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- 1 Interaction of tryptophan-related compounds with ABCG2 transporter:
- 2 comprehensive analysis of tryptophan metabolome in milk
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Abstract

Milk is the main source of the essential amino acid tryptophan and its metabolites for sucklings. The main pathway in tryptophan metabolism is the kynurenine pathway that generates several bioactive metabolites with relevant health effects, including prevention of neurodegenerative disorders. ABCG2 transporter participates in milk secretion of different compounds. Our aim was to provide a comprehensive overview of the *in vitro* and *in vivo* interactions between tryptophan-related compounds and ABCG2. Tryptophan metabolome was analysed in milk and plasma from wild-type and Abcg2-/- mice using liquid chromatography-tandem mass spectrometry. Our results show that Abcg2 mediates the secretion of kynurenic acid, kynurenine, anthranilic acid and xanthurenic acid into milk. Complementary metabolome analysis revealed that the bovine ABCG2 Y581S polymorphism increases kynurenine concentration in cow milk. All these results show the relevance of the interaction of ABCG2 and tryptophan-related compounds present in milk and its potential effect on health, mainly in postnatal development.

1. Introduction

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Milk contains a number of nutrients and bioactive ingredients that play an important role in health. Some of these important nutrients and bioactive ingredients of milk are Ltryptophan (Trp) and its metabolites. It is worth noting that breast milk is the only source of this essential amino acid in breast-fed infants (Schneider, Mutungi, & Cubero, 2018). Trp is an aromatic amino acid critical for protein synthesis; in addition to this essential role, it is the precursor of several bioactive compounds with important physiological roles (Cervenka, Agudelo, & Ruas, 2017). Trp metabolism proceeds mainly by kynurenine (KYN) and serotonin (5HT) pathways (Fig. 1), and the main metabolites generated are related to broad spectrum effects associated with neurodegerative disorders, pain syndromes and autoimmune diseases (Vécsei, Szalárdy, Fülöp, & Toldi, 2013). Probably the best-known pathway is Trp conversion to 5HT and finally to melatonin. Both compounds have important roles in the central nervous system (CNS). 5HT is an important neurotransmitter involved in the control of adaptive responses in the CNS. Melatonin is a neurohormone and a powerful free-radical scavenger, therefore it is an important antioxidant (Reiter et al., 2017). In addition, it plays an essential role in regulating circadian rhythms related to physiological functions such as sleep and blood pressure (Boutin, Audinot, Ferry, & Delagrange, 2005). Nevertheless, more than 95% of Trp is metabolized through the Kynurenine Pathway (KP) in the liver (Badawy, 2019). This pathway generates a range of metabolites, collectively known as kynurenines, involved in immune response, inflammation and neurotransmission (Stone, Stoy, & Darlington, 2013). Most of them are neuroactive and show relevant roles in the regulation of NMDA (N-methyl-D-aspartate) receptor function and the production of free radicals (Nayak & Buttar, 2016). The presence of some of these metabolites in the diet could have important effects on biological processes. For example, an excess of L-KYN in diet could affect neuronal development in young (Pocivavsek, Wu, Elmer, Bruno, & Schwarcz, 2012).

Thus, the content of Trp metabolites in milk might have important effects on human health. The presence of a large number of metabolites and xenobiotics in milk is mediated by two transporter superfamilies: ATP-binding cassette (ABC) and Solute Carrier (SLC) transporters (García-Lino, Álvarez-Fernández, Blanco-Paniagua, Merino, & Álvarez, 2019). In particular, ABCG2/BCRP mediates the active secretion of xenobiotics and several vitamins into milk (van Herwaarden et al., 2007). ABCG2 is present in a multitude of tissues

which affect the body distribution of a wide range of molecules including chemotherapeutic (mitoxantrone, methothexate) and non-chemotherapy drugs (nitrofurantoin, fluoroquinolones), as well as endogenous and dietary compounds such as flavonoids, porphyrins, estrone-3-sulfate and uric acid (Safar, Kis, Erdo, Zolnerciks, & Krajcsi, 2019). However, its role in the regulation of the milk content of Trp metabolites has not been explored.

The function of ABCG2 in regulating milk content can be altered by the presence of several polymorphisms. In cattle, Cohen-Zinder et al. (2005) reported an SNP encoding a substitution of a Ser by Tyr at aminoacidic position 581 (Y581S) described as an *in vitro* and *in vivo* gain-of-function polymorphism (Otero, 2015; Real et al., 2011), which is directly involved in milk quality affecting the presence of ABCG2 substrates in cow milk (García-Lino et al., 2019). In humans, several genetic analyses have demonstrated that single nucleotide polymorphisms (SNPs) leading to ABCG2 deficiency are essential in the pathogenesis of hyperuricemia and gout (Woodward et al., 2009). The Q141K variant yields to decreased ABCG2 protein expression and influences risk of hyperuricemia and gout. Because of this, the Q141k variant has been studied as a potential therapeutic target (Woodward et al., 2009). Although there is some evidence linking hyperuricemia to changes in Trp levels (Liu et al., 2011), and Dankers et al. (Dankers et al., 2013) have suggested that kynurenic acid (KYNA) could interact with human ABCG2, the interaction between ABCG2 and Trp metabolites has not been fully established.

