IDENTIFICATION OF CYP1A2 AS THE MAIN ISOFORM FOR THE PHASE I HYDROXYLATED METABOLISM OF GENISTEIN AND A PRODRUG CONVERTING ENZYME OF METHYLATED ISOFLAVONES

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ABSTRACT:

This study determined the cytochrome P450 (P450) isoforms responsible for metabolism of isoflavones using human liver microsomes (HLM) and expressed P450s. The primary metabolite of genistein is 3'-OH-genistein, as identified with an authentic chemically synthesized standard. CYP1A2 was predominantly responsible for 3'-OH-genistein formation since its formation was inhibited (50%, p < 0.05) by a monoclonal antibody specific for CYP1A2, which was correlated with CYP1A2 activities of HLM, and was catalyzed by expressed CYP1A2. In addition to CYP1A2, CYP2E1 also catalyzed, although to a lesser extent, its formation. The contribution of these P450s to the formation of 3'-OH-genistein was confirmed with a panel of expressed enzymes. Methylated isoflavones biochanin A, prunetin, and formononetin (10–100 μM) were rapidly converted by HLM and expressed CYP1A2 to more active genistein and daidzein. The conversion of biochanin A to genistein appears to be mainly mediated by CYP1A2 because of the strong correlation between the conversion rates and CYP1A2 activities in HLM. Thus, CYP1A2 is an effective prodrug-converting enzyme for less active methylated isoflavones. CYP1A2-catalyzed conversion of biochanin A to genistein (Km, 7.80 μM; Vmax, 903 pmol/min/mg of protein; Vmax/Km, 116 μl/min/mg of protein) was much faster than 3'-hydroxylation of genistein (Km, 12.7 μM and Vmax, 109 pmol/min/mg of protein; Vmax/Km, 8.6 μl/min/mg of protein). The interaction studies showed that genistein inhibited formation of acetaminophen from phenacetin with an IC50 value of 16 μM. Additional studies showed that phenacetin and genistein were mutually inhibitory. In conclusion, CYP1A2 and CYP2E1 metabolized genistein and CYP1A2 acted as prodrug-converting enzymes for other less active methylated isoflavones.

The consumption of an isoflavonoid-rich diet has been associated with a decrease in the incidence of hormone-related cancers (e.g., breast and prostate cancer) (Kurzer and Xu, 1997; Birt et al., 2001; Yang et al., 2001 and references therein). Isoflavonoids, which include aglycons or isoflavones and their glucosides, are generally classified as phytoestrogens. The latter are plant-derived estrogen-like molecules. An important isoflavonoid is genistein, which has been extensively studied and widely publicized in the popular media for its importance for understanding the biological functions of isoflavones. Recently, several phase I metabolites of genistein, which are mainly hydroxylated metabolites, have been identified in vitro and in vivo (Kulling et al., 2000, 2001, 2002). Additionally, formononetin and biochanin A were shown to be converted to daidzein and genistein following oral administration to humans (Setchell et al., 2001). Currently, little is known about the specific
P450s isomers responsible for phase I metabolism of isoflavones. A preliminary studies published by Roberts-Kirchhoff et al. (1999) identified the formation of hydroxylated metabolites as a P450-catalyzed reaction. However, much important information about phase I metabolism of isoflavones is not available. For example, it is not known if P450 convert methylated isoflavones such as formononetin and biochanin A to more active daidzein and genistein. The goals of this study are to characterize the P450 isoform(s) responsible for the hydroxylation of genistein and the O-demethylation of its isoflavone analogs, and to determine the potential for drug-isoflavone interactions.

Materials and Methods

Materials. Human liver microsomes (HLM) were purchased from BD Gentest Corp. (Woburn, MA). Expressed P450s were obtained from BD Gentest Corp. Genistein, daidzein, biochanin A, formononetin, prunetin, and 4′,7-dimethoxyisoflavones (DMIF) (see Table 1 for structures) were purchased from Indofine Chemical Co. Inc. (Somerville, NJ). Hydroxylated metabolites of genistein, 2′-OH-genistein and 3′-OH-genistein were synthesized as described (see Appendix). Inhibitory monoclonal antibodies (mAbs) used in this study have been previously characterized (Gelboin et al., 1995; Krausz et al., 1997; Yang et al., 1998), and were used for single and combinatorial inhibitor studies. Monoclonal antibodies used were mAb1-599-16 (against CYP1A1), mAb26-7-5 (against CYP1A2), mAb151-45-4 (against CYP2A6), mAb49-10-20 (against CYP2B6), mAb281-1-1 (against CYP2C8), mAb1-74-4-8 (against CYP2C9), mAb 1-68-11 (against CYP2C), mAb512-1-8 (against CYP2D6), mAb2-106-12 (against CYP2E1), and mAb3-29-9 (against CYP3A4). The control mAb was Hy Hel raised against egg lysozyme. Other chemicals, generally reagent grade or better, were used as received.

