbic acid. Based on these studies, it is speculated that reduced glutathione might also be efficacious for scavenging OCl⁻ *in vivo*.

4. Conclusion

By using APF as an ROS indicator, an FIA system for the evaluation of antioxidants using DCCA as a source of OCl⁻ was developed. The anti-oxidant activity of several compounds was satisfactorily evaluated by the present FIA system, demonstrating that it can be useful as a screening assay for anti-oxidant activity against OCl⁻.

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Stability-indicating HPLC method for the determination of the stability of oxytocin parenteral solutions prepared in polyolefin bags

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ABSTRACT: Oxytocin is very commonly used in clinical settings and is a nonapeptide hormone that stimulates the contraction of uterine smooth muscles. In this study the stability of extemporaneously compounded oxytocin solutions was investigated in polyolefin bags. The sterile preparations of oxytocin were compounded to the strength of 0.02 U/mL in accordance with United States Pharmacopeia (USP) <797> standards. In order to carry out the stability testing of these parenteral products, the solutions were stored under three different temperature conditions of -20°C (frozen), 2-6°C (refrigerated), and 22-25°C (room temperature). Three solutions from each temperature were withdrawn and were assessed for stability on days 0, 7, 15, 21, and 30 as per the USP guidelines. The assay of oxytocin was examined by an HPLC method at each time point. No precipitation, cloudiness or color change was observed during this study at all temperatures. The assay content by HPLC revealed that oxytocin retains greater than at least 90% of the initial concentrations for 21 days. There was no significant change in pH and absorbance values for 21 days under all the conditions of storage. Oxytocin parenteral solutions in the final concentration of 0.02 U/mL and diluted in normal saline are stable for at least 30 days under frozen and refrigerated conditions for 30 days. At the room temperature, the oxytocin solutions were stable for at least 21 days. The stability analysis results show that the shelf-life of 21 days observed in this study was far better than their recommended expiration dates.

Keywords: Oxytocin, stability, parenteral, compounding, HPLC

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1. Introduction

Oxytocin, a nonapeptide hormone, stimulates the contraction of uterine smooth muscles. It is commonly used in the clinical settings for the induction and augmentation of the first stage of labor, mid trimester abortion, and the cessation or prevention of postpartum hemorrhage (1-4). The recommended doses of oxytocin are as follows; for the induction of labor: 10 units per 1,000 mL, for the control of post-partum bleeding: 10 to 40 units per 1,000 mL, and for the treatment of inevitable abortion: 10 units in 500 mL (5). For these purposes, oxytocin is most often used as parenteral solutions and the recommended solvents are saline or 5% dextrose in saline solution (Product Insert, Teva Parenteral Medicines, 2004). The intravenous forms are diluted from the commercially available oxytocin vials, since the final recommended concentrations are commercially unavailable. These intravenous solutions are prepared using aseptic conditions under the laminar flow hood and should be stored suitably until further used.

There are few studies which provide the guidance regarding the short-term stability of these compounded syringes (3,6). In a study by Boothby *et al.* (7), the stability of oxytocin solution was examined in Ringer's lactate solution at 4 and 25°C for over 31 days. In another study by Trissel *et al.* (8), the stability was done for 90 days at room temperature in 5% dextrose injection, normal saline, and lactated Ringer's solution. However, no stability-indicating study has evaluated the stability of oxytocin stored for a period of 30 days under different temperature conditions in normal saline (the most commonly used diluent for oxytocin). Thus, the hospital pharmacy is placed in a unique situation in determining how to prepare the solutions, how long the product should be stored, and how to improve compounding efficiency while at the same time reducing waste in any health-care model.

The objective of the present study was to determine the physical and chemical stability of extemporaneously compounded oxytocin to 0.02 U/mL in normal saline and stored in polyolefin bags. Three different temperature conditions were used for this study; room temperature (22-25°C), frozen (-20°C) or under refrigeration (2-6°C) for up to 30 days.

2. Materials and Methods

2.1. Materials

All chemicals were of analytical grade. Oxytocin, sodium phosphate, sodium chloride, sodium hydroxide, hydrochloric acid, and all other chemicals were purchased from Sigma Chemical Co. (St. Louis, MO, USA). Acetonitrile (HPLC grade) was purchased from Fisher Chemicals (Fair Lawn, NJ, USA). Oxytocin vials of 100 U/10 mL (10 U/mL), 10 mL Multiple Dose Vial was purchased from Teva Pharmaceuticals (Teva Pharmaceuticals USA, Inc., Sellersville, PA, USA) and normal saline bags were purchased from B. Braun (Bethlehem, PA, USA).