The aim of this work was to study the interaction of Trp-related compounds with ABCG2 transporter through the analysis of Trp metabolome in plasma and milk of wild-type and Abcg2^{-/-} mice. The outcome was corroborated by *in vitro* transport studies with cells overexpressing murine Abcg2. A complementary Trp metabolome analysis was performed in plasma and milk from cows carrying the polymorphism Y581S and non-carrier animals.

2. Materials and methods

2.1. Standards and chemicals

- Reference standards of leucine (Leu), isoleucine (Ile), Trp, 5HT, melatonin, serotonin (5HT), 5-hydroxyindolacetic acid (5HIAA), KYN, KYNA, xanthurenic acid (XA) and anthranilic acid (AA) were purchased from Sigma-Aldrich (St. Louis, MO, USA). Kynurenic acid-d5, phenylalanine-d5 and serotonin-d5 were supplied by Toronto Research Chemicals (Toronto, Canada). Tyrosine-d4, tryptophan-d5, 5-hydroxyindolacetic acid-d4, and kynurenine-13C6 were from Alsachim (Illkirch-Graffenstaden, France). All the other chemicals were analytical grade and obtained from commercial sources.
- 128 2.2. Animals

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- Animals were housed and handled according to institutional guidelines complying with European legislation (2010/63/EU). Experimental procedures were approved by the Animal Care and Use Committee of the University of León and the Junta de Castilla y León ULE_011_2016 and ULE_002_2017.
- Abcg2^{-/-} and wild type female mice aged between 8 and 17 weeks, all of them >99% FVB 133 genetic background (n = 9 - 12), were kindly provided by Dr. A. H. Schinkel (The 134 Netherlands Cancer Institute, Amsterdam, the Netherlands), and were kept in a 135 temperature-controlled environment with a 12-h light / 12-h dark cycle. Pups of 136 approximately 10 days old were separated from their mothers approximately 4 h before milk 137 138 collection. Oxytocin (200 µL of 1 IU/mL solution) was administered subcutaneously to lactating mothers to stimulate milk production 20 min before milk sampling. Milk samples 139 were collected in the morning from the mammary glands by gentle pinching after 140 anesthesia with isoflurane. Blood samples were collected by cardiac puncture under 141 anesthesia with isoflurane and centrifuged immediately at 3000 g for 15 min. At the end of 142 the experiment, the mice were killed by cervical dislocation. Plasma and milk samples were 143 stored at -20 °C until LC-MS/MS analysis. 144
 - Lactating Holstein-breed cows aged between 2 and 5 years and weighing 650 to 1000 kg were used. Their daily milk yield was on average 40 ± 10 kg. The normal milking routine for all the animals involved milk being taken three times each day. The samples were taken at a private farm located at Villalquite, Leon (Spain). The Y581S genotypes were determined

in accordance with the procedure described by Komisarek et al (2009) (Komisarek & Dorynek, 2009). Animals were divided into 2 groups of 8 Y/S 581 heterozygous and 8 Y/Y 581 homozygous cows. No difference was found in age, weight, or milk yield between the two sets of cows. Samples were collected from the first morning milking by mechanical milking. Blood samples were collected from the tail vein and plasma was separated by centrifugation at 3000 x g for 15 min. Plasma and milk samples were stored at -20 °C until LC-MS/MS analysis.

2.3. Sample preparation for LC-MS/MS analysis

An aliquot of plasma (70 μ L for mice and 150 μ L for cows) or milk (between 63 and 180 mg for mice and 150 μ L for cows) was mixed with 300 μ L of acetonitrile in order to precipitate the proteins. After centrifugation, the supernatant was transferred to a clean tube and 50 μ L of the internal standard mixture (containing kynurenic acid-d5, phenylalanine-d5, serotonin-d5, tyrosine-d4, tryptophan-d5, 5-hydroxyindolacetic acid-d4, and kynurenine-13C6) were added. The mixture was evaporated at room temperature under nitrogen stream (<10 psi). After reconstitution with 150 μ L of water, 10 μ L were injected into the system. The standards used for calibration were subjected to the sample procedure.

2.4. Quantification of Trp related compounds by LC-MS/MS

A previously described LC-MS/MS method (Marcos et al., 2016) was used for the determination of Trp-related compounds in milk and plasma from mice and cows. The LC-MS/MS system used to perform the analysis was an Acquity UPLC system (Waters Associates, Milford, MA, USA) coupled to a TQS Micro triple quadrupole mass spectrometer (Waters Associates) equipped with an electrospray ionization interface. Chromatographic separation was carried out on an Acquity BEH C18 column (100 mm x 2.1 mm i.d., 1.7 μm) (Waters Associates) at a flow rate of 300 μL/min. Water and methanol both with ammonium formate (1 mM) and formic acic (0.01% v/v) were used as mobile phase solvents. A gradient elution was used for the chromatographic separation of the analytes. The percentage of the organic solvent linearly changed as follows: 0 min, 1%; 0.5 min, 1%; 7 min, 40%; 8.5 min, 90%; 9 min, 90%; 9.5 min, 1%; 12 min, 1%. Analytes were determined by SRM including 2 ion transitions for each analyte.

179 2.5. Cell cultures

MDCKII cells and their murine Abcg2 transduced subclones were provided by Dr. A.H. Schinkel, Netherlands Cancer Institute, Amsterdam. MDCKII cells stably transduced with both bovine variants (S581 and Y581) of ABCG2 were previously generated by our group (Real et al., 2011). Culture conditions were as previously described (Otero, 2015; Perez et al., 2013).

