Human multidrug and toxic compounds extrusion transporter 1 (hMATE1/SLC47A1) is a H⁺-coupled organic cation exporter responsible for the final step of excretion of various xenobiotics at the kidney and liver. In this study, effects of dietary constituents on hMATE1 mediated drug transport were examined to evaluate possible food–drug interactions. Bergamottin inhibited hMATE1 mediated tetraethyl ammonium transport activity, with a $K_i$ of 98.7 µM. Coumarins, flavonols, and catechin inhibited hMATE1 activity. Among 23 compounds tested, isorhamnetin was the strongest inhibitor of hMATE1 with the $K_i$ of 0.32 µM in a competitive manner. Since isorhamnetin is abundant in Ginkgo biloba that is widely used for herbal supplements, the findings suggest the potential hMATE1 related food–drug interactions.

**Key words**: multidrug and toxic compounds extrusion; food–drug interaction; isorhamnetin; multidrug and toxin extrusion (MATE); Ginkgo biloba

Food–drug interactions are increasing in interest since they have unexpected effects in drug therapy. It has been shown that the concomitant intake of grapefruit juice affects the disposition of various drugs such as calcium channel blockers including nifedipine, nitrendipine and verapamil. Bergamottin is known to be an active compound of grapefruit juice that affects the pharmacokinetics of various drugs through inhibition of CYP3A4. Recent progress of drug transporter studies with bergamottin and flavonoids revealed that multidrug transporters including P-glycoprotein and organic anion transporting polypeptide (OATP) also play important roles in food–drug interactions.

Multidrug and toxin extrusion 1 (MATE1; SLC47A1) is an electroneutral H⁺/organic cation antiporter that functions as an efflux transporter of many organic compounds into the bile and urine. Detailed kinetic studies indicated that hMATE1 recognizes various drugs such as cimetidine, quinidine, verapamil and metformin. Bergamottin is known to be an active compound of grapefruit juice that affects the pharmacokinetics of various drugs through inhibition of CYP3A4. Recent progress of drug transporter studies with bergamottin and flavonoids revealed that multidrug transporters including P-glycoprotein and organic anion transporting polypeptide (OATP) also play important roles in food–drug interactions.

**Materials and Methods**

**Chemicals**: [1-14C]-TEA (118.4 GBq/mmol) was purchased from PerkinElmer Life Sciences Inc. (Waltham, U.S.A.). Coumarins, flavonols and catechins were purchased from Extrasynthese (Lyon, France). Dulbecco’s modified Eagle’s medium (DMEM) and fetal bovine serum were obtained from Invitrogen. All other reagents were of analytical grade and commercially obtained. The chemical structures of the tested compounds are shown in Fig. 1.

**Cell Culture**: Hek293 cells were maintained at 37°C under a humidified atmosphere containing 5% CO$_2$ in DMEM supplemented with 55 mg/L sodium pyruvate, 10% heat-inactivated fetal bovine serum, 10$^5$ units/L penicillin, 100 mg/L streptomycin, and 250 µg/L fungizone. The cells were routinely subcultured by 0.04% trypsinization using trypsin–0.02% ethylenediaminetetraacetic acid (EDTA) solution.

**Establishment of Stable Cell Lines**: A plasmid carrying the cDNA of hMATE1 was described previously. This pcDNA3.1(+) hMATE1 plasmid was used to transfect HEK293 cells by lipofection using TransIT reagent (Mirus). Forty-eight hours after transfection, 800 mg/L geneticin (G418) was added to the medium. Two weeks after the addition of geneticin, a stable clone was isolated by Western blotting using anti hMATE1 antibodies.

