

## Taxonomic, genetic, chemical and estrogenic characteristics of *Epimedium* species

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### Abstract

To understand the factors contributing to estrogenic properties of extracts from the genus *Epimedium* L. (Berberidaceae), we performed taxonomic, genetic and chemical characterization on 37 specimens from 18 species and related these to estrogen receptor (ER $\alpha$  and ER $\beta$ ) bioactivity, as measured by reporter genes in stable human cells. Boot strap values derived from amplified fragment length polymorphisms indicated that specimens of *E. koreanum*, *E. brevicornum*, *E. myrianthum*, *E. leishanense*, and *E. membranaceum* were genetically distinct and this was supported by their very similar ER $\alpha$  activities. In contrast, specimens from *E. pubescens* and *E. sagittatum* were diverse both genetically, chemically and in terms of ER $\alpha$  and ER $\beta$  bioactivities. Strikingly, a genetic cluster comprising six rare *Epimedium* species exhibited strongest ER $\alpha$  and ER $\beta$  activity, and this bioactivity was positively correlated with content of trace flavonoid aglycones (kaempferol, apigenin, quercetin, luteolin and brevisflavone B). In contrast, there was no association between estrogenic activity and the major flavonol glycoside constituents (icariin and epimedin A–C). Although they exhibited equally strong ER $\alpha$  and ER $\beta$  activity, *E. koreanum* can be clearly differentiated from *E. pubescens* and *E. brevicornum* by genetic distance and its significantly lower content of epimedin C. Our morphologic, genetic, chemical and bioactivity profiling provide the basis for the production of extracts with reproducible estrogenic properties. Such reproducibility will be critical for the standardization of *Epimedium*-based products.

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**Keywords:** *Epimedium* species; Berberidaceae; Taxonomy; Genetic; Chemical; Estrogenic activity

### 1. Introduction

The traditional Chinese medicinal herb, *Epimedium* L. (Berberidaceae), is a popular botanical supplement used to improve menopausal symptoms and bone health, amongst other indications. Major *Epimedium* species used for medicinal purposes are *E. koreanum* Nakai, *E. pubes-*

*cens* Maxim., *E. brevicornum* Maxim, *E. sagittatum* (Sieb. Et Zucc) Maxim, and *E. wushanense* T.S. Ying (The State Pharmacopoeia Commission of PR China, 2000). *Epimedium* species have a particularly high content of the prenylated flavonol glycosides (Liang et al., 1997), the most prominent of which is icariin (1), a flavone with two glycoside moieties (Akai, 1935). Icariin and related flavonoids (Fig. 1) were reported to enhance the osteogenic differentiation of rat primary bone marrow stromal cells (Chen et al., 2005), increase osteoblastic proliferation (Meng et al., 2005), reduced osteoclastic bone resorption (Yu et al., 1999) and to increase mineral content, and prevent osteoporosis in ovariectomized rats (Zhang et al., 2006). Because of these properties, icariin and epimedin A (2), B

**Abbreviations:** AFLP, amplified fragment length polymorphisms; ER, estrogen receptor; *E.*, *Epimedium*; PCR, polymerase chain reaction; RE, relative efficacy – the ratio between maximum activity of extract and the maximum activity of estradiol; RP, relative potency – ratio between EC<sub>50</sub> of extract and estradiol.

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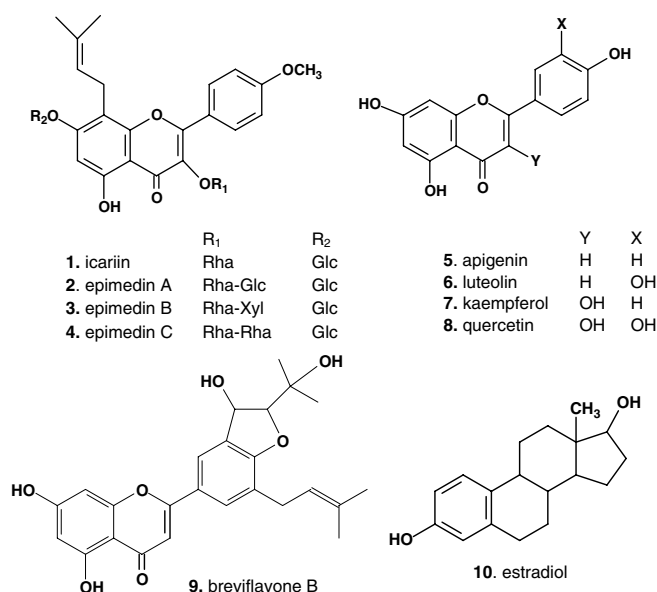


Fig. 1. Structures of major flavonoid glycosides (1–4), bioactive aglycones (5–9) from *Epimedium*, compared to the physiological estrogen, estradiol (10). Rha, rhamnose; Glc, glucose; Xyl, xylose.

(3), and C (4), (all glycosides of anhydroicaritin) are frequently used as marker compounds for the quality control of *Epimedium* and its medicinal extracts (Liu et al., 2006).

Flavonoids can activate the estrogen receptors (ER), part of a 48-member family of transcription factors, which control sexual differentiation and regulate skeletal health in women. The two estrogen receptors, ER $\alpha$  and ER $\beta$ , are similar in their DNA-binding domains but have significant differences in their C-terminal ligand binding domains, sharing only 56% homology (Koehler et al., 2005). These structural differences cause ligands to bind with differing affinity to ER $\alpha$  and ER $\beta$  resulting in different conformations of the activated receptor and differential recruitment of coregulators and selective estrogen modulator activity (McDonnell, 2003). The predominant receptor in the uterus is ER $\alpha$ , while ER $\beta$  is expressed at high levels in the ovary, vascular endothelium, smooth muscle and the central nervous system. When the receptors are expressed in the same tissue, such as the breast, they may cause different actions. Thus ER $\alpha$  stimulates, whereas ER $\beta$  inhibits, cells obtained from lactating mammary glands. (Koehler et al., 2005). Ligands which are selective for either ER $\alpha$  or ER $\beta$  is a subject of intense pharmaceutical research.

Recently extracts of *Epimedium* were found to be potent and specific estrogenic activity (Yap et al., 2005; De Naeyer et al., 2005). Estrogenic activity was not caused by flavonoid glycosides, but was due to the presence of flavonoid aglycones (Yap et al., 2005). These aglycones belong to a recently described class of prenyl flavonoids with potent estrogenic properties (Milligan et al., 2000). Intriguingly prenyl flavones exhibit dose-dependent anti-estrogenic properties in human breast cancer cells (Yap et al., 2005; Pedro et al., 2005) and may be specific inhibitors of the breast cancer resistant protein ABCG2 (Ahmed-Belkacem

et al., 2005). Estrogenic activity in *Epimedium* extracts may also be contributed by known flavonoids such as apigenin (5), luteolin (6), kaempferol (7), and quercetin (8) (Wu et al., 2003). The estrogenic properties of *Epimedium* coupled with these anti-proliferative effects on breast cancer cells suggest its possible utility for menopausal estrogen replacement therapy but without adverse effects on breast health associated with current estrogen–progesterone formulations (Rossouw et al., 2002). Clinical studies are required to establish the role, if any, of *Epimedium* extracts for estrogen-deficient conditions.