2.6. Transport studies

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Transepithelial transport assays using Transwell plates were carried out as described elsewhere (Otero, 2015; Perez et al., 2013) with minor modifications. Cells were grown for 3 days after seeding on microporous polycarbonate membrane filters at a density of 1.0 x 10⁶ cells per well. To check the tightness of the monolayer, transepithelial resistance was measured in each well using a Millicell ERS ohmmeter (Millipore). Transport medium consisted of Hanks' Balanced Salt solution (Sigma-Aldrich) supplemented with 4-(2hydroxyethyl)-1-piperazineethanesulfonic acid, HEPES (25 mM). Two hours before the start of the experiment, medium at both the apical (AP) and basolateral (BL) sides of the monolayer was replaced with 2 mL of transport medium without serum, HANKs solution supplemented, either with or without the specific ABCG2 inhibitor Ko143 (1 µM). The experiment started when medium in either the AP or BL compartment was replaced with fresh medium containing the different compounds at a concentration of 10 µM. Cells were incubated at 37 °C in 5% CO₂ and 100 µL aliquots of culture media were taken at t=1, 2, 3 and 4 h, ours and this volume was replaced with fresh medium. The samples were stored at -20 °C. Concentrations of studied compounds were subsequently determined by HPLC. Active transport across MDCKII monolayers was expressed by the relative transport ratio (R), defined as apically directed transport percentage divided by basolaterally directed translocation percentage, after 4 h.

2.7. High Performance Liquid Cromatography (HPLC) Analysis

Analysis of samples to determine the concentrations of the studied compounds in the transepithelial transport assays were carried out by HPLC analysis. The chromatographic system consisted of a Waters 2695 separation module and a Waters 2998 UV photodiode array detector. Fifty µL of the culture medium was injected directly into the HPLC system.

Separation of the samples was performed on a reverse-phase column (Atlantis® T3 3 μ m, 4.6x150 mm). The mobile phase consisted of 0.14 % trifluoroacetic acid:acetonitrile (80:20), the flow rate of the mobile phase was set to 0.8 mL/min, and UV absorbance was measured at 238 nm. The temperature of the samples was 4 °C. Standard samples were prepared in the appropriate drug-free matrix, yielding a concentration range from 0.039 μ g/mL to 10 μ g/mL, with coefficients of correlation >0.99. The limit of quantification was in the range of 0.02 – 0.03 μ g/mL and the limit of detection was in the range of 0.005 – 0.014 μ g/mL for all the compounds. Both were calculated by the method described by Taverniers et al. (Taverniers, I., De Loose, M., Van Bockstaele, 2004).

2.8. Statistical analysis

Comparisons between groups were performed by the Student's t-test (normal variables) and the Mann-Whitney U test (not normally distributed variables). All analyses were carried out on the assumed significance level of $p \le 0.05$ using SPSS Statistics software (v. 24.0; IBM, Armonk, New York, NY, USA). The results are shown as mean \pm standard deviation (SD).

3. Results

- 3.1. Determination of Trp-related compounds in milk from Abcg2^{-/-}and wild-type mice
- To examine the role of ABCG2 in the transport of Trp metabolites into milk, we collected milk and plasma samples of wild-type and Abcg2^{-/-} female mice.
- The targeted LC-MS/MS method included 10 analytes. These Trp-related analytes were divided into three different classes based on their biological origin (Fig. 1): (a) large neutral amino acids (LNAA) competitors of Trp; (b) metabolites belonging to the serotonin pathway; (c) metabolites from the kynurenine pathway.
 - All the compounds were detected in all plasma and milk samples, with the exception of Trp in milk and melatonin in plasma and milk (Table 1). There were no differences in plasma between wild-type and Abcg2^{-/-} mice for any compound. Nevertheless, a higher secretion into milk concentration of KYN, KYNA, XA, AA and 5HIAA was observed in wild-type compared to Abcg2^{-/-} mice (Table 1). These differences were especially high for XA and KYNA, with a concentration in milk of almost 10-fold higher (75 ± 40 ng/mg versus 7.8 ± 4.7 ng/mg) and 5-fold higher (42 ± 8 ng/mg versus 9.1 ± 3.0 ng/mg) in wild-type and Abcg2^{-/-} mice, respectively. Higher milk-to-plasma ratios were also obtained for these five metabolites in wild-type compared to Abcg2^{-/-} mice, except for 5HIAA. These data indicate that Abcg2 plays a substantial role in the secretion of KYN, KYNA, XA and AA into milk.
- 3.2. Effect of the bovine ABCG2 polymorphism Y581S SNP on secretion into milk of Trprelated compounds
 - A similar analysis was performed on cows carrying the Y581S polymorphism and on non-carrier animals (Table 2). All the analytes were detected in all plasma and milk samples with the exception of AA, which was not detected in any sample, and 5HT, which was not detected in any milk sample. There were no differences in plasma levels between Y/Y 581 and Y/S 581 cows for any compound tested. Significant differences were found only in milk for KYN concentrations (Table 2, Fig. 2). KYN milk concentrations were 2-fold higher in Y/S (4.6 ± 1.8 ng/mL) compared with Y/Y cows (2.4 ± 1.0 ng/mL). The milk-to-plasma ratio for KYN in Y/S cows was 2-fold higher than in Y/Y cows (0.004 ± 0.002 versus 0.002 ± 0.001). This indicates that the polymorphism Y581S affects the *in vivo* active transport of KYN into cow milk.

