

## RESEARCH PAPER

# Resveratrol dimers are novel sphingosine kinase 1 inhibitors and affect sphingosine kinase 1 expression and cancer cell growth and survival

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### Keywords

sphingosine 1-phosphate; sphingosine kinase 1; apoptosis; cancer; proliferation; resveratrol; extracellular signal-regulated kinase; polyADP ribose polymerase; inhibitor kinetics; balanocarpol; ampelopsin A

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## BACKGROUND AND PURPOSE

Sphingosine kinase 1 catalyses formation of the bioactive lipid, sphingosine 1-phosphate, which protects cancer cells from apoptosis. Therefore, sphingosine kinase 1 is a novel target for intervention with anti-cancer agents. We have assessed the effect of the anti-cancer agent, resveratrol and its dimers (ampelopsin A and balanocarpol) on sphingosine kinase 1 activity and on survival of MCF-7 breast cancer cells.

## EXPERIMENTAL APPROACH

Ampelopsin A and balanocarpol were purified from *Hopea dryobalanoides* and their effect on sphingosine kinase 1 activity and expression, [<sup>3</sup>H] thymidine incorporation, ERK-1/2 phosphorylation and PARP activity assessed in MCF-7 cells.

## KEY RESULTS

Resveratrol, ampelopsin A and balanocarpol were novel inhibitors of sphingosine kinase 1 activity. Balanocarpol was a mixed inhibitor (with sphingosine) of sphingosine kinase 1 with a  $K_{ic} = 90 \pm 10 \mu\text{M}$  and a  $K_{iu}$  of  $\sim 500 \mu\text{M}$ . Balanocarpol and ampelopsin A also induced down-regulation of sphingosine kinase 1 expression and reduced DNA synthesis, while balanocarpol stimulated PARP cleavage in MCF-7 breast cancer cells. Resveratrol was a competitive inhibitor (with sphingosine) of sphingosine kinase 1 with a  $K_{ic} = 160 \pm 40 \mu\text{M}$ , reduced sphingosine kinase 1 expression and induced PARP cleavage in MCF-7 cells.

## CONCLUSIONS AND IMPLICATIONS

Each molecule of balanocarpol may bind at least two sphingosine kinase 1 catalytic molecules to reduce the activity of each simultaneously. These findings suggest that resveratrol, ampelopsin A and balanocarpol could perturb sphingosine kinase 1-mediated signalling and this might explain their activity against MCF-7 breast cancer cells.

## LINKED ARTICLE

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## Abbreviations

COSY, correlation spectroscopy; DEPT, distortionless enhancement through polarization transfer; DMS, *N,N*-dimethylsphingosine; HMBC, heteronuclear multiple bond coherence; HMQC, heteronuclear multiple quantum coherence;  $K_{ic}$ , competitive inhibitor constant;  $K_{iu}$ , uncompetitive inhibitor constant; NOESY, nuclear overhauser enhancement spectroscopy; SKi, 2-(*p*-hydroxyanilino)-4-(*p*-chlorophenyl)thiazole; SK1, sphingosine kinase 1; S1P, sphingosine 1-phosphate

## Introduction

Resveratrol (3,4',5-trihydroxy-*trans*-stilbene), isolated as an antifungal agent (Langcake and Pryce, 1977), has several beneficial effects on health including activities against cancer, inflammation, cardiovascular- and age-related diseases (Baur and Sinclair, 2006). Resveratrol exhibits antioxidant activity, which can be attributed to the phenol rings, which are strong scavengers of reactive oxygen species (Leonard *et al.*, 2003). However, many biological effects of resveratrol cannot be explained by its antioxidant properties. For example, the anti-inflammatory effects of resveratrol involve inhibition of COX-2 in human mammary cells (Subbaramaiah *et al.*, 1998). Resveratrol also reduces breast and skin cancer in mice models by blocking COX-1 and COX-1-associated hydroxyperoxidase activity (Jang *et al.*, 1997). Resveratrol has also gained wide attention because of the 'French paradox' where consumption of red wine is associated with a lower incidence of cardiovascular-related deaths, despite patients consuming a high-fat diet (Frémont, 2000). It has been suggested that polyphenols, such as piceatannol and resveratrol may activate human deacetylase (SIRT 1) *in vitro* and sirtuin (SIR 2) *in vivo* to prolong the lifespan of *Saccharomyces cerevisiae*, which may be attributed to the stabilization of rDNA repeats (Howitz *et al.*, 2003). Furthermore, resveratrol has been found to extend the lifespan of *Caenorhabditis elegans* (Wood *et al.*, 2004) and mice fed on a high-calorie diet (Baur *et al.*, 2006), indicating an evolutionary conserved mechanism of SIR2 in regulating metabolism and aging.

Resveratrol has also been shown to be therapeutically useful in reducing the growth and progression of skin, lung and breast cancers (Athar *et al.*, 2007). Resveratrol induces apoptosis in MDA-MB-231 breast cancer cells via a PKC $\delta$ -dependent activation of serine palmitoyltransferase and neutral sphingomyelinase, which results in increased *de novo* synthesis of the pro-apoptotic sphingolipid, ceramide (Scarlatti *et al.*, 2003). Resveratrol has also been shown to inhibit oxidative burst and sphingosine kinase 1 (SK1)-dependent degranulation in human neutrophils (Issuree *et al.*, 2009).

Sphingosine kinase is an enzyme (two isoforms called SK1 and SK2), catalysing the formation of the bioactive lipid, sphingosine 1-phosphate (S1P) and has a central role in cancer progression. For instance, there is increased expression of SK1 mRNA transcript and/or SK1 protein in stomach, lung, brain, colon, kidney and breast cancers and non-Hodgkins lymphoma (Pyne and Pyne, 2010). Moreover, high tumour expression of SK1 is associated with reduced mean survival time and earlier recurrence of tamoxifen resistance in oestrogen receptor positive breast cancers (Long *et al.*, 2010; Watson *et al.*, 2010). Interestingly, resveratrol also reduces SK1 activity by inhibiting the activation of phospholipase D, which is an upstream regulator of SK1 (Issuree *et al.*, 2009) and promotes the down-regulation of SK1 in PC-3 cells (Briuela *et al.*, 2010).

Plants in the Dipterocarpaceae family such as *Hopea dryobalanoides* are known to produce resveratrol oligomers (Sahidin *et al.*, 2005). Even though these secondary metabolites exhibit high biological activities, they have been ignored, largely because of challenges in achieving their isolation in sufficient quantity from natural sources, coupled

with an inability to chemically synthesize these molecules. Hopeaphenol (a resveratrol tetramer) was first isolated from *H. odorata* (Coggon *et al.*, 1965; 1970). This compound is highly active against several cancer cell lines including human epidermoid nasopharynx carcinoma (KB cells), lung carcinoma (A549 cells) and breast cancer (MCF-7 cells) (Ohyama *et al.*, 1999; Muhtadi *et al.*, 2006). Another tetrameric resveratrol, known as vaticanol B was isolated from *H. dryobalanoides* and exhibited moderate cytotoxic activity against P-388 cells (Sahidin *et al.*, 2005; Muhtadi *et al.*, 2006). Therefore, most of the compounds isolated from *H. dryobalanoides* and related species are cytotoxic against several cancer cell lines. However, the exact mechanisms of action and possible molecular targets are unknown.

During our drug discovery programme to identify novel chemical scaffolds, which inhibit SK1 activity, we found that an extract of *H. dryobalanoides* reduced SK1 activity. Therefore, we sought to investigate the biological effects of resveratrol as a novel SK1 inhibitor, and to purify other compounds produced by *H. dryobalanoides* that inhibit SK1 activity, using bioassay-guided fractionation. We present evidence that resveratrol and its dimers, such as ampelopsin A and balanocarpol induce apoptosis of cancer cells, and this is associated with inhibition and down-regulation of SK1 activity and expression.