However the prerequisite for a scientific study is the procurement of high quality, standardized drugs with reproducible pharmacological properties. In common with many traditional medicinal plants, a major problem with *Epimedium* is the absence of a rigorous method to authenticate species. Chinese taxonomists and geneticists have variably reported numbers ranging from 20 to 50 species (Sun et al., 2005). Traditional herbalists do not differentiate among *Epimedium* species, but rather use a mixture of species together as *Herba Epimedii* (Yang, 1985). These species differ significantly in concentrations of major and minor constituents (Wu et al., 2003). Another concern is that *Epimedium*, like the majority of traditional Chinese herbs, is not cultivated but is collected from the wild, which increases the dangers of wrong species identification, genetic diversity, and possible differences in levels of bioactive compounds due to soil and climate (Guo and Xiao, 2003; Chen et al., 1996; Mizuno et al., 1989). Other factors that may cause variations include differences in processing, packaging and storage of raw materials. These multitudes of unknowns render scientific evaluation of *Epimedium* extracts for menopausal and bone health problematic.

The aims of this study are to define the taxonomic, genetic and chemical characteristics of species from *Epimedium*, and determine their relative contributions to estrogenic properties of resultant extracts. Specimens of 18 *Epimedium* species, including the five major medicinal species, were taxonomically identified and subjected to amplified fragment length polymorphism genetic analysis. The content of major flavonol glycosides (1–4) and trace estrogenic aglycones (5–9) (Fig. 1) in various samples of the same species were determined, and compared to ER $\alpha$  and ER $\beta$  activity as measured with reporter genes in stable cell lines. The utility of using such a combined taxonomic, genetic, chemical and bioresponse profiling to obtain reproducible extracts from complex medicinal plants, like *Epimedium* is discussed.

## 2. Results

Leaves from 18 *Epimedium* species, collected from Central and Northern Mainland China, were obtained from the Institute of Medicinal Plant Development, Chinese Academy of Medical Sciences, Beijing (Table 1). These were supplemented with *Epimedium* specimens purchased

Table 1  
Estrogenic effects of bioactive flavonoid aglycones (5–8), and *Epimedium* specimens used in this study

Code	Ligand	ER $\alpha$		ER $\beta$	
		RE <sup>a</sup>	RP <sup>b</sup>	RE <sup>a</sup>	RP <sup>b</sup>
E2	Estradiol	1.00	1.00	1.00	1.00
<i>Bioactive flavonoid aglycones</i>					
A	Apigenin	2.37	4.95E–4	3.33	1.74E–3
K	Kaempferol	1.21	2.36E–4	1.07	1.43E–3
L	Luteolin	1.05	6.83E–4	0.76	1.71E–3
Q	Quercetin	0.15	9.28E–5	0.26	3.87E–4
<i>Epimedium extract</i> <sup>c</sup>					
Code	Botanical name	Location/source			
A1	<i>E. acuminatum</i>	1	5.1E–5	0.04	0.00043
A2	<i>E. acuminatum</i>	0.96	–	0.34	–
SL	<i>E. stelluatum</i>	2.21	2.4E–6	0.42	1.7E–6
FG	<i>E. fargesii</i>	1.49	5.3E–6	1.29	2.8E–6
FH	<i>E. franchetii</i>	2.3	1.8E–6	0.9	1.1E–6
ZH	<i>E. zhushanense</i>	1.76	3.3E–6	0.81	2.1E–6
LI	<i>E. leishanense</i>	1.22	–	0.47	–
L2	<i>E. leishanense</i>	1.7	1.3E–6	0.2	5.5E–7
DO	<i>E. dolichostemen</i>	0.64	6.3E–6	0.71	2.4E–6
S2	<i>E. sagittatum</i>	0.41	2.5E–6	0.21	8.2E–7
M1	<i>E. myrianthum</i>	0.61	–	0.23	–
M2	<i>E. myrianthum</i>	0.99	8.7E–6	0.26	2.2E–6
S3	<i>E. sagittatum</i>	1.18	1.2E–5	0.18	–
LT	<i>E. leptorrhizum</i>	0.77	3.3E–5	0.07	1.1E–6
P1	<i>E. pubescens</i>	0.72	4.7E–5	0.47	2.2E–6
ZB	<i>E. zhenbaense</i>	0.54	8.8E–6	0.37	1.1E–6
WU	<i>E. wushanense</i>	0.33	2.7E–6	0.12	7.4E–7
B1	<i>E. brevicornum</i>	0.65	1.4E–6	0.01	6.6E–7
B3	<i>E. brevicornum</i>	0.7	1.4E–5	0.17	2.5E–6
B2	<i>E. brevicornum</i>	0.52	2.7E–6	0.08	1.6E–6
B4	<i>E. brevicornum</i>	0.73	1.6E–5	0.15	–
P8	<i>E. pubescens</i>	1.1	5.7E–5	0.55	–
P3	<i>E. pubescens</i>	1.37	4.7E–5	1.35	3.8E–6
P2	<i>E. pubescens</i>	1.28	1.2E–5	1.1	2.1E–6
P4	<i>E. pubescens</i>	0.87	5E–5	0.38	–
P5	<i>E. pubescens</i>	0.7	9.9E–6	0.06	–
P6	<i>E. pubescens</i>	0.71	4.8E–5	0.25	–
P7	<i>E. pubescens</i>	0.66	1.7E–5	0.09	–
DA	<i>E. davidii</i>	0.3	3.3E–6	0.19	1.1E–6
PA	<i>E. pauciflorum</i>	0.27	2E–6	0.28	7.5E–7
ME1	<i>E. membranaceum</i>	0.22	2.3E–6	0.06	1E–6
ME2	<i>E. membranaceum</i>	0.29	3.5E–6	0.19	1.3E–6
S1	<i>E. sagittatum</i>	0.26	–	0.08	–
K4	<i>E. koreanum</i>	0.67	9E–6	0.55	–
K2	<i>E. koreanum</i>	0.7	7.5E–6	0.26	1.6E–6
K3	<i>E. koreanum</i>	0.67	6.9E–6	0.11	1.2E–6
K1	<i>E. koreanum</i>	0.67	6.9E–6	0.4	1.9E–6

<sup>a</sup> Relative efficacy (RE): ratio between maximum activity of extract and maximum activity of estradiol.

<sup>b</sup> Relative potency (RP): ratio between EC<sub>50</sub> of extract and EC<sub>50</sub> of estradiol.

<sup>c</sup> Specimens are arranged in according to genetic relatedness as depicted in Fig. 2. All specimens from herbarium of Institute of Medicinal Plant Development, unless otherwise indicated. Other sources: Schwabe, GmbH, Germany; STPT, China; Shenyang Company, China; Yan Bian Company, China; NK, place of collection not known.

from commercial sources in Singapore and PR China (Table 1). A total of 37 *Epimedium* specimens, including eight species with multiple specimens, were taxonomically identified by one of the authors (B.L.G.). All specimens were extracted with 100% ethanol at 37 °C for 7 days, resulting in an average yield of 1.29 ± 0.07 g/100 g (dried wt/wt). All botanical specimens and extracts were subjected to genetic, chemical, and estrogenic bioresponse profiling.

### 2.1. Genetic characterization of *Epimedium* species

To genetically identify *Epimedium* species, fluorescent AFLP analysis was used to examine differences across the entire plant genome. DNA was extracted from leaves, digested with *Eco*RI and *Mse*I, ligated to the adaptor, amplified with polymerase chain reaction (PCR) primers, one of which is end-labeled with FAM fluorescent dye

Table 2  
Similarity for AFLP patterns

	A	B	D0	DA	FG	FH	S	K	L	LT	M	ME	P	PA	SL	WU	ZB	ZH
A(2), <i>E. acuminatum</i>	<b>0.537</b>																	
B(4), <i>E. brevicornum</i>	0.432	<b>0.569</b>																
D0, <i>E. dolichostemen</i>	0.406	0.383	<b>1.000</b>															
DA, <i>E. davidii</i>	0.358	0.382	0.338	<b>1.000</b>														
FG, <i>E. fargesii</i>	0.461	0.413	0.426	0.347	<b>1.000</b>													
FH, <i>E. franchetii</i>	0.450	0.418	0.448	0.372	0.471	<b>1.000</b>												
S(3), <i>E. sagittatum</i>	0.365	0.358	0.357	0.326	0.373	0.374	<b>0.295</b>											
K(4), <i>E. koreanum</i>	0.255	0.268	0.261	0.227	0.277	0.280	0.272	<b>0.500</b>										
L(2), <i>E. leishanense</i>	0.439	0.443	0.420	0.358	0.463	0.412	0.369	0.249	<b>0.603</b>									
LT, <i>E. leptorrhizum</i>	0.414	0.432	0.378	0.332	0.451	0.452	0.354	0.239	0.434	<b>1.000</b>								
M(2), <i>E. myrianthum</i>	0.427	0.393	0.423	0.332	0.425	0.405	0.348	0.246	0.423	0.377	<b>0.485</b>							
ME(2), <i>E. membranaceum</i>	0.435	0.431	0.379	0.397	0.444	0.474	0.365	0.247	0.420	0.417	0.407	<b>0.536</b>						
P(8), <i>E. pubescens</i>	0.420	0.410	0.370	0.368	0.429	0.424	0.349	0.257	0.398	0.411	0.380	0.420	<b>0.458</b>					
PA, <i>E. pauciflorum</i>	0.430	0.432	0.371	0.431	0.413	0.436	0.372	0.242	0.393	0.406	0.381	0.472	0.428	<b>1.000</b>				
SL, <i>E. stelluatum</i>	0.446	0.425	0.412	0.346	0.417	0.461	0.366	0.229	0.409	0.420	0.419	0.404	0.395	0.388	<b>1.000</b>			
WU, <i>E. wushanense</i>	0.432	0.415	0.422	0.368	0.437	0.411	0.361	0.257	0.452	0.399	0.405	0.419	0.388	0.387	0.417	<b>1.000</b>		
ZB, <i>E. zhenbaense</i>	0.418	0.436	0.390	0.415	0.390	0.437	0.361	0.239	0.420	0.373	0.405	0.410	0.415	0.444	0.444	0.462	<b>1.000</b>	
ZH, <i>E. zhushanense</i>	0.475	0.434	0.456	0.359	0.508	0.515	0.379	0.251	0.482	0.451	0.411	0.472	0.419	0.450	0.465	0.459	0.423	<b>1.000</b>

Jaccard values of band sharing were averaged over pairwise comparisons. Figures in parentheses indicate the number of individuals analyzed for that species. Bold indicate intraspecific comparisons.

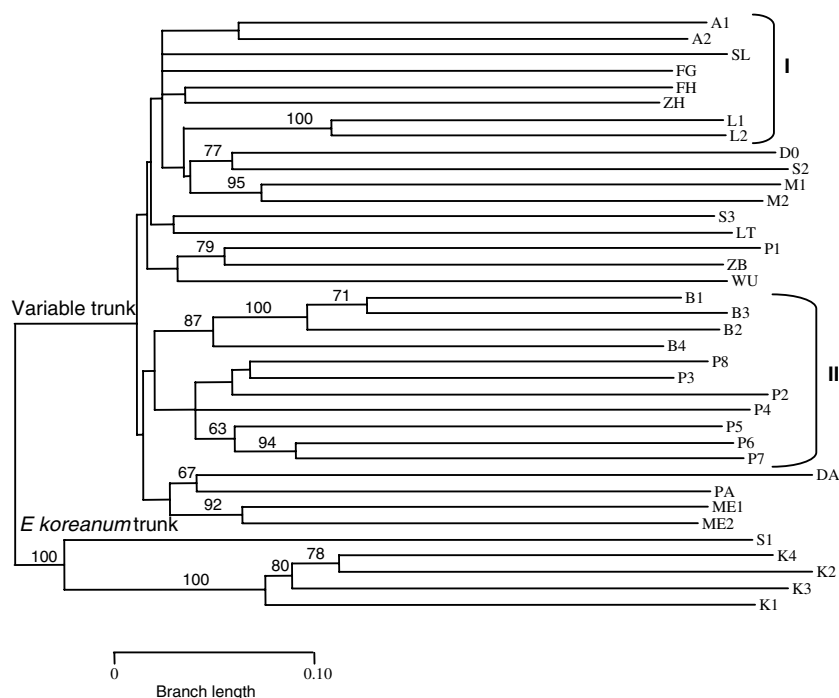


Fig. 2. Dendrogram describing the relationship among the 37 *Epimedium* samples. Shown is the neighboring joining tree based on the similarity of AFLP banding profiles. Numbers indicate the proportion of 1000 bootstrap samples in which a particular node was found. Specimens were collected from locations all over China as listed in Table 1. Scale bar indicates 10% branch length.

