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Doris Cassio, Rocio I. R. Macias, Brigitte Grosse, Jose J. G. Marin, Maria J. Monte. Expression, localization, and inducibility by bile acids of hepatobiliary transporters in the new polarized rat hepatic cell lines, Can 3-1 and Can 10.. *Cell and Tissue Research*, 2007, 330 (3), pp.447-60. 10.1007/s00441-007-0494-3 . inserm-00276792

HAL Id: inserm-00276792

<https://inserm.hal.science/inserm-00276792>

Submitted on 2 May 2008

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Expression, localization, and inducibility by bile acids of hepatobiliary transporters in new polarized rat hepatic cell lines, Can 3-1 and Can 10.

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Abstract

Sinusoidal and apical transporters are responsible for the uptake and biliary elimination of many compounds by hepatocytes. Few in vitro models are available for analyzing such functions. The expression and bile acid-inducibility of thirteen transporters and two nuclear receptors were investigated in new rat polarized lines, Can 3-1 and Can 10, and their unpolarized parent Fao. Relative abundance of mRNA, protein level and localization were examined by real-time quantitative PCR, Western blotting, immunofluorescence and confocal microscopy. Compared to rat liver, mRNA levels of Fao cells were: negligible for Bsep/Abcb11, lower for the uptake transporters (Ntcp, Oatps), similar for SHP, FXR and Bcrp/Abcg2, higher (4-160 fold) for the efflux pumps Mdr1b/Abcb1b, Mdr2/Abcb4, Mrp1/Abcc1, Mrp2/Abcc2, Mrp3/Abcc3, Abcg5, Abcg8. This profile was mostly maintained, and improved for Bsep, in Can 10. Some transporters were less expressed in Can 3-1. Moreover, in both lines sinusoidal (Ntcp, Mrp3) and canalicular transporters (Mdr-P-glycoproteins detected with C219 antibody, Mrp2) were localized at their correct poles. Bile acid-effects on polarity and mRNA levels of transporters were analyzed after a 6 days-treatment with 50 μ M of taurocholic, chenodeoxycholic (CDCA) or ursodeoxycholic acid (UDCA): no polarization of Fao cells was induced; Can 10 and Can 3-1 polarity was maintained; CDCA and UDCA induced marked enhancement of the volume of Can 10 bile canaliculi. CDCA upregulated Bsep, Mdr2, SHP, Mdr1b and Oatp2/1a4 in Can 10 (2-7 fold), and Fao cells. In conclusion, Can 10 constitutes an attractive polarized model for studying vectorial hepatobiliary transport of endogenous and xenobiotic cholephilic compounds.

Key words: hepatobiliary transporters, nuclear receptors, bile acids, hepatocyte polarity, cell models

List of abbreviations: Abc/ABC, ATP-binding cassette; BC, bile canaliculi; Bcrp, breast cancer resistance protein; Bsep/BSEP, bile salt export pump; CDCA, chenodeoxycholic acid; DPPIV, dipeptidylpeptidase IV; FITC, fluorescein isothiocyanate; FXR, farnesoid X activated receptor; Mdr/MDR, multidrug resistance protein; Mrp/MRP, multidrug resistance-associated protein; Ntcp, Na⁺-taurocholate co-transporting polypeptide; PBS: phosphate-buffered saline; Oatp/OATP, organic anion transporting polypeptide; PM: plasma membrane; SHP, small heterodimer partner; TCA, taurocholic acid; UDCA, ursodeoxycholic acid;

Introduction

Bile secretion and biliary elimination of drugs are major functions of hepatocytes. These functions are fulfilled owing to the activity of a large panel of transporters located at the sinusoidal or at the apical pole of the plasma membrane (PM) of hepatocytes (Ito et al. 2005; Trauner and Boyer 2003; St Pierre et al. 2001). Few in vitro models are available for analyzing such functions. Ideally in a pertinent model, transporters should be **produced** at levels similar to those of liver, localized at the appropriate PM pole, regulated as in vivo, and functional.

Few models have been characterized for the **gene** expression and regulation of a large set of transporters, and models that stably express typical hepatocyte polarity are rare. The more complete studies were done with human hepatoma lines, such as Hep G2 (Knowles et al. 1980) and HepaRG (Parent et al. 2004). Although Hep G2 cells express **genes of** some transporters (as MRP2/ABCC2, MRP3/ABCC3, BSEP/ABCB11, ..) (Cantz et al. 2000, Kubitz et al 2004), the expression of others (as OATP2, now termed as OATP1B1), was absent or very weak (Nakai et al. 2001; Le Vee et al. 2006). Moreover Hep G2 is not fully polarized since only 10 to 30% of cells are engaged in bile canaliculi (BC) formation (Kubitz et al 2004, Van Ijzendoorn et al. 2004). HepaRG cells displayed a better **gene** expression pattern of transporters (Le Vee et al. 2006), but their polarity state remains to be precisely characterized, all the more because they are bipotent cells that can differentiate either as hepatocytes or cholangiocytes (Parent et al. 2004).

Starting from the non polarized but well differentiated rat hepatoma line Fao we first isolated polarized rat-human hybrid cells (Cassio et al. 1991), leading to isolation of the WIF-B and WIF-B9 lines (Ihrke et al. 1993; Shanks et al. 1994; Decaens et al. 1996). These cells are able to vectorially transport in BC a fluorescent bile acid analogue (Bravo et al., 1998). They **produce** at a robust level some rat hepatobiliary transporters and at a low level a few human transporters (Nies et al. 1998, Briz et al. 2003; Briz et al. 2007). However, their

complex chromosomal content is very far from that of Fao and they form only simple BC, namely BC between two or three cells (Ihrke et al. 1993). Recently, we isolated by growing transiently Fao cells in spheroids, two new polarized lines, Can 3-1 and Can 10 (Peng et al. 2006). These cells display a chromosomal content similar to that of their rat parental cells. They are fully polarized and form BC, in which bile acid derivatives (such as cholylglycylamido fluorescein), organic cations (such as rhodamine 123), and organic anions (such as fluorescein) are vectorially transported (Peng et al. 2006). Interestingly the BC structures of Can 10 cells are highly developed, as those formed *in vivo*.

The present study was designed to investigate in these new polarized lines and their unpolarized parent the expression and inducibility by bile acids of a large panel of **genes encoding** proteins implicated in bile acid and drug transport, namely thirteen transporters and two nuclear receptors. The localization of some transporters was examined in parallel. Our results designated Can 10 cells as a good model for analyzing several aspects of the physiology and pharmacology of the hepatobiliary transport of endogenous (e.g. bile acids) and xenobiotic (e.g. drugs) cholephilic compounds.

