

RESEARCH PAPER

Mechanisms underlying the metabolic actions of galegine that contribute to weight loss in mice

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Background and purpose: Galegine and guanidine, originally isolated from *Galega officinalis*, led to the development of the biguanides. The weight-reducing effects of galegine have not previously been studied and the present investigation was undertaken to determine its mechanism(s) of action.

Experimental approach: Body weight and food intake were examined in mice. Glucose uptake and acetyl-CoA carboxylase activity were studied in 3T3-L1 adipocytes and L6 myotubes and AMP activated protein kinase (AMPK) activity was examined in cell lines. The gene expression of some enzymes involved in fat metabolism was examined in 3T3-L1 adipocytes.

Key results: Galegine administered in the diet reduced body weight in mice. Pair-feeding indicated that at least part of this effect was independent of reduced food intake. In 3T3-L1 adipocytes and L6 myotubes, galegine (50 µM–3 mM) stimulated glucose uptake. Galegine (1–300 µM) also reduced isoprenaline-mediated lipolysis in 3T3-L1 adipocytes and inhibited acetyl-CoA carboxylase activity in 3T3-L1 adipocytes and L6 myotubes. Galegine (500 µM) down-regulated genes concerned with fatty acid synthesis, including fatty acid synthase and its upstream regulator SREBP. Galegine (10 µM and above) produced a concentration-dependent activation of AMP activated protein kinase (AMPK) in H4IIE rat hepatoma, HEK293 human kidney cells, 3T3-L1 adipocytes and L6 myotubes.

Conclusions and implications: Activation of AMPK can explain many of the effects of galegine, including enhanced glucose uptake and inhibition of acetyl-CoA carboxylase. Inhibition of acetyl-CoA carboxylase both inhibits fatty acid synthesis and stimulates fatty acid oxidation, and this may contribute to the *in vivo* effect of galegine on body weight.

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Keywords: galegine; 3T3-L1 adipocytes; L6 myotubes; glucose uptake; acetyl CoA carboxylase; AMPK

Abbreviations: ACC, acetyl-CoA carboxylase; AICAR, aminoimidazole-4-carboxamide ribonucleotide; AMPK, AMP-activated protein kinase; DMEM, Dulbecco's modified Eagle's media; FASN, fatty acid synthase; FBS, fetal bovine serum

Introduction

Previous studies from this laboratory demonstrated the potential anti-obesity effect of *Galega officinalis*, a plant used in traditional medicine for the treatment of diabetes. Although accompanied by a transient decrease in food intake, pair-feeding studies indicated that the effects on body weight were independent of changes in food intake (Palit *et al.*, 1999). Indeed, food intake in treated mice eventually exceeded that seen in control animals. The main active ingredients of *G. officinalis* are galegine and guanidine, which gave rise to the biguanides (Bailey and Day, 1989). Water-ethanol extracts of *G. officinalis*, which contained significant quantities of galegine, also reduced body weight

and diminished body weight gain in normal mice (unpublished observations). Preliminary studies showed that galegine reduced body weight gain in normal mice and reduced blood glucose and food intake, suggesting that at least part of the effect of *G. officinalis* on body weight was mediated via galegine. The aim of the present work was to investigate the potential mechanisms of action of galegine, as these have not previously been studied in detail, although it was shown to have some insulin-like actions in stimulating glucose uptake in rat diaphragm (Weitzel *et al.*, 1971) and in exerting anti-lipolytic effects in adipocytes (Weitzel *et al.*, 1972).

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Materials and methods

Animals

All animal studies were carried out in accordance with the UK Animals (Scientific Procedures) Act 1986. Adult male

BALB/c mice were obtained from stock at the Biological Procedures Unit within the University of Strathclyde. All animals were housed individually in an air-conditioned environment maintained at $21 \pm 2^\circ\text{C}$ with a 12 h light/12 h dark cycle. Animals were allowed continuous access to tap water and unless indicated otherwise were fed *ad libitum* on standard pellet diet (SDS, Cambridge, UK).

Feeding studies

Before initiation of feeding studies, mice were habituated to individual housing, and individual animal food intake and body weight was monitored daily at 09.30 hours. At the start of the experimental period (day 0), mice were randomly divided into groups ($n=6-8$) and food removed and substituted with standard diet pellets containing indicated concentrations of galegine. Daily food intake and body weight measurements were recorded for the duration of the study period (up to 11 days), and blood glucose determinations were made at the end of the study. In view of the reduction in food intake observed in galegine-treated mice, in some studies, control mice were pair-fed to galegine-treated animals, so that food intake of the two groups did not differ across a 5–11 days period.

Cells and cell culture. 3T3-L1 fibroblasts (CCL-92.1) and L6 muscle cells (CRL-1458) were obtained from the American Type Culture Collection and cultured in Dulbecco's modified Eagle's media (DMEM) containing 10% newborn calf serum and 1% (v/v) antibiotics (100 U ml^{-1} penicillin, 0.1 mg ml^{-1} streptomycin) at 37°C in a humidified atmosphere of 5% CO_2 and 95% air. Confluent 3T3-L1 fibroblasts were induced to differentiate in multi-well plates by culturing for 3 days in 10% fetal bovine serum (FBS)/DMEM containing insulin ($1\text{ }\mu\text{g ml}^{-1}$), 0.5 mM isobutylmethyl xanthine and $0.25\text{ }\mu\text{M}$ dexamethasone. Cells were then cultured in 10% FBS/DMEM containing insulin ($1\text{ }\mu\text{g ml}^{-1}$) for a further 3 days. Subsequently, cells were cultured in 10% FBS/DMEM, which was replaced every 2 days until cells were fully differentiated (8–10 days after the start of the differentiation procedure).