**Expression Test**: Membranes of HEK293 cells were isolated as follows. Cultured cells (2.0×10$^7$ cells) were suspended in ice-cold 20 mM 3-morpholinopropane sulfonic acid (MOPS)-Tris, pH 7.0, containing 0.3 M sucrose, 5 mM EDTA, and protease inhibitors (pepsatin A and leupeptin at 5 µg/L each), homogenized, and then centrifuged at 800×g for 8 min at 4°C. The postnuclear supernatant was centrifuged at 10000×g for 1 h at 4°C. The pellet was suspended in the same buffer, solubilized with dissociation buffer containing sodium dodecyl sulfate (SDS), and then used for polyacrylamide gel electrophoresis in the presence of SDS. Polyacrylamide gel electrophoresis and Western blotting were performed as described previously. Protein concentrations were determined using the Bio-Rad protein assay using bovine serum albumin.
Transport Assay  TEA uptake was measured with mock and stably hMATE1 expressing HEK293 cells grown on BD polystyrene-treated 24-well plates. HEK293 cells (1.0×10^5 cells/well) were grown on 24-well plates containing DMEM supplemented with 10% fetal calf serum, 100 µg/mL penicillin, and 0.25 µg/mL streptomycin at 37°C under 5% CO2. Forty-eight hours later, the cells were washed twice with 800 µL of an assay medium (125 mM NaCl, 4.8 mM KCl, 5.6 mM D-glucose, 1.2 mM CaCl2, 1.2 mM KH2PO4, 1.2 mM MgSO4, and 25 mM Tricine, pH 8.0). Then, the cells were incubated at 37°C for 5 min, transport being initiated by adding [14C]-TEA (0.5 GBq/mmol) to a final concentration of 50 µM. After 10 min incubation at 37°C, the cells were washed twice with 800 µL ice-cold assay medium. The cells were lysed with 400 µL 0.5 M NaOH, and the radioactivity was counted with a liquid scintillation counter.

Data Analysis  For calculation of IC50 values, the percentage of uptake inhibition was calculated from the results of control experiments in the absence of the tested compounds (100% uptake). The corresponding IC50 values for the inhibition of hMATE1-mediated [14C]-TEA uptake by the tested compounds were calculated by fitting to a dose-response curve using KaleidaGraph 3.6 (HULINKS) as previously reported.22) Fitting curve was given by Eq. 1 and IC50 was obtained from Eq. 2.

\[
f(x) = \frac{1}{1 + e^{\alpha x + \beta}}
\]

IC50 = 10^{\alpha/\beta}

(1)  
(2)

All data are presented as means±S.E.M., n=3–9.

To determine the kinetic type of inhibitions (isorhamnetin and bergamottin), the uptake of [14C]-TEA was determined in the presence of increasing concentrations of the inhibitors. The K_i values were calculated using SigmaPlot 12.3 (HULINKS) by fitting data to Eq. 3.

\[
v = \frac{V_{max} \cdot [S]}{[S] + K_m(1+[I]/K_i)}
\]

(3)

Where v is the TEA transport rate; [I] and [S] are inhibitor and substrate concentrations.
Fig. 2. TEA Uptake by hMATE1 Expressing HEK293 Cells

(A) Western blot analysis with an antibody against hMATE1. Membrane fractions of hMATE1 expressing or mock HEK293 cells were analyzed by Western blotting using anti hMATE1 antibodies. Immunoactive proteins were visualized using an ECL detection kit (Amersham Biosciences). The position of hMATE1 is indicated by an arrowhead. (B) Time course of TEA uptake into HEK293 cells stably expressing hMATE1 and into mock cells. The uptake of [14C]-TEA (50 µM) was measured at the indicated time at 37°C, and pH 8.0 in the presence or absence of bergamottin (500 µM). Data are presented as means±S.E.M., n=3.