3.3. Transport of Trp metabolites in MDCKII cell models transduced with murine Abcg2

To corroborate the metabolomic study in mice, a transport assay using parental MDCKII and its subclones transduced with murine Abcg2 was performed to study the role of this protein in the process. KYN, KYNA, XA and AA were tested *in vitro* due to *in vivo* evidence of Abcg2 involved in their milk secretion (Table 2). Trp itself was also included.

In the MDCKII parental cell line, most of the molecules showed similar apically and basolaterally directed translocation (Fig. 3A). However, parental MDCKII cells displayed a high basolaterally directed translocation of KYN and Trp, whereas apically directed translocation was very low, indicating the potential presence of an absorptive KYN and Trp transport process.

In the murine Abcg2-transduced cells (Fig. 3B), we observed for KYN, KYNA and XA increased translocation from the basolateral to the apical compartment, and reduced translocation from the apical to the basolateral compartment compared to parental cells, with high relative transport ratios (AP/BL). For AA, a low transport ratio similar to the parental cells was obtained. The apical transport of KYN, KYNA and XA by Abcg2 was completely inhibited by Ko143, a selective Abcg2 inhibitor (data not shown).

- From this, we can confirm that KYN, KYNA and XA are good *in vitro* substrates of murine Abcg2.
- 3.4. Transport of KYN in MDCKII cell models transduced both bovine variants (Y581 and S581)

To corroborate the role of the bovine Y581S polymorphism in the transport of KYN, transport assays were performed using polarized MDCKII parental cells and their subclones transduced with both bovine ABCG2 variants (S581 and Y581) (Fig. 4). The relative transport ratio of KYN was significantly higher in the S581 expressing cells versus parental cells (1.08 ± 0.25 vs. 0.59 ± 0.19) since apical transport increased and basolateral transport decreased compared with parental cells. However, in the case of cells expressing Y581 cells, no changes were observed in the transport ratio compared with parental cells, indicating that KYN is not transported by this variant. Statistically significant differences were found between transport ratios of both variants of bovine ABCG2, Y581 and S581 (0.42 ± 0.25 vs. 1.08 ± 0.25). Therefore, the differences found between two bovine variants

showed that the Y581S polymorphism affects the *in vitro* transport of KYN, corroborating the differences found *in vivo* between carrier and non-carrier animals.

4. Discussion

In this study a targeted metabolomic analysis was performed to elucidate the role of ABCG2 transport in the active secretion of Trp-related compounds in plasma and milk. The relevance of the bovine Y581S polymorphism on the transport of these compounds present in cow milk was also studied.

Due to the complication in controlling all the animal factors, such as age, nutrition and time of lactation, among others, a study of the association between *in vitro* and *in vivo* results is difficult, especially in cows (Otero, 2015). However, this LC-MS/MS metabolomic study permits a comprehensive estimation of the effect of ABCG2 in Trp-related pathways in plasma and milk in mice and cows. What is of relevance, and an advantage over other studies analysing a limited number of metabolites (Cubero et al., 2005; Laeger, Görs, Metges, & Kuhla, 2012), is the fact that a large number of compounds were determined.

The amount of L-Trp in humans is essential for maintaining health, and one of the most common sources of Trp in the human diet is milk (Richard et al., 2009). Furthermore, the biological activities of its derivative compounds, mainly as neuroactive molecules, adds to the importance of this complete study. Focussing, therefore, on the relevance of these neuroactive compounds in the diet, we performed the metabolome analysis to elucidate the role of ABCG2 transport in active secretion into milk of Abcg2^{-/-} and wild-type lactating mice. Differences in the concentrations of KYN, KYNA, XA and AA were observed in the milk-to-plasma ratio between wild-type and Abcg2^{-/-} mice, which suggests the involvement of Abcg2 in the milk secretion of these compounds (Table 1). These results were corroborated by transport assays showing that KYN, KYNA and XA are *in vitro* substrates of Abcg2 (Fig. 3). Only KYNA had been previously described as a potential Abcg2 substrate in humans (Dankers et al., 2013). Conversely, AA was not confirmed as an *in vitro* substrate of ABCG2, and due to the positive results observed in the *in vivo* study, we cannot exclude its *in vitro* interaction with the Abcg2 transporter in other experimental conditions.

Binding of substrate to ABCG2 has been shown to be dependent of hydrophobic interactions, mainly between hydrogen bond acceptors (HBAs) present in substrates and hydrogen bond donors (HBDs) present in the transmembrane region of the transporter, among other physicochemical features (Matsson et al., 2007). In adition, Xu *et al.* (Xu et al., 2015) demonstrated that binding of substrates with the ABCG2 transporter increases with the number of HBAs present in the potential substrates. In our case, KYN, KYNA and XA, identified as Abcg2 substrates, gave a higher number of HBAs than Trp, 5HIAA and 5HT and AA (non Abcg2 substrates) (Supplementary material, Table S1).