(blue). PCR fragments were resolved in an Applied Biosystems 3730xl DNA analyzer. Fragment sizes were calculated by mixing samples with DNA size markers in the range of 75–500 bp end labeled with Rox (red). The four primer combinations yielded a total number of 840 polymorphic markers among the samples analyzed. Table 2 shows pairwise Jaccard genetic similarity indices which were used to construct a neighbor joining phylogenetic tree (Fig. 2). Bootstrap values >50 were labeled along the nodes that they support. Genus *Epimedium* was divided clearly into two main genetic trunks; a well defined trunk consisting of mainly one *Epimedium* species, *E. koreanum*; and a more variable trunk comprising all the other species (Fig. 2). The eight *Epimedium* species with multiple samples were subjected to bootstrap analysis to evaluate intraspecies diversity (Efron et al., 1996). Jaccard values of samples from *E. myrianthum*, *E. leishanense*, *E. brevicornum*, *E. membranaceum* and *E. koreanum* strongly supported their grouping into the same species. For these species, average value of intraspecies similarity was higher than any of interspecies similarity, suggesting higher level of variation between species (Table 2). Clear identification of herbal materials for *E. myrianthum*, *E. leishanense*, *E. brevicornum*, *E. membranaceum* and *E. koreanum* through comparison of AFLP patterns was therefore more straightforward. However samples for *E. acuminatum*, *E. sagittatum* and *E. pubescens*, could not form a group per species, suggesting the possibility that the majority of genetic variation resides within species. For these species, intraspecies similarity is close or even lower than interspecies similarity and hence AFLP pattern may not be suitable for species identification. For the species with only one sample analyzed, no strong conclusion can be reached despite generally low level of interspecies similarity.

## 2.2. Chemical profiling of *Epimedium*

Major compounds used for standardization of *Epimedium* products and extracts are the flavonol glycosides, icariin, epimedin A, B and C (Fig. 1) (Yap and Yong, 2004). The relative abundance of these flavonol glycosides in all 37 plant extracts were determined by liquid chromatography tandem mass spectrometry. The major flavonoid glycosides in order of abundance were epimedin C ( $7.41 \pm 0.91$  g/100 g), icariin ( $3.04 \pm 0.30$  g/100 g), epimedin B ( $1.91 \pm 1.0$  g/100 g), and epimedin A ( $1.88 \pm 0.33$  g/100 g) (Table 3). These four flavonoid glycosides constitute an average of  $10.4 \pm 1.0$  g/100 g of *Epimedium* ethanol extracts. The average sum of major constituents in *E. koreanum* was significantly lower than *E. brevicornum* and *E. pubescens* ( $4.66 \pm 0.55$  versus  $9.09 \pm 1.21$  and  $13.30 \pm 1.05$  g/100 g respectively,  $P < 0.01$ ), due mainly to lower levels of the most abundant glycoside, epimedin C ( $1.31 \pm 0.28$  versus  $5.17 \pm 0.81$  and  $9.35 \pm 0.63$  g/100 g respectively,  $P < 0.01$ ). In contrast, concentrations of the second most abundant glycoside, icariin, were similar in *E. koreanum*, *E. brevicornum* and *E.*

Table 3

Major flavonoid glycosides (1–4) in *Epimedium* specimens<sup>a</sup>

Code	Epimedin A <sup>b</sup>	Epimedin B <sup>b</sup>	Epimedin C <sup>b</sup>	Icariin <sup>b</sup>
A1	1.2758	1.4027	6.8413	3.1154
A2	0.8671	1.0324	6.3863	2.6273
SL	0.0520	0.1960	0.2071	0.1137
FG	2.2304	2.1474	10.5556	3.9104
FH	0.0474	0.5178	0.1440	0.0932
ZH	0.0439	0.0401	0.1770	0.0376
LI	1.4047	1.4504	16.6478	1.9580
L2	1.1182	1.4022	17.9569	1.3670
DO	0.7129	0.8472	5.7090	2.2029
S2	0.4397	0.5374	15.7854	6.2445
M1	6.1859	3.9292	11.5630	2.2282
M2	2.5819	1.1806	22.1945	0.8395
S3	0.2902	0.4303	16.6333	0.8557
LT	0.0633	0.1458	0.4512	0.2300
P1	2.3832	2.8676	6.3187	4.0355
ZB	0.9069	1.1287	7.5597	3.0058
WU	0.5382	0.6731	6.4458	0.7365
B1	1.5024	5.6980	3.2119	2.7782
B3	1.3286	5.0952	5.3778	3.9265
B2	1.5556	4.5492	4.9236	4.2382
B4	1.4876	2.0251	7.1848	4.7455
P8	2.0552	3.3589	9.6943	2.6149
P3	1.4620	2.1029	8.2134	2.3273
P2	1.4318	2.0737	9.9739	3.6700
P4	1.6242	2.0561	9.2314	2.9885
P5	3.1375	2.8485	12.0893	3.4422
P6	4.0989	4.0738	9.6384	8.6503
P7	2.6874	3.0065	6.6212	3.9868
DA	9.4539	1.1183	4.4325	3.9508
PA	7.6014	2.5513	11.4250	5.0796
ME1	1.0279	1.2288	7.5661	6.5034
ME2	1.3437	1.4148	7.1720	3.7776
S1	0.4728	0.3446	0.6598	3.0959
K4	1.8507	2.0609	1.5569	3.4961
K2	1.9802	2.2844	1.8966	4.0939
K3	1.6269	1.8296	1.2204	2.9763
K1	0.9336	1.1829	0.5719	2.8297

<sup>a</sup> Specimens were arranged in order of genetics relatedness as depicted in Fig. 2.

<sup>b</sup> Concentrations expressed as g/100 g of dried ethanol extracts.

*pubescens* ( $3.34 \pm 0.28$ ,  $3.92 \pm 0.41$ , and  $3.95 \pm 0.81$  g/100 g respectively).

Concentrations of bioactive minor compounds, the flavonoid aglycones apigenin, kaempferol, luteolin, quercetin and brevivflavone B (9) (Fig. 1) were also determined (Table 4). These five flavonoid aglycones form an average of  $0.038 \pm 0.0061$  g/100 g of the ethanol extract. The average concentrations of flavonoids in *Epimedium* species were in the order quercetin > luteolin > brevivflavone B > apigenin > kaempferol with concentrations of  $0.013 \pm 0.003$ ,  $0.0080 \pm 0.002$ ,  $0.0075 \pm 0.0010$ ,  $0.0062 \pm 0.0019$ , and  $0.0035 \pm 0.0009$  g/100 g, respectively.

## 2.3. Estrogenic activity of *Epimedium* extracts and correlations with genetic clusters

In view of *Epimedium* being advertized as a drug for male impotence, we proceeded to examine the effects of



Table 4  
Minor bioactive flavonoid aglycones (5–9) in *Epimedium* specimens<sup>a</sup>