2.2. HPLC sample preparation using solid phase extraction (SPE) method

All the samples for the HPLC analysis were prepared using Oasis HLB cartridges (60 mg-3 mL; Product No. WAT094226) obtained from Waters Corporation (Milford, MA, USA). The cartridges were pre-washed with 3 mL of acetonitrile and 1.5 mL of water, then conditioned with 1.5 mL of 10 mM phosphate buffer (pH 7.5). The cartridge was not permitted to run dry during the whole conditioning procedure. The sample (3 mL) of oxytocin solution was loaded on the cartridge and was eluted with 0.75 mL 50% of 10 mM phosphate buffer (pH 7.5) and 50% acetonitrile. This sample was immediately centrifuged at $9,000 \times g$ for 3.5 min and 500 μ L of supernatant was transferred to an HPLC vial with a micro-insert.

2.3. HPLC system

All the chromatographic studies were performed on a Dionex Ultimate 3000 HPLC system connected with an absorbance detector. The separations were done on X Bridge™ C8 cartridge column (250 \times 4.6 mm I.D., Waters Associates, Milford, MA, USA) with the column particle diameter of 3.5 μ m. The column effluents were monitored at wavelength of 214 nm for a run time of 16 min at the temperature of 30°C. The absorbance value was selected based on the full range scan of an oxytocin sample that was carried under the same conditions in the HPLC system. The mobile phase was 50% of 10 mM phosphate buffer (pH 7.5) and 50% acetonitrile. The mobile phase was filtered and degassed before use. The flow rate was 0.2 mL/min and the injection volume was 50 μ L.

2.4. Stability-indicating HPLC assay

The suitability of the present HPLC conditions to be used as a stability-indicating method was tested by accelerating the degradation of oxytocin. The stability-indicating capability of the assay was examined by degradation of 0.02 U/mL oxytocin in 1 M HCl and 1

M NaOH solutions and these solutions were heated at 90°C for 1 h. The samples were withdrawn before and after heating each of the solutions and were subjected to SPE. Each sample was analysed by HPLC using the conditions as explained above.

2.5. Oxytocin standard curve

The calibration of HPLC system was performed by constructing a standard curve using five known concentrations of oxytocin. In order to prepare standard curve, fixed amounts (0.01, 0.015, 0.02, 0.03, and 0.04 U/mL) of standard oxytocin were respectively added to the normal saline solutions and these samples were analyzed by subjecting them to SPE and then using the standardized HPLC conditions.

All the samples were analyzed three times (intra-day variation) on three different days (inter-days variation). The accuracy was calculated at each concentration as the ratio of the measured concentration to the nominal concentration multiplied by 100%.

The limit of quantitation (LOQ) of the method was defined as the lowest concentration of that could be quantitatively determined with acceptable precision and accuracy. Acceptance limits were defined as accuracy of 80-120% and precision of $\leq 20\%$.

2.6. Preparation of oxytocin parenteral solutions

Dilutions of the oxytocin parenteral solutions to nominal concentrations of 0.02 U/mL were performed by the Charleston Area Medical Center (CAMC Solutions, Charleston, WV, USA) under aseptic conditions. All the solutions were prepared on the same day. Briefly, 2 mL of solution was withdrawn from a 1,000 mL normal saline bag and was replaced by 2 mL of oxytocin stock (10 U/mL). The bag was then agitated for couple of seconds to ensure uniform mixing.

The solutions were frozen (-20°C), refrigerated ($2-6^{\circ}\text{C}$), or stored at room temperature ($22-25^{\circ}\text{C}$). The solutions at room temperature were exposed to normal fluorescent light. Frozen and refrigerated samples were kept in a dark refrigerator and were exposed to light only during sampling. These samples were allowed to warm at room temperature and no external source of heat was used to warm these parenteral solution bags. Three bags for each of the storage conditions were assessed for physical and chemical stability over 30 days. Stability was assessed on days 0, 7, 15, 21, and 30. The storage temperatures were closely monitored throughout the study.

2.7. Physical evaluation

Physical stability of oxytocin parenteral solutions was assessed by visual examination. The solutions were evaluated against black and white backgrounds for visible particulate matter, cloudiness, and color changes.

