Impact of fatty acids on human UDP-glucuronosyltransferase 1A1 activity and its expression in neonatal hyperbilirubinemia

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While breast milk has been known as a cause of neonatal hyperbilirubinemia, the underlying mechanism of breast milk-induced jaundice has not been clarified. Here, the impact of fatty acids on human UDP-glucuronosyltransferase (UGT) 1A1 – the sole enzyme that can metabolize bilirubin – were examined. Oleic acid, linoleic acid, and docosahexaenoic acid (DHA) strongly inhibited UGT1A1 activity. Forty-eight hours after a treatment with a lower concentration of DHA (10 mg/kg), total bilirubin significantly increased in neonatal hUGT1 mice, which are human neonatal jaundice models. In contrast, treatments with higher concentrations of fatty acids (0.1–10 g/kg) resulted in a decrease in serum bilirubin in hUGT1 mice. It was further demonstrated that the treatment with higher concentrations of fatty acids induced UGT1A1, possibly by activation of peroxisome proliferator-activated receptors. Our data indicates that activation of peroxisome proliferator-activated receptors would increase UGT1A1 expression, resulting in reduction of serum bilirubin levels in human infants.

UDP-glucuronosyltransferases (UGTs) are phase II drug-metabolizing enzymes that catalyze glucuronidation of compounds by transferring glucuronic acid from a co-substrate, UDP-glucuronic acid, to substrates. UGTs are super-family enzymes that are mainly classified into two subfamilies, UGT1 and UGT2, on the basis of evolutionary divergence. The human UGT1 gene encodes multiple unique exons 1 and common exons 2 to 5, producing nine functional UGT1A isoforms, UGT1A1, UGT1A3, UGT1A4, UGT1A5, UGT1A6, UGT1A7, UGT1A8, UGT1A9, and UGT1A10. These isoforms exhibit substrate specificity and are expressed tissue specifically. In the liver, which is known as the most important tissue for detoxification, UGT1A1, UGT1A3, UGT1A4, UGT1A6, and UGT1A9 are expressed. Accumulating evidence indicates that in addition to the liver, extrahepatic tissues such as the small intestine and kidneys potentially contribute to metabolism of a wide variety of compounds. While UGTs metabolize various clinically used drugs, they are also involved in metabolism of endogenous compounds, such as bilirubin – a specific substrate of UGT1A1.

Increased or decreased expressions of drug-metabolizing enzymes or the presence of factors that can activate or inhibit drug-metabolizing enzymes can cause accelerated or delayed metabolism of administered drugs or endogenous substrates, which can result in severe adverse effects of the drugs or hormone imbalance. Therefore, it is important to understand the factors that affect the expression and activity of drug-metabolizing enzymes to avoid potential adverse effects. It has been known that breast-fed infants have a 3- to 6-fold greater probability of developing elevated total serum bilirubin levels than formula-fed infants. Though controversial findings were subsequently reported, Arias et al. originally found an inhibitory effect of 5β-pregnane-3α,20β-diol – an endogenous compound present in breast milk – on glucuronosyltransferase activity in vitro. A recent study by Ota et al. revealed that 5β-pregnane-3α,20β-diol inhibited enzyme activity, as well as transcriptional activity of human UGT1A1. In addition to β-pregnane-3α,20β-diol, fatty acids included in breast milk have also been reported as inhibitors of bilirubin glucuronosyltransferase activity. In 1972, Bevan and Holton demonstrated that oleic acid – one of the fatty acids highly present in breast milk – inhibited bilirubin conjugation in rat liver slices. Subsequently, Hargreaves demonstrated that while saturated fatty acids did not, C18 and C20 unsaturated fatty acids inhibited bilirubin conjugation in rat liver slices. While several studies revealed that mammalian UGTs, such as monkey UGT2B9 and human UGT2B7, possessed glucuronidation activities toward saturated and
unsaturated fatty acids in the 1990s to early 2000s\textsuperscript{14–20}, the inhibitory effects of fatty acids on human UGT enzymes were not investigated until 2004. A study by Tsoutsikos et al. demonstrated that certain fatty acids strongly inhibited 4-methylumbelliferone (4-MU) glucuronidation in human kidney microsomes, as well as in recombinant human UGT1A9 and UGT2B7\textsuperscript{21}. Inhibitory effects of unsaturated fatty acids, oleic acid and linoleic acid, on recombinant human UGT1A9-catalyzed 4-MU and propofol glucuronidation were subsequently demonstrated in 2008\textsuperscript{22}. However, inhibitory effects of fatty acids on human UGT1A1-catalyzed glucuronidation have not been investigated.