Furthermore, we examined the importance of the bovine Y581S polymorphism in the transport of Trp-related compounds present in cow milk, as previously reported for milk secretion of fluoroquinolone drugs, antiinflammatory drugs and endogenous and dietary compounds (García-Lino et al., 2019; Otero, 2015). In fact uric acid, which is related to Trp levels (Dankers et al., 2013), has been previously reported as an endogenous compound actively secreted into milk with a 2-fold increase in the milk plasma ratio for carrier animals (Otero et al., 2016). Our data show that the transfer of KYN into milk is influenced by the bovine Y581S polymorphism with an almost 2-fold increase in the concentration and milk-to-plasma ratio of KYN in the Y/S animals (Fig. 2). This is the first time that differences between both variants of cows, carriers and non-carriers of the Y581S polymorphism has been observed for Trp-related compounds with neuroactive activity. Furthermore, these results are in agreement with *in vitro* transport studies showing that cells overexpressing the S581 variant transported KYN in the basolateral to apical direction more efficiently than the cells with the Y581 variant (Fig. 4), indicating a higher transport capacity *in vitro* for this variant.

Our results could have great relevance in the field of nutrition since all of these metabolites show important effects on health and some of them are present in consumed dairy milk. Furthermore, our results could be relevant in the early stages of development where milk is the main food consumed. ABCG2 inhibitors such as drugs (Barrera et al., 2013) and dietary compouds (Miguel et al., 2014) can alter the transfer of these compounds into milk. In fact, these kinds of interactions mediated by ABCG2 have been observed for other ABCG2 substrates. For instance, soy-enriched diet and flaxeed-enriched diet modify *in vivo* milk secretion mediated by ABCG2 of the antimicrobial danofloxacin in sheep (Otero et al., 2018; Perez et al., 2013). Therefore, potentially different concentrations of Trp-related

compounds in consumed milk due to polymorphisms or inhibition of this transporter may affect the intake of these compounds by the offspring or the dairy consumer.

Trp-related compounds generated by KP have an important role in health due to their effects on the CNS since they have been linked to several psychiatric and mental health disorders such as depression and schizophrenia (Chen & Guillemin, 2009). This pathway is also currently involved in immunological processes (Badawy, 2019). KYNA has proved to be a neuroprotective molecule and its neuroprotective role in various experimental models of neurotoxicity has been reported (Majláth, Török, Toldi, & Vécsei, 2016). There is evidence in preclinical models that changes in KYNA concentrations in the brain could influence the development of Huntington's disease (Okamoto et al., 2009) and schizophrenia (Beggiato, Notarangelo, Sathyasaikumar, Giorgini, & Schwarcz, 2018). These compounds may also have effects in other organs. Reports suggest that KYNA could have a protective role in the intestinal mucosa in the framework of obstructive jaundice and may protect from induced ulcers (Glavin, Bose, & Pinsky, 1989). Since we demonstrated that KYNA is an ABCG2 substrate, different milk intake of KYNA due to differences in ABCG2 function may affect the protective role of KYNA in several pathological conditions.

KYN is the main precursor of KYNA. Due to the lower permeability of KYNA in the blood-brain-barrier than presented by KYN, KYN is used as a protective compound in models of Huntington's disease induced by malonate and quinolinic acid (Santamaría et al., 1996). However, despite the benefits in the neurodegenerative diseases provided by KYN administration, several studies report that an exposure to KYN during a vulnerable period of brain development causes cognitive deficits in adulthood related to schizophrenia and other psychiatric diseases (Pocivavsek et al., 2012). In addition, it has been shown that a continuous KYN administration during the prenatal period causes learning and memory deficits in adults rats (Pocivavsek, Thomas, Elmer, Bruno, & Schwarcz, 2014). Since our results show the role of ABCG2 in the presence of KYN in milk, this interaction may play an important role in cognitive development of the offspring.

Our data may have relevant effects in other health fields. Some studies have demonstrated that KYN and KYNA are endogenous ligands of the aryl hydrogen receptor (AhR) (Julliard, Fechner, & Mezrich, 2014). The coupling between AhR and these compounds increases concentration of TGF- β and IL-6 in early sepsis (Wirthgen & Hoeflich,

2015). Also, recent studies suggest that Trp-containing diet might regulate epithelial homeostasis through AhR in ulcerative colitis (Islam et al., 2017).

In conclusion, this work demonstrates that several Trp metabolites are secreted into milk by the Abcg2 transporter. It shows, in addition, that lactating dairy cows carrying the Y581S polymorphism produced milk with higher amounts of KYN compared with non-carriers.

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Figure captions

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Fig. 1. Tryptophan-related metabolites included in the target LC-MS/MS method. (a) Large neutral amino acids (LNAA) competitors of Trp; (b) Kynurenine pathway and (c) Serotonin pathway. Isoleucine (Ile), leucine (Leu), tryptophan (Trp), kynurenine (KYN), kynurenic acid (KYNA), xanthurenic acid (XA) and anthranilic acid (AA), 5-hydroxyindolacetic acid (5HIAA), serotonin (5HT).

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Fig. 2. Plasma concentration (A), milk concentration (B) and milk/plasma ratio (C) of KYN in Y/Y 581 and Y/S 581 cows. Samples were collected at the first routine morning milking. Results are mean; error bars indicated S.D (n=8; *p<0.05).