To complement the visual inspection of the syringes, absorbance was also measured at the wavelength of 600 nm using spectrophotometer (Hitachi High Technologies America, Inc., CA, USA).

The pH was measured at each sampling point with Corning Pinnacle pH meter (Model 530, Cole-Parmer, IL, USA) fitted with Orion 9102BNWP electrode (single junction Ag/AgCl refillable combination pH), using three-point standardization with buffer solutions (pH of 4.0, 7.0, and 10.0). The pH of each sample was read at least two times. pH is defined as an important parameter that governs the stability of the product as the change in pH can cause the precipitation of the product. Thus pH along with spectroscopy can be a valuable tool for evaluating the stability.

2.8. Oxytocin stability assay

Chromatogram peak heights were used to determine oxytocin concentrations in the parenteral solutions. Three samples were withdrawn on each of the sampling day from each of the storage temperature and each sample was injected at least two times. The percentage of oxytocin remaining in each of the infusion after day zero was calculated based on the oxytocin content at day zero. The drug concentration was considered stable if its concentration was more than 90% of the initial concentration.

2.9. Data Analysis

Student's *t*-test was used to compare the difference between the data of interest with $\alpha = 0.05$ as the minimal level of significance. Wherever possible, the data is presented as mean \pm standard deviation.

3. Results

3.1. Standard curve and method validation

The chromatogram of oxytocin standards in normal saline showed a peak at retention time of 11.6 min (Figure 1). A

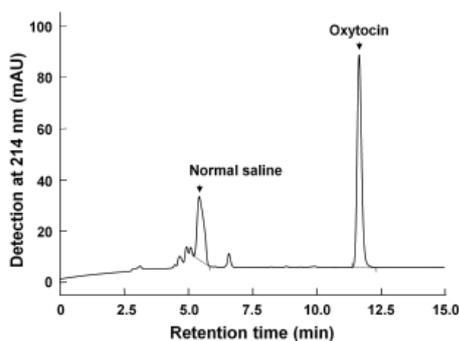


Figure 1. A representative HPLC chromatogram of oxytocin sample (0.02 U/mL) in the saline solution. The chromatogram showed a displacement peak originating from the saline solution at 5.4 min and oxytocin peak at 11.6 min.

blank sample (normal saline only) was also injected to the HPLC system and a peak at 5.4 min was observed from this sample. No peak was observed when sterile water or mobile phase was injected to the HPLC system. Thus the first peak in the chromatogram is because of normal saline and the second peak at 11.6 min is for oxytocin. As shown in Figure 2, a good linearity was exhibited in the concentration range (0.01-0.04 U/mL) by using the presently developed HPLC method. The average coefficient of determination of 0.99 was observed for the standard curve. The slopes of the curves illustrated an excellent agreement with coefficient of variability.

The intra- and inter-day relative standard deviations (RSD) were calculated to be 1.83% and 1.67%, respectively. For each concentration studied, a relative error (RE) of less than 10% was obtained. An acceptable precision and accuracy was acquired by this method for all the standards and quality controls based on the recommended criteria (9). The percentage recovery of oxytocin using the present HPLC method was also calculated from the peak areas obtained. As shown from the data in Table 1, an admirable recovery was accomplished at each of the added concentration. In accordance to the official requirements the LOQ was 0.005 U/mL.

3.2. Stability-indicating HPLC method characterization

Figures 3A and 3B show the degradation of oxytocin after heating at 90°C for 1 h in the presence of 1 M HCl and 1 M NaOH, respectively. As is clear from the chromatograms, multiple degradation product peaks were observed. Under both the conditions, acidic and alkaline, there was significant loss of

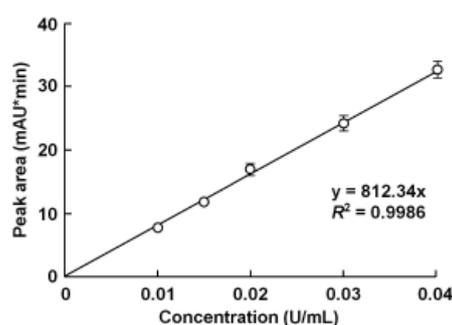


Figure 2. Standard curve of oxytocin assay. The samples were processed by HPLC as described in Section 2.3. For each concentration point 3 set of samples were prepared and this study was done on three different days.