Miscroscopic Incubation Experiments. A typical assay contained 25 to 50 pmol expressed P450 or 150 to 300 pmol P450 in HLM in 0.5 ml 100 mM, pH 7.5, K2 HPO4 buffer. For incubations with mAbs, 15–30 μl (10–30 μg of ascites protein) of ascites fluid was added to the enzyme and allowed to incubate for 5 min at 37°C. Monoclonal antibody Hy Hel, anti egg lysozyme (Gelboin et al., 1998), was used as a control. For combinatorial antibody inhibition studies, total protein content was kept at approximately the same level by using appropriate amounts of mAb Hy Hel. Substrate (60 μM final concentration, unless otherwise indicated) in 10 μl of methanol was added with additional buffer to a final volume of 0.5 ml. The oxidative metabolism reaction was initiated with the addition of NADPH (1 mM final concentration) or an NADPH-regenerating system (BD Gentest Corp.). The reaction proceeded for 30 min at 37°C in a shaking water bath (100 rpm). 7-OH-comarin (10 nmol), used as an internal standard, was added and the metabolites were extracted with 8 ml of CH2 Cl2 which ensured >85% recovery. For 3′-OH-genistein, an additional extraction with 4 ml of CH2 Cl2 yielded >85% recovery.

The conditions for microsomal incubation were described in the text under Materials and Methods. Briefly, the microsomes were incubated with pooled human liver microsomes purchased from BD Gentest Corp. for 45 min at 37°C in the presence of NADPH. The sample was extracted with methylene chloride, the organic phase was dried, and the dried sample was reconstituted and injected into a HP 1050 system and eluted with a gradient method as described in the text. M1 and M2 are two metabolites of genistein, and the internal standard was 7-OH-comarin.

HPLC Analysis. The metabolic samples were analyzed by HPLC using Agilent 1050 system running ChemStation software that was consisted of a quaternary pump, a diode array or variable wavelength detector and an autosampler (Agilent Technologies Inc., Wilmington, DE). The metabolites were separated using a Zorbax SB-C18 (4.6 mm × 150 mm) column (Agilent) with the following conditions: flow rate, 1 ml/min; wavelength, 260 nm; mobile phase A, acetic acid in water (pH 3.3); mobile phase B, acetonitrile; gradient, 0 to 20 min, mobile phase B 15 to 35%, linear gradient; 20 to 30 min, mobile phase B 35–45%, linear gradient. We allowed 5-min re-equilibrium time between injections. The retention time of these compounds together with their chemical structures are listed in Table 1.

Results

Metabolism of Genistein by HLM. The metabolic pattern of genistein was determined with HLM, which resulted in two major peaks that eluted earlier than genistein (designated as M1-M2) in the HPLC diagram (Fig. 1). This pattern is similar to those reported by earlier studies (Roberts-Kirchhoff et al., 1999; Kulling et al., 2000; 2001). Additional hydroxylated metabolites of genistein, which elutes earlier than genistein, have been previously reported (Kulling et al., 2000, 2001). Therefore, we did not further pursue the identities of these minor metabolites. Instead, we focused on the enzymatic and kinetic characterization of the P450 isomers responsible for the formation of the primary phase I metabolite of genistein, M1.

Confirmation of Structures of M1. The M1 peak was purified from HPLC, and injected into a liquid chromatographer/mass spectrometer with atmospheric pressure ion-spray (not shown). The metabolite was found to be either 2′-OH or 3′-OH genistein (not shown). Authentic standards of 2′-OH-genistein and 3′-OH-genistein were synthesized (see Appendix). Comparison of the metabolites formed using HLM with the metabolites synthesized chemically indicated that M1 is 3′-OH genistein and not 2′-OH genistein. An earlier study of the hydroxylation of genistein did not show which metabolite (3′-OH genistein or 2′-OH genistein) was formed by CYP1A2 (Roberts-Kirchhoff et al., 1999).