Fig. 3. Inhibition of hMATE1 Mediated TEA Uptake by Various Compounds

The uptake of [14C]-TEA (50µM) was measured for 5 min at 37°C and pH 8.0 in the presence or absence of the indicated compounds (0.1–500µM). The averaged TEA uptake for the control was 0.10±0.01 (nmol·mg protein−1·min−1). Inhibitions by bergamottin (A), imperatorin (B), quercetin (C), isorhamnetin (D), kaempferol (E) and (−)-EGCG (F) are shown. Data are expressed as percentages of the control. All data are presented as means±S.E.M., n=3–9.
RESULTS

**Bergamottin Is a Natural hMATE1 Inhibitor** To examine the effects of dietary constituents on hMATE1 activity, a stable hMATE1 expression line was established by genetic screening. The expression of hMATE1 in HEK293 cells derived from an isolated colony was examined by Western blot analysis. An immunoreactive band at 62kDa which corresponded to the molecular weight of hMATE1, was observed for the membranes of HEK293 cells expressing hMATE1 (Fig. 2A). However mock cells did not show immunoreactivity except for minor nonspecific band at ca. 80kDa. Immunoreactivity at 128kDa may correspond to hMATE1 dimer. hMATE1 activity was measured as [14C]-TEA (50µM) uptake by HEK293 cells. When cells were incubated at an alkaline pH, which generates a downhill H+ gradient across the plasma membrane, hMATE1 expressing HEK293 cells took up TEA in a time dependent manner up to 30 min (Fig. 2B). On the other hand, mock cells did not show significant TEA uptake. These results indicate that the established HEK293 cell line is useful for measuring hMATE1 activity. Initially we tested the effect of bergamottin, which is a major ingredient of grapefruit juice, regarding food–drug interactions. When 500µM bergamottin was included in the medium, TEA uptake was greatly inhibited, suggesting hMATE1 is sensitive to this compound (Fig. 2B).

**Inhibitory Effects of Natural Compounds on hMATE1 Transport Activity** Since bergamottin is a furanocoumarin, we examined the effects of various coumarins on hMATE1-mediated TEA uptake (Figs. 1, 3). Most coumarins caused dose-dependent inhibition of TEA uptake, the exceptions being dicoumarol and bergapten. These two coumarins were insoluble at higher than 10µM and thus inhibition could not be measured at higher concentrations. Bergamottin, psoralen and xanthotoxin inhibited TEA uptake to similar extents. Among the tested coumarins, imperatorin caused the strongest inhibition, the apparent IC50 being 6µM (Figs. 1, 3B).

We also tested 12 flavonols and 3 catechins. As shown in Figs. 1 and 3, isorhamnetin showed strong inhibition of TEA uptake, whereas myricetin, ombuin and quercetin tetramethylether at up to 100µM failed to inhibit TEA uptake (Figs. 1, 3). Other flavonols i.e., kaempferol, quercetin, kaempferide, galangin and morin, caused moderate inhibition (Figs. 1, 3). For catechins, TEA uptake was moderately inhibited by epigallocatechin gallate (EGCG) (Figs. 1, 3F). On the other hand, catechin did not inhibit TEA uptake at the tested concentrations.

The apparent IC50 values for these compounds are summarized in Fig. 1. Among the tested compounds, isorhamnetin was the most potent inhibitor of hMATE1 although it did not show complete inhibition even at high concentrations (Fig. 3D).

**Kinetic Analysis of the hMATE1 Inhibition by Bergamottin and Isorhamnetin** In order to elucidate the mechanism underlying the hMATE1 inhibition by bergamottin and isorhamnetin, the inhibitory effects of these compounds were measured with various concentrations of TEA and inhibitors (Fig. 4). Clearly, both bergamottin and isorhamnetin caused competitive type inhibition with K values of 98.7 and 0.32µM, respectively. This suggests these compounds share the same binding site with TEA.