To understand the potential inhibitory effects of fatty acids on human UGT1A1, in the present study, inhibitory effects of 15 saturated and unsaturated fatty acids on the enzymatic activity of human UGT1A1 were investigated. Estradiol 3-O-glucuronide formation is one of the reactions that are selectively catalyzed by UGT1A1\textsuperscript{26}. In our study, the control activity of Estradiol 3-O-glucuronidation in UGT1A1 Supersomes was 468 pmol/min/mg at 20 $\mu$M estradiol, which was in agreement with the reported catalytic rate. Out of the 15 tested fatty acids, oleic acid, linoleic acid, DHA, EPA, palmitoleic acid, arachidonic acid, and alpha-linolenic acid strongly inhibited UGT1A1 activity with IC\textsubscript{50} values of 31.6 $\mu$M, 33.1 $\mu$M, 37.1 $\mu$M, 11.6 $\mu$M, 19.9 $\mu$M, 22.7 $\mu$M, and 26.1 $\mu$M, respectively (Fig. 1 and Table 1).

**Table 1 | IC\textsubscript{50} values of fatty acids for estradiol 3-O-glucuronidation**

| Fatty acids      | IC\textsubscript{50} values |
|------------------|----------------------------|
| Oleic acid       | 31.6 $\mu$M                |
| Linoleic acid    | 33.1 $\mu$M                |
| Palmitoleic acid | 37.1 $\mu$M                |
| Alpha-linolenic acid | 26.1 $\mu$M        |
| Arachidonic acid | 22.7 $\mu$M                |
| DHA              | 11.6 $\mu$M                |
| EPA              | 19.9 $\mu$M                |

IC\textsubscript{50} values were directly determined in the UGT1A1 recombinant system from linear regression (Fig. 1). IC\textsubscript{50} values of stearic acid, myristic acid, lauric acid, and palmitic acid are the fatty acids highly present in human breast milk\textsuperscript{23}. In the present study, the inhibitory effects of these fatty acids and 9 other fatty acids, which are moderately present in human breast milk, on the enzymatic activity of human UGT1A1 were investigated. Estradiol 3-O-glucuronide formation is one of the reactions that are selectively catalyzed by UGT1A1\textsuperscript{26}. In our study, the control activity of Estradiol 3-O glucuronidation in UGT1A1 Supersomes was 468 pmol/min/mg at 20 $\mu$M estradiol, which was in agreement with the reported catalytic rate. Out of the 15 tested fatty acids, oleic acid, linoleic acid, DHA, EPA, palmitoleic acid, arachidonic acid, and alpha-linolenic acid strongly inhibited UGT1A1 activity with IC\textsubscript{50} values of 31.6 $\mu$M, 33.1 $\mu$M, 11.6 $\mu$M, 19.9 $\mu$M, 37.1 $\mu$M, 22.7 $\mu$M, and 26.1 $\mu$M, respectively (Fig. 1 and Table 1).

**Results**

**Inhibitory effects of free fatty acids on UGT1A1-catalyzed glucuronidation.** Fatty acids present in breast milk have been considered to be the causes of breast milk-induced neonatal hyperbilirubinemia. However, inhibitory effects of fatty acids on human UGT1A1-catalyzed glucuronidation have not been investigated. Oleic acid, linoleic acid, stearic acid, myristic acid, lauric acid, and palmitic acid are the fatty acids highly present in human breast milk\textsuperscript{23}. In the present study, the inhibitory effects of these fatty acids and 9 other fatty acids, which are moderately present in human breast milk, on the enzymatic activity of human UGT1A1 were investigated. Estradiol 3-O glucuronidation is one of the reactions that are selectively catalyzed by UGT1A1\textsuperscript{26}. In our study, the control activity of Estradiol 3-O glucuronidation in UGT1A1 Supersomes was 468 pmol/min/mg at 20 $\mu$M estradiol, which was in agreement with the reported catalytic rate. Out of the 15 tested fatty acids, oleic acid, linoleic acid, DHA, EPA, palmitoleic acid, arachidonic acid, and alpha-linolenic acid strongly inhibited UGT1A1 activity with IC\textsubscript{50} values of 31.6 $\mu$M, 33.1 $\mu$M, 11.6 $\mu$M, 19.9 $\mu$M, 37.1 $\mu$M, 22.7 $\mu$M, and 26.1 $\mu$M, respectively (Fig. 1 and Table 1).