Effect of mAbs on Genistein Metabolism by HLM. The contribution of each P450 isomar was determined with a panel of well characterized inhibitory monoclonal antibodies or mAbs (Gelboin et al., 1999) against various human P450 isomers (Fig. 2). When they
were added singly, it was found that mAb 26-7-5 against CYP1A2 was the most effective inhibitor of 3'-OH-genistein formation \((p < 0.05)\). Monoclonal antibodies against CYP2E1 also significantly inhibited \((p < 0.05)\) the formation of 3'-OH-genistein but to a lesser extent than mAb 26-7-5 against CYP1A2 (Fig. 2). Monoclonal antibodies against CYP2C8, CYP2B6, CYP2A6, and CYP3A4 caused minor inhibition of formation of 3'-OH-genistein, whereas mAbs against CYP1A1, CYP2C19, and CYP2D6 did not inhibit the metabolism of genistein into 3'-OH-genistein.

A combinatorial mAb approach was used to determine the contribution of liver P450 isoforms (Fig. 3). The results indicated that mAb against CYP1A2 inhibited metabolism by about 50% \((p < 0.0005)\). Addition of mAb against CYP2E1 further inhibited remaining metabolism by about 50%, which represented 75% decrease when compared with the control \((p < 0.05)\). Addition of mAb against CYP2A6 or CYP2C8 did not result in additive inhibition \((p > 0.05)\). The addition of mAbs against both CYP2C8 and CYP3A4 further inhibited the remaining metabolism by about 36% \((p < 0.05)\), which was 87% decrease when compared with the control. Therefore, the combinatorial addition of the four relevant mAbs inhibited metabolism of genistein to 3'-OH-genistein by 87% \((p < 0.01)\).

Metabolism of Genistein by Expressed Human CYP1A2. Expressed CYP1A2 formed 3'-OH-genistein as the primary metabolite formed. Formation of 3'-OH-genistein was measured as a function of time (0, 15, 30, 45, 60, and 90 min) at a genistein concentration of 2 \(\mu\)M, and the results indicated the formation was approximately linear with time for up to 60 min (not shown).

Formation of 3'-OH-Genistein via Other Expressed P450 Isoforms. We also determined the formation of 3'-OH-genistein using other expressed P450 isoforms (25–50 pmol each). Expressed CYP1A1, CYP2E1, and CYP1B1 formed substantial amounts of 3'-OH-genistein (Fig. 4). Expressed CYP2D6, CYP2C8, and CYP2C19 formed small amounts of 3'-OH-genistein, whereas other expressed P450s including CYP3A4 and CYP3A5, the major P450 in the liver and intestine, did not form 3'-OH-genistein (Fig. 4).

Correlation between HLM 1A2 Activity and Formation Rate of 3'-OH-Genistein. Further definition of the contribution of CYP1A2 in the formation of genistein metabolites was accomplished with 11 preparations of liver microsomes [HLM pool (lot 1 and lot 2), HG32, HG42, HG89, HG93, HK21, HK23, HK43, HL56, HL112 from BD Gentest Corp.] containing different CYP1A2 activities. We found a clear linear relationship \((r^2 = 0.92)\) between CYP1A2 activity of the liver microsomes and the formation rate of 3'-OH-genistein (Fig. 5). We also plotted formation rates of 3'-OH-genistein against CYP2E1, CYP2A6, CYP3A4, CYP2C8, and CYP2C9 activities. None of these activities correlated with formation rates of 3'-OH-genistein except there was a weak correlation between formation rates of 3'-OH-genistein and CYP2E1 activities \((r^2 = 0.23)\). Further studies were focused on CYP1A2-catalyzed metabolism of genistein and its isoflavone analogs.

Determination of the Kinetic Parameters of 3'-OH-Genistein Formation by CYP1A2. We determined initial rates of 3'-OH-genistein and CYP2E1 activities \((r^2 = 0.23)\). Further studies were focused on CYP1A2-catalyzed metabolism of genistein and its isoflavone analogs.
CYP1A2-MEDIATED METABOLISM OF ISOFLAVONES