Table 1. Recovery of oxytocin from the standard samples (n = 3)

Concentration added (U/mL)	Concentration obtained (U/mL)	Recovery (%)
0.01	0.012 \pm 0.0008	100.2
0.03	0.032 \pm 0.0011	100.2
0.05	0.053 \pm 0.002	100.3

oxytocin after heating at 90°C for 1 h. There was no interference of the degradation products as these peaks were seen at retention times other than the retention time for oxytocin (11.6 min, arrowhead in Figure 3). These results suggest that the present method is sufficiently specific to the drug and that, based on the chromatograms of forced degradation of oxytocin, this method can simultaneously analyze oxytocin and its degradation products in a sample.

3.3. Physical evaluation

The oxytocin parenteral solution remained clear throughout the study when kept frozen, under refrigeration, or at room temperature. No precipitation was observed. No significant increase ($p > 0.05$) in absorbance was observed as compared to the freshly read sample of the distilled water at the wavelength of 600 nm (data not shown).

The pH value of normal saline before addition of oxytocin was around 4.40. After the addition of oxytocin this value increases to 5.48 ± 0.18 . As shown in Table 2, there was no significant increase or decrease ($p > 0.05$) in this pH value under all the three temperatures used in this study over a period of 30 days.

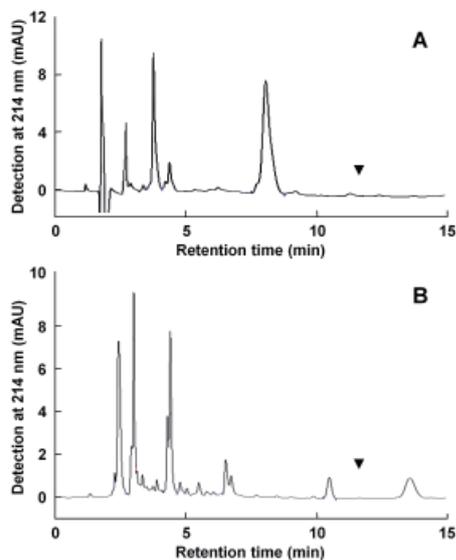


Figure 3. Chromatograms of oxytocin after accelerated degradation. Oxytocin was treated under (A) acidic and (B) alkaline conditions and the samples were processed by HPLC as described in Section 2.3. Arrowhead: expected elution position of oxytocin (11.6 min).

3.4. Drug content

As shown in Figure 4, all the oxytocin parenteral solutions were in the range of 90-110% of the labeled drug amount under frozen (-20°C) and refrigerated ($2-6^{\circ}\text{C}$) conditions for 30 days; at room temperature ($22-25^{\circ}\text{C}$) the oxytocin solutions were stable for at least 21 days. Minimal to no degradation was observed and no new degradation peaks were observed in the stable samples (data not shown).

4. Discussion

The stability data of the routinely used large volume parenteral in the final working condition are unavailable as they are commercially not available. Knowing the stability of these preparations would minimize compounding them just before the administration. A concentration of 0.02 U/mL is commonly used in the clinical setting.

The parent drug stability test guideline Q1A (R2) issued by the International Conference on Harmonization (ICH) suggest that stress studies should be carried out on a drug to ascertain its inherent stability characteristics (10). A proper identification of degradation products would hence support the suitability of the proposed analytical procedures. It also requires that analytical test procedures for stability samples should be stability-indicating and be fully validated. In order to analyze the concentration of oxytocin, a stability-indicating HPLC method was validated and used for the present study.

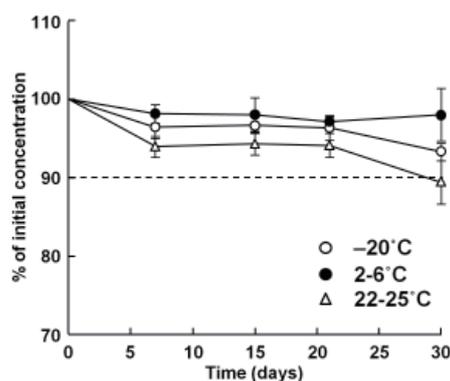


Figure 4. Percentage of initial concentration versus time profile for oxytocin parenteral solutions ($n = 9$). Open circle, -20°C ; closed circle, $2-6^{\circ}\text{C}$; open triangle, $22-25^{\circ}\text{C}$. Dashed line indicates 90% of initial concentration of oxytocin.