**Figure 1 | Inhibitory effects of fatty acids on estradiol 3-O-glucuronidation.** Estradiol 3-O-glucuronidation was determined in the UGT1A1 recombinant system in both the presence and absence of fatty acids. Results are duplicate determinations.
Inhibitory pattern of oleic acid, linoleic acid, and DHA on UGT1A1 activity in the recombinant system. We further determined the inhibitory pattern of oleic acid, linoleic acid, and DHA against estradiol 3-O-glucuronidation by UGT1A1. Dixon-plot analyses indicated that oleic acid and linoleic acid non-competitively inhibited UGT1A1 activity (Fig. 2A and B), showing their $K_i$ values of 23.4 μM and 22.1 μM, respectively (Table 2). DHA more potently inhibited UGT1A1 activity than oleic acid and linoleic acid, showing its $K_i$ value of 1.8 μM. The inhibitory pattern of DHA against estradiol 3-O-glucuronidation by UGT1A1 was non-competitive, the same as the patterns of oleic acid and linoleic acid (Fig. 2C and Table 2). These data indicate that oleic acid, linoleic acid, and DHA present in breast milk might be the causes of breast milk-induced neonatal hyperbilirubinemia.

Inhibitory pattern of oleic acid, linoleic acid, and DHA on UGT1A1 activity in human liver and small intestine microsomes. While recombinant systems are convenient tools to determine enzymatic activities of single drug-metabolizing enzymes, differences in the enzymatic properties between recombinant systems and other in vitro tools, such as human liver microsomes or hepatocytes, have been reported. Investigations have speculated that those differences might be attributed to the differences in composition of endoplasmic reticulum membrane, topology of enzymes, or ratio of active/inactive proteins. Although UGT1A1 is extensively expressed in various tissues, its expression levels are dramatically high in the endoplasmic reticulum membrane, topology of enzymes, or ratio of active/inactive proteins. Although UGT1A1 is extensively expressed in various tissues, its expression levels are dramatically high in the liver and small intestine. Indeed, previous studies have revealed that in addition to the liver, the small intestine plays an important role in bilirubin metabolism. Therefore, in the present study, the inhibitory effects of oleic acid, linoleic acid, and DHA on UGT1A1 activity in human liver and small intestine microsomes were examined. In human liver microsomes, Dixon-plot analyses revealed that oleic acid and linoleic acid non-competitively inhibited UGT1A1 activity with $K_i$ values of 29.3 μM and 24.0 μM, respectively (Fig. 3A-B and Table 2). DHA more potently inhibited UGT1A1 activity than oleic acid and linoleic acid, showing its $K_i$ value of 4.3 μM with a non-competitive inhibition pattern (Fig. 3C and Table 2). These data are in agreement with the finding observed in the UGT1A1 recombinant system. In contrast, oleic acid and linoleic acid slightly inhibited estradiol 3-O-glucuronidation in human small intestine microsomes (Fig. 4A). At 50 μM oleic acid and linoleic acid, the residual activities of estradiol 3-O-glucuronidation were more than 80%. However, DHA still potently inhibited intestinal estradiol 3-O-glucuronidation, showing its $K_i$ value of 5.2 μM with a non-competitive inhibition pattern (Fig. 4B and Table 2). This indicates that DHA has a potency to inhibit both hepatic and small intestinal UGT1A1 activities.