## Methods

### *Extraction and isolation of ampelopsin A and balanocarpol*

Dried and ground *H. dryobalanoides* leaf (50g) was placed in a brown jar with 500 ml of methanol and 5g of polyvinylpyrrolidone (PVP). The jar was sealed and left at room temperature for at least 48 h. The jar was then shaken twice daily for 1–2 min. The extract was filtered and dried by rotary evaporation to produce a gummy dark greenish extract (2 g). A sample (0.5 g) of this extract was fractionated by Flash chromatography using a 20 g ISOLUTE® Flash Si II cartridge in Biotage Flash Master. The flow rate was set at 20 ml min<sup>-1</sup> for gradient elution using hexane, dichloromethane, butan-2-ol and methanol. The volumes of the fractions were then reduced by rotary evaporation. Subsequently, all fractions were freeze-dried. 12 major fractions were obtained by monitoring their UV-VIS spectra (F1: 2 mg; F2: 5 mg; F3: 3 mg; F4: 3 mg; F4A: 15 mg; F5: 11 mg; F 5B: 2.5 mg; F6: 7.7 mg; F7: 46 mg; F8: 68 mg; F9: 3.4 mg; F10: 2 mg). Ampelopsin A was obtained from the methanol phase after extracting F5 with 2,2,4-trimethylpentane:methanol (1:1). The yield of ampelopsin A (5 mg) was 0.01% (based on dried weight of plant material). Additionally, 700 g of dried and ground stem bark of *H. dryobalanoides* was successively extracted with 3.5–5 L of hexane, ethyl acetate and methanol at their respective boiling points for 48–72 h. The solvent was removed from the ethyl acetate extract by rotary evaporation to yield 5 g of residue after freeze-drying; this was further processed by vacuum liquid chromatography (Coll and Bowden, 1986). Balanocarpol was then purified from this residue on a Sephadex LH-20 column (5 g) using methanol as the only eluent. Typically, 20 fractions were collected per

column bed volumes and monitored by TLC, NMR and MS. The yield of balanocarpol (300 mg) was 0.043%, based on dried weight of plant material.

### Structure elucidation using NMR and MS

All NMR experiments were performed with a JEOL (JNM LA400) operating at 400 ( $^1\text{H}$ ) and 100 ( $^{13}\text{C}$ ) MHz using deuterated and residual solvent peaks as internal reference.  $^1\text{H}$ -NMR was performed on all samples to establish identity and yield of compounds present in the sample. Further structure elucidation was performed using 2-D NMR experiments such as correlation spectroscopy (COSY), heteronuclear multiple quantum coherence (HMQC) and heteronuclear multiple bond coherence (HMBC).  $^{13}\text{C}$  and distortionless enhancement through polarization transfer (DEPT) NMR experiments were performed when samples were sufficiently pure. Spatial structural information was obtained with nuclear overhauser enhancement spectroscopy (NOESY). MS was used to establish the molecular weights and molecular formulae of selected samples. The sample (1 mg) was dissolved in appropriate solvents and separated by HPLC before being ionized in a ThermoFinnigan LCQ-Decaiontrap or Orbitrap HRESI mass spectrometer. Negative or positive mode electrospray ionization (ESI) analysis was used dependent on the nature of the compounds. Samples were also run in Agilent 6130, LC/MS (Agilent Technologies, Palo Alto, CA, USA) using atmospheric pressure chemical ionization (APCI).

### Cell culture

MCF-7 breast cancer cells (either Neo cells expressing a vector containing a neomycin resistance gene or parental cells) were grown in a monolayer culture in high glucose Dulbecco's modified Eagle's medium (DMEM) with 10% European fetal calf serum (EFCS) and 100 U·mL<sup>-1</sup> penicillin, 100 µg·mL<sup>-1</sup> streptomycin, 0.4% Geneticin (for MCF-7 Neo cells), and 15 µg·mL<sup>-1</sup> insulin at 37°C with 5% CO<sub>2</sub>. HEK 293 cells stably over-expressing SK1 were cultured in DMEM supplemented with 10% EFCS, 100 U·mL<sup>-1</sup> penicillin, 100 µg·mL<sup>-1</sup> streptomycin, 1% non-essential amino acids and 0.8% Geneticin at 37°C in 5% CO<sub>2</sub>.

### Western blotting

Cell lysates were prepared in sample buffer containing 125 mM Tris, pH 6.7, 0.5 mM Na<sub>4</sub>P<sub>2</sub>O<sub>7</sub>, 1.25 mM EDTA, 0.5% w/v SDS containing 1.25% (v/v) glycerol, 0.06% (w/v) bromophenol blue and 50 mM dithiothreitol. Equal amount of proteins were subjected to SDS-PAGE and Western blotting. Resolved proteins were immunoblotted with anti-SK1, anti-ERK2, anti-phospho ERK-1/2, anti-PARP or anti-actin antibodies.

### Sphingosine kinase (SK) activity assay

SK1 activity was assayed as described previously (Lim *et al.*, 2011). Briefly, sphingosine was solubilized in Triton X-100 (final concentration 0.063% w/v) and combined with buffer 1 containing 20 mM Tris (pH 7.4), 1 mM EDTA, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 40 mM β-glycerophosphate, 1 mM NaF, 0.007% (v/v) β-mercaptoethanol, 20% (v/v) glycerol, 10 µg·mL<sup>-1</sup> aprotinin, 10 µg·mL<sup>-1</sup> soybean trypsin inhibitor, 1 mM PMSF