CYP1A2-Activation of Methylated Isoflavones. We determined whether CYP1A2 transform methylated isoflavones (Table 1) to genistein or daidzein through metabolic activation by CYP1A2. We used biochanin A, prunetin, formononetin, daidzein, 4',5-dimethoxyisoflavone to incubate with expressed CYP1A2. We found rapid conversion of biochanin A to genistein (see Fig. 6) and very slow formation (barely detectable but not quantifiable) of genistein from daidzein (100 μM). Prunetin (40 μM) was also rapidly converted and its rate of conversion (340 ± 47 pmol/min/mg of protein) was about 60% less than the conversion rates of biochanin A (40 μM) to genistein. At lower concentrations, the rate of conversion of prunetin to genistein (231 ± 5 pmol/min/mg of protein) was similar to that of biochanin A to genistein (202 ± 4 pmol/min/mg of protein). Formononetin was rapidly converted to daidzein and genistein, whereas 5,4'-dimethoxyisoflavone was rapidly converted to the daidzein but poorly converted to genistein. The rate of conversion from formononetin (100 μM) to genistein (295 ± 27 pmol/min/mg of protein) was similar to its conversion to daidzein (393 ± 68 pmol/min/mg of protein). The latter was approximately 40% slower than the conversion rate of biochanin A to genistein but was approximately 80 times greater than the conversion rate of 5,4'-dimethoxyisoflavone (100 μM) to daidzein. 5,4'-dimethoxyisoflavone (100 μM) was mainly converted to formononetin at a rate 4 times faster than the rate of conversion from biochanin A to genistein.

HLM-Activation of Methylated Isoflavones. Biochanin A, formononetin, and 5,4'-dimethoxyisoflavone at a high concentration (100 μM each) and at a low concentration (10 μM) were also incubated with HLM. At high isoflavone concentration, and the rate of metabolism was found in descending order as 5,4'-dimethoxyisoflavone (to formononetin) > biochanin A (to genistein) > 5,4'-dimethoxyisoflavone (to daidzein) > formononetin (to daidzein) ~ formononetin (to genistein). The only change in the order, with comparison to using an expressed CYP1A2, was that 5,4'-dimethoxyisoflavone was metabolized to daidzein faster than formononetin to daidzein in HLM. At low isoflavone concentration, biochanin A, prunetin, and formononetin were converted to genistein, with the rates of conversion of 232 ± 5, 202 ± 4, and 52 ± 1 pmol/min/mg of protein for prunetin, biochanin A, and formononetin, respectively. Formononetin was also converted to daidzein (40 ± 2 pmol/min/mg of protein) at a rate comparable with its conversion rate to genistein.

We then determined the rates of biochanin A conversion to genistein in a panel of nine human liver microsomes and plot them against the CYP1A2 activities supplied by the vendor (Fig. 7). The results indicated that the rates of conversion strongly correlated with the CYP1A2 activities (r² = 0.84, Fig. 7), but weakly correlated with CYP2E1 (r² = 0.45) and CYP2C9 (r² = 0.34) activities (not shown). These rates of conversion did not correlate with other P450 activities supplied by the vendor (BD Gentest Corp.).

Determination of Kinetic Parameters of CYP1A2-Catalyzed Metabolism of Biochanin A. We observed rapid conversion of biochanin A to genistein by CYP1A2. Similar to metabolism of genistein to 3'-OH-genistein, the metabolic rate of biochanin A (measured at 2, 5, 10, 20, 40, 70, and 100 μM) into genistein was also concentration dependent with a Kₘ value of 7.80 μM and a Vₘₐₓ value of 903 pmol/min/mg of protein (Fig. 6). The ratio of Vₘₐₓ/Kₘ, which represents intrinsic clearance, is 116 μl/min/mg of protein. The Eadie-Hofstee plot shows that the involvement of one major enzyme (linear regression generated r² of 0.89).

Interactions between CYP1A2-Catalyzed Metabolism of Genistein and Phenacetin. We determined the potential for drug interaction involving genistein and phenacetin, a typical substrate of CYP1A2. Formation rates of acetaminophen, the major metabolite produced by metabolism of phenacetin using expressed CYP1A2, was decreased in the presence of increasing concentration of genistein (Fig. 8). The IC₅₀ value was approximately 16 μM (Fig. 8). The interaction between phenacetin and genistein is mutually inhibitive in that increasing concentration of phenacetin also inhibited hydroxylation of genistein (10 μM). At a phenacetin concentration of 10 μM, the formation rate of 3'-OH-genistein was inhibited 56 ± 5%; at 20 μM, it was inhibited 93 ± 3%; and at 40 μM, it was inhibited 96 ± 4%.

Discussion

Phase I metabolism of isoflavones is an attractive target for metabolism studies since phase I biotransformation may generate metabolites that are more potent than the parent compounds (van Acker et al., 2000 and references therein). Recently, it has been shown that O-demethylation and hydroxylation were viable in vivo pathways for isoflavone activation, because less active biochanin A and formononetin were converted into genistein and daidzein (Setchell et al., 2001; Kulling et al., 2002).