Code	Apigenin <sup>b</sup>	Kaempferol <sup>b</sup>	Luteolin <sup>b</sup>	Quercetin <sup>b</sup>	Breviflavone B <sup>b</sup>
A1	0.0043	0.0000	0.0000	0.0020	0.0105
A2	0.0103	0.0002	0.0002	0.0036	0.0030
SL	0.0655	0.0270	0.0338	0.0385	0.0084
FG	0.0129	0.0021	0.0026	0.0075	0.0000
FH	0.0124	0.0114	0.0114	0.0888	0.0106
ZH	0.0096	0.0092	0.0011	0.0114	0.0102
LI	0.0102	0.0056	0.0190	0.0055	0.0107
L2	0.0014	0.0000	0.0031	0.0018	0.0137
DO	0.0120	0.0034	0.0226	0.0033	0.0216
S2	0.0081	0.0010	0.0036	0.0026	0.0074
M1	0.0044	0.0025	0.0026	0.0015	0.0011
M2	0.0049	0.0062	0.0015	0.0013	0.0064
S3	0.0057	0.0001	0.0026	0.0025	0.0055
LT	0.0012	0.0014	0.0094	0.0195	0.0000
P1	0.0225	0.0063	0.0020	0.0060	0.0034
ZB	0.0242	0.0011	0.0019	0.0025	0.0000
WU	0.0045	0.0007	0.0000	0.0033	0.0042
B1	0.0017	0.0001	0.0131	0.0051	0.0086
B3	0.0011	0.0000	0.0060	0.0062	0.0249
B2	0.0010	0.0000	0.0013	0.0424	0.0252
B4	0.0016	0.0002	0.0066	0.0031	0.0096
P8	0.0014	0.0023	0.0326	0.0030	0.0091
P3	0.0000	0.0000	0.0690	0.0566	0.0042
P2	0.0014	0.0061	0.0177	0.0352	0.0091
P4	0.0000	0.0000	0.0004	0.0038	0.0000
P5	0.0008	0.0027	0.0004	0.0039	0.0111
P6	0.0000	0.0020	0.0000	0.0014	0.0000
P7	0.0000	0.0168	0.0010	0.0038	0.0118
DA	0.0013	0.0000	0.0000	0.0045	0.0095
PA	0.0022	0.0132	0.0000	0.0223	0.0111
ME1	0.0000	0.0000	0.0109	0.0175	0.0000
ME2	0.0010	0.0011	0.0013	0.0103	0.0047
S1	0.0000	0.0000	0.0000	0.0165	0.0000
K4	0.0000	0.0000	0.0000	0.0020	0.0030
K2	0.0000	0.0003	0.0015	0.0175	0.0108
K3	0.0009	0.0002	0.0015	0.0022	0.0026
K1	0.0008	0.0046	0.0137	0.0183	0.0060

<sup>a</sup> Specimens were arranged in order of genetic relatedness as depicted in Fig. 2.

<sup>b</sup> Concentrations expressed as g/100 g of dried ethanol extracts.

the extract on the androgen receptor and related members of the steroid receptor family. Despite its colloquial name of “horny goat weed”, extracts of *Epimedium* did not increase androgen, glucocorticoid or progesterone receptor activities; but instead dose-dependently increased ER $\alpha$  activity in transient transfections (Fig. 3). To increase the accuracy and reproducibility of bioassays, we constructed permanently transfected cells incorporating either ER $\alpha$  or ER $\beta$ , and an estrogen responsive reporter gene in its chromosomes (pERE4-Luc<sub>hygro</sub>). Dose-response studies were performed for the *Epimedium* extracts and estradiol (10) (Fig. 4). For easy comparability, estrogenic activity of each extract was expressed as relative efficacy (RE), the ratio between maximum activity of extract and the maximum activity of estradiol; and relative potency (RP), the ratio between EC<sub>50</sub> of extract and estradiol. Generally *Epimedium* species were ER $\alpha$ -selective in terms of potency (Table

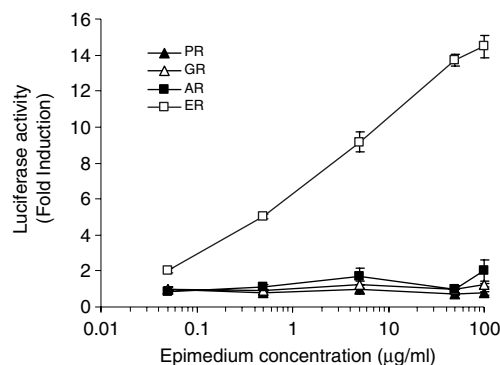


Fig. 3. Effect of *Epimedium* extracts on estrogen, androgen, progesterone and glucocorticoid receptor activities. HeLa cells, transiently transfected with plasmids encoding for ER $\alpha$ , androgen (AR), progesterone (PR) and glucocorticoid (GR) receptors and their cognate response elements driving luciferase-reporter genes, were exposed to an *E. pubescens* extract at the indicated doses. Data (mean  $\pm$  s.e. of at least three replicates) represent increase in luciferase activity over replicates exposed to vehicle only.

1). Thus whereas the phytoestrogens apigenin, kaempferol, luteolin and quercetin had RP which were higher for ER $\beta$  compared to ER $\alpha$ , the reverse was generally observed for *Epimedium* extracts. Extracts also displayed higher maximal activity (higher RE) for ER $\alpha$  compared to ER $\beta$ .

The four samples from the most genetically distinct group, *E. koreanum*, exhibited remarkably similar ER $\alpha$  activity (mean RE of  $0.67 \pm 0.007$ , and mean RP of  $7.5 \pm 0.4$ ) (Table 1). In contrast their ER $\beta$  activities were relatively weaker with mean RE of  $0.33 \pm 0.094$ . In the same way, the four samples from the closely linked *E. Brevicornum* specimens exhibited similar and strong ER $\alpha$  (mean RE of  $0.65 \pm 0.046$ ) and relatively weaker ER $\beta$  (mean RE of  $0.10 \pm 0.036$ ) activities. Although having only two specimens each, ER $\alpha$  activities of *E. leishanense*, *E. myrianthum*, and *E. membranaceum* specimens were very similar suggesting that estrogenic activity may have utility to differentiate these *Epimedium* species.

In contrast, the three specimens from the genetically diverse species, *E. sagittatum*, exhibited correspondingly divergent estrogenic activities with RE for ER $\alpha$  ranging from 1.18 to 0.26 (mean:  $0.61 \pm 0.28$ ) (Table 1). Similarly the eight specimens from the genetically divergent *E. pubescens* species displayed ER $\alpha$  maximum activities ranging from RE of 1.37 to 0.66. In contrast the three *E. pubescens* specimens that were genetically linked (P5, P6, P7) exhibited similar RE ranging from 0.66 to 0.71. This was remarkable as they were collected from different locations at different times.

Intriguingly *E. acuminatum*, *E. stellatum*, *E. fargesii*, *E. franchetii*, *E. zhushanense* and *E. leishanense* clustered at one end (genetic cluster I) of the phylogenetic tree (Fig. 2) exhibited highest ER $\alpha$  (mean RE:  $1.58 \pm 0.25$ ) and ER $\beta$  (mean RE:  $0.55 \pm 0.20$ ) activity. These species exhibited maximum ER $\alpha$  activities that was higher than that of estradiol itself, suggesting the presence of compounds which may augment estrogenic activity.