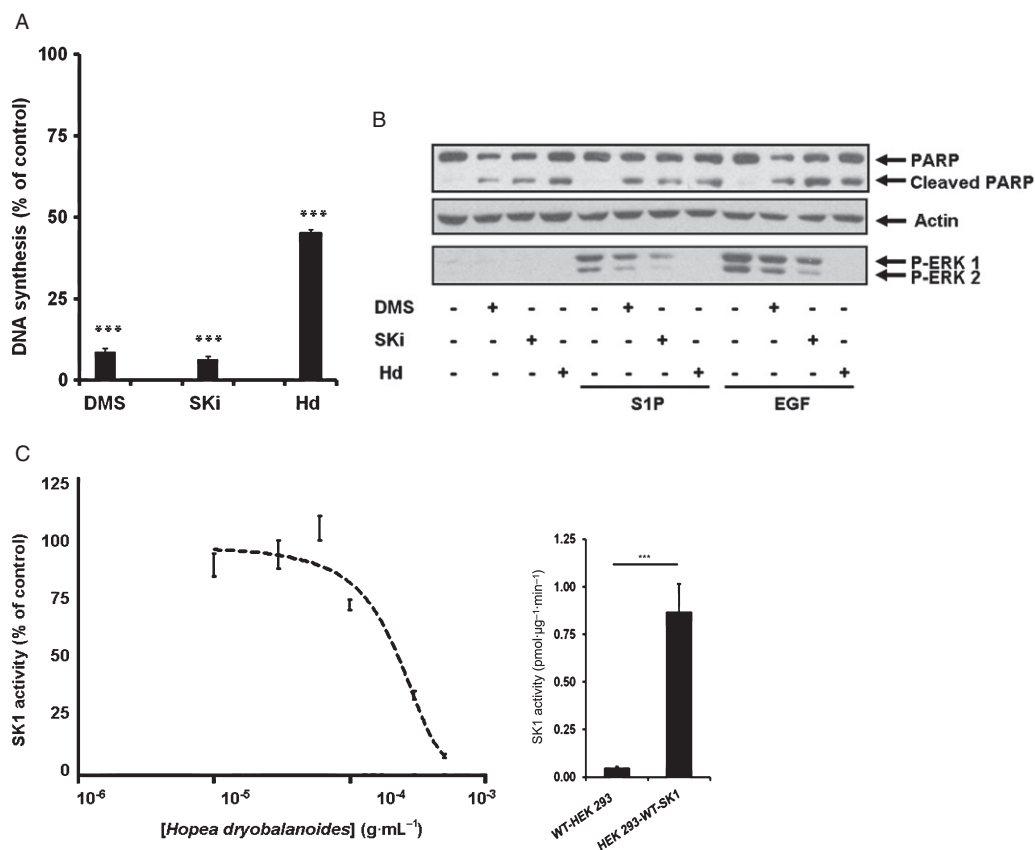
and 0.5 mM 4-deoxy pyridoxine. For SK2 assays, the sphingosine was complexed with bovine serum albumin (final concentration 0.2 mg·mL<sup>-1</sup>) in buffer 1 supplemented with 400 mM KCl. SK activity was determined by incubating purified SK1 or SK2 (15 ng) or cell lysate (15 µg protein, determined by Bradford protein assay) from HEK 293 cells containing stably over-expressed recombinant SK1, for 15–20 min at 30°C, in the presence of sphingosine (0.5 to 20 µM), [ $^{32}\text{P}$ ]ATP (250 µM,  $4.4 \times 10^4$  cpm·nmol<sup>-1</sup> in 10 mM MgCl<sub>2</sub>), and varying concentrations of inhibitors or control vehicle (5% (v/v) DMSO final concentration). The total assay volume per sample was 200 µL. For ATP kinetics, 20 µM sphingosine was combined with 25–1000 µM [ $\gamma$ - $^{32}\text{P}$ ]ATP ( $2.2 \times 10^6$  cpm/assay in 100 mM MgCl<sub>2</sub>), and varying concentrations of inhibitors or control vehicle (5% (v/v) DMSO final concentration). Reactions were stopped by the addition of 500 µL 1-butanol and mixed with 1 mL of 2 M KCl. [ $^{32}\text{P}$ ]-S1P was extracted from the organic phase by washing twice with 1 mL of 2 M KCl before quantification by Cerenkov counting. Kinetic parameters were obtained using the graph plotting and curve fitting programs Biograph (University of Strathclyde, Glasgow UK) and Prism 4.03 (GraphPad). Substrate kinetics were analysed according to the Michaelis-Menten equation and the inhibition constants ( $K_{ic}$  and  $K_{iu}$ ) were determined using Dixon and Cornish-Bowden plots (Cortés *et al.*, 2001).

### [ $^3\text{H}$ ] thymidine assay

MCF-7 cells were plated in 24-well plates ( $2.0\text{--}4.0 \times 10^4$  cells per well) and maintained in complete medium. After 24 h, cells were treated with varying concentrations of inhibitor or vehicle control for a further 15–72 h, as indicated. [ $^3\text{H}$ ]-thymidine (0.5 µCi·mL<sup>-1</sup>) was added for the final 5 h. Samples were terminated by washing 3 times with 1 mL ice cold 10% (w/v) trichloroacetic acid. [ $^3\text{H}$ ]-thymidine incorporated into DNA were harvested with 0.25 mL 0.1% (w/v) SDS, 0.3 M NaOH and quantified by liquid scintillation counting with 2 mL scintillation cocktail.

### Materials

All general biochemicals and anti-actin antibody were from Sigma (Poole, UK). High glucose (DMEM), minimum essential medium (MEM), penicillin-streptomycin (10 000 U·mL<sup>-1</sup> penicillin and 10 000 µg·mL<sup>-1</sup> streptomycin) and Lipofectamine 2000™ were from Invitrogen (Paisley, UK). MCF-7 cells were a gift from R. Schiff (Baylor College of Medicine, Houston, TX, USA). Anti-ERK2 antibody was from BD Transduction Laboratories (Oxford, UK), and anti-SK1 antibody was a gift from A. Huwiler (University of Bern, Switzerland). Sphingosine and S1P were from Avanti Polar Lipids (Alabaster, AL, USA). Purified SK1 and SK2 were from Enzo Life Sciences (Exeter, UK). The SK inhibitor, 2-(*p*-hydroxyanilino)-4-(*p*-chlorophenyl)thiazole (SKi) was from Merck Biosciences (Nottingham, UK). Dried and ground leaves of *H. dryobalanoides* were originally collected by the Forest Institute of Malaysia (FRIM) and stored and extracted at Strathclyde Innovations in Drug Research. 1–2 kg of dried and ground leaves, bark and twigs of *H. dryobalanoides* were obtained.



**Figure 1**

Biological activities of *Hopea dryobalanoides* extract. (A) Bar graph showing the effect of SK1 inhibitors and *H. dryobalanoides* leaf extract on [<sup>3</sup>H]-thymidine incorporation in MCF-7 Neo cells. Quiescent MCF-7 Neo cells were treated with 10 µM *N,N*-dimethylsphingosine (DMS), 10 µM SKi, 5 µg·mL<sup>-1</sup> *H. dryobalanoides* leaf (Hd) extract or control (0.05% v/v DMSO final concentration). Cells were then incubated for a further 15 h and then with [<sup>3</sup>H]-thymidine (0.5 µCi·mL<sup>-1</sup>) for 5 h. Data are expressed as percentage of control and represent means and SDs of triplicate determinations, \*\*\**P* < 0.001, significantly different from control. (B) Western blot showing the effect of SK1 inhibitors and Hd extract on ERK-1/2 activation and PARP cleavage. Quiescent MCF-7 Neo cells were incubated with control (0.1% (v/v) DMSO final concentration) or indicated inhibitors (10 µM DMS, 10 µM SKi and 500 µg·mL<sup>-1</sup> *H. dryobalanoides* leaf extract (Hd)) for 48 h before being stimulated by 25 ng·mL<sup>-1</sup> EGF or 1 µM S1P for 5 min. Cell lysates were separated by SDS-PAGE and immunoblotted with anti-PARP or anti-phospho-ERK1/2 antibodies. Blots were then stripped and reprobed with anti-actin antibody to ensure comparable protein loading. Results are representative of three independent experiments. (C) Graph showing the concentration-dependent inhibitory effect of Hd extract on SK1 activity. Enzyme activity was measured using 10 µM sphingosine and 250 µM [<sup>32</sup>P]-ATP as the substrates and using lysates of HEK 293 cells over-expressing recombinant SK1. Data are expressed as means ± SD of triplicate determinations. Also shown is a histogram comparing SK1 activity, using 10 µM sphingosine and 250 µM [<sup>32</sup>P]-ATP, in lysates of HEK293 cells stably transfected with vector (WT-HEK293) or from HEK293 cells stably transfected with SK1 (HEK293-WT-SK1). *P* < 0.001.