The objectives of this study were to determine the primary P450 isoform responsible for the oxidative metabolism of isoflavones in vitro and to determine the potential for drug-isoflavone interactions. The results indicated that human liver microsomal P450 and expressed CYP1A2 were capable of metabolizing genistein into 3'-OH-genistein and converting methylated isoflavones into more biologically active genistein and daidzein. The results also indicated mutual inhibition of metabolism between genistein and phenacetin. Linear correlations between CYP1A2 activities and 3'-OH-genistein formation rates and between CYP1A2 activities and conversion rates of biochanin A to genistein using a panel of human liver microsomal samples supported this conclusion. Substantial inhibition of 3'-OH-genistein formation by specific mAbs validated the role played by P450 and CYP1A2 in the in vitro metabolism of genistein.

This study demonstrated that oxidative metabolism of genistein to 3'-OH-genistein is mainly by CYP1A2, and to a less extent by CYP2E1 in human liver microsomes. CYP2C8 and CYP3A4 may provide a minor contribution to its formation (Fig. 2). This con-
conclusion is supported by the fact that the combined use of a panel of mAb to CYP1A2, CYP2E1, CYP2C8, and CYP3A4 inhibited the formation of 3'-OH-genistein by 87% (Fig. 3). This conclusion is also supported by the fact that the metabolism rate of genistein (into 3'-OH-genistein) correlated with the CYP1A2 activities in the human liver microsomes (Fig. 5). Expressed CYP1A2-catalyzed metabolism of genistein also confirmed the hypothesis that CYP1A2 is primarily responsible for the formation of 3'-OH-genistein, the major phase I metabolites of genistein in human liver microsomes (Fig. 4, Fig. 6). Secondary role played by CYP2E1 is also well supported in that expressed CYP2E1 formed 3'-OH-genistein, and inhibitory mAb against CYP2E1 inhibited formation of 3'-OH-genistein. There was also a significant albeit weak correlation between CYP2E1 activities and formation rates of 3'-OH-genistein (not shown). In addition to CYP1A2 and CYP2E1, CYP1A1 and CYP1B1, which are poorly expressed in human liver, may also contribute to the formation of 3'-OH-genistein in vivo and perhaps at the target sites other than liver, but further study is necessary to confirm their roles. Roles played by other P450 isoforms such as CYP2D6 and CYP2A6 may also need to be defined. For example, expressed CYP2A6 also did not form 3'-OH-genistein, but inhibitory mAb against CYP2A6 showed minor inhibition of 3'-OH-genistein formation when used alone. In another instance, expressed 2D6 formed small amounts of 3'-OH-genistein, but the inhibitory mAb against CYP2D6 did not inhibit the formation of 3'-OH-genistein. Previously, these two mAbs were shown to be specific against these two P450 isoforms with minimal cross inhibition (Gelboin et al., 1997, 1999). Although further studies are needed to definitively resolve these apparent inconsistencies, it is likely that one of the reasons is due to difference in how substrates are presented to the enzyme. In an expressed system, isoflavone has only one P450 pathway, whereas in the HLM, multiple P450 pathways could exist for the same isoflavone. Furthermore, in the HLM some P450 isoform may be polymorphic, and there could be large variations in individual P450 content between different HLM samples. The use of the mAbs as metabolism inhibitors is able to more accurately identify and quantify the individual P450 isoforms involved in the metab-

Fig. 6. Kinetic characterization of CYP1A2 catalyzed metabolism of genistein to 3'-OH-genistein (panel A) and biochanin A to genistein (panel B).

$K_m$ and $V_{max}$ values for hydroxylation of genistein were 12.7 μM and 109 pmol/min/mg of protein, respectively. The $K_m$ and $V_{max}$ values for O-demethylation of biochanin A were 7.80 μM and 903 pmol/min/mg of protein, respectively. The insert in each panel shows the Eadie-Hofstee plot for each metabolic reaction. Each point represents the mean value of two determinations.
olism of isoflavones and additionally can demonstrate interindividual difference due to polymorphism and/or P450 content variations. Therefore, their use in the inhibition studies may provide a more accurate isoform contribution to metabolism, compared with single expressed P450 enzyme studies.