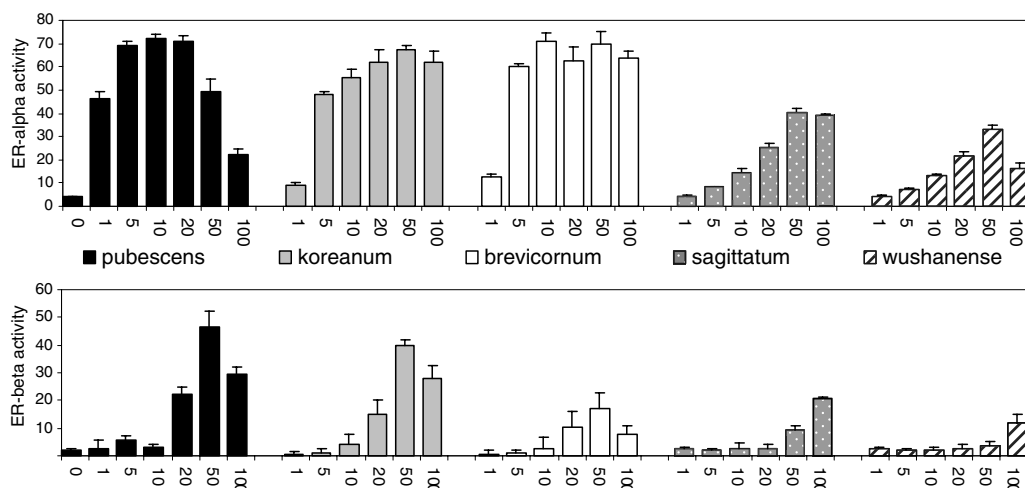


Fig. 4. Estrogenic activity of major *Epimedium* species. HeLa cells stably transfected with ER $\alpha$  (upper panel) or ER $\beta$  (lower panel) and an estrogen-responsive reporter gene were exposed to increasing doses (1–100 µg/ml) of ethanol extracts of indicated *Epimedium* species. Relative estrogenic activity (mean  $\pm$  s.e.) was expressed as percentages of maximal estradiol activity.

## 2.4. Relationship between chemical constituents, genetic clusters and bioactivity

### 2.4.1. Major constituents

There were no simple correlations between the abundance of major flavonoid glycosides and estrogenic activity (Tables 1 and 3). This was not unexpected as the major glycosides (icariin, epimedin A, B, C) were inactive in our estrogenic assays (data not shown).

### 2.4.2. Minor bioactive flavonoid aglycones

In contrast, dose–response studies indicated that minor flavonoid aglycones such as apigenin, kaempferol, luteolin and quercetin exerted significant estrogenic effects (Fig. 5). As expected, phytoestrogens were several fold more ER $\beta$  selective in terms of relative potency (Table 1). However *Epimedium* species appear ER $\alpha$  selective, suggesting that combined activity of extracts was different from that expected from their flavonoid constituents. However within species, some correlations between ER $\alpha$  activity and sum

of the concentrations of bioactive flavonoids (apigenin, kaempferol, luteolin, quercetin, and brevivflavone B) (Fig. 1) were observed (Table 4). The eight specimens from genetic cluster I with highest ER $\alpha$  activity had the highest concentration of bioactive flavonoids ( $0.059 \pm 0.02\%$ ). In this cluster, ER $\alpha$  activity correlated positively ( $R_2 = 0.78$ ) with the sum of minor bioactive flavonoids (Fig. 6A). The *E. pubescens* group had the next higher concentration of bioactive flavonoids ( $0.044 \pm 0.016\%$ ) and reflecting their greater genetic diversity had a lesser degree of correlation ( $R_2 = 0.55$ ) between minor components and ER $\alpha$  activity (Fig. 6B). No definite correlations were observed between ER $\beta$  activity and flavonoid content.

## 3. Discussion

This systematic study of 37 taxonomically-identified specimens revealed differences in the genetic, chemical, and ER $\alpha$ /ER $\beta$  activity profiles that may be used to stan-

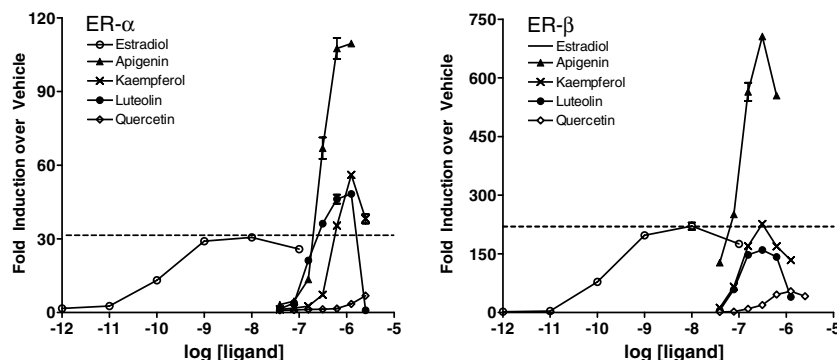


Fig. 5. Estrogenic effects of estradiol and bioactive flavonoid aglycones present in *Epimedium* extracts. Stable cells, permanently transfected with plasmids encoding ER $\alpha$  and ER $\beta$  and the pERE $_4$ -Luc $_{hygro}$  estrogen-responsive reporter gene, were exposed to indicated doses of estradiol, apigenin, kaempferol, luteolin and quercetin. Dotted line indicates maximum activity of estradiol. Data (mean  $\pm$  s.e. of at least three replicates) represent increase in luciferase activity over replicates exposed to vehicle only.

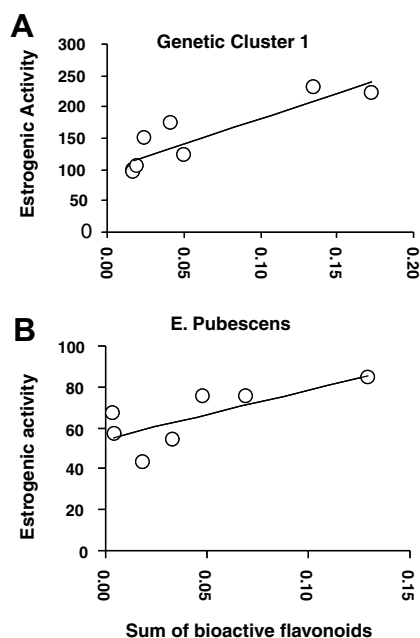


Fig. 6. Relationship between minor bioactive flavonoid aglycones and estrogenic activity. Scatter plot relating bioactivity (ER $\alpha$  activity, Table 1) of ethanol extracts of specimens from, (A) genetic cluster I and, (B) *E. pubescens* to the sum of the concentrations of bioactive flavonoids (5–9). Relative estrogenic activity (mean  $\pm$  s.e.) was expressed as percentages of maximal estradiol activity.

dardize *Epimedium* species, and medicines derived from their extracts.