## Results

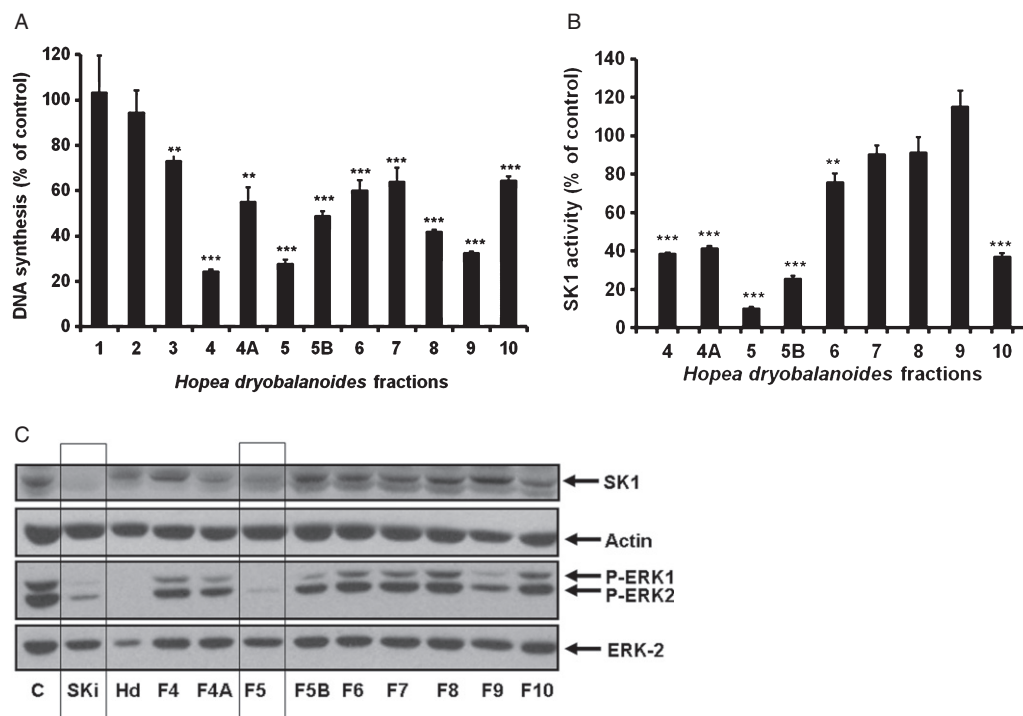
### Isolation of ampelopsin A and balanocarpol as bioactive SK1 inhibitors

We previously demonstrated that MCF-7 (parental) and MCF-7 Neo breast cancer cells express SK1 (GenBank™ number: NM\_001142601), which has a molecular mass of 42 kDa (Loveridge *et al.*, 2010; Lim *et al.*, 2011). MCF-7 Neo cells were used because the action of SK1 inhibitors on SK1 and apoptosis have been extensively characterized in these cancer cells (Loveridge *et al.*, 2010; Lim *et al.*, 2011). Initial screening of plant extracts for bioactivity against SK1 demonstrated that the *H. dryobalanoides* leaf extract reduced [<sup>3</sup>H]thymidine incorporation into DNA (Figure 1A), abolished

S1P- and EGF-stimulated activation of ERK-1/2, promoted PARP cleavage (Figure 1B) and inhibited SK1 activity [SK1 stably over-expressed in HEK 293 cells 10–30-fold increase (dependent on passage number) of SK1 activity vs. lysate from vector-transfected cells] (Figure 1C). These effects were reproduced with the SK1 inhibitors, SKi and *N,N*-dimethylsphingosine (Figure 1A and B).

Therefore, the *H. dryobalanoides* extract was fractionated using Flash chromatography to isolate compounds with SK1 inhibitor activity. Fractions 4, 5 and 9 were the most active fractions in reducing [<sup>3</sup>H] thymidine incorporation into DNA in MCF-7 cells (Figure 2A), while fraction 5 was the most active fraction in inhibiting SK1 activity (~90% inhibition) (Figure 2B). We have previously shown that the SK1 inhibitor, SKi reduces SK1 expression in cancer cells via an inhibitor-





**Figure 2**

Biological activities of *Hopea dryobalanoides* fractions from Flash chromatography. (A) Bar graph showing the effect of purified *H. dryobalanoides* fractions on [<sup>3</sup>H]-thymidine incorporation in MCF-7 Neo cells. Quiescent MCF-7 Neo cells were treated with 5  $\mu\text{g}\cdot\text{mL}^{-1}$  of the solid residue from each *H. dryobalanoides* fractions (1–10) or control (0.05% v/v DMSO final concentration). The cells were then incubated for 15 h and then with [<sup>3</sup>H]-thymidine (0.5  $\mu\text{Ci}\cdot\text{mL}^{-1}$ ) added for 5 h. Data are expressed as percentage of control and represent means  $\pm$  SD of triplicate determinations, \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , significantly different from control. (B) Bar graph showing the effect of purified *H. dryobalanoides* fractions (500  $\mu\text{g}\cdot\text{mL}^{-1}$ ) on recombinant SK1 activity. Data are expressed as percentage of control and represent means  $\pm$  SD of triplicate determinations, \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , significantly different from control. (C) Western blots showing the effect of purified *H. dryobalanoides* fractions on SK1 and phosphorylated ERK-1/2 levels. MCF-7 Neo cells were treated with control (0.2% v/v DMSO), C; 10  $\mu\text{M}$  SKi, 200  $\mu\text{g}\cdot\text{mL}^{-1}$  *H. dryobalanoides* leaf extract (Hd), 20  $\mu\text{g}\cdot\text{mL}^{-1}$  Fractions 4–10 (F4–F10) for 48 h (inhibitors were replenished after 24 h). EGF 25  $\text{ng}\cdot\text{mL}^{-1}$  was added for 5 min before harvesting the cells. Cells lysates were analysed by Western blotting using anti-SK1 antibody. Blots were also probed with anti-phospho ERK1/2 antibody and then stripped and re-probed with anti-actin or anti-ERK-2 antibodies to ensure comparable protein loading. Results are representative of three independent experiments.

