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cmgh ORIGINAL RESEARCH



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SUMMARY

A fluorescent resonance energy transfer–based, highthroughput screen was established to identify inhibitors of organic solute transporter (OST) α -OST β -mediated bile acid efflux. We identified clofazimine as an inhibitor of OST α -OST β and showed that such inhibition enhanced intestinal farnesoid X receptor activation.

BACKGROUND & AIMS: The organic solute transporter α-β (OSTα-OSTβ) mainly facilitates transport of bile acids across the basolateral membrane of ileal enterocytes. Therefore, inhibition of OSTα-OSTβ might have similar beneficial metabolic effects as intestine-specific agonists of major nuclear
receptor for bile acids, the farnesoid X receptor (FXR). However, no OSTα-OSTβ inhibitors have yet been identified.

METHODS: Here, we developed a screen to identify specific inhibitors of $OST\alpha$ - $OST\beta$ using a genetically encoded Förster Resonance Energy Transfer (FRET)-bile acid sensor that enables rapid visualization of bile acid efflux in living cells.

RESULTS: As proof of concept, we screened 1280 Food and Drug Administration–approved drugs of the Prestwick chemical library. Clofazimine was the most specific hit for $OST\alpha$ - $OST\beta$ and reduced transcellular transport of taurocholate across Madin–Darby canine kidney epithelial cell monolayers expressing apical sodium bile acid transporter and $OST\alpha$ - $OST\beta$ in a dose-dependent manner. Moreover, pharmacologic inhibition of $OST\alpha$ - $OST\beta$ also moderately increased intracellular taurocholate levels and increased activation of intestinal FXR target genes. Oral administration of clofazimine in mice (transiently) increased intestinal FXR target gene expression, confirming $OST\alpha$ - $OST\beta$ inhibition in vivo.

CONCLUSIONS: This study identifies clofazimine as an inhibitor Q^{696} of OST α -OST β in vitro and in vivo, validates OST α -OST β as a 97 drug target to enhance intestinal bile acid signaling, and 98 confirmed the applicability of the Förster Resonance Energy 99 Transfer-bile acid sensor to screen for inhibitors of bile acid efflux pathways. (*Cell Mol Gastroenterol Hepatol 2017*; \blacksquare - \blacksquare ; 101 *https://doi.org/10.1016/j.jcmgh.2017.11.011*) 102

Keywords: Fluorescence Resonance Energy Transfer (FRET); FXR; $OST\alpha$ - $OST\beta$; Bile Acids.

Abbreviations used in this paper: ASBT, apical sodium-dependent bile acid transporter; BAS, bile acid sensor; FACS, fluorescence-activated cell sorting; FDA, Food and Drug Administration; FGF15/19, fibroblast growth factor 15/19; FRET, fluorescent resonance energy transfer; FXR, farnesoid X receptor; MDCKII, Madin-Darby canine kidney epithelial cells; mRNA, messenger RNA; nucleoBAS, nucleus-localized bile acid sensor; OST α -OST β , organic solute transporter α - β ; TCDCA, taurochenodeoxycholic acid; TICE, transintestinal cholesterol excretion; U2OS, human bone osteosarcoma epithelial cells © 2017 The Authors. Published by Elsevier Inc. on behalf of the AGA Institute. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/). 2352-345X

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2 van de Wiel et al

B ile acids are released in the duodenum during a meal, where they act as digestive detergents crucial 117**Q7** 118<mark>0</mark>8 119<mark>0</mark>9 for intestinal absorption of lipids, fat-soluble vitamins, and 120010 other lipophilic nutrients.¹ After the meal, most bile acids 121 are reabsorbed from the ileum and return to the liver via the 122 hepatic portal vein. In the hepatocyte, the bile acid pool is 123 replenished via de novo bile acid synthesis from choles-124 terol.² Subsequently, bile acids are conjugated and stored in 125 the gallbladder until the next meal. The meal-regulated 126 dynamics renders bile acids as potent signaling molecules 127 that modulate triglyceride, lipid, glucose, and energy homeostasis, making bile acid signaling an interesting target 128 129 for metabolic diseases.³ Activation of the farnesoid X 130 receptor (FXR), a major nuclear receptor involved in bile 131 acid signaling, has been suggested to be beneficial in many 132 metabolic disorders. In particular, selective intestinal acti-133 vation of FXR led to metabolic improvements such as 134 enhanced glucose tolerance, reduced diet-induced weight gain, and reduced inflammation,⁴ but also enhanced tran-135 136 sintestinal cholesterol excretion (TICE).5

137 The organic solute transporter α - β (OST α -OST β) mainly 138 facilitates transport of bile acids across the basolateral 139 membrane of ileal enterocytes.^{6,7} It consists of 2 proteins 140 forming a heterodimer, OST α (encoded by *SLC51A*) and its 141 subunit OST β (encoded by *SLC51B*), which are both 142 required for normal trafficking and function of OST α -OST β .