DHA inhibits bilirubin glucuronidation in vivo. Our in vitro studies revealed that DHA could be a potent inhibitor of UGT1A1 activity. To understand the impact of the fatty acid in bilirubin glucuronidation in vivo, DHA was orally administered to hUGT1 mice, which are human breast milk-induced jaundice models. HUGT1 mice develop severe hyperbilirubinemia and their serum bilirubin levels reach a peak at 14 days after birth. The model mice were orally treated with DHA (10 mg/kg) at 12 days after birth and their bilirubin levels were measured at 14 days after birth, 48 hours after the treatment. Corn oil is usually used to dissolve hydrophobic compounds; however, it was reported that treatment of hUGT1 mice with corn oil decreased serum bilirubin levels, possibly by an inhibition of enterohepatic circulation of bilirubin. Rapeseed oil, on the other hand, did not affect the serum bilirubin levels of neonatal hUGT1 mice (data not shown). Therefore, in the present study, rapeseed oil was used to dissolve DHA and also to treat mice in the control group. In accordance with the previous report, serum

Table 2 | $K_i$ values of fatty acids for estradiol 3-O-glucuronidation

| Fatty acids            | $K_i$ values |
|------------------------|--------------|
| Recombinant UGT1A1     |              |
| Oleic acid             | 23.4 μM      |
| Linoleic acid          | 22.1 μM      |
| DHA                    | 1.8 μM       |
| Human liver microsomes |              |
| Oleic acid             | 29.3 μM      |
| Linoleic acid          | 24.0 μM      |
| DHA                    | 4.3 μM       |
| Human small intestine  |              |
| Oleic acid             | -            |
| Linoleic acid          | -            |
| DHA                    | 5.2 μM       |

$K_i$ values were determined in the UGT1A1 recombinant system and liver microsomes of the human liver and small intestine (Fig. 2-4).

3.2903 | DOI: 10.1038/srep02903
bilirubin levels of neonatal hUGT1 mice in the control group increased by 2.3 ± 3.2 mg/dL from 12 days to 14 days after birth (13.6 mg/dL to 15.9 mg/dL). When DHA was orally given to the mice, the serum bilirubin levels increased by 6.6 mg/dL ± 1.3 from 12 days to 14 days after birth, which was statistically greater than the increased amount in the control group (Fig. 5). In hUGT1 mice, it was demonstrated that intestinal UGT1A1 played an important role in bilirubin metabolism. Thus, the data indicates that DHA inhibited bilirubin glucuronidation in the small intestine, causing increased serum bilirubin levels in the human neonatal jaundice model.

High concentration of fatty acids induce UGT1A1 in the liver and small intestine. To investigate the concentration-dependent effects of fatty acids on bilirubin clearance in vivo, hUGT1 mice were further treated with high doses of oleic acid, linoleic acid, and DHA. Contrary to our expectation, the fatty acids concentration-dependently decreased serum bilirubin levels (Fig. 5 and 6). Especially, oleic acid and linoleic acid dramatically decreased serum bilirubin levels, from 13.6 mg/dL to 2.4 and 1.9 mg/dL, respectively. Fatty acids have been known to activate peroxisome proliferator-activated receptors (PPARs), which are transcriptional factors of human UGTs. Indeed, a PPAR agonist, 4-Chloro-6-(2,3-xylidino)-2-pyrimidinylthioacetate Acid (WY-14643), has been shown to induce human UGT1A1. UGT1A1 mRNA expression was subsequently determined in fatty acids- and vehicle-treated mice to understand the effects of oleic acid and linoleic acid on the UGT1A1 expression. In the small intestine, oleic acid treatment resulted in 20- to 40-fold increases in UGT1A1 mRNA expression (Fig. 7A). Oleic acid also induced UGT1A1 in the liver, in which the UGT1A1 levels increased 10- to 100-fold (Fig. 7B). While linoleic acid similarly decreased serum bilirubin levels in hUGT1 mice, the induction of UGT1A1 was specifically observed in the small intestine, in which UGT1A1 levels increased 20- to 30-fold (Fig. 7C and D). These data indicate that induction of UGT1A1 in the small intestine might be responsible for the decrease of serum bilirubin in fatty acid-treated hUGT1A1 mice.

**Discussion**

Breast milk has been recognized as a leading cause of neonatal hyperbilirubinemia. A recent study demonstrated that suppression of the gastrointestinal UGT1A1 expression by breast milk led to the development of neonatal jaundice. However, inhibition of the UGT1A1-catalyzed bilirubin glucuronidation by the components of breast milk has still been considered the underlying mechanism.
for breast milk–induced jaundice. To our knowledge, this is the first study examining the inhibitory effects of fatty acids on human UGT1A1 activity, although fatty acids were reported to inhibit mammalian UGT enzymes. In the present study, seven fatty acids out of 15 tested fatty acids inhibited UGT1A1 activity with IC\textsubscript{50} values of 11.6 \( \mu \)M to 37.1 \( \mu \)M in the recombinant UGT1A1 system. Oleic acid and linoleic acid also inhibited UGT1A1 activity in the human liver microsomes. More than 30% and, to a lesser amount, 7% of the free fatty acids present in human breast milk are oleic acid and linoleic acid, respectively\textsuperscript{27}. Therefore, those fatty acids could inhibit UGT1A1 activity in breast-fed infants, contributing to elevation of serum bilirubin levels in infants. Interestingly, DHA significantly inhibited the UGT1A1 activity not only in human liver microsomes, but also in small intestine microsomes, with \( K_m \) values of 4.3 \( \mu \)M and 5.2 \( \mu \)M, which were much lower than the values of oleic acid and linoleic acid. In the human neonatal hyperbilirubinemia model, intestinal UGT1A1 plays an important role in bilirubin metabolism\textsuperscript{23}. The inhibitory effects of DHA in UGT1A1-catalyzed bilirubin glucuronidation was further confirmed in our \textit{in vivo} study with \textit{hUGT1} mice (Fig. 5). The amount of DHA in human breast milk is less than 1% of the total free fatty acids in breast milk\textsuperscript{27}. Given the strong inhibitory effects of DHA toward intestinal UGT1A1 activity, however, not only oleic acid and linoleic acid, but DHA could also be a factor contributing to the onset of breast milk–induced neonatal jaundice.

To examine whether our observation of the \textit{in vivo} study was valid or not, the inhibitory effects of DHA on bilirubin clearance in the \textit{hUGT1} mice were estimated using equations described in the methods section (Eq 1–4). The \( K_m \) value of bilirubin glucuronidation by human UGT1A1 was reported to be around 2 \( \mu \)M\textsuperscript{25}. Because the serum free bilirubin concentration is much lower than the \( K_m \) value, Equation 4 can be used to estimate the inhibitory effects of DHA on bilirubin clearance. With the parameters of \( K_m \) value (5.2 \( \mu \)M) and concentration of DHA used for the oral treatment (0.5 mg/mL), the R value was calculated to be 0.0034 (Equation 4). This maximum inhibitory effect of DHA on bilirubin clearance would be temporal due to the rapid absorption of DHA into the blood stream. However, the increased serum bilirubin levels by DHA in \textit{hUGT1} mice was considered to be attributed to the inhibition of bilirubin glucuronidation in the small intestine, because in the \textit{hUGT1} mice, hepatic UGT1A1 is not functional due to its very low expression in the liver\textsuperscript{24}. In humans, breast milk of mothers contains on average 3.2% total fat and the DHA concentration of human breast milk was reported to be 0.06% of total fat\textsuperscript{24}. This indicates that the concentration of DHA in human breast milk is around 60 \( \mu \)M, giving an R value of 0.08 (Equation 4). Breast-fed human infants constantly drink breast milk from their mothers; therefore, inhibition of bilirubin glucuronidation in the small intestine by DHA would be the cause of breast milk–induced jaundice. Interindividual variability in the DHA content in breast milk\textsuperscript{24} might explain the difference of serum bilirubin levels in human infants.

A key question was why higher concentrations of DHA or other fatty acids did not increase, but rather decreased serum bilirubin levels in \textit{hUGT1} mice. Our \textit{in vivo} study demonstrated that UGT1A1 expression was significantly induced with treatment of a high concentration of fatty acids, possibly by activating PPARs. Therefore, it is considered that in conditions in which excessive amounts of fatty acids were treated, the inducible effect of fatty acids on UGT1A1 overcame their inhibitory effects on UGT1A1 activity. In physiological conditions, however, DHA in breast milk would not be involved in induction of UGT1A1, because it was demonstrated that breast-fed \textit{hUGT1} mice fed with formula in which DHA was not included\textsuperscript{23}. Clinical intervention to treat episodes of severe hyperbilirubinemia calls for extended phototherapy treatment or even blood transfusion. While exchange transfusions using the umbilical vein and phenobarbital treatments are now becoming extinct, phototherapy also has disadvantages; for example, increased insensible water loss, skin rashes, pyrexia, decreased maternal-infant interaction, and lack of visual sensory input are potential disadvantages of this treatment. Unlike the constitutive androstane receptor, which is activated by phenobarbital, PPARs can be activated by various natural compounds such as saturated fatty acids\textsuperscript{36}, which do not function as UGT inhibitors (Table 1). Therefore, supplemental treatment of mothers or direct treatment of newborn infants with natural ligands for PPAR would be an alternative method to increase UGT1A1 expression to accelerate bilirubin conjugation in infants.

It should be noted that oleic acid and linoleic acid did not inhibit estradiol 3-\( \beta \)-glucuronidation in human small intestine microsomes, although they inhibited the reaction in the UGT1A1 recombinant system and in human liver microsomes. Among functional human UGT enzymes, UGT1A1, UGT1A3, UGT1A8, UGT1A10, UGT2A1, UGT2A2, and UGT2B15 exhibit estradiol 3-\( \beta \)-glucuronide formation activities\textsuperscript{40}. Due to the high expression levels of UGT1A1 among those UGT isoforms in the human liver, 3-\( \beta \)-glucuronidation of estradiol has been used as a probe reaction of UGT1A1 in the human liver\textsuperscript{40}. Because oleic acid and linoleic acid inhibited the 3-\( \beta \)-glucuronidation
of estradiol in the UGT1A1 recombinant system, the inhibitory effects of oleic acid and linoleic acid against estradiol 3-O-glucuronidation in liver microsomes were considered to be attributed to the inhibition of UGT1A1. However, the expression pattern of UGT enzymes in other tissues is different from that in the liver. In the small intestine, not only UGT1A1, but also other UGT enzymes such as UGT1A8 and UGT1A10 are highly expressed. UGT1A8 shows slightly less 3-O-glucuronidation activity toward estradiol compared with UGT1A1. On the other hand, UGT1A10 shows a nearly 10-fold greater V_{max} of estradiol 3-O-glucuronidation than UGT1A1. These suggest that estradiol 3-O-glucuronidation might be mainly catalyzed by UGT1A10 in the small intestine, in which oleic acid and linoleic acid weakly inhibit UGT1A10. Bilirubin, in contrast, is a specific substrate of UGT1A1. Therefore, it is still possible that oleic acid and linoleic acid inhibit UGT1A1-catalyzed bilirubin glucuronidation in the small intestine, contributing to breast milk-induced neonatal hyperbilirubinemia. It will be important to determine whether oleic acid and linoleic acid inhibit UGT1A1 in the human small intestine with a UGT1A1 specific substrate, such as bilirubin. In this study, however, we were unable to conduct UGT1A1 enzyme assay with bilirubin due to its lack of stability against light. Although inhibition study with bilirubin as an inhibitor is also promising to understand the contribution ratio of UGT1A1 to estradiol 3-O-glucuronidation in the human small intestine microsomes, it has been demonstrated that bilirubin could inhibit not only UGT1A1, but also other UGT isoforms due to their high amino acid sequence similarities.

Interestingly, all of the fatty acids that strongly inhibited UGT1A1 activity were unsaturated fatty acids, and fatty acids that did not inhibit UGT1A1 were all saturated fatty acids. This finding is in accordance with previously reported data that most unsaturated fatty acids dramatically inhibited human UGT1A9- and UGT2B7-catalyzed 4-MU glucuronidation in their recombinant systems. Therefore, it can be speculated that unsaturated fatty acids have the potency to inhibit a wide variety of human UGT enzymes. Unsaturated fatty acids are abundant in food such as vegetable oil and fish oil. Additionally, nowadays certain unsaturated fatty acids, such as DHA and EPA, are widely used as daily supplements, as DHA and fish oil. Additionally, nowadays certain unsaturated fatty acids, such as DHA and EPA, are widely used as daily supplements, as DHA and fish oil. Furthermore, certain unsaturated fatty acids, such as DHA and EPA, are widely used as daily supplements, as DHA and fish oil. Further research is needed to elucidate the mechanisms by which these fatty acids inhibit UGT1A1 activity.

Figure 7 | Effects of oral oleic acid (A) and linoleic acid (B) treatments on UGT1A1 expressions in hUGT1 mice. Neonatal hUGT1 mice were orally treated with oleic acid and linoleic acid (10 g/kg). Forty-eight hours after the treatment, livers and small intestines were isolated and UGT1A1 expression levels were determined. Fold changes of UGT1A1 levels are shown. Error bars show SD, n = 3. **, P < 0.01 compared to the vehicle treated group.

### Methods

#### Chemicals and reagents

UDP-glucuronic acid, estradiol, estradiol 3-O-glucuronide, and deconjugase were purchased from Sigma-Aldrich (St Louis, MO). The UGT1A1 recombinant system, UGT1A1 Supersomes, human liver microsomes, and human small intestine microsomes were purchased from BD Gentest (Woburn, MA). Oleic acid, linoleic acid, and DHA were purchased from Cayman Chemical Company (Ann Arbor, MI). Palmitic acid, stearic acid, myristic acid, and palmitoleic acid were purchased from MP Biomedicals (Solon, OH). Lauric acid was purchased from Tokyo Chemical Industry (Tokyo, Japan) and arachidonic acid, palmitoleic acid, alpha-linolenic acid, arachidonic acid, eicosapentaenoic acid (EPA), and DHA were purchased from Cayman Chemical Company (Ann Arbor, MI). All other chemicals and solvents were of analytical grade or the highest grade commercially available.

#### Enzyme assays

Estradiol 3-O-glucuronide formation was determined according to the method of Fujikawa et al. (2007) with slight modifications. Briefly, a typical incubation mixture (200 µl of total volume) contained 50 mM Tris-HCl (pH 7.4), 10 mM MgCl₂, 2 mM UDPGA, 25 µg/ml alamethicin, 0.25 mg/ml UGT1A1 Supersomes or microsomes and 10 µM estradiol. For an inhibition study, 1 to 50 µM of inhibitors were included in the reaction mixtures. To estimate the inhibition type of oleic acid, linoleic acid, and DHA for substrate binding, analyses were conducted using a fixed concentration of 2 mM UDPGA and varied concentrations of estradiol (5, 10, and 20 µM). The reaction was initiated by the addition of UDPGA after a 3-min incubation at 37°C. The reaction was terminated by the addition of 5 mM unlabeled estradiol. The glucuronide was separated by reverse-phase liquid chromatography. The peak area of the glucuronide was quantified by liquid chromatography/mass spectrometry. The degree of inhibition was calculated as the percentage change in the amount of glucuronide. The inhibition type was determined by comparing the degree of inhibition with the calculated values for the inhibition type of oleic acid, linoleic acid, and DHA.
preincubation at 37°C. After incubation at 37°C for 30 min, the reaction was terminated by addition of 200 μL of cold methanol. After removal of the protein by centrifugation at 15,000 rpm for 5 min, a 50 μL portion of the sample was subjected to HPLC. The type of inhibition and the Ki value were determined by a Dixon plot analysis.

HPLC conditions. HPLC was performed using an LC-10AD pump (Shimadzu, Kyoto, Japan), a FP-2020 fluorescence detector (JASCO, Tokyo Japan), a SIL-10A autosampler (Shimadzu), a SCL-10A system controller (Shimadzu), and a Mightysil RP-18 GP column (4.6 × 150 mm, 5 μm; Kanto Chemical, Tokyo, Japan). The flow rate was 1.0 mL/min, and detection was accomplished with a fluorescence detector at 280-nm excitation and 310-nm emission. The mobile phase was 10 mM H3PO4, methanol (45:55, v/v). Quantification of estradiol 3-β-glucuronide was performed by comparing the HPLC peak height to that of the authentic standard.

Animals and treatments. Tg(UGT1A1<sup>-/-</sup>/Ugt1A<sup></sup>/<sup>-/-</sup>) (b/UGT1) mice were developed previously in a C57BL/6 background<sup>14,15</sup>. All animals received food and water ad libitum, and mouse handling and experimental procedures were conducted in accordance with our animal care protocol, which was previously approved by Kitasato University. Fatty acids dissolved in rapeseed oil were orally administered to 12-day-old mice. Blood was obtained before and after treatment and the serum total bilirubin levels were measured. Rapeseed oil was orally administered to mice in the control group. For tissue collections, mice were anesthetized by diethyl ether inhalation, and the liver was perfused with ice-cold 1.15% KCl. The small intestine and liver were rinsed in cold 1.15% KCl and stored at −80°C.

Bilirubin measurements. Blood was obtained from the submandibular vein and centrifuged at 2000 × g for 5 minutes. Serum samples (20 μL) were measured for total serum bilirubin using a Bilirubinometer (B-105N, Erma, Tokyo, Japan). For each serum sample, the bilirubin value was measured three times and the mean value was used for data analysis. To avoid hemolysis of the serum and photolysis of the bilirubin, serum bilirubin using a Bilirubinometer (B-105N, Erma, Tokyo, Japan). For each serum sample, the bilirubin value was measured three times and the mean value was used for data analysis. To avoid hemolysis of the serum and photolysis of the bilirubin, serum bilirubin using a Bilirubinometer (B-105N, Erma, Tokyo, Japan).

Semi-quantitative and quantitative (Q) reverse transcription (RT)-PCR. Total RNA was extracted from tissues with Trizol reagent (Invitrogen). The cDNA was synthesized from total RNA using ReverTra Ace (TOYOBO, Osaka, Japan) according to the manufacturer’s protocol. Primers for human UGT1A1 and GAPDH were used for data analysis. To avoid hemolysis of the serum and photolysis of the bilirubin, serum bilirubin using a Bilirubinometer (B-105N, Erma, Tokyo, Japan).

Equations. When an inhibitor exhibits noncompetitive inhibition against enzymes, the velocity of the metabolic reaction can be described by the following equation (Eq 1):

\[
V = \frac{V_{\text{max}}}{1 + \frac{I}{K_i}} \cdot C_n + C_u
\]

where \(V\) is the velocity of the metabolic reaction, \(C_n\) is the free substrate concentration, and \(I\) is the free inhibitor concentration. The \(V_{\text{max}}\) is the maximum rate of metabolism and \(K_m\) is the Michaelis constant, which is defined as the substrate concentration at 1/2 the maximum velocity. \(K_i\) is the inhibition constant. When \(C_u\) is much lower than the \(K_m\) value, the equation can be described by the following equation (Eq 2):

\[
V = \frac{V_{\text{max}}}{K_m} \cdot \left(1 + \frac{I}{K_i}\right)
\]

In this case, intrinsic clearance (\(CL_{in}\)) can be described by the following equation (Eq 3):

\[
CL_{in} = \frac{V_{\text{max}}}{K_m} \cdot \left(1 + \frac{I}{K_i}\right)
\]

Therefore, the ratio of intrinsic clearance (\(R\)) in the presence (\(V^+\)) and absence (\(V^-\)) can be described by the following equation (Eq 4):

\[
R = \frac{V^+}{V^-} = \frac{1}{1 + \frac{I}{K_i}}
\]

This equation was used to estimate the ratio of intrinsic clearance in the presence and absence of an inhibitor.

Statistical analyses. Statistical significances were determined by analysis of variance followed by Dunnett’s test. A value of \(P<0.05\) was considered statistically significant.
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Acknowledgements
This work was supported by a Grant-in-Aid for Research Activity Start-up [24890224].

Author contributions
T.L., R.H.T. and R.F. wrote the main manuscript text and A.S. and R.F. conducted experiments and prepared figures 1–7. All authors reviewed the manuscript.

Additional information
Competing financial interests: The authors declare no competing financial interests.

How to cite this article: Shibuya, A., Itoh, T., Tukey, R.H. & Fujiwara, R. Impact of fatty acids on human UDP-glucuronosyltransferase 1A1 activity and its expression in neonatal hyperbilirubinemia. Sci. Rep. 3, 2903; DOI:10.1038/srep02903 (2013).

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