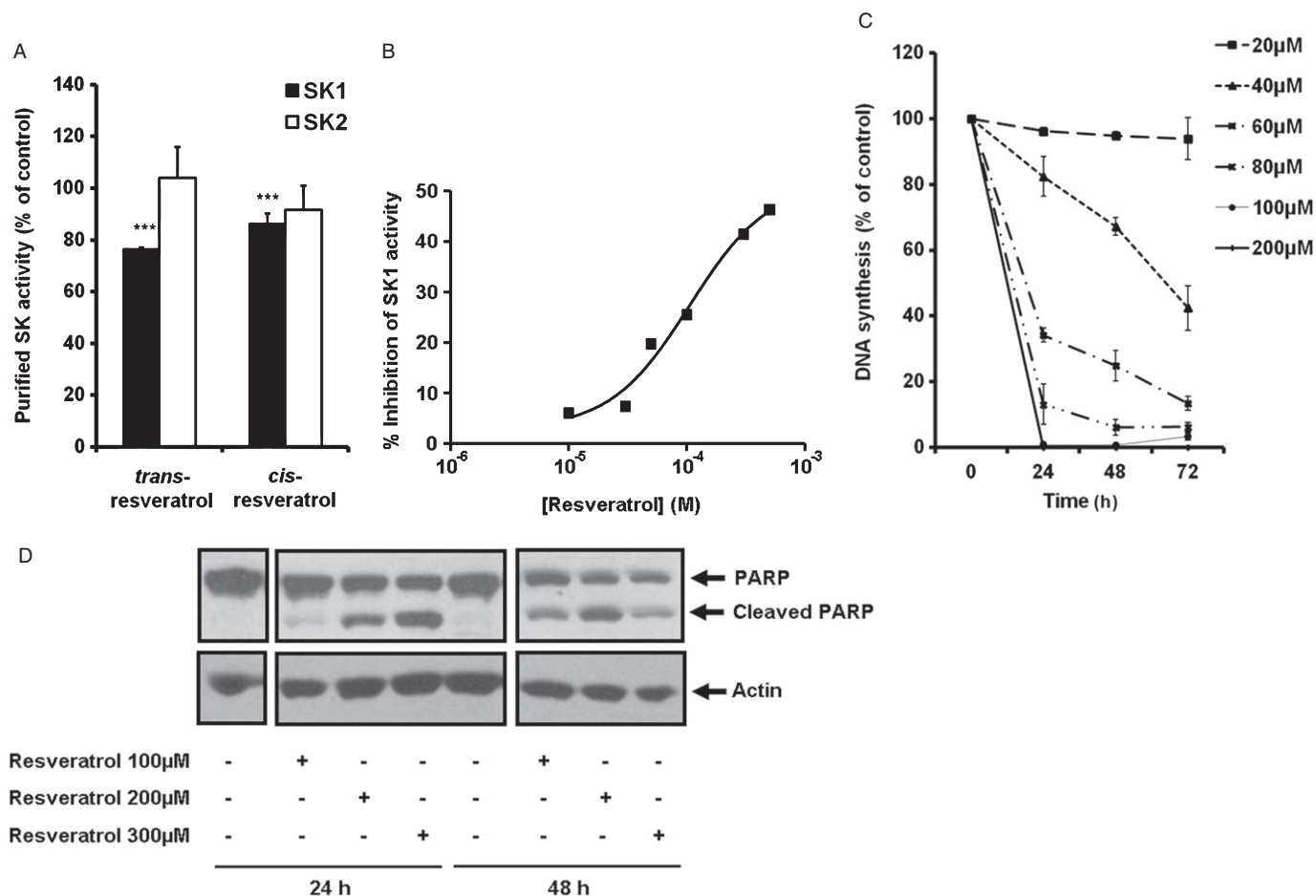
induced ubiquitin-proteasomal degradation pathway (Loveridge *et al.*, 2010; Lim *et al.*, 2011). This effect was reproduced here, along with the finding that SKi also reduced EGF-stimulated ERK-1/2 phosphorylation (Figure 2C). Similarly, the treatment of MCF-7 Neo cells with fraction 5 reduced the expression of SK1 and inhibited EGF-stimulated ERK-1/2 activation (Figure 2C). This was also observed using parental MCF-7 cells (data not shown). Interestingly, other fractions exhibited different effects. For instance, Fraction 4 and 5B inhibited SK1 activity (Figure 2B), but did not induce down-regulation of SK1 expression (Figure 2C). Fraction 9 reduced ERK1/2 activation but did not affect SK1 expression, whereas Fraction 10 reduced SK1 activity (Figure 2B) and expression, but had less effect on ERK1/2 activation (Figure 2C). The total plant extract also reduced SK1 expression, used as a positive control (Figure 2C). These data show that Fraction 5 exhibits properties typical of an SK1 inhibitor and this fraction was therefore subjected to NMR and MS analysis. These analyses demonstrated that Fraction 5 contained one major compound, ampelopsin A which is a resveratrol dimer (Supporting Information Figure S1, Table S1). The structure of

ampelopsin A was consistent with that isolated from *H. parviflora* (Tanaka *et al.*, 2000).

The aromatic region ( $\delta$  6–8) of <sup>1</sup>H-NMR of the bark extract of *H. dryobalanoides* indicated the presence of phenolic compounds (data not shown). Upon fractionation of the bark extract and subsequent purification using Sephadex LH-20, balanocarpol was isolated. Structural elucidation with NMR revealed that balanocarpol is an isomer (7a-epimer) of ampelopsin A (Supporting Information Figure S1, Table S1). The structure of balanocarpol is consistent with that isolated from *H. parviflora* (Tanaka *et al.*, 2000). The molecular weights of ampelopsin A and balanocarpol were established by MS; both compounds have the same molecular formula of C<sub>28</sub>H<sub>22</sub>O<sub>7</sub>.

### Resveratrol is an SK1 inhibitor

Because the isomers ampelopsin A and balanocarpol are resveratrol dimers formed by the fusion of *cis*- and *trans*-isomers of resveratrol (Supporting Information Figure S2), we investigated whether resveratrol itself is an SK1 inhibitor. In this