Materials and methods

Materials

Fetal bovine serum was from Gibco BRL (Invitrogen, Cergy Pontoise, France). Culture medium, medium supplements, dimethyl sulfoxide (DMSO), bile acids and α tubulin antibody, were from Sigma Chemical Company (St. Louis, MO, USA). Rabbit polyclonal antibodies against Mrp2/Abcc2 (Buchler et al. 1996) and guinea pig polyclonal antibodies against Mrp3/Abcc3 (Donner and Keppler 2001) were given by D. Keppler. Rabbit polyclonal antibodies directed against Na⁺-taurocholate cotransporting polypeptide, Ntcp, was a gift from B. Stieger (Stieger et al. 1994). The mouse monoclonal anti-P-glycoprotein C219 was purchased from Dako, (Carpinteria, CA, USA); it recognizes Mdr1a, Mdr1b and Mdr2. Rabbit polyclonal antibodies against dipeptidylpeptidase IV (DPPIV) and rat monoclonal antibodies against the tight junction associated protein, ZO-1 were provided by A. Hubbard (Shanks et al. 1994) and by B.R. Stevenson (Stevenson et al. 1986), respectively. Fluorescent Alexa dye-conjugated and horseradish peroxidase-linked secondary antibodies were from Molecular Probes (Eugene, OR, USA) and Amersham Biosciences Europe (Orsay, France), respectively. FITC-conjugated goat antibodies directed against guinea pig IgG were from Sigma.

Cell lines

The unpolarized Fao line (Deschatrette and Weiss 1974) is a subclone of the rat hepatoma H4IIEC3 line. Can 3-1 and Can 10 are stable highly polarized cell lines obtained by transient spheroid culture of Fao cells (Peng et al., 2006). **Their karyotype, growth rate and production of epithelial markers (such as E cadherin) and tight junctional proteins (such as occludin) have been determined (Peng et al., 2006). These lines are available (INSERM, U757, University of Orsay, 91400, Orsay, France) for further studies on hepatocyte differentiation.**

Cell Culture

Cell culture was performed in Coon's modified F-12 medium exactly as described (Cassio et al. 1991). Fao and Can 10 cells were plated at $3.5 \times 10^3/\text{cm}^2$. Can 3-1 cells were plated at $10^5/\text{cm}^2$; this high density is necessary for Can 3-1 polarization (Peng et al. 2006). Cells were cultured in 10 cm Petri dishes or glass coverslips for seven days. Cells grown in dishes were used for RNA extraction or protein lysate preparation. Cells on coverslips were used for immunolocalization and for FITC-dextran paracellular diffusion studies.

Treatment with bile acids

Stock solutions (50 mM) of bile acids, taurocholic (TCA), chenodeoxycholic (CDCA) and ursodeoxycholic acid (UDCA) were prepared in DMSO. Cells were cultured for 7 days. They were plated in normal medium and after one day they were cultured without (control) and with TCA, CDCA, UDCA (50 μM) or DMSO, the final concentration of DMSO being 0.1%. Those media were renewed twice (at day 3 and 6).

Determination of gene expression by real-time RT-PCR

Total RNA was isolated from cells using RNeasy spin columns from Qiagen (Izasa, Barcelona, Spain), treated with DNase I (Roche, Barcelona, Spain), and measured with the RiboGreen RNA-Quantitation kit (Molecular Probes, Leiden, The Netherlands). RT was carried out using random nonamers and the Enhanced Avian RT-PCR kit (Sigma-Genosys, Cambridge, UK). Real-time quantitative PCR was performed using Amplitaq Gold polymerase (Perkin-Elmer Applied Biosystems, Madrid, Spain) in an ABI Prism 5700 Sequence Detection System (Perkin-Elmer Applied Biosystems) with the following thermal conditions: a single cycle of 95°C for 10 min followed by 45 cycles of 95°C for 15 s and 60°C

for 60 s. Primer oligonucleotides obtained from Sigma-Genosys (Table 1) were designed with the assistance of Primer Express software (Perkin-Elmer Applied Biosystems) for cDNA fragments in described sequences, and their specificity was checked using BLAST. Since amplification products were detected using SYBR Green I, it was ascertained previously that no non-specific products were formed during PCR in any case (data not shown). Total liver RNA from a healthy male adult rat was used in all determinations as an external calibrator. The results obtained from each sample were normalized using both polyubiquitin-C and 18S rRNA to correct the differences of RNA extraction in each sample. The latter was measured using the TaqMan® Ribosomal RNA Control Reagents kit (Perkin-Elmer Applied Biosystems).

Immunolocalization of proteins

Cells on glass coverslips were rinsed three times in PBS and then fixed. They were fixed in 3% formaldehyde (4°C, 1 min) and permeabilized with methanol (4°C, 15 min) for Mrp2 and DPPIV, or with acetone (4°C, 2 min) for Mdr or 5 min at -20°C in methanol followed by 1 min at -20°C in acetone for Mrp3. For Ntcp localization cells were fixed in 4% formaldehyde (4°C, 30 min), permeabilized with Triton 0,1% in PBS (room temperature, 3 min) and then incubated with 10% serum in PBS (room temperature, 30 min). ZO-1 was localized using any of these fixations. After fixation, indirect immunofluorescence was performed (Bender et al. 1998) and cells were analyzed with a Zeiss Axioskop fluorescence microscope. Confocal analysis was performed with a Zeiss LSM 510 confocal microscope, with a x63 objective. Series of xy sections were taken in 0.3 µm steps.

Western blotting

Cells cultured for 7 days were lysed as described (Bender et al. 1998). Proteins (25 µg) from lysate aliquots were separated by 7.5% SDS-PAGE (Laemmli 1970), then transferred to a nitrocellulose membrane (Hybond ECL, Amersham). Immunoblotting was performed using

anti-P-glycoprotein C219 with anti α -tubulin antibodies, and the appropriate peroxidase-conjugated secondary antibodies. Peroxidase activity was detected by enhanced chemiluminescence (kit from Amersham Biosciences Europe), and NIH Image 1.62 software was used for densitometric analysis.

Tight junction paracellular permeability

Paracellular permeability of tight junctions was evaluated by measuring the paracellular diffusion into the BC of FITC-dextran (molecular weight 4,400). Cells grown on glass coverslips were washed once with serum-free medium, incubated exactly as previously described (Ihrke et al. 1993) with 3 mg/ml FITC-dextran for 30 min at 2°C and washed two times with cold serum-free medium before observation. Each coverslip was examined within 5-10 min after mounting on a glass microscope slide. Fluorescent BC were counted and expressed as a percent of the total number of BC observed in phase contrast. For each culture, 2-3 coverslips were examined (4 fields per coverslip and at least 200 BC observed).

Statistical analysis

Effects of bile acid treatment were analyzed using a Student's t-test. The criterion of significance was $p < 0.05$.

Results

Expression and localization of hepatobiliary transporters

The steady state of relative abundance of mRNA of thirteen transporters and two nuclear receptors involved in hepatobiliary function was measured by real-time quantitative PCR in the unpolarized parental cell line Fao and in the polarized Can 3-1 and Can 10. Fao cells displayed a **gene** expression pattern near to that of liver (Fig. 1). Ten over the fifteen genes examined were expressed at 60% or more of the liver level. The expression of Ntcp was low (3%) as well as that of the three isoforms of organic anion transporters involved in the uptake of bile acids by the rat liver, i.e., Oatp1/1a1 (0.7 %), Oatp2/1a4 (23%), Oatp4/1b2 (16%). In contrast, high **gene** expression of most ABC transporters was found. Among the nine ABC **genes** examined, seven (Mdr1b, Mdr2, Mrp1, Mrp2, Mrp3, Abcg5 and Abcg8) were expressed at a higher level than in liver. This was particularly striking for Abcg5 and Abcg8, their mRNA being 80 and 160 times more abundant than in liver, respectively. The level of Bcrp was similar to that in normal rat liver, whereas that of Bsep was negligible (0.04 %). The **gene** expression of the nuclear receptor FXR and the transcription factor SHP was of the same order in Fao cells and liver.

When we examined the polarized lines, Can 3-1 and Can 10 (Fig. 2), we found that the **gene** expression pattern of Fao was almost maintained in these subclones, although some differences were observed. Can 3-1 cells displayed a significant lower level for Oatp4/1b2, Mdr2, Mrp2, Abcg5, Abcg8. Moreover in these cells Bsep mRNA could not be detected. Can 10 cells **produced** lower amounts of Oatp2/1a4 and Oatp4/1b2 **mRNA** whereas that of Mrp3, Bcrp and Bsep was increased. However Bsep **mRNA level** remained still very low (0.53% of liver level). In both sublines **mRNA level** of FXR and SHP was reduced but attained at least 30 % of the liver level.

The next step of this study was to ascertain that hepatobiliary transporters were localized at their appropriate poles in polarized Can lines. Mdr proteins (detected with the C219 antibody, that recognizes Mdr1a, Mdr1b, Mdr2) and Mrp2 were chosen as typical apical transporters; Mrp3 and Ntcp as basolateral ones. In non-polarized Fao cells, apical transporters were exclusively localized in an intracellular compartment (Fig. 3b,k), basolateral transporters were mostly present on the whole plasma membrane (Fig. 4b,k) and ZO-1 was discontinuously distributed on this membrane (Figs. 3c,l, 4c,l). In Can 3-1 and Can10 cells these proteins displayed a polarized localization (Figs. 3, 4). Apical transporters were confined in the BC delineated by the tight junction-associated protein ZO-1 (Fig. 3e,h,q). Basolateral transporters were excluded from BC and present on the rest of the PM and in the cytoplasm (Fig. 4e,h,n,q).

Thus, these two polarized sublines, in particular Can 10 cells expressed at a high level several hepatobiliary transporters, which are localized in similar PM domains to those of normal polarized hepatocytes.

Effects of bile acids on transporters expression, targeting, and polarity

To investigate the effects of bile acids, cells were cultured for 6 days with taurocholic acid (TCA), chenodeoxycholic acid (CDCA) or ursodeoxycholic acid (UDCA). TCA was tested because it is the most abundant natural bile acid in rat bile (Monte et al. 1997), whereas CDCA, which is also abundant in this species, is a very strong natural ligand for the nuclear receptor FXR (Lew et al. 2004, Makishima et al. 1999). UDCA was included as comparison because it is a poor ligand for FXR (Lew et al. 2004, Makishima et al. 1999). However, this bile acid has very interesting properties regarding its ability to stimulate bile formation (Scharschmidt & Lake, 1989). In each case 50 μ M was used because it is high enough to be considered as similar to serum total bile acid concentrations reaching the liver parenchyma in situations of mild cholestasis or transiently in postprandial periods. Moreover this

concentration is low enough not to induce any toxic effect and it is far below the critical micellar concentration for most bile acid species. Indeed we have verified that it was not toxic for the three cell lines examined here.

No significant change in **gene** expression was observed in TCA- and UDCA-treated cells, whatever the line examined, except for Bsep in UDCA-treated Fao cells (Fig. 5g). In contrast the **gene** expression of some hepatobiliary transporters, namely Mdr2, Bsep, and to a lesser extent Mdr1b and Oatp2/1a4, together with that of SHP, was upregulated by CDCA in Fao and Can 10 cells (Fig. 5f,g,e,c,o). Bsep was the most markedly induced (280 fold in Fao and 7 fold in Can 10), but it was also by far the less expressed in non-treated cells. Its level reached 12% and 4% of liver level in Fao and Can 10 treated by CDCA, respectively. For the other CDCA-induced genes, a 2-4 fold increase was observed in CDCA-treated cells. In Can 10 cells cultured for only 3 days with CDCA, upregulation of Bsep, Mdr2 and SHP was already induced, but at a slightly lower level than after a 6 days-treatment (results not shown). No significant changes in gene expression were induced by treatment of Can 3-1 with CDCA.

The effect of CDCA on Mdr was examined at the protein level: Western blotting was performed and the protein was quantified (Fig. 6a,b). We effectively found a 4 fold increase in Fao and Can 10 cells cultured with CDCA, and no change in Can 3-1 cells.

Finally, the effects of bile acids on the polarized state of the three cell lines and on the targeting of transporters were investigated. No polarization of Fao cells was induced whatever the bile acid used (results not shown). In the case of Can 10 cells that form long tubular BC, enhancements of the volume of BC were induced by UDCA, CDCA, but not frequently by TCA (Fig. 7). However, polarity was well maintained in cells treated with bile acids, as illustrated by the localization of the apical marker DPPIV (Fig. 7b,d,f,h) and the confocal analysis of the localization of Mrp2 with ZO-1 (Fig. 7i-x) and of Mdr with ZO-1 (Fig. 8e-h). Can 3-1 polarity was also maintained after treatment with bile acids. Besides Mdr proteins were more concentrated in the BC, and consequently better targeted, in CDCA-treated Can 3-

1 cells (Fig. 8c,d). In the case of Can 10, these apical proteins were already correctly targeted to BC in non-treated cells (Fig. 8e,f).

The effect of bile acids on the paracellular permeability of tight junctions was evaluated using FITC-labeled dextrans. As shown in Fig. 9 paracellular permeability of both lines was enhanced in cells treated with bile acids.

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Discussion

The present study was designed to investigate in new polarized rat cell lines the **gene** expression, localization, and inducibility by bile acids of a large panel of proteins implicated in hepatobiliary transport. This analysis was undertaken for two reasons. Firstly, whereas most in vivo experiments on bile transport and biliary elimination of drugs are performed on rats, in vitro experiments are rarely performed with rat cell lines. Secondly, human hepatic cell lines, such as Huh7, Hep G2, Hep3B, PLC/PRF/5 are the most popular systems for in vitro studies, but most of them are poor models for transport studies, and are not or badly polarized.

Comparison of the expression and localization of transporters in hepatic cell lines

In the non-polarized parental Fao cells the majority of the thirteen transporters examined and the two nuclear receptors FXR and SHP are expressed at a level, similar and often higher than in liver. Strikingly, efflux transporters, whether they are apical or sinusoidal, are well expressed, except Bsep, whereas influx transporters expression is rather low. This expression profil is well maintained and even improved for some transporters, in Can 10 cells. In contrast several transporters, in particular Mrp2, are badly expressed in Can 3-1.

We recently analyzed the expression of seven transporters in the polarized rat-human hybrid WIF-B9 clone (Briz et al. 2007). Some transporters (Mdr1b, Mrp1, Mrp2) are expressed at similar levels in WIF-B9 and Can 10, but others (Bcrp, Mdr2, and to a lesser extent Bsep and Mrp3) are expressed at much lower levels in WIF-B9. So among the polarized clones established from Fao, Can 10 has the better **gene** expression pattern of transporters.

Two human cell lines Hep G2 and HepaRG were previously characterized for the **gene** expression of a dozen of transporters (Le Vee et al. 2006). As observed here for rat cell lines, these human lines are characterized by a very low expression of uptake transporters, and a

better expression of efflux pumps. In this study mRNA levels were compared to levels of primary hepatocytes (and not to liver levels as in our study). Assuming that expression levels are not very different in liver and primary hepatocytes, the comparison between Can 10 and human lines levels shows that **genes** of transporters are better expressed in Can 10 than in Hep G2 and even HepaRG, except for Bsep that is well expressed in HepaRG cells.

Compared to Fao and human cell lines both Can sub-lines present the advantage to correctly localized transporters: apical transporters are present at the BC and basolateral transporters are absent from BC and present on the rest of the PM. These localizations were observed for all cells, attesting to the homogeneity and stability of the polarized phenotype of both Can clones. In the case of Mdr proteins detected with the C219 antibody, we observed a higher enrichment in BC with a weaker cytoplasmic staining in Can 10, suggesting a better targeting of these apical transporters in Can 10 than in Can 3-1 cells.

Comparison of the effects induced by bile acids in hepatic cell lines

The consequences of treatment of Fao, Can 3-1 and Can 10 cells by various bile acids were evaluated. CDCA is by far the most efficient of the three bile acids tested. As the other bile acids, CDCA induces increase of tight junction paracellular permeability of Can 3-1 and Can 10 cells. It also promotes, as UDCA, a marked enhancement in the volume of Can 10 BC. Such changes could result from the accumulation of bile acids in BC during the 6 days treatment. In contrast to TCA and UDCA, CDCA induces also in Fao and Can 10 cells **upregulation** of a same subset of transporters. CDCA is known to be the most efficient FXR activator (Lew et al. 2004) and the genes overexpressed in CDCA-treated Fao and Can 10 cells have been shown to be activated by bile acid-FXR-dependent mechanisms (Lee et al. 2006; Eloranta and Kullak-Ubrick, 2005). FXR response elements have been identified on the promoters of the most induced genes, Bsep, Mdr2 and SHP (Ananthanarayan et al. 2001; Huang et al. 2003; Goodwin et al. 2000; Lu et al. 2000). Consequently we attribute the

overexpression of transporters occurring in CDCA-treated cells to FXR-mediated transcriptional activation.

Our data demonstrate that induction of **gene** expression and targeting of transporters are dissociable. Indeed in CDCA-treated Fao cells, some transporters are overexpressed but their localization is not improved and the cells remain unpolarized. Conversely a better targeting of Mdr proteins occurs in CDCA-treated Can 3-1 cells without any change in the **mRNA** level of these transporters. This improvement in targeting could result from some changes in cytoskeletal structures or components known to be involved in the targeting of apical transporters to the BC, as for example myosin II light regulatory chain (Chan et al. 2005).

It has been previously shown that CDCA can reversibly induce polarization of Fao cells and Fao-derived revertant C2rev7 cells (Ng et al. 2000). In our study, no sign of polarization was found in CDCA-treated Fao cells. This discrepancy might have been due to the fact that we used a lower and non-toxic CDCA concentration and a shorter period of treatment. Indeed, when we applied the drastic conditions described by Ng et al. (2000), no polarized cells were obtained. In the same way, when plated at low density, Can 3-1 cells did not polarize even in presence of CDCA.

Information about the effects of bile acids in other hepatic cell lines is scarce. The level of Ntcp is hardly increased (25%) and its targeting to the BC slightly induced in TC-treated WIF-B9 cells (Konieczko et al. 1998; Sai et al. 1999). BSEP expression is down-regulated by lithocholate and up-regulated by CDCA in Hep G2 cells (Yu et al. 2002; Lew et al. 2004). It is also highly induced by CDCA in HepaRG cells (Le Vee et al. 2006), but the effect of CDCA on other transporters was not examined. In none of the previous related studies, the effects of several bile acids on the **gene** expression of a large panel of transporters were analyzed.

Usefulness of Can 10 cells

The aim of this study was to evaluate if the new rat polarized lines, Can 3-1 and Can 10, could be pertinent rat hepatic models for further *in vitro* studies on vectorial transport processes involved in the biliary secretion of bile acids and other cholephilic compounds. The most interesting line is undoubtedly Can 10. Compared to their parental Fao cells, Can 10 cells have retained the same **gene** expression profile of transporters, many of these transporters being expressed at a level, similar and even higher than in the liver. Moreover the localization of these transporters was improved in Can 10 cells. Finally in Can 10 and Fao the expression of the same subset of transporters was induced by CDCA. This implies that Can 10 cells, while acquiring a polarized phenotype, have retained regulatory pathways controlling transporter levels. Another advantage of Can 10 is the formation of highly organized BC in which bile acids, organic anions and cations are vectorially transported (Peng et al. 2006).

In summary Can 10 is an interesting rat cell line for studying **synthesis**, regulation, and targeting of many hepatobiliary transporters. For example, since it expresses high levels of Abcg5 and Abcg8, it could be very useful to study how these transporters interact, traffic and promote biliary sterol excretion. Moreover, because of its functional polarity and its capacity of transport, Can 10 represents an attractive model for *in vitro* pharmacological and toxicological studies.

Acknowledgments

We greatly thank A. Hubbard, D. Keppler, B.R. Stevenson, and B. Stieger for kindly providing antibodies. We also thank V. Nicolas (IFR141-ITFM, Chatenay Malabry, France) and Mickael Bourge (INSERM U757, Orsay, France) for help with confocal analysis and figure artwork, respectively. This work was supported by a grant from Egide (PAI Picasso) and the Acción Integrada Hispano-Francesa (HF2003-0089). The group is member of the Network for Cooperative Research on Membrane Transport Proteins (REIT), co-funded by the Ministerio de Educación y Ciencia, Spain and the European Regional Development Fund (ERDF) (Grant BFU2005-24983-E/BFI) and belongs to the “Centro de Investigación Biomédica en Red” for Hepatology and Gastroenterology Research (CIBERehd), Instituto de Salud Carlos III, Spain.

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Table 1. Oligonucleotide sequences of primers used in real-time quantitative RT-PCR.

Target sequence	Primer Sequence (5'-3')		Amplicon Position	(bp)	GenBank Accession number.
	Forward	Reverse			
Ntcp	CGTTGCCGGAATGTTTGTCT	TGCCCTCTGTCTCAGTTCATG	1253-1327	75	NM_017047
Oatp1/Oatp1a1	CTACTGCCCTGTTCAAGGCC	ATTGTATCTCTCAGGATCCGAGG	2108-2175	68	L19031
Oatp2/Oatp1a4	TGCGGAGATGAAGCTCACC	TCCTCCGTCACCTTCGACCTT	2032-2100	69	U88036
Oatp4/Oatp1b2	AGACGTCCCATCACACCAC	GCCTCTGCAGCTTTCCTTGA	86-153	68	AJ271682
Mdr1b/Abcb1b	AACCCCATGGCTGGAACA	TGGGCTCCTGGGACACAAT	3288-3387	100	NM_012623
Mdr2/Abcb4	ACTGTCCGGAATGCAGATGTC	TCTTTATCAGCTCACTGTGGCTT	1800-1881	82	NM_012690
Bsep/Abcb11	GCCATTGTGCGAGATCCTAAA	TGCAGGTCCGACCCTCTCT	3956-4073	118	NM_031760
Mrp1/Abcc1	GTGAAGCTTTCGGTACTGGAA	TACATGATTGCACAGGAAAAGGA	742-831	90	AJ 277881
Mrp2/Abcc2	TGATCGGTTTGCTGAAGAGCT	CGCACATCCCAACACAAA	1119-1257	139	NM_012833
Mrp3/Abcc3	TCCCACTTCTCGGAGACAGTAACT	CACCTTAGCATCACTGAGGACCTT	3481-3570	90	NM_080581
Bcrp/Abcg2	ATGATGCTCTTTTCTGGCCTCT	AAGCCATATCGAGGAATGCTAAA	1723-1814	92	NM_181381
Abcg5	GGCTCGGCACAGCTTAGG	CTGGCATGATTTGATGTTCCA	216-300	85	NM_053754
Abcg8	CCCTGATCCGTCGTCAGATT	AGAGACATCAGGCAGGCTTCT	1309-1393	85	AF351785
FXR	GTGACAAAGAAGCCGCGAAT	GCAGGTGAGCGGTTGTAAT	481-594	114	U18374
SHP	CTCGGTTTGATACAGTGTGAC	GCATATTGGCCTGGAGGTTTT	920-994	75	D86580
UBQ-C	TGGCTATTAATTCTTCAGTCTGCATTC	GGCAAATGGCTAGAGTGCAGAG	972-1042	71	D16554

Figures legends

Figure 1 Expression of thirteen hepatobiliary transporters and two nuclear receptors in Fao cells. Steady state mRNA levels were determined by real-time RT-PCR. They are expressed using a logarithmic scale as % of mRNA level in rat liver and correspond to the mean of values from three independent cultures. The results shown were normalized using 18S rRNA (a similar pattern of expression was obtained using polyubiquitin-C). Liver level is visualized by a dotted line.

Figure 2 Expression of hepatobiliary transporters and nuclear receptors in Can 3-1 and Can 10 cells. Steady state mRNA levels were determined by real-time RT-PCR. They are expressed as % of corresponding mRNA level in parental Fao cells and are the mean of values from 3 and 4 independent cultures for Can 3-1 and Can 10 lines, respectively. The results shown were normalized using 18S rRNA (a similar pattern of expression was obtained using polyubiquitin-C). Fao level is visualized by a dotted line.

Figure 3 Immunolocalization of apical hepatobiliary transporters in Fao (a-c; j-l) and Can clones (d-i; m-r). Each transporter was colocalized with the tight junction-associated protein ZO-1. In each case the corresponding phase-contrast image is given on the left. Some BC formed by Can 3-1 and Can 10 cells were indicated by short (simple BC formed between two cells) or by long arrows (complex BC). In non-polarized Fao cells, apical transporters (b,k) were exclusively localized in an intracellular compartment and ZO-1 was discontinuously distributed on the plasma membrane (c,l). In Can 3-1 and Can 10 cells apical transporters were confined in the bile canaliculi (e,h,q) delimited by ZO-1 (f,i,r). The anti-Mdr-P-glycoprotein C219 antibody detects Mdr1a, Mdr1b and Mdr2. Due its very low level of expression (Fig. 2), Mrp2 was not detected in Can 3-1 cells (n). Scales: 10 μ m.

Figure 4 Immunolocalization of basolateral hepatobiliary transporters in Fao (a-c; j-l) and Can clones (d-i; m-r). Each transporter was colocalized with the tight junction-associated protein ZO-1. In each case the corresponding phase-contrast image is given on the left. Some BC formed by Can 3-1 and Can 10 cells were indicated by short (simple BC formed between two cells) or long arrows (complex BC). In non-polarized Fao cells basolateral transporters (b,k) were mostly present on the whole plasma membrane and ZO-1 was discontinuously distributed on this membrane (c,l). In Can 3-1 and Can10 cells basolateral transporters (e,h,n,q) were excluded from the canaliculi sealed by ZO-1 (f,i,o,r) and present on the rest of the PM and in the cytoplasm. Scales: 10 μ m.

Figure 5 Effects of bile acids on the expression of hepatobiliary transporters and nuclear receptors in Fao and Can 10 cells (a-o). Steady state mRNA levels, determined by real-time RT-PCR, were expressed as % of corresponding mRNA level in cells cultured with DMSO and without bile acids. They are the mean of values from 2-3 independent cell cultures. The results shown were normalized using 18S rRNA or polyubiquitin-C; similar results were obtained in both cases. * $p < 0.05$ when compared to cells cultured with DMSO and without bile acids.

Figure 6 Effects of bile acids on Mdr protein level in Fao and Can clones. Western blot (a). Each lane was loaded with a cell lysate aliquot (25 μ g of total protein). Mdr proteins (detected with the C219 antibody that recognizes Mdr1a, Mdr1b, Mdr2) were codetected with α -tubulin. Mdr level was normalized using α -tubulin as the internal control and expressed as % of the level of DMSO treated cells (b). The experiment was performed three times with lysates from 2-3 independent cultures. One representative experiment is shown. In all cases a significant increase in Mdr level was obtained in Fao and Can 10 cells treated with CDCA.

Compared to liver, Fao and both Can clones expressed a higher Mdr protein level; this result is in agreement with measurement of Mdr mRNA level (Fig. 1, Fig. 2).

Figure 7 Effects of bile acids on the polarity of Can 10 cells.

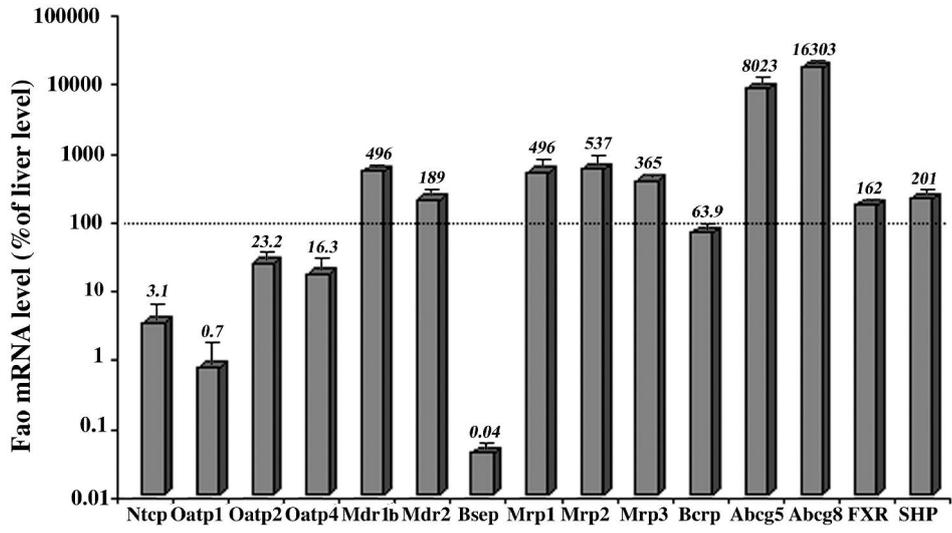
Immunolocalization of the apical marker DPPIV in cells treated by bile acids (a-h). In each case the corresponding phase contrast image is given on the left. DPPIV is present at the membrane of BC, that are greatly inflated (*) in CDCA and UDCA-treated cells. Confocal analysis of the localization of the apical transporter Mrp2 with the tight junction-associated protein ZO-1 in cells treated by bile acids (i-x). In each case, the localization of Mrp2, ZO-1, and the corresponding merge image (compilation of 20-30 xy sections taken in 0.3 μm steps) are shown (left to right) followed by one xy section taken in the middle of the cell layer. Note the marked enhancement of the volume of BC in CDCA and UDCA-treated cells, particularly well visible in the xy sections. Scales: 10 μm .

Figure 8 Confocal analysis of the localization of apical Mdr transporters (detected with the C219 antibody) with the tight junction-associated protein ZO-1 in DMSO and CDCA treated Can 3-1 (a-d) and Can 10 cells (e-h). For each culture an image corresponding to a compilation of 18-21 xy sections and a graph of the distribution of Mdr along the dotted line indicated on the corresponding compilation are shown. Note that Mdr proteins are well concentrated at the BC of Can 3-1 only after CDCA treatment (c,d). In contrast these apical markers are already highly concentrated at the BC of Can 10 cells cultured without CDCA (e,f). Scales: 10 μm .

Figure 9 Effects of bile acids on tight junction paracellular permeability of Can clones, evaluated by measuring the paracellular diffusion of FITC-dextran (molecular weight 4,400)

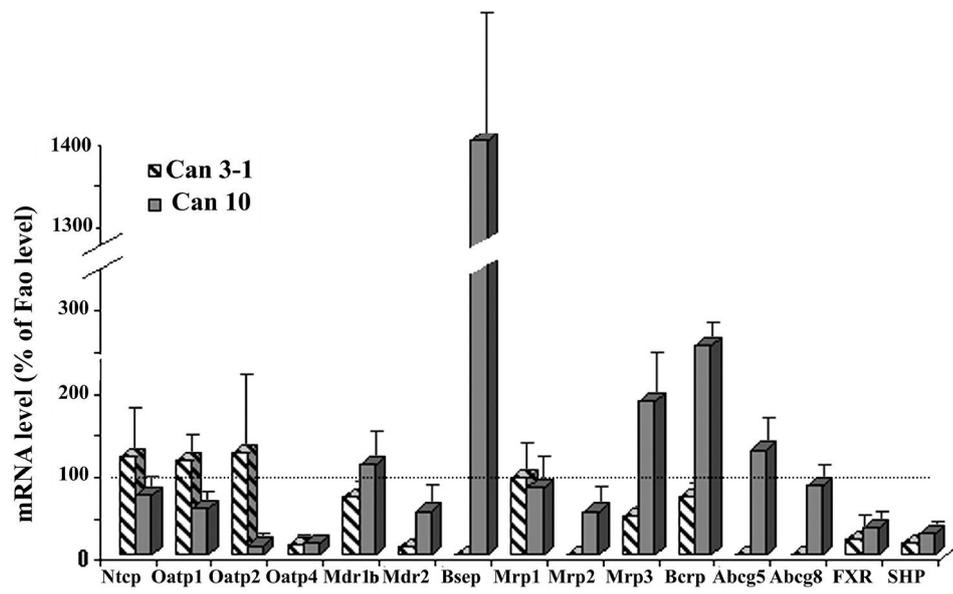
into the BC. The values are means from 2-3 independent cell cultures. * $p < 0.05$ when compared to untreated cells.

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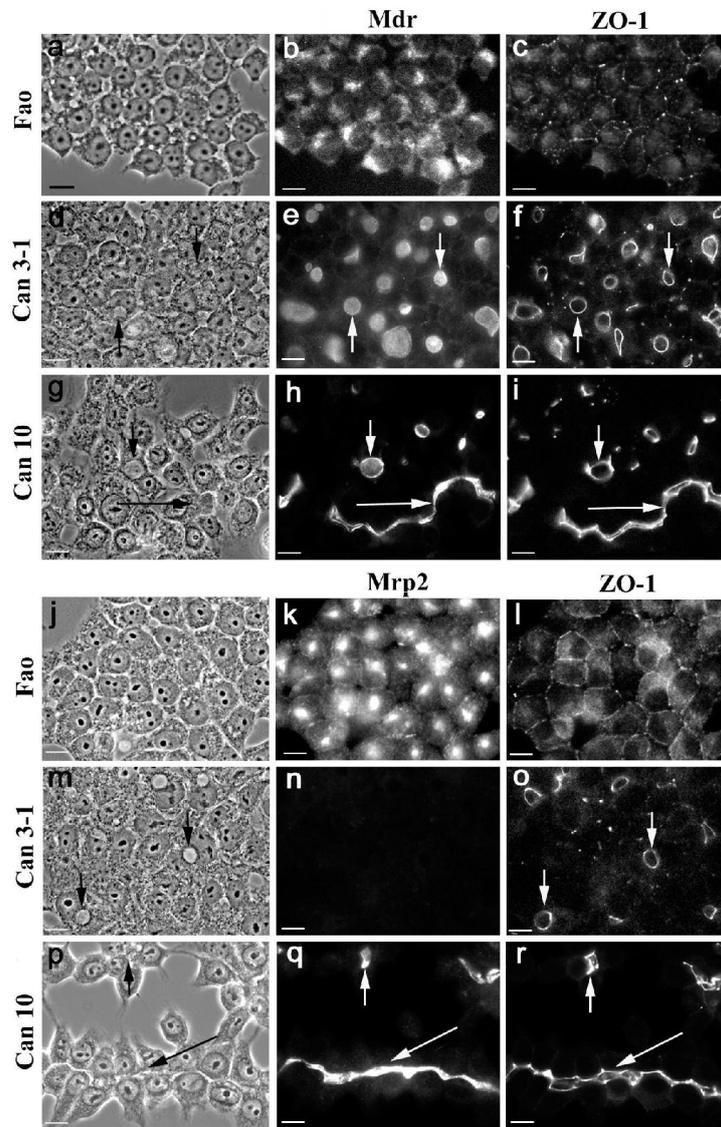
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Review

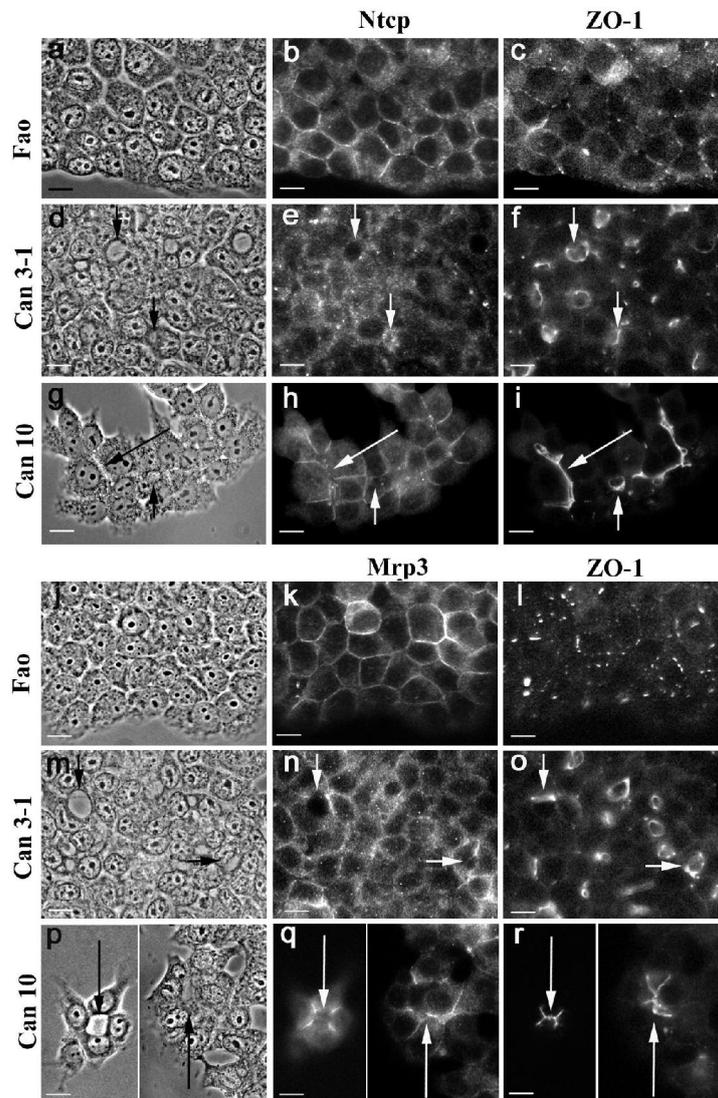


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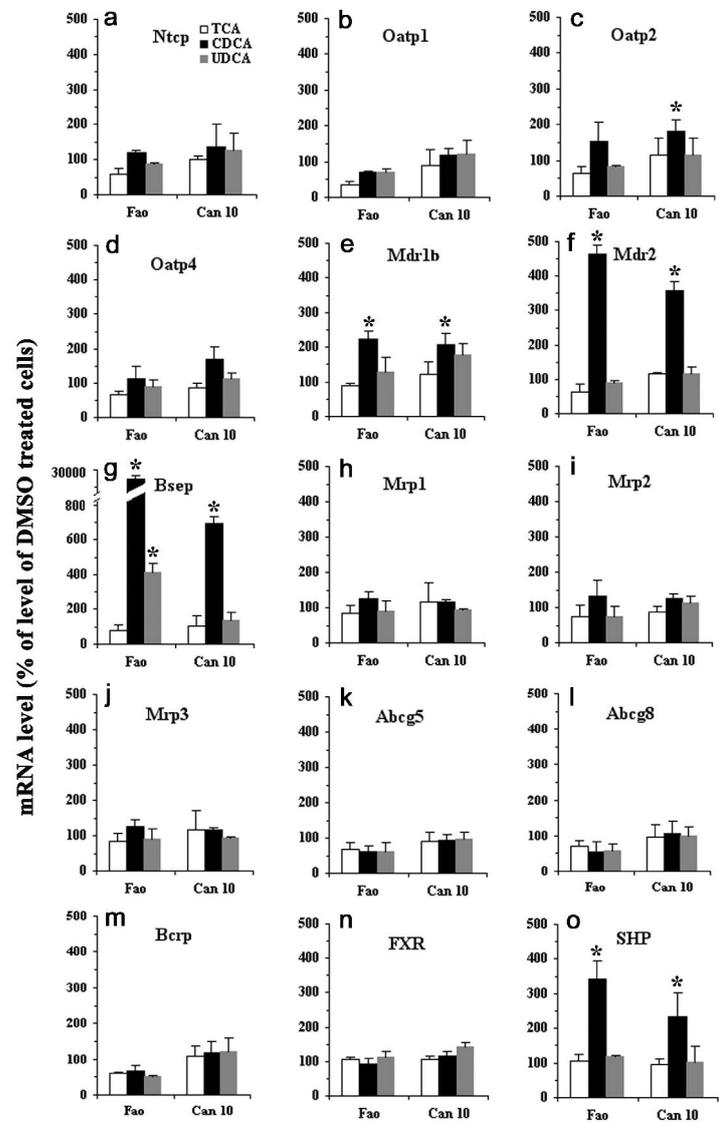
Review



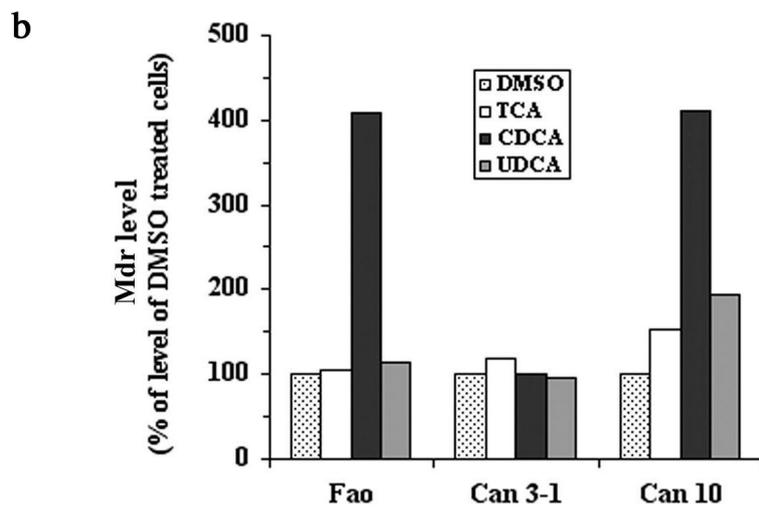
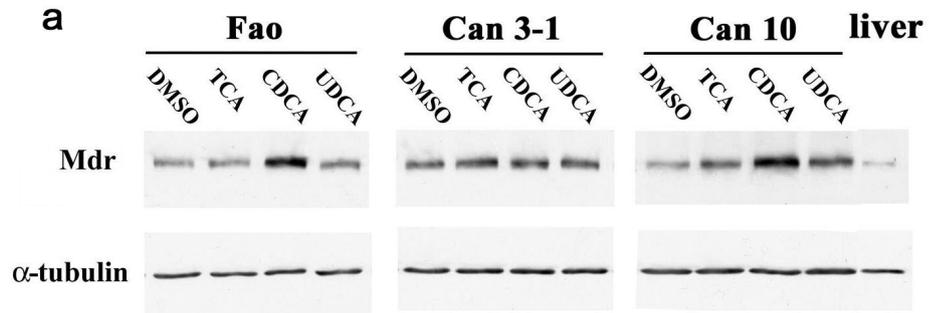
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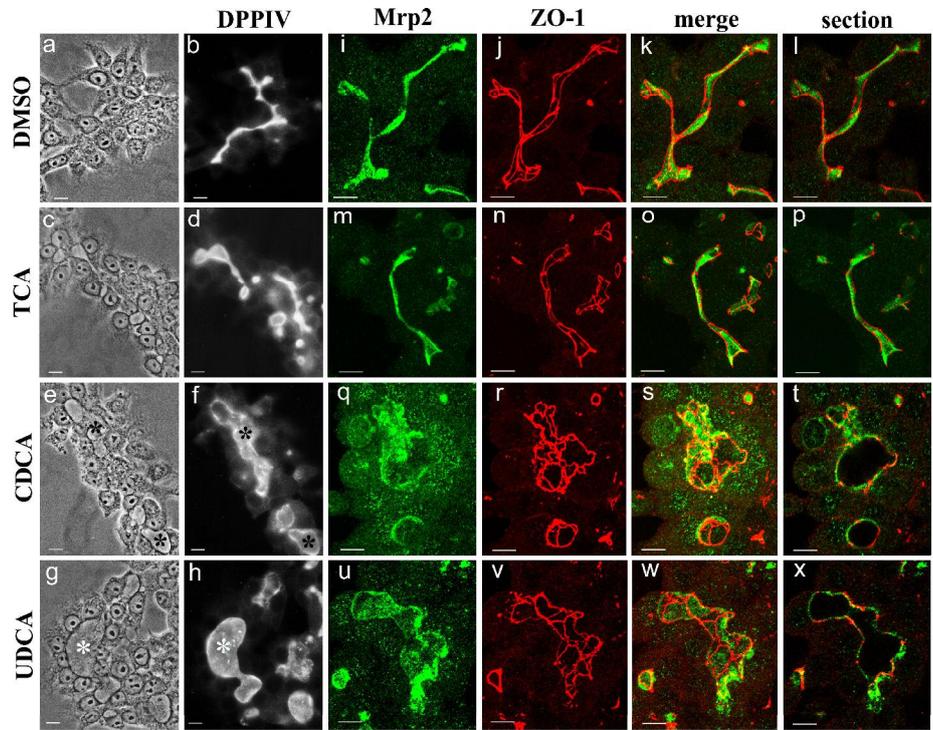


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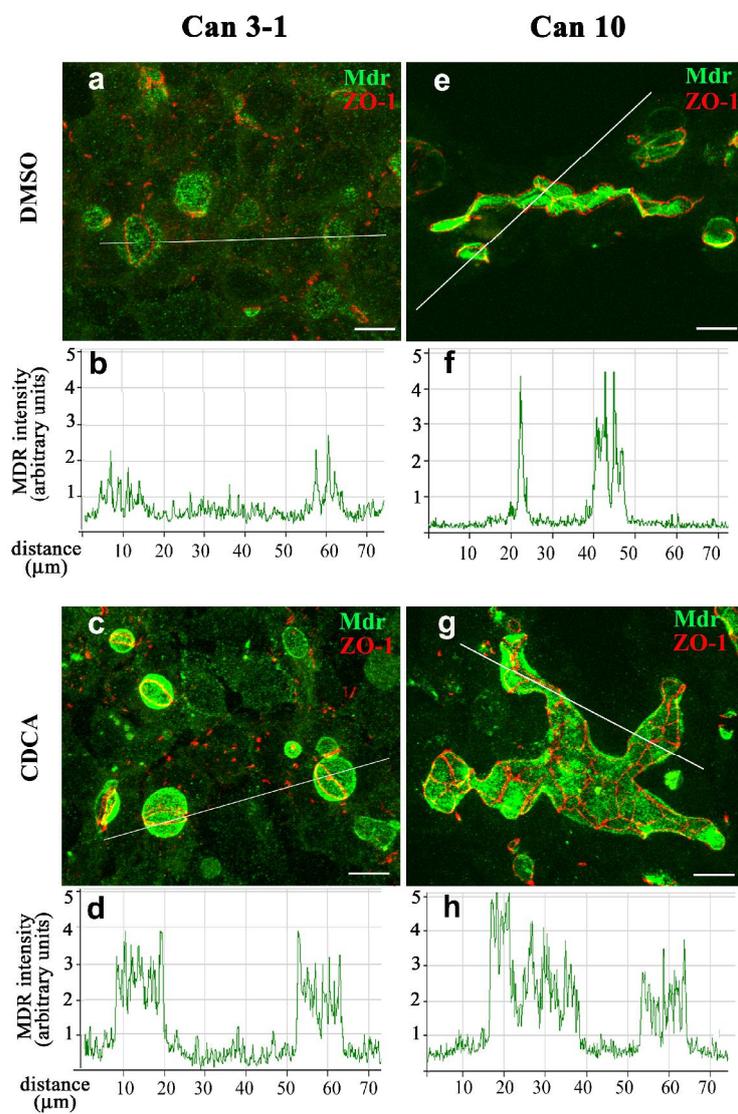
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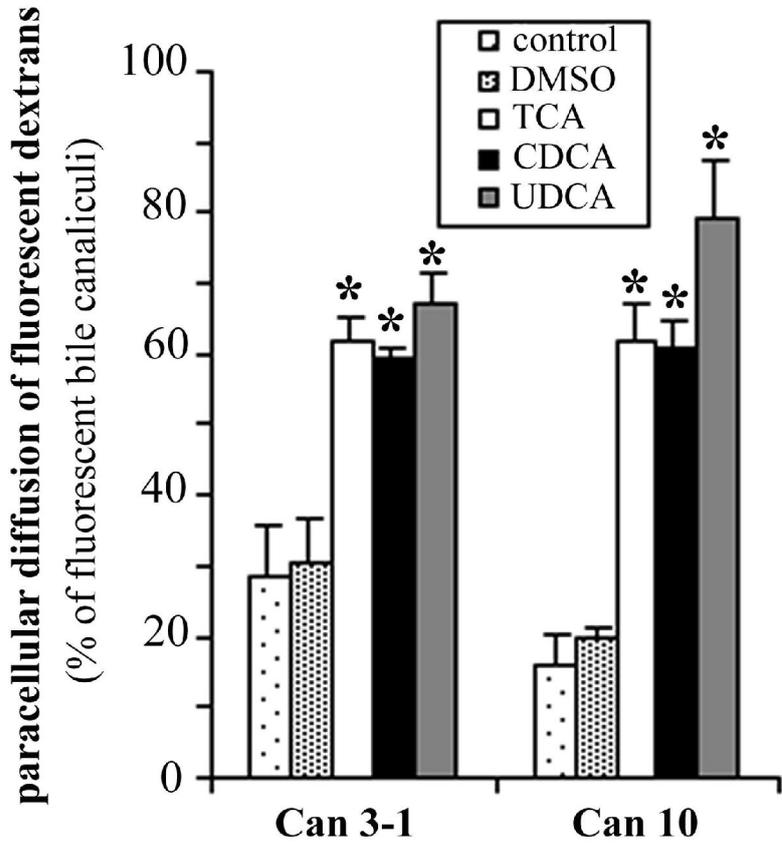


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