For DNA analysis, the use of 840 AFLP bands covering the entire plant genome provides rich genetic data adding to previous phylogenetic analyses based on limited specimens (Nakai et al., 1996) or less informative ribosomal DNA sites (Rossouw et al., 2002). Dendrogram analyses confirm the uniqueness of *E. koreanum*, a plant with an exclusive geographical footprint in the colder regions of Northern China and Korea (Guo and Xiao, 2003). In addition, *E. myrianthum*, *E. leishanense*, *E. brevicornum*, and *E. membranaceum* had samples of the same species consistently grouped together. Their average value of intraspecies similarity is higher than interspecies similarity. For these five species, AFLP can be reliably used for identification with Jaccard genetic similarity  $>50\%$  and  $>90$  bootstrap value. On the other hand, specimens of *E. acuminatum*, *E. sagittatum* and *E. pubescens* demonstrated greater intra-genetic variability, suggesting that conclusive identification of this species, and medicines derived from it, may be problematic. Differences in genomic DNA are regarded as definitive means of botanical identification compared to morphology. Genetic profiles like that generated by AFLP analyses may help in the difficult task of standardization and quality assurance for wildcrafted medicinal plants (like *Epimedium*) with many species, a situation that is not uncommon for many herbs in the Chinese herbal pharmacopoeia.

To examine whether marker compounds may aid standardization of extracts, we measured the concentrations

of four major flavonoid glycosides and five minor bioactive flavonoid aglycones (Fig. 1) using chromatographic tandem mass spectrometry. The content of the major glycoside epimedin C was significantly lower in *E. koreanum* compared to *E. brevicornum* and *E. pubescens*. Thus *E. koreanum* can be differentiated from other major *Epimedium* species by its AFLP profiles and its content of Epimedin C. In contrast no differences were observed in the content of icariin, the next most abundant glycoside.

To explore mechanism(s) underlying relationships between genetic, chemical profiles and estrogenic activity, we devised cellular tools to capture summated estrogenic effects of plant extracts. In our bioassay, coding sequences for ER $\alpha$  or ER $\beta$  and a reporter gene, containing a promoter with estrogen-response elements driving a luciferase gene, were stably incorporated into the chromosomes of ER-negative HeLa lines. Such permanent cell lines provide accurate, sensitive, and reproducible biomarkers for the action of estrogens. They have near-linear responses over several orders of magnitude (Wang et al., 2005) and have been shown to correlate with estrogen-driven outcome parameters such as increases in thickness of uterine lining in rats (Sonneveld et al., 2006). Generally *Epimedium* species exhibited high ER $\alpha$  activity with maximal activity that was about 70% of that observed with estradiol. The estrogenicity of *Epimedium* extracts were consistent with findings from rat studies where *Epimedium* flavonoids enhances the osteogenic differentiation of rat primary bone marrow stromal cells (Chen et al., 2005) and can prevent OVX-induced osteoporosis independent of its enhancement of intestinal calcium absorption (Zhang et al., 2006).

Unlike common phytoestrogens, *Epimedium* extracts were ER $\alpha$ -selective, probably reflecting their content of prenyl-flavonoids (Wu et al., 2003; Yap and Yong, 2004), a new class of phytoestrogens with potent ER $\alpha$  activity isolated from hop (*Humulus lupulus* L.) extracts. Furthermore, the ER $\alpha$  activity of genetically related *Epimedium* specimens was consistent, more so than was observed with ER $\beta$ . Specimens that were genetically distinct such as *E. koreanum* and *E. brevicornum* induced remarkably similar ER $\alpha$  activities. In contrast divergent species such as *E. sagittatum* induced correspondingly diverse estrogenic activities. The remarkable correlation between bioassay and genetic classification suggest that their combined use may result in greater accuracy for identification of specimens. The validity of this conclusion has to be tested in a larger study.

Interestingly a cluster comprising six relatively rare *Epimedium* species (*E. acuminatum*, *E. stelluatum*, *E. fargesii*, *E. franchetii*, *E. zhushanense* and *E. leishanense*), exhibited ER $\alpha$  activity that was up to twofold higher than that observed with saturating doses of the physiological estrogen, estradiol. Further studies are needed to investigate whether such “super-agonist” activity may be due to the presence of unknown compounds that activate pathways that synergize steroid receptor signaling, such as the mitogen-activated protein kinase pathway (Jansen et al., 2004).



Although ER $\beta$  generally correlated with ER $\alpha$  activity, some striking differences were also observed. The two specimens of *E. acuminatum* showed high ER $\alpha$  activity, but almost negligible ER $\beta$  activity. In addition to its strong ER $\alpha$  activity (RE: 1.49), *E. fargesii* displayed exceptionally strong ER $\beta$  activity (RE: 1.29), suggesting the possibility that ER $\beta$ -specific compounds may be isolated from this *Epimedium* species. Whether these differences between ER $\alpha$  and ER $\beta$  activities can aid species identification of *E. acuminatum* and *E. fargesii* needs to be determined with larger numbers of specimens. Among major *Epimedium* species, specimens of *E. koreanum*, *E. pubescens* and *E. brevicornum* exhibited high ER $\alpha$  and ER $\beta$  activities. However *E. koreanum* was the most distinct genetically, chemically and in their ER $\alpha$  and ER $\beta$  activities, suggesting that this well-defined species may have value as the source of estrogenic medicines with consistent activity.

#### 4. Concluding remarks

Legend has it that *Epimedium* received its colloquial name, “horny goat weed”, when goats grazing on the herb were observed to have excessive copulating behaviors. In contrast to reports that *Epimedium* extracts may elicit penile erections in rats (Chen and Chiu, 2006), we found extracts of *Epimedium* to be strong activators of ER $\alpha$  and ER $\beta$  but not the androgen receptor. Like other dietary supplements, *Epimedium* is currently sold to consumers in dozens of different preparations. In many preparations, the source materials, chemical contents and bioactivities are not known. We have examined the estrogenicity of ethanol extracts of taxonomically identified *Epimedium* species prepared under standardized conditions, established a reproducible hierarchy of ER $\alpha$  and ER $\beta$  bioactivities, and defined their relationships to phylogenetic origins and chemical content. Although more specimens from each of the *Epimedium* species would be required to completely validate our conclusions, nevertheless our approach, combining taxonomic, genetic chemical and bioresponse profiling, may be generally relevant for the standardization of botanical extracts affecting processes mediated by steroid/nuclear receptors. This integrated approach to the standardization and authentication of herbal raw materials, and products derived from them, may lead to botanical medicines of pharmaceutical quality.

#### 5. Experimental

##### 5.1. General experimental procedures

Genomic DNA was isolated using the DNeasy Plant Mini Kit (QIAGEN, Hilden, Germany). Pre-amplification and amplification steps were performed according to instructions of AFLP Plant Mapping Kit (Applied Biosystems, Foster City, USA). Capillary electrophoresis was per-

formed with an ABI PRISM 3730xl and size of DNA fragments determined with GeneMapper 3.7 software (Applied Biosystems, Singapore). Chromatographic separation was performed on a Cadenza CD-C18 column (150  $\times$  2 mm, Imatakt, Japan). Concentrations of major constituents and minor bioactive components were quantified using an Agilent 1100 LC/MSD system (Ion-trap, Agilent, Germany). Major flavonol glycosides icariin, epimedin A–C were purchased from Chromadex, USA. Bioactive flavonoids, luteolin, apigenin, kaempferol were purchased from Fluka Chemie AG, Germany; and quercetin from Sigma, Switzerland. Isolation and characterization of brevivflavone B have been described (Yap et al., 2005).