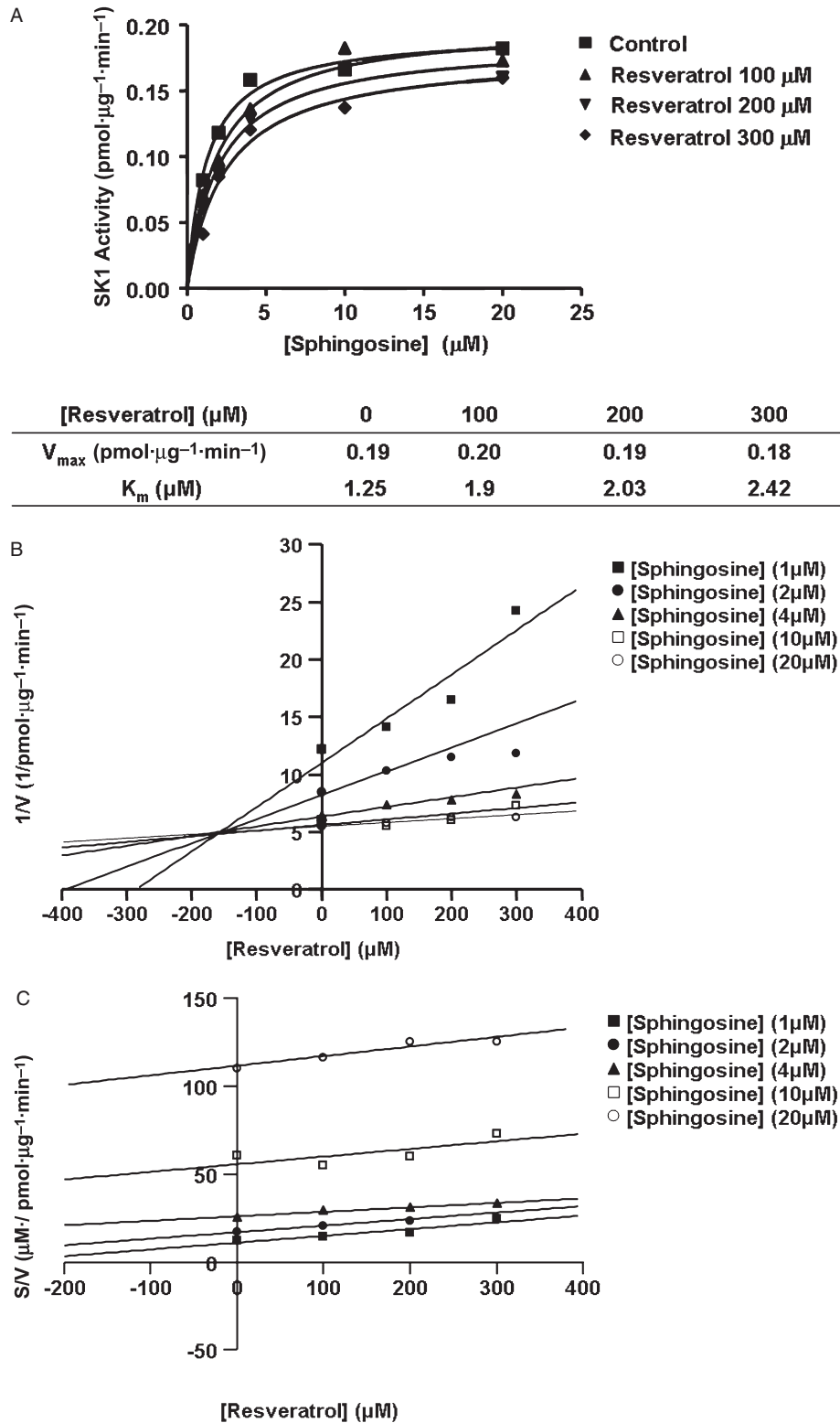
**Figure 3**

Biological activities of resveratrol. (A) Bar graph showing the effect of *trans*- and *cis*-resveratrol (both at 500 μM) on purified SK1 or SK2 activity. SK1/2 was assayed using 10 μM sphingosine and 250 μM [<sup>32</sup>P]-ATP as the substrates. Data are expressed as percentage of control and represent mean and SDs of six determinations, \*\*\**P* < 0.001, significantly different from control. (B) Graph showing the concentration-dependent inhibitory effect of *trans*-resveratrol on SK1 activity in lysates from SK1 stably over-expressed HEK 293 cells. (C) Graph showing the effect of *trans*-resveratrol on [<sup>3</sup>H]-thymidine incorporation in MCF-7 cells. MCF-7 cells were treated with *trans*-resveratrol at indicated concentrations or control (0.05% v/v DMSO final concentration) for 24–72 h and then with 0.5 μCi·mL<sup>-1</sup> of [<sup>3</sup>H]-thymidine added for 5 h. Data are expressed as percentage of control and represent means and SDs of triplicate determinations. (D) Western blot showing the effect of *trans*-resveratrol on PARP cleavage. MCF-7 cells were treated with *trans*-resveratrol (100, 200, 300 μM) or control (0.3% v/v DMSO final concentration) for 24 and 48 h. Cell lysates were prepared and analysed by Western blotting using anti-PARP antibody and then stripped and reprobed with anti-actin antibody to ensure comparable protein loading. The images shown were taken from the same Western blot. Results are representative of three independent experiments.

regard, both *cis*- and *trans*-resveratrol inhibited purified SK1 but not purified SK2 activity (Figure 3A). *Trans*-resveratrol inhibited purified SK1 activity and SK1 stably over-expressed in HEK 293 cells to a similar extent, for example ~30–40% (Figure 3A and B). Calculation of the % inhibition of SK1 activity using the Michaelis–Menten equation, 10 μM sphingosine, the kinetic constants determined using lysates from HEK 293 cells stably over-expressing SK1 (see later) and 500 μM resveratrol yielded a 25% inhibition of SK1 activity, which is also close to the level of inhibition determined experimentally for the purified enzyme (~30%, Figure 3A).

*Trans*-resveratrol inhibited SK1 activity in a concentration-dependent manner in lysates from cells stably over-expressing SK1 (Figure 3B). The treatment of MCF-7 cells with *trans*-resveratrol also reduced [<sup>3</sup>H] thymidine incorpora-

tion into DNA in a time and concentration-dependent manner (Figure 3C) and induced PARP cleavage in MCF-7 cells (Figure 3D). Kinetic analysis of lysates from HEK 293 cells stably over-expressing SK1 demonstrated that SK1 obeyed Michaelis–Menten kinetics and that *trans*-resveratrol (thereafter referred to as resveratrol) was a competitive inhibitor (with sphingosine) of SK1 with a  $K_{ic} = 160 \pm 40 \mu\text{M}$  (Figure 4A–C). When ATP was varied and the sphingosine concentration kept constant (20 μM), resveratrol reduced the  $V_{max}$  without affecting the  $K_m$  for ATP indicating that it is non-competitive with ATP (control,  $K_m = 42.3 \mu\text{M}$ ,  $V_{max} = 0.25 \text{ pmol} \cdot \mu\text{g}^{-1} \cdot \text{min}^{-1}$ ; 100 μM resveratrol,  $K_m = 36.1 \mu\text{M}$ ,  $V_{max} = 0.21 \text{ pmol} \cdot \mu\text{g}^{-1} \cdot \text{min}^{-1}$ ; 300 μM resveratrol,  $K_m = 41 \mu\text{M}$ ,  $V_{max} = 0.18 \text{ pmol} \cdot \mu\text{g}^{-1} \cdot \text{min}^{-1}$ ; 500 μM resveratrol,  $K_m = 34.5 \mu\text{M}$ ,  $V_{max} = 0.15 \text{ pmol} \cdot \mu\text{g}^{-1} \cdot \text{min}^{-1}$ ,  $n = 4$ ).



**Figure 4**

Inhibitor kinetic analysis of resveratrol for stably over-expressed recombinant SK1 in lysates from HEK 293 cells. (A) Non-linear regression analysis. (B) Dixon plot. (C)  $S/V$  versus resveratrol concentration plot. Results are representative of three independent experiments.

### Kinetic and functional studies of balanocarpol

In contrast with resveratrol, balanocarpol is a mixed competitive inhibitor (with sphingosine) of SK1 with a  $K_{ic} = 90 \pm 10 \mu\text{M}$  and a  $K_{iu} \sim 500 \mu\text{M}$  (Figure 5A–C). These findings demonstrate that balanocarpol is more potent (approximately twofold) compared with resveratrol in inhibiting SK1 activity. When ATP was varied and the sphingosine concentration kept constant ( $20 \mu\text{M}$ ), balanocarpol reduced the  $V_{\text{max}}$  without affecting the  $K_m$  for ATP indicating that it is non-competitive with ATP (control,  $K_m = 36.6 \mu\text{M}$ ,  $V_{\text{max}} = 0.24 \text{ pmol}\cdot\mu\text{g}^{-1}\cdot\text{min}^{-1}$ ;  $300 \mu\text{M}$  balanocarpol,  $K_m = 31.1 \mu\text{M}$ ,  $V_{\text{max}} = 0.18 \text{ pmol}\cdot\mu\text{g}^{-1}\cdot\text{min}^{-1}$ ,  $n = 4$ ). Moreover, balanocarpol is more potent in inhibiting [ $^3\text{H}$ ] thymidine incorporation into DNA in MCF-7 cells; the  $\text{IC}_{50}$  for inhibition of DNA synthesis is  $\sim 50 \mu\text{M}$  for resveratrol (Figure 3C) and  $\sim 10 \mu\text{M}$  for balanocarpol (Figure 6A). In common with resveratrol and SKi, balanocarpol also induced PARP cleavage indicating that it is an inducer of apoptosis (Figure 6B). Resveratrol and balanocarpol also exhibited properties typical of SK1 inhibitors by inducing the down-regulation of SK1 expression in MCF-7 Neo cells (Figure 6C).

## Discussion

The major finding of our study is that resveratrol and its dimers, ampelopsin A and balanocarpol are SK1 inhibitors. This conclusion is based on the finding that these compounds inhibited SK1 catalytic activity and induced the down-regulation of SK1 expression, properties similar to that observed for the known inhibitor, SKi. Moreover, these compounds also inhibit proliferation and induce apoptosis of MCF-7 breast cancer cells. Indeed, the effect of these compounds on apoptosis can be reproduced by siRNA knock-down of SK1 in MCF-7 breast cancer cells; this involving caspase activation associated with increased ceramide formation and Bax oligomerization (Taha *et al.*, 2006). In addition, our findings concerning the effect of resveratrol, ampelopsin A and balanocarpol on SK1 expression are supported by studies that demonstrate that dietary polyphenols including resveratrol induce down-regulation of SK1 activity and expression in prostate cancer cells (Brizuela *et al.*, 2010). The mechanism of down-regulation of SK1 expression might involve changes in SK1 protein turnover, as demonstrated for other SK1 inhibitors that induce the ubiquitin-proteasomal degradation of SK1 (Tonelli *et al.*, 2010; Loveridge *et al.*, 2010) or lysosomal-cathepsin B catalysed proteolysis (Ren *et al.*, 2010) or changes in gene promoter activity. Further studies are required to establish the precise mechanism by which resveratrol and balanocarpol reduce the expression of SK1.