143 $OST\alpha$ knockout mice show a reduction in the bile acid 144 pool and serum levels combined with increased FXR activation in ileal enterocytes.⁸ In addition, deficiency of OST α is 145 protective for liver injury during obstructive cholestasis, 146 147 and leads to decreased body fat and lipid accumulation and improved insulin sensitivity.⁹⁻¹¹ Furthermore, $OST\alpha$ 148 149 knockout mice show increased elimination of cholesterol in 150 the feces and decreased levels of cholesterol and triglyceride in serum.^{8,12} These data point to $OST\alpha$ - $OST\beta$ as a novel 151 target to treat diabetes and obesity, but also lipid and 152 cholesterol disorders, for instance, by inducing the TICE 153 154 pathway via intestinal FXR activation.¹³ However, no inhibitors for $OST\alpha$ -OST β have yet been identified and 155 techniques available for measuring bile acid efflux are 156 157 limited. Therefore, we developed a novel assay for cell-158 based high-throughput screening to identify specific 159 inhibitors for OST α -OST β , making use of a fluorescent 160 resonance energy transfer (FRET)-based bile acid sensor that enables rapid visualization of bile acid efflux in living 161 cells.¹⁴ This screen specifically measures bile acid efflux, 162011 which simultaneously provides a readout based on 163 164 increased FXR activation as a consequence of $OST\alpha$ - $OST\beta$ 165 inhibition. We screened 1280 Food and Drug Administration 166 (FDA)-approved drugs of the Prestwick chemical library and 167 confirmed several positive hits. Here, we show that inhibi-168 tion of $OST\alpha$ - $OST\beta$ leads to intestinal FXR activation.

170 171 Materials and Methods

172 Reagents

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The Prestwick chemical library (1280 FDA approvedcompounds in 96-well plates) was purchased from

Prestwick Chemical. Bifonazole, bromhexine HCl, clofazi- ⁰¹²176 mine, lovastatin, meclozine 2HCL, and simvastatin were 177 purchased from Sigma-Aldrich. The fibroblast growth factor 178 (FGF)19 human enzyme-linked immunosorbent assay kit 179 was purchased from BioVendor R&D. [³H]-taurocholate and 180 [¹⁴C]-inulin were purchased from Perkin Elmer. 181

Cell Culture

184 Human bone osteosarcoma epithelial cells (U2OS) wild-185 type cells (HTB-96; American Type Culture Collection) 186 were cultured in Dulbecco's modified Eagle medium 187 (high glucose) supplemented with 10% fetal bovine 188 serum, 1% penicillin/streptomycin, and 1% L-glutamine. 189 NucleoBAS (bile acid sensor localized in the nucleus),¹⁴ Q14 190 NA⁺-taurocholate co-transporting polypeptide, and 191 $OST\alpha$ - $OST\beta$ -expressing U2OS cells were engineered by 192 transfecting cells using polyethylenimine. Stable cell lines 193 were generated by colony picking using cloning rings over 194 well-separated colonies. Apical sodium-dependent bile 195 acid transporter (ASBT) and mouse $OST\alpha$ - $OST\beta$ 196 co-expressing Madin-Darby canine kidney epithelial cells 197 (MDCKII) cells were a gift from Paul Dawson (Emory 198 University School of Medicine, Atlanta, GA).¹⁵ All cells 199 were cultured at 5% CO₂ at 37°C. 200

Fluorescence-Activated Cell Sorting

203 Two days before the experiments, wild-type and trans-204 fected U2OS cells were cultured in 5% charcoal-treated fetal 205 bovine serum to prevent bile acid overload of the sensor. 206 The adherent cell layer was trypsinized by 5 mmol/L EDTA 207 to create a suspension of single cells for fluorescence-208 activated cell sorting (FACS) analysis. Cells were harvested 209 by centrifugation and the resulting pellet was suspended in 210 FACS uptake buffer (0.3 mmol/L EDTA, 0.5% bovine serum 211 albumin, 0.01% NaN₃, and 10 mmol/L D-glucose), plated in 212 96-wells and subsequently incubated with 10 μ mol/L of 1 of 213 1280 compounds of the Prestwick Chemical library. After 5 214 minutes, 3 μ mol/L TCDCA was added and incubated for 30 215 minutes at room temperature while shaking. Citrine and 216 cerulean intensity was measured by a violet 405-nm laser. 217 Spectral range for emission detections were as follows: 218 cerulean, 450/40 nm; and citrine, 525/20 nm. 219

Taurocholate Uptake Assay in MDCKII–ASBT Cells

Cells were cultured in a 24-well plate format at 223 50%-60% confluency. The next day, cells were washed 224 with uptake buffer and incubated with compounds at 37°C 225 for 30 minutes. Uptake buffer was aspirated and a mix of 226 trace amounts of $[{}^{3}H]$ -taurocholate and 20 μ mol/L 227 nonradiolabeled trichloroacetic acid was administered. Q16228 After 2 minutes of incubation at 37°C, the cells 229 were washed using ice-cold phosphate-buffered saline and 230 lysed in 0.05% sodium dodecyl sulfate in distilled water. 231 Tritium signal of each sample was measured in 3 mL of 232 scintillation liquid. 233

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