Table 2. The results of pH analysis ($n = 9$)

Storage temperature	Day 0	Day 7	Day 15	Day 21	Day 30
-20°C	5.48 ± 0.18	5.17 ± 0.15	5.16 ± 0.07	5.16 ± 0.06	5.10 ± 0.13
$2-6^{\circ}\text{C}$	5.48 ± 0.18	5.22 ± 0.10	5.21 ± 0.04	5.20 ± 0.13	5.08 ± 0.03
$22-25^{\circ}\text{C}$	5.48 ± 0.18	5.36 ± 0.16	5.33 ± 0.06	5.20 ± 0.04	5.28 ± 0.17

The present HPLC method met with all the acceptance criteria and was sensitive and reproducible enough for the acceptable study of oxytocin in unknown samples. As reported by Trissel (11), the failure to recognize the degradation products is the most common point that leads to erroneous reporting of the data on the stability studies.

There are some published studies in the literature that do report the short-term stability of oxytocin (3,6). Gard *et al.* (3) have reported oxytocin is stable in lactated Ringer's solution and lactated Ringer's-dextrose 5% solution over a 24-hour period at 25°C and over a 7-day period at 5°C. In an attempt to measure long term stability, Boothby *et al.* (7) published that oxytocin in lactated Ringer's solution is stable for 31 days at 4 and 25°C. Another comprehensive study done by Trissel *et al.* (8) have examined the extended stability of oxytocin in 5% dextrose injection, 0.9% sodium chloride injection, lactated Ringer's injection bags for 90 days at room temperature. Based on their results they have reported that oxytocin diluted in 5% dextrose or 0.9% sodium chloride is stable to be used for 90 days at room temperature. However, when diluted in lactated Ringer's solution it is advisable to be used only for 28 days when stored at room temperature. The most commonly recommended solvent for the dilution of oxytocin for clinical use is 0.9% sodium chloride. It is very pertinent to establish the stability of commonly used concentration of oxytocin solution at three normally used storage temperatures for parenteral products in the hospitals. Thus for the present study we have used all the three different temperature conditions of room temperature (22-25°C), refrigeration (2-6°C), and frozen (-20°C) and done the stability studies for 30 days.

Based on our results, oxytocin parenteral solutions are stable for at least 30 days when stored under refrigerated or frozen conditions. At room temperature, these solutions were observed to be stable for at least 21 days. Trissel *et al.* (8) have recommended that the use of oxytocin parenteral solutions should be restricted to no greater than 28 days at room temperature, regardless of the diluent used. The recommendation based on our present study is reasonably more than what is presently used.

A major driver of operations within the pharmacy department in a hospital setting is increasing efficiency and reducing waste. Therefore, Charleston Area Medical Center (CAMC) utilizes a consolidated intravenous medication preparation model where a central, USP <797> compliant clean room prepares many compounded sterile products in batches for use within our organization. Thus, the pharmacy is able to mass produce extemporaneously compound parenteral solutions from this facility and ship the finished products to each hospital of our health system, thus allowing the organization to free up its pharmacy team to provide more direct patient care while improving operational efficiencies. One of the challenges to this

model is expiration dating of medications. The aim of the compounding pharmacy is to prepare sterile products that can be compounded in large batches and with long expiration dates. Based on the present study, oxytocin solutions are stable for at least 21 days under different temperature conditions. By consolidating with the pharmacy in a hospital, workload can be removed from each facility thus shortening the turnaround time for the pharmacy to send medications to the patients.

The results of this study show that extended stability for oxytocin parenteral solutions can be assigned. However, assigned stability data should not exceed the mandates from USP <797> in the absence of end product sterility testing according to USP <71>.

5. Conclusion

A stability-indicating method was developed, which separates all the degradation products formed. The present HPLC method meets all the acceptance criteria and was sensitive and reproducible enough for the acceptable study of oxytocin in unknown samples. A good linearity of the standard curves with reproducible slopes was observed with this method. Hence it is recommended for analysis of the drug and degradation products in stability samples. The present method proved to be rapid and simple, and its precision is sufficient for routine quality control of oxytocin.

The oxytocin solutions were stable for at least 21 days under all the temperatures studied in the concentrations normally used in clinical setting when prepared in 0.9% sodium chloride (normal saline) for parenteral use. The extended stability of this preparation can improve operational efficiency, reduce waste, and reduce medication delivery delays.

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