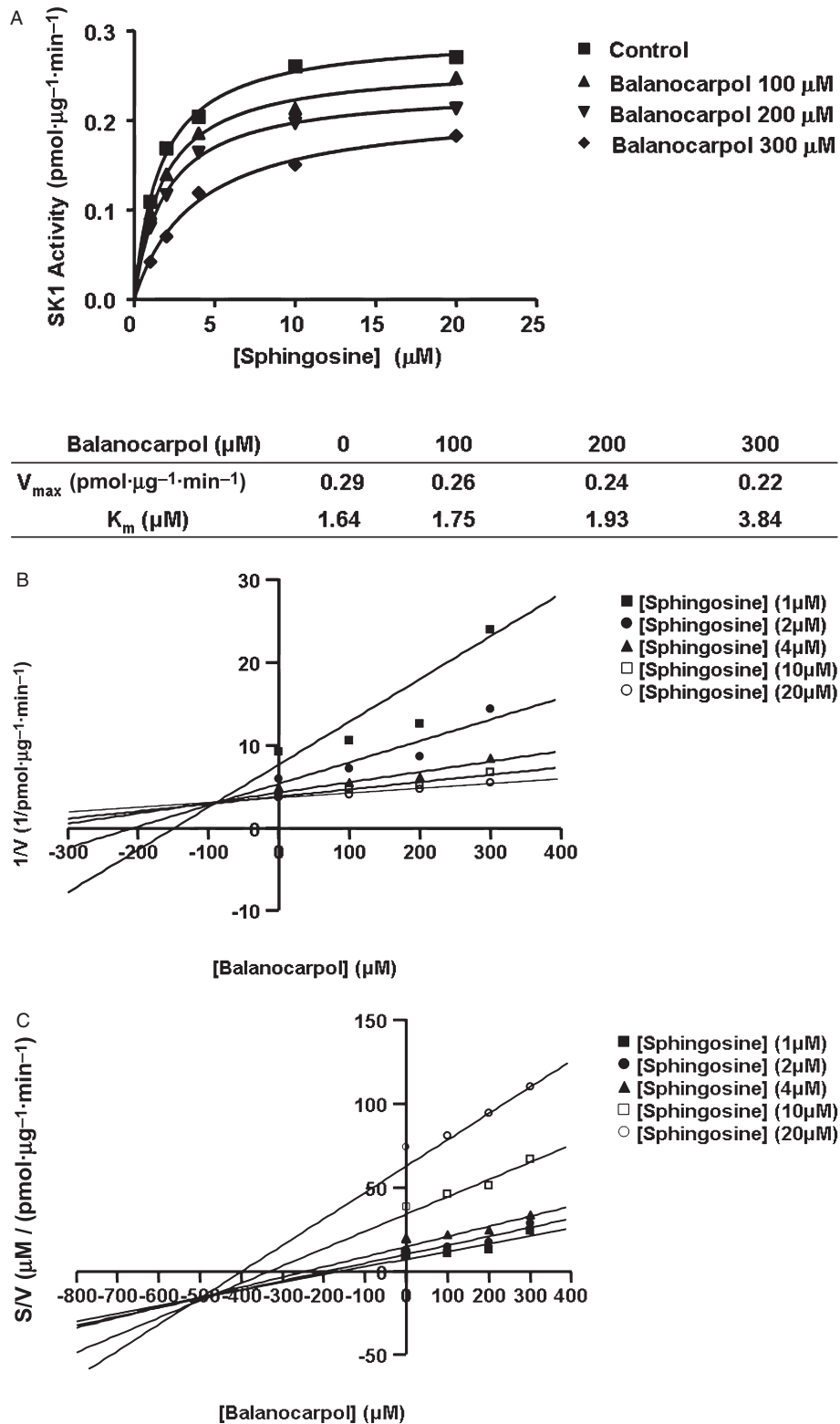
Previous studies using red grape skin polyphenolic extracts (SGE) to study the 'French paradox', linked the anti-angiogenic effects of resveratrol (which is present in SGE) to effects on S1P and VEGF signalling (Barthomeuf *et al.*, 2006). Thus, pretreatment of endothelial cells with SGE reduced S1P-induced migration, ERK1/2 activation and platelet activating factor synthesis (Barthomeuf *et al.*, 2006). However, SGE contains other polyphenolic compounds including sig-

nificant quantities of resveratrol oligomers. Based on the finding that balanocarpol is a more potent inhibitor of SK1 compared with resveratrol, the attribution of the anti-angiogenic effect of SGE to resveratrol alone should be exercised with caution. Our findings also raise the question of whether the anti-cancer effect of resveratrol might be due to direct inhibition of SK1 and/or down-regulation of SK1 expression leading to alterations in sphingolipid metabolism and signalling. Indeed, perturbation of sphingolipids has been demonstrated before by resveratrol, which induces an increase in the synthesis of ceramide and which, in turn, stimulates the apoptosis of MDA-MB-231 cells (Scarlatti *et al.*, 2003). Our findings also suggest that some of the effects of resveratrol on the inflammatory response to C5a, previously described by Issuree *et al.* (2009), and on prostate cancer survival (Brizuela *et al.*, 2010) might be a consequence of the direct inhibition of SK1 activity and down-regulation of SK1 expression.

In the current study, we also provide a novel insight into the enzymatic mechanism by which resveratrol inhibits SK1 activity. Kinetic analysis established that resveratrol is a competitive inhibitor (with sphingosine) of SK1. The  $K_{ic}$  for inhibition of SK1 activity is  $160 \pm 40 \mu\text{M}$ . Although this represents a relatively low potency, it is consistent with the concentration of resveratrol required for induction of apoptosis of cancer cells (Nakagawa *et al.*, 2001; Pozo-Guisado *et al.*, 2002; Scarlatti *et al.*, 2003). The maximum serum concentration of resveratrol is  $2 \mu\text{M}$  using relevant dietary concentrations in humans (Walle *et al.*, 2004). Although this concentration is considerably lower than the  $K_i$  for inhibition of SK1 activity, it is possible that the intracellular concentration obtained from a serum concentration of  $2 \mu\text{M}$  is considerably higher. Low serum concentrations of resveratrol might also be increased by higher dose administration. Interestingly, resveratrol is also an oestrogen receptor antagonist and can therefore reduce oestrogen-induced MCF-7 cell growth (Lu and Serrero, 1999). In this regard, SKi is also an oestrogen receptor antagonist and exhibits structural similarity with resveratrol (Antoon *et al.*, 2011). Therefore, both resveratrol and SKi could be usefully exploited to launch a two-pronged attack on cancer.