Further studies of the metabolism of several additional isoflavones by CYP1A2 indicated rapid O-demethylation of biochanin A and prunetin to genistein, of formononetin to genistein and daidzein, and of 5,4'-dimethoxyisoflavone to formononetin and daidzein, respectively. These results suggest that CYP1A2 may be a prodrug-converting enzyme for other less active isoflavones by metabolizing them to more active genistein and daidzein in vivo. We then determined the kinetics of CYP1A2-catalyzed 3'-hydroxylation of genistein and O-demethylation of biochanin A (Fig. 6). The resulting $K_m$ value of 12.7 $\mu$M for hydroxylation of genistein was 63% higher than a $K_m$ value of 7.80 $\mu$M for O-demethylation of biochanin A. Furthermore, the $V_{max}$ value of 109 pmol/min/mg of protein for hydroxylation of genistein was 12% of a $V_{max}$ value of 903 pmol/min/mg of protein for O-demethylation of biochanin A. Taken together, the difference in intrinsic clearance (or $V_{max}/K_m$) value (116 versus 8.6 pmol/min/mg of protein) indicates that O-demethylation of biochanin A would be more than 10 times faster than hydroxylation of genistein at concentrations of 5 $\mu$M or less. In other words, CYP1A2 is capable of activating other methylated isoflavones to genistein, which suggests that biochanin A and other isoflavones may serve as natural prodrugs of genistein. These in vitro results are consistent with in vivo results obtained by Setchell et al. (2001), and with our observations that the conversion rates of biochanin A to genistein were correlated with CYP1A2 activities in a panel of HLMs (Fig. 7). Interestingly, a popular product sold today as an herbal supplement called red clover extracts contains high percentages of biochanin A and formononetin.

We determined the potential for drug-isoflavone interactions since CYP1A2 is a major P450 isofor (13% total P450) in human liver and responsible for the metabolism of many drugs (Rendic and DiCarlo, 1997). The results indicated that genistein is a competitive inhibitor of CYP1A2 catalyzed metabolism of phenacetin to acetaminophen (Fig. 8), with an IC$_{50}$ value (16 $\mu$M) close to its $K_m$ value (13 $\mu$M) (Fig. 6). Furthermore, phenacetin also inhibited the metabolism of genistein with its IC$_{50}$ value (<10 $\mu$M) since 10 $\mu$M of phenacetin inhibited the metabolism of genistein (10 $\mu$M) by more than 50%. The potential for drug-isoflavone interaction is clearly present since the metabolism rate of phenacetin was decreased even at the lowest genistein concentrations (e.g., 5 $\mu$M). A genistein concentration in the low micromolar range is achievable in vivo (Setchell et al., 2001). Although the vast majority of the genistein is conjugated in vivo when measured at the systemic level (Kurzer and Xu, 1997; Setchell et al., 2001; Yang et al., 2001), it is entirely possible that the concentration of unconjugated genistein during its first-pass through the liver could reach that level. This is because genistein is rapidly absorbed and 40 to 60% absorbed genistein may reach liver as aglycone (Chen et al., 2003).

In conclusion, this study indicates that CYP1A2 is the main P450 isoform responsible for the formation of major phase I metabolites of genistein and for activation of methylated isoflavones to genistein and daidzein via the O-demethylation pathway. We have demonstrated the potential for drug-isoflavone interaction, but further studies are needed to show whether these interactions are pharmacokinetically relevant in vivo. The latter studies are important because many people are supplementing their diets with herbal supplements containing small amounts of isoflavones.

APPENDIX

Synthesis of 2'-OH-genistein and 3'-OH-genistein. The synthetic pathways of these two compounds were similar, and only the synthesis of 3'-OH-genistein (see Fig. 9) is shown here. This synthetic scheme was developed based on earlier work of Robertson and Suckling (1949), Helena et al. (1999), and Nabaei-Bidhendi and Banneree (1991).

4,6-Di(methoxym ethoxy)-2-hydroxyacetophenone (I). Anhydrous potassium carbonate (61.2 g, 0.45 mol) was added to chloroacetophenone (12.6 g, 0.0875 mol) in anhydrous acetonitrile (400 ml), followed by rapidly dropping a solution of methoxymethyl chloride (23 ml, 0.3 mol) in acetonitrile (70 ml). The mixture reacted at room temperature for 2 h. After the mixture was filtered to remove potassium carbonate, the filtrate was evaporated under vacuum. The residue was loaded into a column and eluted with EtOAc-petroleum ether to obtain white nee-
Fig. 9. A synthetic scheme for metabolic standards used in the present studies.

Example shown is the synthesis of 3'-OH-genistein, and a similar method was adapted to synthesize other isoflavones.

References