L6 muscle cells were grown as monolayers in multi-well plates, and after reaching confluency were cultured in 1% newborn calf serum/DMEM to limit cell growth and aid the formation of multinucleated myotubes. Media was replaced every 2 days, and cells were used 4–6 days after reduction in the FBS media content.

Cell viability

The viability of 3T3-L1 adipocyte and L6 myotube cells following treatment during various incubations with galegine was determined using the alamarBlue assay technique (Nociari *et al.*, 1998).

Glucose uptake

Glucose uptake studies were performed on fully differentiated 3T3-L1 adipocytes and L6 myotubes by measuring the transport of 2-deoxy- ^3H D-glucose (final concentration

$50\text{ }\mu\text{M}$, $0.5\text{ }\mu\text{Ci}$). (Hayes *et al.*, 1993), following incubation at 37°C with increasing concentrations of galegine for 5 h.

Lipolysis

Lipolysis in fully differentiated 3T3-L1 cells incubated at 37°C for 4 h, with or without $10\text{ }\mu\text{M}$ isoprenaline, was determined by measurement of glycerol release (Wieland, 1974).

Acetyl-CoA carboxylase enzyme activity

The acetyl-CoA carboxylase (ACC) activity within 3T3-L1 adipocytes and L6 myotubes was assayed using the method as described by Inoue and Lowenstein (1975).

Determination of activity of lipogenic enzymes

Epididymal fat deposits were excised from young adult male Sprague–Dawley rats (220–260 g) and placed on ice. Tissue was then homogenized with 1.5 volumes homogenizing buffer (70 mM KHCO_3 , $85\text{ mM K}_2\text{HPO}_4$, $9\text{ mM KH}_2\text{PO}_4$, 1 mM dithiothreitol, pH 8.0) by a Janke and Turrax T-25 homogenizer for 20 s. The homogenate was centrifuged at $20\text{ }000\text{ g}$ for 10 min and supernatant removed and centrifuged again at $105\text{ }000\text{ g}$ for 60 min. The resulting enzyme extract supernatant was stored at -20°C and used in previously described continuous spectrophotometric assays (based on the oxidation or formation of NADPH at 37°C) to assess the activity of fatty acid synthase (FASN) (Nepokroeff *et al.*, 1975), ATP-citrate lyase (Takeda *et al.*, 1969) and malic enzyme (Hsu and Lardy, 1969). Activities of enzymes were measured following incubation of enzyme extracts with concentrations of galegine of up to $500\text{ }\mu\text{M}$.

AMP-activated protein kinase activity. AMP-activated protein kinase (AMPK) activity assays were performed on lysates of H4IIE hepatocyte cells, HEK293 cells, 3T3-L1 adipocytes and L6 myotubes cultured in 10 cm well plates with 10% FBS/DMEM supplemented with various concentrations of galegine for 1 h by measurement of phosphorylation of a synthetic peptide substrate, SAMS (HMRSAMSGHLHLVRR) (Davies *et al.*, 1989). AMPK activity was expressed as nmol of phosphate incorporated into the peptide substrate per min per mg of lysate ($\text{nmol min}^{-1}\text{ mg}^{-1}$).

Gene expression and reverse transcription-PCR

Fully differentiated 3T3-L1 adipocytes were incubated with $500\text{ }\mu\text{M}$ galegine for 24 h at 37°C . At the end of the incubation period, media was aspirated and cells washed twice with phosphate-buffered saline at 37°C .

Total RNA was extracted from the cells using a RNeasy kit (Qiagen, Crawley, UK) with an on-column DNase digestion step to remove genomic DNA. The integrity and concentration of extracted RNA was determined using a BioAnalyzer (Agilent, Wokingham, UK) and then subjected to microarray analysis using Mouse Expression 430 GeneChips (A and B chips, Affymetrix, High Wycombe, UK) at the Sir Henry Wellcome Functional Genomics Facility (SHWFGF, University of Glasgow).

Quantitative real-time PCR (QPCR) analysis was undertaken to validate data obtained from the microarray studies. Several key genes from these processes were selected for QPCR analysis. Where possible, the QPCR primers for these genes were designed to recognize the same region of the target transcripts to which the Affymetrix GeneChip probes were hybridized. The expression of two housekeeping genes, β -2-microglobulin (B2M), and hypoxanthine guanine phosphoribosyl transferase, were examined for normalization purposes. All necessary reactions were carried out in single runs on a 96-well Chromo4 QPCR thermocycler (GRI, Essex, UK). To confirm their identity, QPCR reactions were run on 2% agarose gels, the amplicons gel purified (GFX PCR DNA and gel band purification kit, Amersham Biosciences, Little Chalfont, UK), and their sequence confirmed by automated DNA sequencing.

Data analysis

All data shown are the mean \pm standard error of the mean (mean \pm s.e.m.). The differences in food intake and body weight between control and galegine-treated groups were analysed by two-way analysis of variance for repeated measures. For *in vitro* assays, statistical significance was tested with one-way analysis of variance with Dunnett's *post hoc* test. Mean fold changes in gene transcript expression in 3T3-L1 adipocytes treated *in vitro* with galegine were determined using SHWFGF in-house FunAllyse v1.2 Microarray Data software. Only genes exhibiting at least a two-fold up- or downregulation following treatment were entered into the Gene Ontology Tree Machine web server (Zhang *et al.*, 2004). This was used to identify gene ontology categories with significantly enriched gene numbers ($P < 0.001$) in the galegine-treated cell transcript sets. The threshold cycle values (C_t) of data produced via QPCR analysis were determined using Opticon Monitor v2.0 software provided with the Chromo4 QPCR Thermocycler. These values were then entered into a Microsoft Excel QPCR analysis macro, REST (relative expression software tool; Pfaffl *et al.*, 2002). This enabled group-wise comparison and statistical analysis of relative expression results from the QPCR of control and galegine-treated adipocyte cDNA, and normalization using housekeeping gene data.