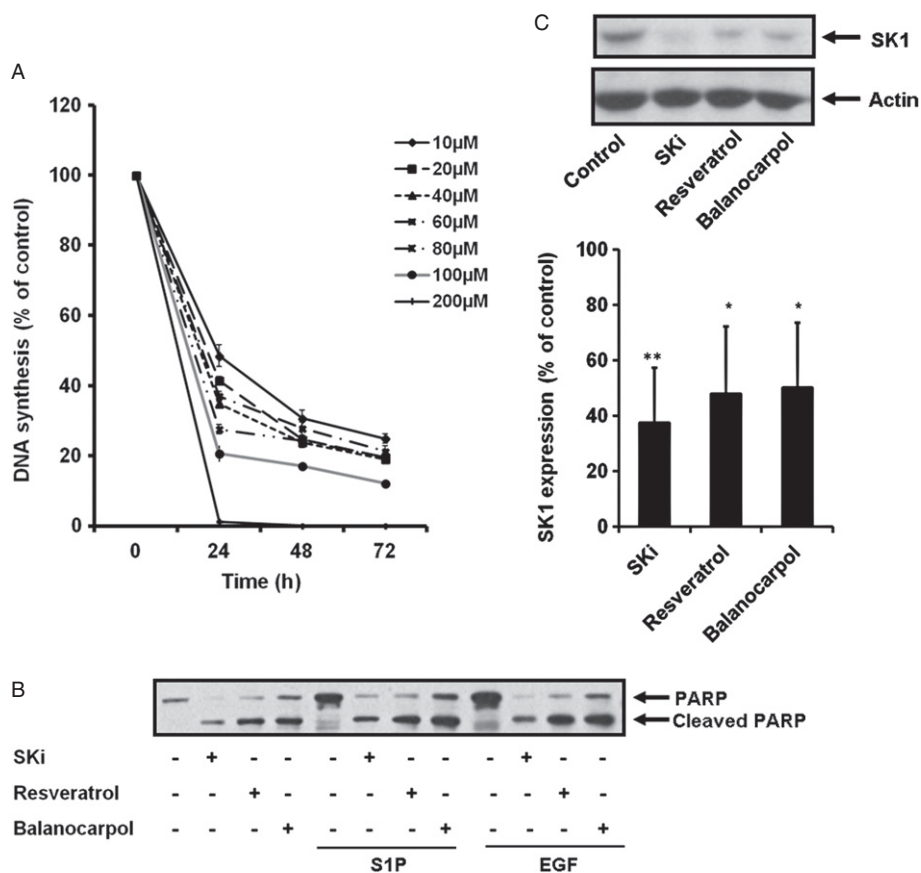
High concentrations (usually  $>50 \mu\text{M}$ ) of resveratrol induce apoptosis whereas low concentrations ( $10\text{--}40 \mu\text{M}$ ) suppress cancer cell growth (Nakagawa *et al.*, 2001; Pozo-Guisado *et al.*, 2002; Scarlatti *et al.*, 2003). This is in line with our findings, where apoptosis was induced by high concentrations of resveratrol ( $100\text{--}300 \mu\text{M}$ ) and this was correlated with inhibition/down-regulation of SK1 expression. In contrast, the  $\text{IC}_{50}$  for inhibition of MCF-7 cell growth is  $\sim 50 \mu\text{M}$  and might therefore, be SK1-independent although, as the intracellular concentration of inhibitor attained is unknown, the inhibition of growth by targeting SK1 cannot be entirely excluded. This is relevant if, for instance, different levels of SK1 inhibition are required to observe effects on growth, as distinct from apoptosis. Alternatively, the mechanism for growth inhibition might be via a different target where reported  $\text{IC}_{50}$  values are closer to those determined here, such as COX-1 ( $15 \mu\text{M}$ ), COX-2 ( $32.2 \mu\text{M}$ ), antioxidant/free radical scavenging ( $27 \mu\text{M}$ ) and oestrogen receptors ( $5 \mu\text{M}$ ) (Jang *et al.*, 1997; Subbaramaiah *et al.*, 1998; Lu and Serrero, 1999; Frémont, 2000).





**Figure 5**

Inhibitor kinetic analysis of balanocarpol for stably over-expressed recombinant SK1 in lysates from HEK 293 cells. (A) Non-linear regression analysis. (B) Dixon plot. (C)  $S/V$  versus balanocarpol concentration plot. Results are representative of three independent experiments.



**Figure 6**

Biological activities of balanocarpol. (A) Graphs showing the effect of balanocarpol on [<sup>3</sup>H]-thymidine incorporation in MCF-7 cells. MCF-7 cells were treated with balanocarpol at indicated concentrations or control (0.05% v/v DMSO final concentration) for 24–72 h and then with 0.5 μCi·mL<sup>-1</sup> of [<sup>3</sup>H]-thymidine added for 5 h. Data are expressed as percentage of control and represent means and SDs of triplicate determinations. (B) Western blots showing the effect of balanocarpol, SKi or resveratrol on PARP cleavage. Quiescent MCF-7 Neo cells were incubated with control (0.1% v/v DMSO final concentration) or indicated inhibitors (10 μM SKi, 200 μM resveratrol or 200 μM balanocarpol for 48 h before being stimulated with 25 ng·mL<sup>-1</sup> EGF or 1 μM S1P for 5 min. Cell lysates were separated by SDS-PAGE and immunoblotted with anti-PARP antibody. Results are representative of three independent experiments. (C) Western blots showing the effect of balanocarpol, SKi or resveratrol on SK1 expression. Quiescent MCF-7 Neo cells were incubated with control (0.1% v/v DMSO final concentration), 10 μM SKi, 200 μM resveratrol or 200 μM balanocarpol for 48 h. Cells were harvested and analysed by Western blotting using anti-SK1 antibody. Blots were then stripped and reprobbed with anti-actin antibody to ensure comparable protein loading. The bar graph represents densitometric quantification of the effects of inhibitors on SK1 expression (SK1: actin ratio). Data are expressed as a percentage of control (\**P* < 0.05, \*\**P* < 0.01, significantly different from control, *n* = 3). Results are representative of three independent experiments.

The fact that secondary metabolites are produced via the biogenesis of resveratrol monomers raises the question whether oligomerization produces compounds which exhibit better potency as anti-cancer agents. Biosynthesis of resveratrol oligomers might therefore be regarded as a diversity-oriented approach. This is supported by the finding that balanocarpol is about twice as potent as resveratrol with respect to inhibition of SK1 activity. Therefore, it is possible that a single molecule of balanocarpol might bind at least two SK1 catalytic molecules to inhibit the activity of each simultaneously. Indeed, we have previously demonstrated that SK1 is a minimal dimer in which the catalytic sites are non-cooperative and therefore function independently of each other (Lim *et al.*, 2011). Therefore, each catalytic site in the SK1 dimer might bind balanocarpol simultaneously, while resveratrol might bind to and inhibit only one catalytic site in

the SK1 dimer. Indeed, this possible generic mechanism of action, where potency is augmented by dimerization of resveratrol is supported by the finding that the resveratrol tetramer, dibalanocarpol is more potent than resveratrol and its dimers in inhibiting the proliferation of P-388 cells (Sahidin *et al.*, 2005). The proposed action of balanocarpol as an SK1 inhibitor bears some similarity with rapamycin, which inhibits two different proteins (FKBP-12 and mTOR) in a complex concomitantly (Choi *et al.*, 1996).

In summary, we have provided the first evidence that resveratrol and balanocarpol directly inhibit SK1 activity and induce down-regulation of the enzyme in cancer cells. We also speculate that oligomerization of resveratrol might provide a means to improving inhibitor potency against SK1. The study of resveratrol oligomers has been hampered by difficulty in isolation from natural sources and intractable

chemical synthesis. Nevertheless, a recent study reported a programmable, controlled and potentially scalable synthesis of the resveratrol family via a three-stage design (Snyder *et al.*, 2011). It is anticipated that this will accelerate the development of novel therapeutics and translation to the clinic.

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## Conflict of interest

The authors declare that there are no conflicts of interest.

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## Supporting information

Additional Supporting Information may be found in the online version of this article:

**Figure S1** Molecular structures of ampelopsin A and balanocarpol.

**Figure S2** Proposed biosynthetic route of ampelopsin A or balanocarpol. Adapted from Sotheeswaran *et al.* (1993).

**Table S1** Chemical shifts of ampelopsin A and balanocarpol

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