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Fig. 3. Transepithelial transport of tested compounds (10 µM) in (A) parental MDCKII cells 585 and (B) their murine Abcg2 transduced derivative. (o) translocation from the apical to the 586 basolateral compartment; (•) translocation from the basolateral to the apical compartment. 587 The experiment was started with the addition of each compound to one compartment 588 (basolateral or apical). Aliquots of 100 µL were taken from the opposite compartment at t=1, 589 2, 3 and 4 h, and measured by HPLC. The fraction of each compound transported to the 590 acceptor compartment was presented as a percentage of the total amount of compound 591 added to the donor compartment at the beginning of the experiment. Results are means, 592 error bars indicated SD (n=3-8). Ratio represents the relative transport ratio (i.e. the apical 593 directed translocation divided by the basolateral directed translocation) at t=4h. 594

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Fig. 4. Transepithelial transport of KYN (10 µM) in polarized MDCKII parental (nonmonolayers. MDCKII-S581-ABCG2, and MDCKII-Y581-ABCG2 transduced), The experimental was started (t=0) by replacing the medium in either the apical or basolateral compartment with fresh Hanks supplemented medium containing 10 µM KYN. Aliquots of 100 µL were taken from the opposite compartment at t=1, 2, 3 and 4 h, and measured by HPLC. The fraction of KYN transported to the acceptor compartment was presented as a percentage of the total amount of KYN added to the donor compartment at the beginning of the experiment. Results are means, error bars indicated SD (n=5). Ratio represents the relative transport ratio (i.e. the apical directed translocation divided by the basolateral directed translocation) at t=4h. (o) translocation from the apical to the basolateral compartment; (•) translocation from the basolateral to the apical compartment.

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Table 1. Levels of Trp-related compounds (ng/mL) in plasma and milk samples from wild-type and Abcg2^{-/-} female mice (n=9-12). Milk to plasma ratio is also represented.

		Wild-type	Abcg2 ^{-/-}	p-value
	Leu	24400 ± 7229	20937 ± 7925	0.292
Plasma	lle	25209 ± 7670	21507 ± 8275	0.285
	Trp	88660 ± 11302	89831 ± 11302	0.911
	KYN	1628 ± 363	1451 ± 344	0.259
	KYNA	7.1 ± 2.5	6.6 ± 2.2	0.647
	XA	73 ± 37	69 ± 30	0.792
	AA	155 ± 98	124 ± 66	0.404
	5HT	11937 ± 5080	8643 ± 1959	0.079
	5HIAA	702 ± 138	581 ± 143	0.060
	Leu	497 ± 238	386 ± 282	0.292
	lle	1645 ± 515	2051 ± 684	0.250
	Trp	< LOD	<lod< td=""><td>-</td></lod<>	-
	KYN	32 ± 17	15 ± 7	0.039*
	KYNA	42 ± 8	9.1 ± 3.0	< 0.001*
Milk	XA	75 ± 40	7.8 ± 4.7	0.001*
	AA	36 ± 10	17 ± 7	0.01*
	5HT	41 ± 22	40 ± 19	0.585
	5HIAA	118 ± 24	84 ± 14	0.001*
	Leu	0.02 ± 0.01	0.08 ± 0.18	0.268
	lle	0.08 ± 0.02	0.53 ± 1.35	0.265
	Trp	< LOD	< LOD	-
	KYN	0.02 ± 0.01	0.01 ± 0.00	0.012*
Milk to plasma	KYNA	6.9 ± 2.6	1.47 ± 0.51	< 0.001*
	XA	0.97 ± 0.48	0.22 ± 0.33	0.001*
	AA	0.23 ± 0.10	0.12 ± 0.04	0.008*
	5HT	0.004 ± 0.00	0.004 ± 0.00	0.794
	5HIAA	0.17 ± 0.04	0.15 ± 0.05	0.322

Results are expressed as means of concentration (ng/mL) ± S.D. * p<0.05 versus wild-type

Table 2. Levels of Trp-related compounds (ng/mL) in plasma and milk samples from non-carrier (Y/Y) and carrier (Y/S 581) cows (n=4). Milk to plasma ratio is also represented.

		Y/Y	Y/S	p-value
Plasma	Leu	38067 ± 9127	39692 ± 10952	0.751
	lle	13559 ± 3439	13315 ± 3283	0.886
	Trp	11283 ± 3053	12348 ± 2443	0.453
	KYN	1054 ± 381	1067 ± 256	0.935
	KYNA	8.1 ± 3.4	7.1 ± 0.5	0.407
	XA	95 ± 26	108 ± 18	0.263
	5-HT	0.09 ± 0.07	0.14 ± 0.08	0.197
	5HIAA	0.011 ± 0.003	0.011 ± 0.003	0.724
	Melatonin	0.006 ± 0.004	0.005 ± 0.001	0.284
	Leu	432 ± 184	569 ± 291	0.280
	lle	342 ± 148	441 ± 180	0.248
	Tyr	41 ± 11	64 ± 31	0.071
	Trp	192 ± 74	252 ± 90	0.170
	KYN	2.4 ± 1.0	4.6 ± 1.8	0.012*
Milk	KYNA	7.9 ± 3.8	8.2 ± 2.6	0.840
	XA	0.32 ± 0.12	0.31 ± 0.08	0.816
	5HT	< LOD	<lod< td=""><td>-</td></lod<>	-
	5HIAA	0.56 ± 0.27	0.81 ± 1.14	0.593
	Melatonin	0.003 ± 0.002	0.003 ± 0.001	0.713
Milk to plasma	Leu	0.01 ± 0.00	0.01 ± 0.00	0.361
	lle	0.03 ± 0.01	0.03 ± 0.01	0.174
	Tyr	0.009 ± 0.004	0.013 ± 0.006	0.152
	Trp	0.018 ± 0.008	0.020 ± 0.007	0.492
	KYN	0.002 ± 0.001	0.004 ± 0.002	0.012*
	KYNA	1.0 ± 0.4	1.2 ± 0.4	0.405
	XA	0.004 ± 0.001	0.003 ± 0.001	0.816
	5HT	< LOD	<lod< td=""><td>-</td></lod<>	-
	5HIAA	0.46 ± 0.17	0.67 ± 0.26	0.697
	Melatonin	0.46 ± 0.17	1.7 ± 0.8	0.157