Reagents

Synthesis and purification of galegine. Galegine is not commercially available and was synthesized by reaction of benzyl amine with 2-methylpseudourea sulphate. The white solid formed after re-crystallization was identified as galegine using ^1H and ^{13}C nuclear magnetic resonance spectroscopy, infrared spectroscopy and elemental analysis techniques, and had a purity greater than 99%.

Other reagents. Insulin, isobutylmethyl xanthine, dexamethasone, wortmannin and isoprenaline were sourced from Sigma (Poole, Dorset, UK). DMEM tissue culture medium, FBS, newborn calf serum, penicillin and streptomycin were all purchased from Gibco (Paisley, Strathclyde, Scotland). 2-Deoxy- ^3H D-glucose, $\text{NaH}^{14}\text{O}_3$ and $[\gamma\text{-}^{32}\text{P}]$ ATP were obtained from Amersham Pharmacia Biotech (Buckinghamshire, UK). All water used in these experiments

was purified using a Milli-Q Water Purification System (Millipore, Milford, MA, USA). All other chemicals used were of the highest available purity.

Results

Feeding studies

Galegine ($3.41 \text{ mmol kg}^{-1}$ feed that is 600 mg kg^{-1} feed; approximate daily dose of $0.5 \text{ mmol galegine per kg body weight}$) produced a decrease in the body weight and food intake of mice relative to controls (Figure 1). Blood glucose was also reduced at the end of the experiment ($3.2 \pm 0.4 \text{ mmol l}^{-1}$ compared with control value of $6.0 \pm 0.5 \text{ mmol l}^{-1}$; $n = 5$; $P < 0.001$). In pair-fed mice, where the food intake was not significantly different from controls, galegine ($3.41 \text{ mmol kg}^{-1}$ feed) also produced a significant reduction in body weight, which was sustained across 11 days (Figure 2). In this study, however, the fall in blood glucose did not reach statistical significance ($4.7 \pm 0.9 \text{ mmol l}^{-1}$ compared with control of $6.0 \pm 1.1 \text{ mmol l}^{-1}$; $n = 7$). Galegine in this dose produced no toxicity when administered over 28 days.

Effect of galegine on *in vitro* glucose uptake by 3T3-L1 adipocytes and L6 myotubes

Pre-treatment with galegine ($10 \mu\text{M}$ – 3 mM ; 5 h) produced a concentration-dependent stimulation of insulin-independent

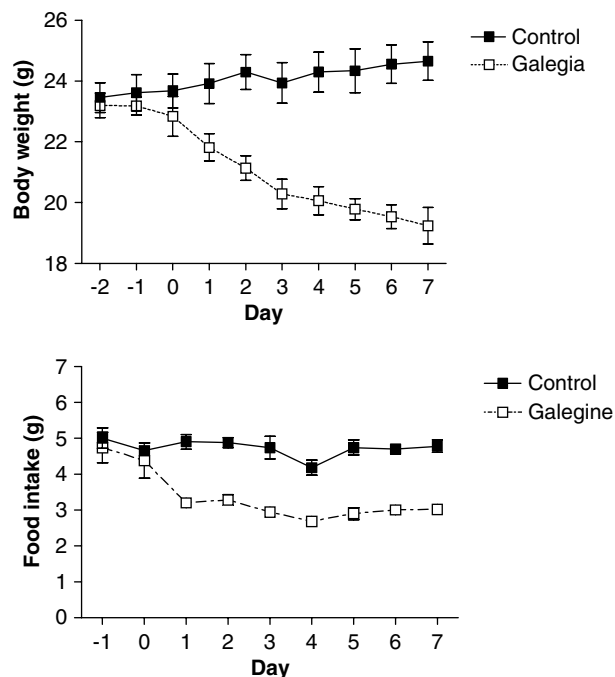


Figure 1 Effect of galegine incorporation ($3.41 \text{ mmol kg}^{-1}$ feed) into the diet of male BALB/c mice on body weight (upper panel) and food intake (lower panel) after 7 days treatment. Values shown are mean \pm s.e.m. ($n = 6$ – 8 per group). Two-way analysis of variance showed a significant difference between galegine-treated and control mice for both body weight ($P < 0.0001$) and food intake ($P < 0.0001$). In each case, there was a significant effect of 'time' ($P < 0.005$), but no significant interaction between time and treatment.

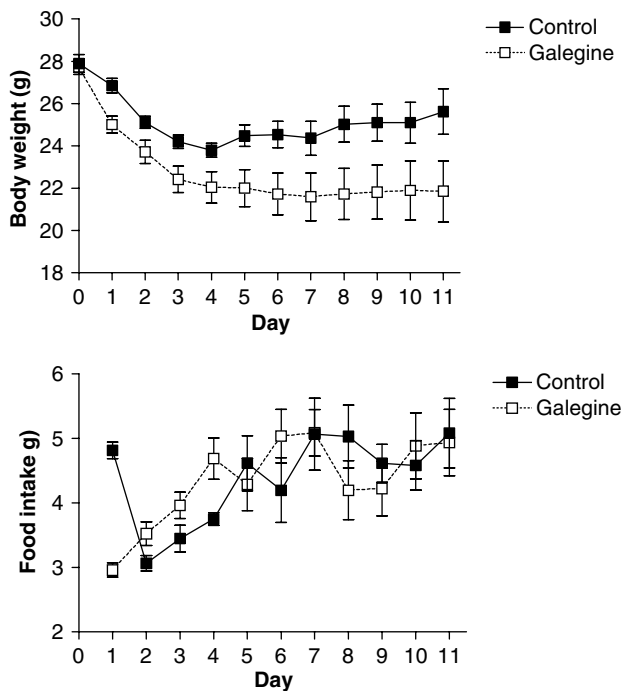


Figure 2 Body weight (upper panel) and food intake (lower panel) in mice receiving galegine ($3.41 \text{ mmol kg}^{-1}$ feed) compared with that in pair-fed controls. Values shown are mean \pm s.e.m. ($n=8$). Two-way analysis of variance showed a significant difference between body weight of galegine-treated and control mice ($P<0.001$), but no significant difference for food intake. In each case, there was a significant effect of time ($P<0.0001$), but no significant interaction between time and treatment.

glucose uptake by 3T3-L1 adipocytes (Figure 3a) without any effect on cell viability (data not shown). The PI3 kinase inhibitor LY294002 ($100 \mu\text{M}$) prevented the effect of galegine on glucose uptake even at maximal galegine concentrations. Another PI3 kinase inhibitor, wortmannin ($1 \mu\text{M}$), reduced, but did not abolish, galegine-stimulated glucose uptake (Figure 3a). The stimulatory effect of insulin ($1 \mu\text{M}$) on glucose uptake by 3T3-L1 adipocytes was prevented by either wortmannin or LY294002 (Figure 3b). Treatment of cells with $10 \mu\text{M}$ cytochalasin B completely inhibited both galegine- and insulin-stimulated glucose uptake into 3T3-L1 adipocytes (Figures 3a and b).

Incubation with galegine ($1 \mu\text{M}$ – 1 mM , 5 h) produced a concentration-dependent stimulation of glucose uptake into L6 myotubes (Figure 4a), again without any effect on cell viability (data not shown). Treatment of L6 myotubes with $100 \mu\text{M}$ LY294002 inhibited this effect of galegine, although wortmannin ($1 \mu\text{M}$) produced only a small reduction in galegine-stimulated glucose uptake. The stimulatory action of $1 \mu\text{M}$ insulin on glucose uptake by L6 myotubes was prevented by both wortmannin and LY294002 (Figure 4b). Treatment of L6 myotubes with $10 \mu\text{M}$ cytochalasin abolished both galegine- (Figure 4a) and insulin-stimulated (Figure 4b) glucose uptake. A 24-h cell incubation with phenformin stimulated 2-deoxy-D-glucose uptake in both 3T3-L1 adipocytes (Figure 5a) and L6 myotubes (Figure 5b) with a broadly similar potency and maximal response to galegine. Metformin

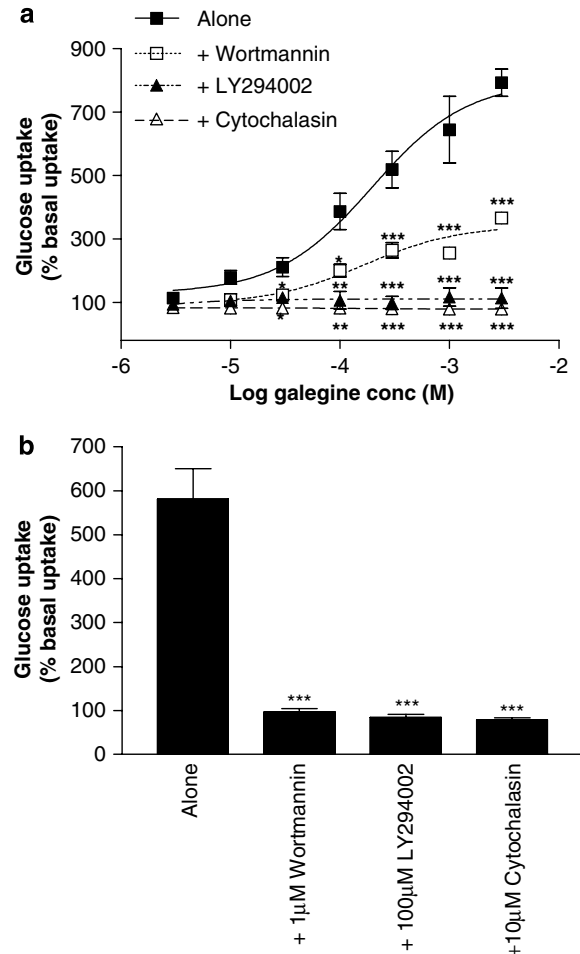


Figure 3 Effects of 5 h treatment with galegine (a, upper panel) and insulin (b, lower panel) on 2-deoxy-D-glucose uptake by 3T3-L1 adipocytes in the absence or presence of $1 \mu\text{M}$ wortmannin, $100 \mu\text{M}$ LY294002 or $10 \mu\text{M}$ cytochalasin B. Results are expressed as a percentage of basal glucose uptake and values shown are mean \pm s.e.m. of observations from two individual experiments of three replicates per experiment. * $P<0.05$, ** $P<0.01$, *** $P<0.001$ vs galegine alone.

was much less potent in 3T3-L1 adipocytes and showed no activity in L6 myotubes up to $100 \mu\text{M}$.

Effect of galegine on lipolysis in 3T3-L1 adipocytes

A 24-h incubation with galegine (0.3 – $300 \mu\text{M}$) produced a slight reduction in basal glycerol release and a more marked reduction in isoprenaline-stimulated glycerol release from 3T3-L1 adipocytes (Figure 6).

Effect of galegine on acetyl-CoA carboxylase activity

Galegine (0.3 – $30 \mu\text{M}$, 24 h) produced a concentration-dependent reduction in ACC activity in 3T3-L1 adipocytes. The known activator of AMPK aminoimidazole-4-carboxamide ribonucleotide (AICAR) also significantly reduced ACC activity in 3T3-L1 adipocytes, although it was less potent than galegine (Figure 7a). Galegine and AICAR also reduced acetyl CoA carboxylase activity in L6 myotubes (Figure 7b), galegine appearing less potent in this cell line with effects reaching statistical significance only at concentrations of $30 \mu\text{M}$ and above.

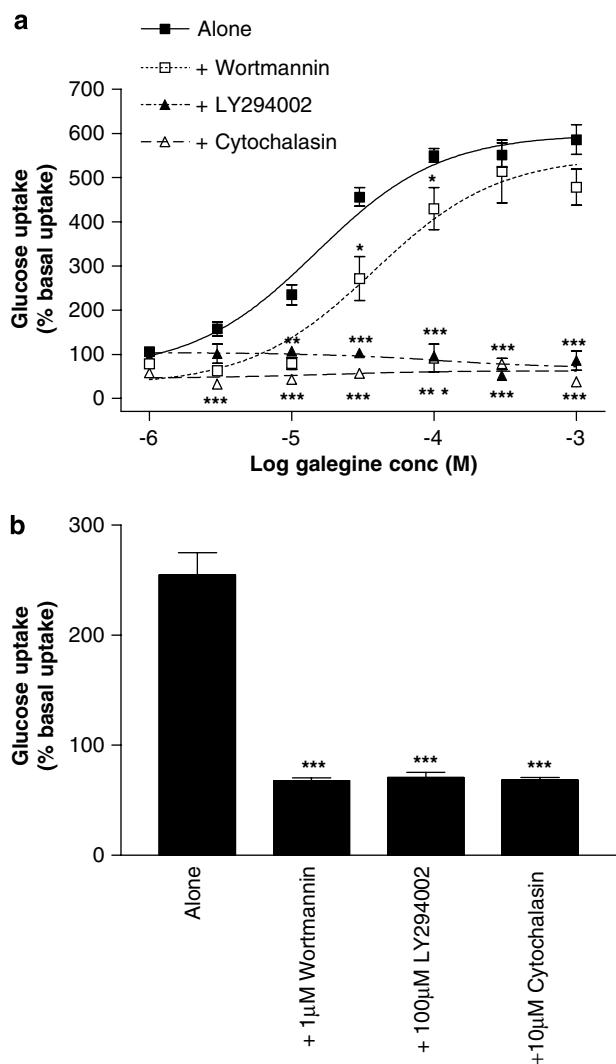


Figure 4 Effects of 5 h treatment with galegine (a, upper panel) and insulin (b, lower panel) on 2-deoxy-D-glucose uptake by L6 myotubes in the absence or presence of 1 μM wortmannin, 100 μM LY294002 or 10 μM cytochalasin B. Results are expressed as a percentage of basal glucose uptake and values shown are mean ± s.e.m. of observations from two individual experiments with three replicates per experiment. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ vs galegine alone.

Direct effect of galegine on lipogenic enzymes

Direct incubation of galegine (500 μM) with homogenates of rat epididymal adipose had no effect on the activities of FASN, malic enzyme or ATP citrate lyase (Table 1).

Effect of galegine on AMPK activity

Incubation of H4IIE rat hepatoma cells with galegine (10 or 300 μM) for up to 6 h produced a time-dependent activation of AMPK measured in cell lysates, with maximal activation at 360 min and twofold activation still evident at 24 h (data not shown) in the presence of 300 μM galegine. The effect of 300 μM galegine was markedly greater than that of 10 μM (Figure 8). Incubation with galegine for 1 h produced a concentration-dependent activation of AMPK in both 3T3L-1

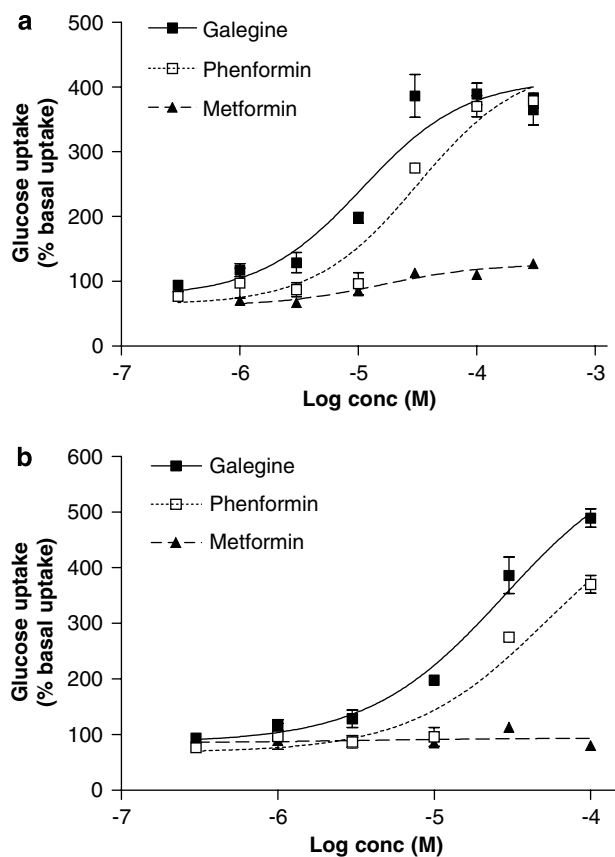


Figure 5 Comparison of the effects of 24 h treatment with galegine, phenformin and metformin on 2-deoxy-D-glucose uptake by 3T3-L1 adipocytes (a, upper panel) and L6 myotubes (b, lower panel). Results are expressed as a percentage of basal glucose uptake and are mean ± s.e.m. of observations from two individual experiments with three replicates per experiment.

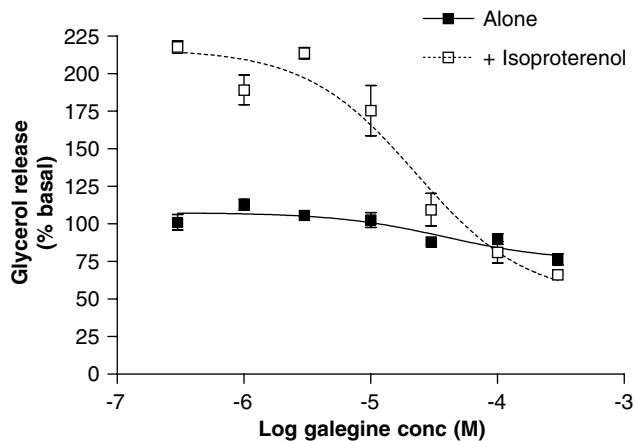


Figure 6 Effect of 24 h treatment with galegine on basal lipolysis and 10 μM isoprenaline-stimulated lipolysis in 3T3-L1 adipocytes. Results are expressed as a percentage of basal glycerol release and are mean ± s.e.m. of observations from three individual experiments with three replicates per experiment.

adipocytes and L-6 myotubes (Figure 8). Galegine also produced a concentration-dependent activation of the enzyme in a human kidney cell line (HEK293) (data not shown).

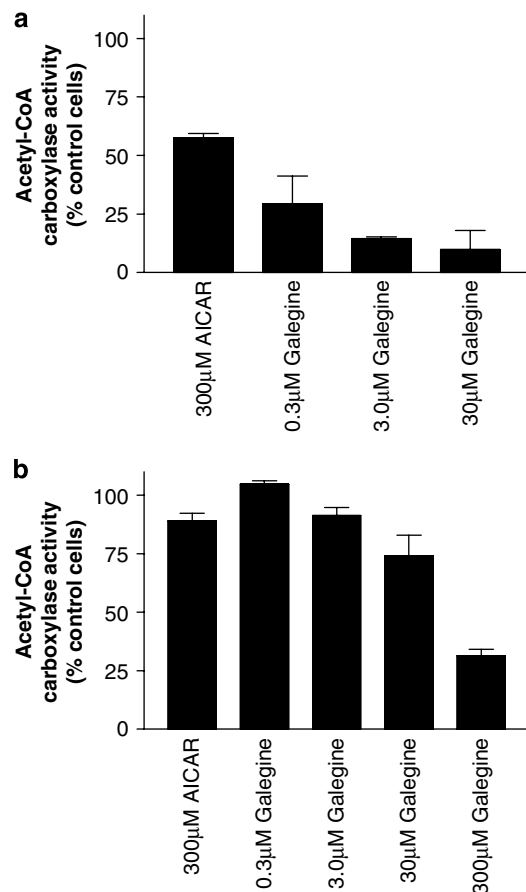


Figure 7 Effect of 24 h galegine treatment on acetyl-CoA carboxylase activity in 3T3-L1 adipocytes (a, upper panel) and L6 myotubes (b, lower panel). Results are mean \pm s.e.m. ($n=3$) and are expressed as a percentage of activity measured within control cells. In 3T3-L1 adipocytes (upper panel), all values are statistically significantly different from control (ANOVA followed by Dunnett's $P<0.01$), whereas there is no significant effect of galegine on acetyl-CoA carboxylase in L6 myotubes. ANOVA, analysis of variance.

Table 1 The effect of 500 μ M galegine on the activity of fatty acid synthase, ATP-citrate lyase and malic enzyme extracted from fat tissue

Treatment	Enzyme activity ($\text{nmol mg}^{-1} \text{min}^{-1}$)		
	Fatty acid synthase	ATP-citrate lyase	Malic enzyme
Control	1.48 \pm 0.01	2.19 \pm 0.01	487.6 \pm 10.0
Galegine	1.50 \pm 0.05	2.18 \pm 0.1	514.5 \pm 3.2

The activities of fatty acid synthase and ATP-citrate lyase are expressed as nmol of NADPH oxidized per mg protein per min and the activity of malic enzyme expressed as nmol of NADPH formed per mg protein per min. Data are expressed as mean \pm s.e.m., $n=3$.

Effects of galegine on gene expression

Microarray analysis of 3T3-L1 adipocytes exposed for 24 h to galegine (500 μ M) revealed changes in expression of a group of lipid and insulin signalling pathway genes (Table 2). QPCR studies were carried out on adipocytes to validate data obtained from microarray analysis. REST analysis found that changes in expression of the genes analysed by QPCR, in general, matched the direction and approximate mean fold changes observed using the Affymetrix Mouse Expression

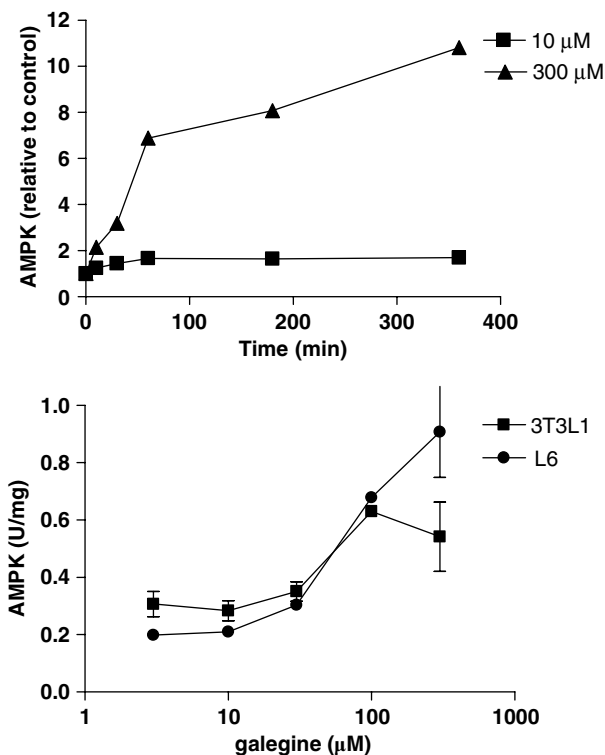


Figure 8 Effect of galegine treatment on AMPK activity of H4IIE rat hepatoma cells (upper panel) or 3T3-L1 adipocytes and L6 myotubes (1 h incubation, lower panel). AMPK activities are expressed relative to activity detected in control cells (upper panel) or in absolute units (lower panel). Each point is the mean (\pm s.e.m.) of three separate experiments, with each assay being done in duplicate. AMPK, AMP-activated protein kinase.

GeneChip microarray (Table 2). These genes included FASN and sterol-regulatory-element-binding protein (SREBP), which exhibited 2.5-fold and 2.0-fold down-regulation in expression, respectively. Hormone-sensitive lipase and acyl-CoA synthetase also showed altered expression in adipocytes, where they both exhibited significant downregulation, although these observations have not been validated by QPCR.

The transcriptional co-activator, peroxisome proliferator-activated receptor- γ co-activator-1 α was upregulated by 1.87-fold in adipocytes incubated with galegine. There was a significant downregulation (-1.8 fold, $P=0.001$) of phosphoenolpyruvate carboxykinase-1 in galegine-treated adipocytes.

Tribbles-3 expression was increased in 3T3-L1 adipocytes (4.09-fold) incubated with galegine, although this has not been validated by QPCR.

Discussion

Galegine produced weight loss, reduction in weight gain and reduced blood glucose, accompanied by a decrease in food intake. A significant part of this weight loss is due to reduction in food intake, but a pair-feeding experiment demonstrated that it could at least partly be attributed to other effects, as suggested for the plant *G. officinalis* administered in the same way (Palit *et al.*, 1999). The *in vivo* effect of galegine may depend on initial body weight, as

Table 2 Galegine-induced change in expression of genes in mouse liver and 3T3-L1 adipocytes

Gene	3T3-L1 adipocyte microarray Fold change	3T3-L1 adipocyte QPCR Fold change	P-value
Acyl-CoA synthetase long-chain family member 1 (ACSL1)	-2.80	NA	NA
Lipase, Hormone sensitive (HSL)	-2.25	NA	NA
Peroxisome proliferator activated receptor gamma coactivator 1 alpha (PGC-1)	+1.87	NA	NA
Tribbles homolog 3 (TRB3)	+4.09	NA	NA
Fatty acid synthase (FASN)	-2.31	-2.458	0.001
Phosphoenolpyruvate carboxykinase 1, cytosolic (PCK1)	-2.56	-1.809	0.001
Sterol regulatory element binding protein 1 (SREBP)	-1.89	-1.991	0.001

Abbreviations: NA, not assayed; QPCR, real-time quantitative PCR.

Mice were fed a diet as detailed in Materials and methods. Adipocytes were treated for 24 h with 0.5 mM galegine relative to expression in control non-treated cells. The microarray results were from a single microarray study where four pooled control adipocyte cultures were compared with four pooled galegine-treated adipocyte cultures. The QPCR results were obtained from three pairs of control and galegine-treated adipocytes, with three replicate readings per gene of interest, with the exception of PCK1 data, which were represent three replicate readings from two pairs of treated and control adipocytes. QPCR values were normalized by HPRT expression. QPCR data analysis was carried out using REST (Relative Expression Software Tool). P-values are for QPCR results only.

another pair-feeding experiment (data not shown), using mice with a markedly lower initial body weight (23 g compared with 28 g in data presented in Figure 2), showed no dissociation of body weight loss from reduced food intake. Nevertheless, the *in vitro* pharmacology of galegine demonstrated actions compatible with *in vivo* metabolic activities, independent of reduced food intake. Thus, galegine stimulated glucose uptake by both 3T3-L1 adipocytes and L6 myotubes, which could contribute to the *in vivo* hypoglycaemic action of galegine. The stimulation of glucose uptake is not due to cell toxicity, as evidenced by the lack of cytotoxicity in the concentration range used. Moreover, cytochalasin B, in addition to inhibiting the effect of insulin, also inhibited the stimulatory effect of galegine on glucose uptake by both cell types, suggesting a role for glucose transporter GLUT4 (Keller and Mueckler, 1990; Sarabia *et al.*, 1992) and other glucose transporters such as GLUT1, which is also expressed in L6 myotubes and 3T3-L1 adipocytes, in mediating the effect of galegine. Like the effects of insulin, the stimulatory effects of galegine on glucose uptake by 3T3-L1 adipocytes were reduced by the PI3 kinase inhibitor wortmannin (Ui *et al.*, 1995) and abolished by LY294002, another PI3 kinase inhibitor (Stein, 2001). Although LY294002 also markedly reduced the effects of galegine in L6 myotubes, wortmannin was much less effective, despite being just as effective in reducing the action of insulin. The reason for this discrepancy is unclear, but the balance of evidence supports some contribution from the PI3 kinase pathway in the stimulatory action of galegine on glucose uptake. However, effects of wortmannin and LY294002 on other kinases cannot be excluded.

Galegine is a guanidine derivative, and guanidine was the compound derived from *G. officinalis*, which gave rise to the biguanides, metformin and phenformin (Bailey and Day, 1989; see Figure 9 for structure of galegine and guanidine compared with the biguanides). Metformin and phenformin both activate AMPK (Zhou *et al.*, 2001), and the use of mice with a liver-specific knockout of the kinase LKB1 suggests that activation of AMPK in the liver (which requires the upstream kinase LKB1) is largely responsible for the glucose-lowering effect of metformin (Shaw *et al.*, 2005). The effects of biguanides on AMPK are probably mediated by their ability to inhibit complex I of the respiratory chain (El-Mir

et al., 2000; Owen *et al.*, 2000), and thus increase the cellular AMP:ATP ratio. Increases in AMP:ATP have indeed been demonstrated for phenformin (Hawley *et al.*, 2005), although they have been more difficult to demonstrate for the less potent biguanide, metformin.

In the present study, phenformin had similar effects to those of galegine in stimulating glucose uptake in 3T3-L1 adipocytes and L6 myotubes, although metformin was much less effective and less potent. Activation of AMPK stimulates glucose uptake in skeletal muscle (Merrill *et al.*, 1997; Hayashi *et al.*, 1998) and mediates the effect of AICAR (Sakamoto *et al.*, 2005). Unlike the effects of galegine in L6 myotubes in the present study, the effects of AICAR in stimulating glucose uptake in skeletal muscle were not affected by wortmannin, although LY294002 was not tested (Hayashi *et al.*, 1998; Bergeron *et al.*, 1999). There is considerable evidence that the effects of AICAR in stimulating glucose uptake involve glucose transporters (Kurth-Kraczek *et al.*, 1999; Yamaguchi *et al.*, 2005), as suggested in the present study for galegine. The mode of activation of glucose transporters remains to be determined, but there were no changes in the expression of GLUT1 (+1.2-fold) or GLUT4 (no change) genes (data not shown).

The effects of metformin *in vivo* appear to be primarily on the liver, possibly because hepatocytes express high levels of the organic cation transport, OCT1, which appears to facilitate the uptake of metformin (Wang *et al.*, 2002; Shu *et al.*, 2007). Whether OCT1 is involved in the uptake of galegine is not clear at present.

Although we show that galegine had no direct effect on adipose tissue enzymes (malic enzyme, FASN, citrate lyase) that are important in the synthesis of fat, it produced a potent inhibition of acetyl CoA carboxylase when incubated for 24 h with 3T3L-1 adipocytes. ACC exists as two isoforms, with the ACC1 (ACC- α) isoform predominating in adipocytes where it functions in fatty acid synthesis, and the ACC2 (ACC- β) isoform predominating in muscle where its inhibition stimulates fatty acid oxidation (Abu-Elheiga *et al.*, 2001). Both ACC1 (Davies *et al.*, 1992) and ACC2 (Winder *et al.*, 1997) are phosphorylated and activated by AMPK. In the present experiments, we found that the AMPK activator AICAR inhibited ACC in 3T3-L1 adipocytes and, albeit much less markedly, in L6 myotubes. Galegine also inhibited

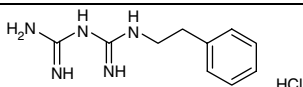
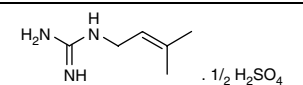
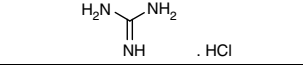
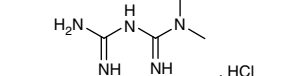
	Phenformin hydrochloride
	Galegine hemi-sulphate
	Guanidine hydrochloride
	Metformin hydrochloride

Figure 9 Structure of galegine compared with that of guanidine, metformin and phenformin.

isoprenaline-stimulated lipolysis, another action associated with AMPK activation (Sullivan *et al.*, 1994; Corton *et al.*, 1995; Watt *et al.*, 2006). The observation that galegine activates AMPK in 3T3-L1 adipocytes and L6 myotubes, as well as in the H4IIE rat hepatoma and HEK293 human kidney cell lines, supports the hypothesis that galegine works, at least in part, through activation of AMPK. In the liver, activation of AMPK leads to inhibition of lipogenesis and glucose production and stimulation of fatty acid oxidation (Viollet *et al.*, 2006).

Finally, in 3T3-L1 adipocytes, galegine downregulated the expression of mRNAs encoding several genes involved in lipid metabolism, including FASN, sterol-regulatory-element-binding protein, hormone-sensitive lipase and acyl-CoA synthetase, although the changes in the last two must be interpreted cautiously as they have not been validated using QPCR. Activation of AMPK is known to reduce the expression of proteins involved in lipogenesis, including the upstream regulator SREBP, at least in the liver (Zhou *et al.*, 2001; Foretz *et al.*, 2005). Hence, activation of AMPK may provide a unifying hypothesis explaining the mechanism of action of galegine. This has already been proposed for the biguanides (Zhou *et al.*, 2001; Hawley *et al.*, 2002) which, like galegine, have been found to have an anti-obesity action (Pasquali *et al.*, 2000; Kay *et al.*, 2001; Kim *et al.*, 2006). The upregulation of the transcriptional co-activator PGC-1 α (peroxisome proliferator-activated receptor- γ co-activator-1 α) by galegine is also compatible with previous studies demonstrating increased PGC-1 α mRNA expression in skeletal muscle in response to AMPK activation (Terada *et al.*, 2002; Zong *et al.*, 2002). PGC-1 α has important roles in promoting cold-induced thermogenesis and fatty acid oxidation, and its expression is reduced in morbidly obese subjects (Semple *et al.*, 2004). The observed downregulation of hormone-sensitive lipase may be an additional explanation for the inhibition of isoprenaline-induced lipolysis. AMPK-mediated inhibition of hormone-sensitive lipase is accompanied by reduced phosphorylation of the lipase (Garton *et al.*, 1989; Watt *et al.*, 2006), but the role of downregulation of protein expression has not been investigated. The significance of the upregulation of Tribbles-3 (which has not been validated using QPCR) is unclear, but this gene was found recently to be upregulated in 3T3-L1 adipocytes in response to reductions in available glucose and was suggested to be involved in cell-survival in response to

glucose deprivation (Yacoub Wasef *et al.*, 2006). This may relate to the previously demonstrated inhibitory effect of galegine on mitochondrial respiration (Lotina *et al.*, 1973), which could be the mechanism whereby it activates AMPK, as indicated above for the biguanides. In this context, however, we were unable to detect any significant effect of galegine on the expression of genes concerned with oxidative phosphorylation (for example, NADPH-ubiquinone dehydrogenase, succinate dehydrogenase complex, ATP synthase) in 3T3-L1 adipocytes.

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Conflict of interest

The authors state no conflict of interest.

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