##### 5.2. *Epimedium* raw materials

Specimens of *Epimedium* were collected under the supervision of one of the coauthors (B.L.G.). Reference specimens were archived at Institute for Medicinal Plant Development, Chinese Academy of Medical Sciences, Beijing (Guo and Xiao, 2003). Sample collection was done at spring time, when the corolla characteristics of flower petals can be used to aid identification. Some samples were obtained from commercial sources and subsequently taxonomically identified (B.L.G.). Dried *Epimedium* leaves were grounded into powder and soaked in 100% ethanol (1:10, wt/vol) at 37 °C for 7 days, filtered and the supernatant dried *in vacuo*. The dried ethanol residue were weighed and re-dissolved in DMSO to give various concentrations for chemical and bioactivity profiling.

##### 5.3. Genomic DNA amplification and fluorescent AFLP analyses

Genomic DNA was isolated from 2 to 5 g of dried leaves and DNA quantity and quality verified using 1% agarose gel electrophoresis. Plant DNA ( $\sim$ 0.5  $\mu$ g) was digested with *Eco*RI and *Mse*I and ligated to adaptor. For each sample, four specific PCR amplifications were performed with *Eco*RI-ACA/*Mse*I-CACC (B2 + C), *Eco*RI-AGG/*Mse*I-CACC (B7 + C), *Eco*RI-ACT/*Mse*I-CAC (B1) and *Eco*RI-ACT/*Mse*I-CTG (G1). The *Eco*RI primers were labeled with the fluorescent dye FAM (blue) or HEX (green) at the 5' end. After mixing with DNA size standards (50–500 bp) labeled with ROX (red), amplified fragments were resolved in an Applied Biosystems 3730xl DNA analyser. Fragment data were collected by using GENEMAPPER 3.7 software.

##### 5.4. Chemical profiling

The concentrations of major constituents and minor bioactive components were quantified with liquid chromatography tandem mass spectrometry. Following chromatographic separation, concentrations of major flavonoid glycosides (icariin, epimedin A, B, C) were determined under full scan mode using 4-hydroxybenzophenone as

internal standard (RSD < 5%). Minor bioactive flavonoid aglycones were determined by multiple-reaction monitoring ( $m/z$  269  $\rightarrow$  225 for apigenin,  $m/z$  285  $\rightarrow$  185, 239 for kaempferol,  $m/z$  285  $\rightarrow$  175 for luteolin,  $m/z$  301  $\rightarrow$  151, 179 for quercetin,  $m/z$  437  $\rightarrow$  351 for brevipflavone B) (RSD < 8%). The MSD settings for the maximum signal of the selected ion pairs for the each of the standard compounds were optimized by flow injection analysis. Concentrations for each bioactive compound were quantified using a six-point calibration curve of peak area ratio for compound to internal standard against the concentration. The concentrations of each compound were determined on three separate occasions. Results are the mean of these determinations and concentrations expressed as g per 100 g of original dried ethanol extract.

### 5.5. Measurement of estrogenic activity using stably transfected ER $\alpha$ and ER $\beta$ -responsive cell lines

#### 5.5.1. Plasmids

pEGFP-ER $\alpha_{\text{neo}}$  and pEGFP-ER $\beta_{\text{neo}}$  were constructed by excising the full length hER $\alpha$  or hER $\beta$ , and inserting into the eukaryotic pEGFP-N2 expression plasmid (Clontech, Palo Alto, CA) with neomycin resistance gene. The pERE $_4$ -Luc $_{\text{hygro}}$  reporter gene consists of four tandem copies of consensus vitellogenin ERE cDNA (5'-AGGTCA-CAGTGACCT-3') cloned into a pGL3-basic plasmid (Promega, Madison, WI), upstream of the luciferase reporter gene. The hygromycin resistance gene was inserted into the vector as selection marker. All plasmids were sequenced to check fidelity of construction. Transient transfections for ER $\alpha$ , ER $\beta$ , androgen, progesterone and glucocorticoid receptors were performed as described previously (Lim et al., 2006).

#### 5.5.2. Determination of LD $_{90}$ for hygromycin and neomycin

HeLa (ATCC) cells plated in 96-well plates with a cell density of 10,000 cells/well were exposed to increasing concentrations of hygromycin B (Stratagene, CA) or neomycin (G418 sulfate, Stratagene). HeLa cells exposed to vehicle alone served as controls. After 10 days of incubation in antibiotics containing media, the number of viable cells was determined by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide reduction assay to draw the kill curves and determine the LD $_{90}$  for hygromycin B and neomycin.

#### 5.5.3. Stable transfection and clone selection

HeLa cells plated in 24-well plates were sequentially transfected with pERE $_4$ -Luc $_{\text{hygro}}$  and pEGFP-ER $\alpha_{\text{neo}}$ . Selection of stably transfected clones was done with LD $_{90}$  concentration of hygromycin B and neomycin respectively. Hygromycin/neomycin-resistant foci were identified, subcultured and screened for estradiol-induced luciferase activity. A highly inducible clone  $\alpha$ C3, stably expressing ER $\alpha$ , and  $\beta$ C3, stably expressing ER $\beta$  were obtained and used for subsequent experiments.

#### 5.5.4. ER $\alpha$ and ER $\beta$ -responsive bioassays

ER $\alpha$  and ER $\beta$  stable cells were plated in 24-well plates at an optimized density of  $4 \times 10^4$  cells in RMPI medium supplemented with 10% dextran-coated charcoal-stripped fetal bovine serum for 12 h. After 30 h of incubation with test samples, luciferase activity of lysates was measured with luminometer. Each data point was mean  $\pm$  s.e. of at least triplicate samples. Dose–response curves were performed for each extract to determine maximal and EC $_{50}$  values.

#### 5.6. Data and statistical analyses

For AFLP data analysis, electrophoretic patterns were converted into binary matrixes (1 for presence, 0 for absence of a band). Jaccard similarity matrix (number of shared bands/total number of bands) was calculated for each pair-wise comparison with the software package NTSYSpc 2.1 (Exeter software, NY). The same software was used to generate a neighbor joining phylogenetic tree. Phylogenetic consistency was evaluated by bootstrapping analysis of 1000 replicates with neighbor-joining search by using PAUP\* 4.0b10 (Sinauer, MA). Comparisons of differences in chemical contents between genetic groups were performed with Student's *t*-test after ANOVA.  $P < 0.05$  was regarded as significant. Regression analysis was performed using SPSS (SPSS Inc., Chicago, IL). Maximum activity ( $C_{\text{max}}$ ) and the dose that elicited half maximum activity (EC $_{50}$ ) were calculated from dose–response curves using Graphpad prism V4. In those instances where there was insufficient extract for a full dose response curve, maximum activity was measured at a dose of 50  $\mu\text{g}$  /ml.

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