Results are expressed as means of concentration (ng/mL) ± S.D. * p<0.05 vs wild-type

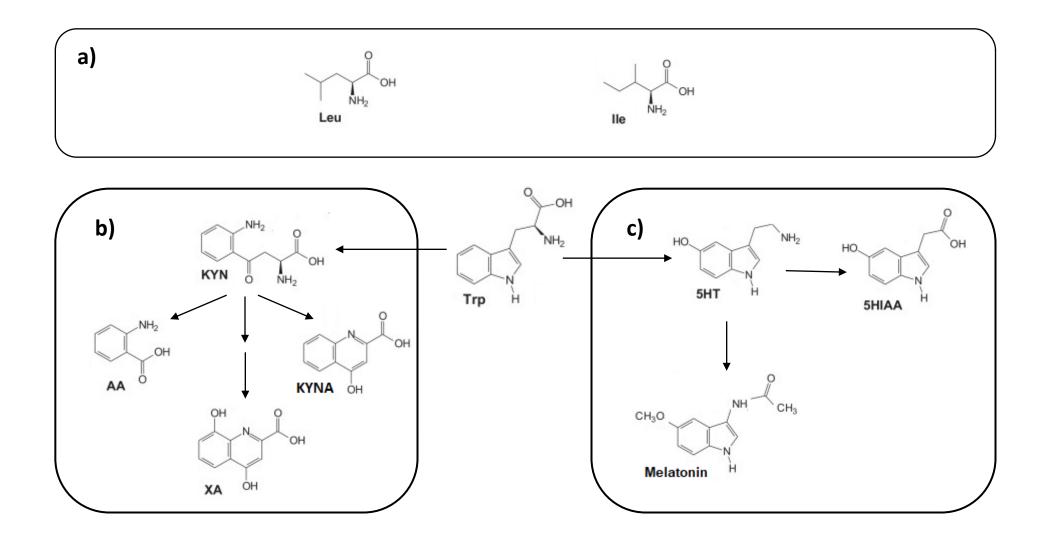


Fig. 1 Garcia-Lino et al.

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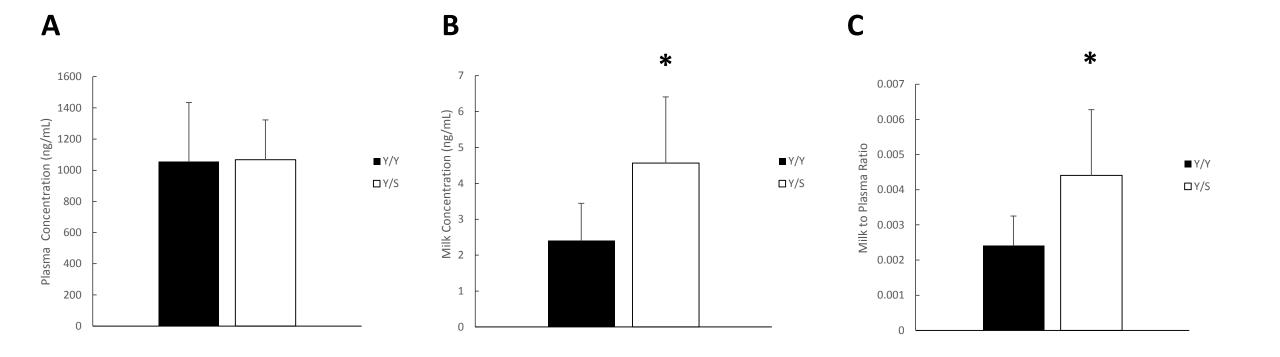


Fig. 2 Garcia-Lino et al.