**DISCUSSION**

MATE1 is a cationic drug exporter located in the apical membranes of renal tubular cells and hepatocytes. Recent pharmacokinetic studies revealed that pyrimethamine is a specific inhibitor of MATE1, the K being ca. 90 nM. Sugiyma and co-workers used pyrimethamine as a tool to determine MATE1 dependent drug efflux and showed that MATE1 is responsible for metformin efflux in human and mouse. Interestingly, they found pyrimethamine treatment increased the hepatic or plasma metformin concentrations, and caused greater gluconeogenesis inhibition, indicating a metformin–pyrimethamine interaction through MATE1. These studies indicated the importance of MATE1 in the efflux of cationic drugs and in the drug–drug interactions.

In the present study, we showed the inhibition of hMATE1 by bergamottin. Bergamottin is well known as an ingredient of grapefruit juice and causes food–drug interactions by interfering with CYP activity. Plasma bergamottin concentration after grapefruit juice intake is reported as 2–10 nM, far
lower than the observed $K_i$ value.\textsuperscript{3} Our study suggests bergamottin in food may not interfere with the pharmacokinetics of cationic drugs through hMATE1.

Inhibition of hMATE1 by bergamottin also suggests the possibility that hMATE1 is inhibited by other compounds in food. As expected, many flavonoids and coumarins inhibited hMATE1. Among 23 compounds tested, isorhamnetin was the strongest inhibitor of hMATE1. Isorhamnetin is the major flavonoid in Ginkgo biloba L.\textsuperscript{24} It is also well known as a metabolite of quercetin.\textsuperscript{25,26}\textsuperscript{26} de Pascual-Teresa and co-workers reported that the plasma isorhamnetin concentration after oral administration of quercetin-rich food is ca. 0.36 $\mu$M.\textsuperscript{27} This value is comparable to the IC\textsubscript{50} and $K_i$ values for isorhamnetin, indicating special caution is required when taking G. biloba supplements and/or quercetin-rich foods. G. biloba is used to treat mild cognitive impairment and dementia although it effect is still controversial.\textsuperscript{28,29} Since G. biloba leaves are among the most widely used herbal remedies in the United States, further investigation of G. biloba and its interaction with metabolic targets are required.\textsuperscript{28}

Although isorhamnetin and bergamottin have different structures, they inhibited hMATE1 mediated TEA uptake in a typical competitive manner. This indicates these two compounds and TEA share the same binding site.

The inhibitory effects of the tested compounds were highly dependent on the positions of the hydroxyl and methoxy groups on the flavonol. For instance, isorhamnetin and geraldol are only different in the 5-hydroxy group. However, hMATE1 is not sensitive to geraldol, indicating the importance of the 5-hydroxy group for inhibition. Isorhamnetin, kaempferol and quercetin have the same structure except for the functional group at the 3' position. Isorhamnetin has a methoxy group at this position, and shows a ca. 100 fold lower IC\textsubscript{50} than kaempferol and quercetin. When quercetin and rhamnetin are compared, it can be seen they have the same structure except for the functional group at position 7. Quercetin has a hydroxyl group at this position, whereas rhamnetin has a methoxy group and this group causes more than 10 fold weaker inhibition. Also, the importance of a hydroxyl group at the 4' position was indicated by quercetin and tamarixetin inhibition. These results indicate that hydroxy groups at positions 5, 7 and 4', and a methoxy group at position 3' are important for inhibition of hMATE1. Interestingly, isorhamnetin has hydroxy groups at positions 3, 5, 7 and 4', and a methoxy group at position 3', and thus fulfills this inhibition constraint.

CONCLUSION

In conclusion, some coumarins, EGCG, and flavonoids especially for isorhamnetin inhibited hMATE1. These compounds may affect pharmacokinetics through inhibition of food excretion. Although further in vivo and clinical studies are necessary, our findings point out the hMATE1 related risk of food–drug interactions.

Acknowledgments This study was supported in part by Grants-in-Aid for scientific research from the Ministry of Education, Culture, Sports, Science and Technology of Japan and the Smoking Research Foundation to H.I. and H.O. We thank Dr. Y. Moriyama for his help and discussion.

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