A) PARENTAL

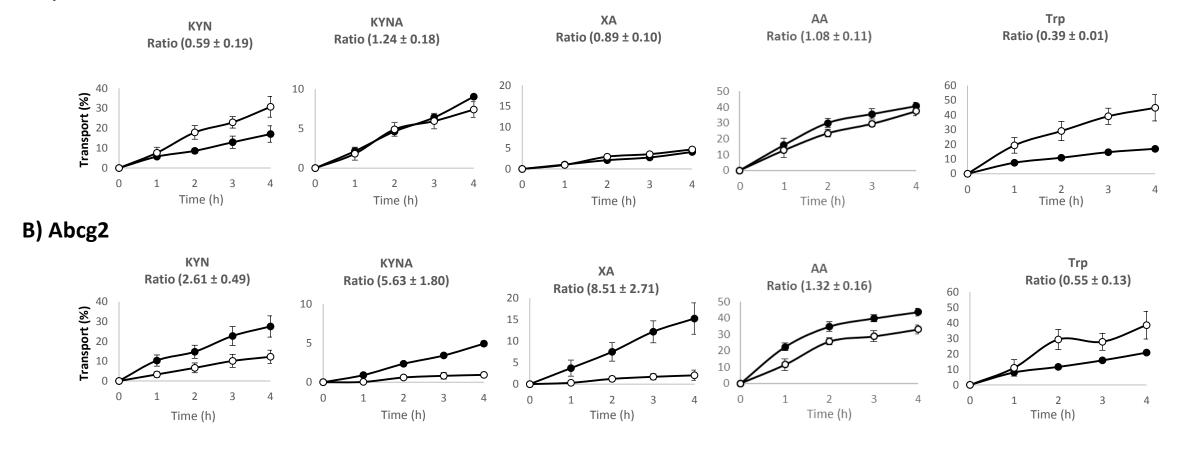


Fig. 3 Garcia-Lino et al.

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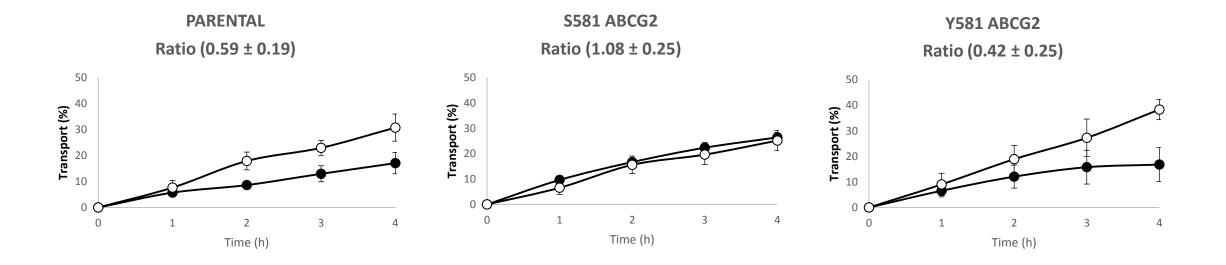
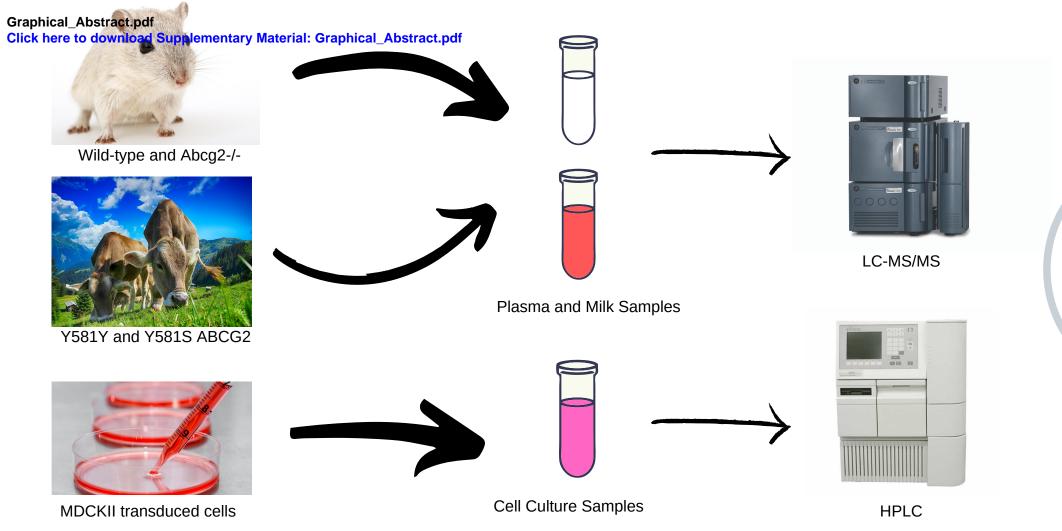


Fig. 4 Garcia-Lino et al.

Supplementary material

Table S1. Hydrogen bond acceptors (HBA) count and hydrogen bond donors (HBD) count for Trp metabolites detected in mouse samples. Data were obtained from U.S. National Library of Medicine (PubChem).

	Hydrogen bond	Hydrogen bond donors
	acceptors (HBA) count	(HBD) count
KYNA	5	3
XA	5	3
KYN	4	2
AA	3	2
Trp	3	3
5HT	2	3
5HIAA	2	2



Interactions between
ABCG2 and
Tryptophan-related
compounds

*Highlights (for review)

HIGHLIGHTS

LC-MS/MS analysis of tryptophan metabolome in plasma and milk shows ABCG2 interaction

Abcg2 mediates milk secretion of metabolites of the kynurenine pathway in mice

Bovine ABCG2 Y581S polymorphism increases kynurenine concentration in milk

Kynurenine and kynurenic and anthranilic acid are *in vitro* substrates of murine Abcg2

The bovine ABCG2 Y581S polymorphism affects the *in vitro* transport of kynurenine

*Declaration of Interest Statement

Declaration of interests
\boxtimes The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.
☐ The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: