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Abstract

Oxidative stress is a common feature in most hepatopathies. Accumulating evidences indicate that reactive oxygen species (ROS) induce a number of functional changes either deleterious or adaptive in the capability of the hepatocytes to produce bile and to secrete exogenous and endogenous compounds. This review is aimed to describe the mechanisms involved in these changes. For this purpose, we will summarize:

1.	The current evidence that acutely induced oxidative stress is cholestatic,
	by describing the mechanisms underlying the hepatocyte secretory failure,
	including the disorganization of the actin cytoskeleton and its most
	noticeable consequences, that is, the impairment of tight-junctional
	structures and the endocytic internalization of canalicular transporters
	relevant to bile formation.
2.	The role for oxidative-stress-activated signalling pathways in the
	pathomechanisms described above, particularly those involving Ca^{2+}
	elevation and its consequent activation via Ca ²⁺ of "classical" and "novel"
	PKC isoforms.
3.	The mechanisms involved in the adaptive response against oxidative stress
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3. The mechanisms involved in the adaptive response against oxidative stress mediated by ROS-responsive transcription factors, such as upregulation of GSH synthesis pathway, antioxidant enzymes, and hepatocellular efflux pumps.

4. The consequences on hepatocellular secretory function when this adaptive response can be surpassed by the sustained/high production of ROS. This deleterious effects include transcriptional and posttranscriptional changes in the expression of transporters relevant to bile formation, as has been shown to occur, for example, after long-term administration of aluminum to rats, in the Long-Evans Cinnamon rat (a model of chronic hepatic copper accumulation mimicking Wilson's disease), and in ischemia-reperfusion injury.

Keywords	Actin - Bile secretion - Calcium - Cholestasis - Oxidative stress -
(separated by "-")	Protein kinases - Signalling - Tight junctions

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Abstract

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45 Introduction

Due to its multiple energy-dependent functions, liver has a high mitochondrial 46 metabolic rate and is heavily engaged in detoxification mechanisms that involve 47 redox-enzyme systems. Since these are major sources of endogenous radical oxy-48 gen species (ROS), production of these highly reactive, cytotoxic compounds is 49 higher in liver as compared with most organs. Hence, hepatocytes are rich in 50 antioxidant defenses to counterbalance this oxidative challenge (Cesaratto et al. 51 2004). However, this borderline equilibrium makes liver highly susceptible to the 52 pro-oxidant injury induced by pathological conditions (Kaplowitz and Tsukamoto 53 1996). Oxidative stress (OS) is a common feature in most hepatopathies including 54 hepatic ischemia-reperfusion injury following hepatectomy or liver transplantation 55 (Czubkowski et al. 2011; Galaris et al. 2006); obstructive cholestasis (Vendemiale 56 et al. 2002); chronic cholestatic liver diseases (Copple et al. 2010; Salunga et al. 57 2007); sepsis-induced cholestasis (Sakaguchi and Furusawa 2006); viral (Simula 58 59 and De V 2010), toxic (Stehbens 2003), and autoimmune (Pemberton et al. 2004) hepatitis; alcoholic (Wu and Cederbaum 2009) and nonalcoholic (Koek et al. 2011) 60 steatohepatitis; and pathologies leading to hepatic accumulation of heavy metals, 61 such as iron (hemochromatosis, iron-loading anemia) (Alla and Bonkovsky 2005) 62 or copper (Wilson's disease) (Dalgic et al. 2005). 63

In recent years, evidence has accumulated that OS is cholestatic. Functional changes involve impairment of biliary secretion through both direct oxidative damage of cellular structures involved in this process or, more significantly, via modification of intracellular signal transduction pathways sensitive to changes in the intracellular redox state. We will summarize here the mechanisms involved in these alterations. Au1

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Fig. 140.1 Main transport systems and metabolic events involved in bile flow formation and the hepatic handling of the endogenous and exogenous cholephilic organic anions bilirubin and sulfobromophthalein (BSP), respectively. For details, see section "Normal Mechanisms of Bile Formation"

70 Normal Mechanisms of Bile Formation

Bile formation is an osmotic process driven by the vectorial transport of certain 72 osmotic solutes into bile, mainly bile salts and both oxidized (GSSG) and reduced 73 (GSH) glutathione (see Fig. 140.1). For these solutes to drive blood-to-bile water 74 transport, they need to be concentrated and retained into a confined space (the bile 75 canaliculus), sealed by tight-junctional structures localized in the paracellular 76 pathways. Once secreted, these solutes induce osmotic water movement, mainly 77 through aquaporins type 9 and 8, located at the basolateral and apical membranes, 78 respectively (Marinelli et al. 2011). 79

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This primary (canalicular) secretion is further modified by cholangiocytes dur-80 ing its transit along bile ducts, as a result of a balance between hormone-dependent 81 water and electrolyte secretion and, on the other hand, the obligatory absorption 82 of water, electrolytes and organic solutes (Elsing et al. 1996; Marinelli and 83 LaRusso 1996). 84

Bile salts are the predominant organic solutes in bile. The main sinusoidal 85 transport system for bile-salt uptake is the Na^+ -taurocholate cotransporting poly-86 *peptide*, which has been cloned from both rat (Ntcp, *Slc10a1*) (Hagenbuch et al. 87 1991) and human liver (NTCP, SLC10A1) (Hagenbuch and Meier 1994). Ntcp/ 88 NTCP is driven by a transmembrane Na⁺ gradient maintained by the Na⁺/K⁺-89 ATPase pump, which is also localized in the sinusoidal membrane (Bohan and 90 Boyer 2002). Ntcp/NTCP accounts for the transport of more than 80 % of amidated 91 bile salts (the major circulating bile salts) and only 40 % of their unconjugated, 92 parent compounds (Kouzuki et al. 1998). The remaining fraction of circulating bile 93 salts is taken up by a non-electrogenic, Na⁺-independent transport system, formed 94 by a family of transporters collectively named organic anion-transporting poly-95 peptides (Oatps/OATPs for rat and human, respectively) (Kullak-Ublick et al. 96 2000). Apart from bile salts, Oatps/OATPs accept a wide range of amphipathic, 97 organic compounds, including bilirubin, bilirubin glucuronides, leukotrienes, 98 estrogens, "type II" organic cations, and several exogenous organic anions, the 99 cholephilic dye sulfobromophthalein (BSP) being a prototypical example of the 100 latter one (Hagenbuch and Meier 2003). 101

After traversing the cell by Fick's diffusion bound to high-affinity cytosolic pro-102 teins, monoanionic bile salts (C24 amides conjugated with glycine or taurine) are 103 excreted in the canalicular pole by the *bile-salt export pump* (BSEP/Bsep; ABCB11/ 104 Abcb11), an ATP-binding cassette transporter (Suchy and Ananthanarayanan 2006). 105 In contrast, canalicular efflux of divalent, bipolar sulfated or glucuronidated bile salts 106 is mediated by the *multidrug resistance-associated protein 2* (MRP2/Mrp2; ABCC2/ 107 Abcc2). This carrier also transfers endogenous and exogenous non-bile-acid 108 organic anions conjugated with glutathione and glucuronic acid, including bilirubin 109 glucuronides and BSP both in its unconjugated and conjugated forms (Nies and 110 Keppler 2007). 111

The bile-salt-independent fraction of the bile flow depends on glutathione 112 excretion, mainly in its reduced form (~80 %) (Ballatori and Truong 1992). 113 Hepatocellular glutathione transport is poorly understood. The liver is the main 114 site of glutathione synthesis, through a pathway involving two consecutive steps, 115 catalyzed by the enzymes γ -glutamyl-cysteinyl synthetase (γ -GCS) and glutathione 116 synthetase (GS). The tripeptide is then exported into both blood and bile, and all 117 biliary glutathione comes from this intracellular source (Garcia-Ruiz et al. 1992). 118 However, a high-affinity, electrogenic carrier has been functionally characterized, 119 but not cloned as yet (Ballatori and Dutczak 1994), which exports actively GSH 120 into bile, and can transfer with low-affinity GSSG and GSH conjugates as well. 121 Another transporter likely involved in glutathione canalicular transport is Mrp2. 122 However, this carrier bears low affinity towards GSH, although it can transfer 123 124 GSSG and GSH conjugates with high affinity (Yang and Hill 2001).

Canalicular bile flow is further modified during its transit along bile ducts by 125 both secretory and absorptive processes (Bogert and LaRusso 2007). Ductular fluid 126 secretion is mainly driven by the secretin-regulated, cAMP-dependent output of 127 a HCO₃⁻-rich fluid secreted via the Cl⁻/HCO₃⁻-exchange system, anion 128 exchanger 2 (Ae2/AE2). Exchange is dependent on the out-to-in Cl⁻-concentration 129 gradient, which is maintained by the Cl^- efflux across the apical membrane via the 130 ATP-dependent, secretin-activated, cystic fibrosis transmembrane regulator 131 (CFTR). Blood-to-bile water movement at the ductular level is facilitated by 132 constitutive AQP4 in the basolateral membrane and secretin-stimulated AQP1 in 133 the apical membrane (Marinelli et al. 2011). On the other hand, absorption of 134 ductular water and electrolytes is driven by the osmotic gradients created by bile-135 to-plasma transport of electrolytes and organic solutes. They comprise (i) gluta-136 mate, transported by as yet unidentified carriers; (ii) glucose, transported by SGLT1 137 and GLUT1 at the apical and basolateral domains, respectively; and (*iii*) bile salts, 138 taken up by the apical Na⁺-dependent bile-salt transporter, ASBT/Asbt (SLC10A2/ 139 *slc10a2*), and extruded by both the basolateral export pump, MRP3/Mrp3 (ABCC3/ 140 Abcc3), and the heterodimeric organic solute transporter, OST a-OST B/Ost a-Ost B 141 (Marinelli and LaRusso 1996; Xia et al. 2006). 142

¹⁴³ Changes in Hepatobiliary Secretory Function Induced by OS

Compelling evidence in the literature indicates that oxidative challenge affects the
hepatocyte secretory machinery by impairing both bile flow (hepatocellular cholestasis) and the biliary excretion of both endo- and xenobiotics.

OS-induced impairment of bile flow generation has been demonstrated to occur 148 after exposure to a number of pro-oxidant agents, including tertsoon 149 butylhydroperoxide (tBOOH) (Ballatori and Truong 1989; Schmitt et al. 2000), 150 hydrogen peroxide (Akerboom et al. 1984; Ballatori and Truong 1989), menadione 151 (te Koppele et al. 1991), allyl alcohol (te Koppele et al. 1991), ethylhexanol 152 (te Koppele et al. 1991), chloro-dinitrobenzene (Schmitt et al. 2000), CCl₄ (Eipel 153 et al. 2007), ethacrynic acid (Ji et al. 2004), and lindane (Barros et al. 1988), among 154 others. Some pharmacological agents, such as cyclosporine A (Bramow et al. 155 2001), dapsone (Veggi et al. 2002, 2005), and nitrofuran derivatives (Hoener 156 1988), also induce cholestasis due, at least in part, to their pro-oxidant properties. 157 Finally, maneuvers leading to hepatic OS, such as hepatic (Accatino et al. 2003; 158 Bowers et al. 1987; Lee et al. 2000) and intestinal (Turnage et al. 1991) ischemia-159 reperfusion or aluminum intoxication (Gonzalez et al. 2004, 2007), also induce bile 160 flow impairment. 161

The classical view to interpret the cholestasis associated to pro-oxidant conditions is based upon the following pathomechanisms:

i. Reduction of the number of living parenchymal liver cells by necrosis and
 apoptosis depending on the severity of the oxidative injury (Czaja 2007).

- ii. Impairment of the bile-salt-dependent fraction of the bile flow due to compet-
- itive inhibition of bile-salt transport by the intracellular GSSG formed in excess

during the oxidative challenge (Akerboom et al. 1984; Ballatori and Truong 168 1989). Indeed, GSSG cis-inhibits the transport bile salts in liver canalicular 169 membrane vesicles (Griffiths et al. 1987), and mirror curves showing an inverse 170 relationship between GSSG and bile-salt biliary excretions have been obtained 171 when different pro-oxidizing compounds were administered in the isolated rat 172 perfused liver, such as hydrogen peroxide (Akerboom et al. 1984), menadione 173 (Akerboom et al. 1988), and tBOOH (Akerboom et al. 1984; Ballatori and 174 Truong 1989). 175

iii. Impairment of the bile-salt-independent bile flow due to a decrease in the biliary
excretion of total glutathione (GSH plus GSSG). This occurs due to depletion of
the hepatic levels of glutathione due to the sustained plasmatic and biliary
exportation from the cell as GSSG to maintain the GSH/GSSG ratio (Koeppel
et al. 1998).

The above-mentioned mechanisms may be predominant under strong oxidizing 181 conditions. However, under mild or even low oxidizing conditions, bile secretory 182 failure still occurs. For example, upon administration of different pro-oxidant 183 compounds to isolated perfused rat livers, drop in bile flow and/or decrease in 184 bile-salt secretion occurs before leakage of cytosolic hepatocellular enzymes or 185 increments in intracellular GSSG become apparent (Ballatori and Truong 1989; Ji 186 et al. 2004). Likewise, in isolated rat hepatocyte couplets, the apical secretion of 187 fluorescent bile-salt analogues was impaired by low concentrations of the pro-188 oxidant compounds tBOOH and 2,3-dimethoxy-1,4-naphthoquinone, even when 189 cell viability and intracellular GSSG levels remained unaffected (Pérez et al. 190 2006a). Overall, these results suggest that more subtle changes in the machinery 191 involved in bile formation occur under mild OS conditions. Among them, the actin-192 cytoskeletal disruption, which occurs even at very low OS levels, seems to be 193 a crucial causal factor, as discussed next. 194

195 Cytoskeletal Integrity and Hepatocanalicular Function

The actin cytoskeleton is a dynamic network of filamentous actin (F-actin), formed
by the reversible assembly of monomeric actin (G-actin), spatially distributed as
a belt around the bile canaliculus.

Actin cytoskeleton is one of the primary targets of ROS. The oxidative challenge promotes the oxidation of actin at a sulfhydryl group of a cysteine in position 374 (Dalle-Donne et al. 2001). This induces conspicuous changes in actin-spatial distribution, resulting in marked changes in the cellular topology (plasma membrane blebbing) (Dalle-Donne et al. 2001; Mirabelli et al. 1988).

Most of the above-mentioned roles of actin in cellular biology apply to hepatocytes, and many of them are involved in the biliary secretory processes. It is therefore not surprising that disorganization of the actin cytoskeleton induced by ROS has several deleterious effects on hepatobiliary function.

The interrelationship between ROS, Ca^{2+} elevations, actin-cytoskeletal integrity, and hepatocanalicular secretory function was exhaustively investigated in the

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1990s by our group, using the hepatocyte couplet model. These studies revealed 210 that, under mild OS conditions with preserved hepatocellular viability, 211 a close relationship exists between the disarrangement of the pericanalicular actin 212 cytoskeleton and the impairment in the capability of the couplets to accumulate 213 apically and retain in their canalicular vacuoles fluorescent bile-salt analogues 214 induced by the oxidizing compounds tBOOH (Ahmed-Choudhury et al. 1998; 215 Pérez et al. 2006a, b; Roma et al. 1997) and menadione (Stone et al. 1994, 1996). 216 These two independent tests indicated that both the apical secretion of bile salts and 217 their further tight-junctional-dependent retention in the bile canaliculus are impaired 218 early under OS conditions. Impairment of tight-junctional permeability was 219 also observed in isolated perfused rat livers exposed to tBOOH (Ballatori and 220 Truong 1989). 221

These functional alterations to secrete and retain bile salts in the biliary space 222 seem to have a structural correlate. tBOOH induces disorganization of the tight-223 junctional complex in hepatocyte couplets, as suggested by the redistribution of the 224 tight-junctional-associated protein, ZO-1 (Pérez et al. 2006a); F-actin is anchored to 225 zonula occludens-associated proteins thus regulating paracellular permeability 226 (Anderson and Van Itallie 1995). In addition, the canalicular bile-salt transporter 227 Bsep suffers endocytic internalization into intracellular vesicles in hepatocyte 228 couplets (Pérez et al. 2006b), which reduces dramatically the density of transporters 229 230 properly located at the membrane domain. A similar phenomenon has been described for the canalicular transporter Mrp2, which suffered endocytic internali-231 zation after exposure of isolated perfused rat livers to pro-oxidant insult, such as 232 exposure to tBOOH (Schmitt et al. 2000), chloro-dinitrobenzene (Schmitt et al. 233 2000), ethacrynic acid (Ji et al. 2004; Sekine et al. 2006), and lipopolysaccharide 234 (LPS) (Sekine et al. 2010), or after hepatic ischemia-reperfusion (Yu et al. 2007). 235 Mrp2 relocalization has a clear-cut functional correlate. Experiments in isolated 236 perfused rat livers indicated that a high, sustained exposure to ethacrynic acid has an 237 inhibitory effect on the excretion of both unchanged and conjugated forms of the 238 model cholephilic dye and Mrp2 substrate BSP (James and Ahokas 1992). Mrp2 239 relocalization under OS conditions is reversible in nature. When the OS induced by 240 tBOOH in isolated perfused rat livers was reverted by replenishment of GSH with 241 the cell-permeable form, GSH-ethyl ester internalized Mrp2 was relocalized back to 242 the canalicular membrane in a microtubule-dependent manner (Sekine et al. 2008). 243

The exact mechanisms that link OS-induced actin disorganization with tight-244 junctional impairment and transporter internalization are unknown, but previous 245 studies in the literature provide some clues. Hepatic tight-junctional permeability 246 increases following administration of the actin-disrupting agent phalloidin (Elias 247 et al. 1980). F-actin is anchored to tight-junctional-associated proteins (e.g., ZO-1), 248 and it is likely that F-actin disorganization induces relocalization of zonula 249 occludens intermediary proteins or even proteins forming the tight-junctional 250 strands, such as occludin and claudin. 251

Phalloidin-induced F-actin disorganization also induces internalization of cana-252 licular transporters, such as Mrp2 (Rost et al. 1999). The retrieval of canalicular 253 transporters under OS conditions (Ji et al. 2004; Pérez et al. 2006b; Schmitt et al. 254

2000; Sekine et al. 2006) is therefore also likely due to the simultaneous F-actin 255 disarrangement. The molecular bases to understand this causal relationship are just 256 emerging. Mice lacking radixin, which cross-links actin filaments and plasma 257 membrane proteins, develop conjugated hyperbilirubinemia associated to retrieval 258 of Mrp2 (Kocher et al. 1999), and the same holds true for obstructive and estrogen-259 induced cholestasis, where a disturbed colocalization of Mrp2 and radixin is 260 associated with Mrp2 endocytic internalization (Kojima et al. 2008). Interestingly, 261 the internalization of Mrp2 that occurs after hepatic ischemia-reperfusión is coin-262 cident with a virtual loss of radixin expression (Shu et al. 2007), and ROS-mediated 263 dephosphorylation and relocalization of radixin has been proposed to account for 264 LPS-induced Mrp2 internalization (Saeki et al. 2011). This latter phenomenon has 265 been associated with the OS induced by this cytokine (Sekine et al. 2010) and seems 266 to involve a decrease in the total amount of the active, phosphorylated form radixin 267 and its degree of interaction with Mrp2 (Saeki et al. 2011). 268

Mediation of Signal Transduction Pathways in OS-Induced Acute Hepatocanalicular Dysfunction

Cytosolic Ca2+ elevations occur under OS conditions, due to both the entry of 271 extracellular Ca²⁺ via plasma membrane receptor-operated Ca²⁺ channels and the 272 release of Ca²⁺ from intracellular Ca²⁺ storages, particularly in the endoplasmic-273 reticulum (calciosome) (Reed 1990). Ca²⁺ elevations are a major determinant of the 274 impairment in bile secretion following the oxidative injury. The intracellular Ca²⁺ 275 chelator BAPTA/AM fully prevents the impairment induced by low levels of tBOOH 276 in the capability of the hepatocyte couplets to accumulate and retain in their cana-277 licular vacuoles bile-salt-fluorescent analogues (Stone et al. 1994). Suggestively, the 278 associated actin-cytoskeletal disarrangement is also prevented by BAPTA/AM, 279 further supporting a causal relationship between both phenomena. Furthermore, 280 Ca^{2+} -elevating agents, such as the Ca^{2+} ionophore A23187 (Stone et al. 1994) or 281 the inhibitor of endoplasmic-reticulum Ca²⁺-ATPase thapsigargin (Ballatori and 282 Truong 1989), mimic the deleterious effects of ROS on both actin-cytoskeleton 283 integrity and hepatocanalicular function. 284

A number of signal pathways downstream of Ca²⁺ are involved in this phenom-285 enon. Activation of Ca²⁺-dependent, "classical" protein kinase C isoforms (cPKCs) 286 seems to be one of the most important ones. Our group demonstrated that the 287 pro-oxidant agent *t*BOOH induces cytosolic-Ca²⁺ elevations and translocation of 288 the cPKC isoform, PKCa, from the cytosol to the plasma membrane in isolated 289 hepatocytes, even at concentrations low enough to only affect the biliary secretory 290 machinery (Pérez et al. 2006a). Furthermore, several previous findings showed 291 strong similarities between the effect of ROS and those induced by Ca²⁺ and PKC 292 agonists, namely: (i) Both cytosolic Ca^{2+} elevations (Nathanson et al. 1992a) and 293 PKC activation (Corasanti et al. 1989) impair bile flow generation in the isolated 294 perfused rat liver (Corasanti et al. 1989), in part by increasing paracellular perme-295 ability (Kan and Coleman 1988; Llopis et al. 1991); we and others further 296

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characterized these effects in the hepatocyte couplet model and showed that cytosolic Ca²⁺ elevations impair the couplet capability to secrete and retain in their
canalicular vacuoles fluorescent bile-salt analogues by activating cPKC (Roma et al.
1999) and that PKC activation by vasopressin and phorbol esters reproduced these
effects (Nathanson et al. 1992b; Roma et al. 1997, 1998). (*ii*) PKC agonists induce
F-actin-cytoskeletal disarrangements (Roma et al. 1998), and Ca²⁺-elevating agents
reproduced these effects by a PKC-dependent mechanism (Roma et al. 1999).

Final confirmation of a crucial role for cPKC activation in actin disorganization and hepatocanalicular dysfunction induced by ROS was provided by recent studies in hepatocyte couplets. ROS-mediated actin-cytoskeleton disarrangements were fully prevented by both PKC-pan-specific and cPKC-specific inhibitors (Pérez et al. 2006a). More relevant from the therapeutic point of view, both cytoskeleton disruption and canalicular dysfunction were reversed within 1 h by these inhibitors (Pérez et al. 2006a).

The retrieval of the bile-salt transporter Bsep from the canalicular membrane 311 was also fully prevented by PKC antagonists (Pérez et al. 2006b). The same holds 312 true for the impairment of the tight-junctional-retentive properties, another possible 313 consequence of the actin disassembly induced by exposure to pro-oxidant 314 agents (Pérez et al. 2006a). The mechanisms that explain the harmful effect 315 of PKC on actin integrity and, by extension, on the hepatocanalicular function as 316 whole remain unclear. PKC phosphorylates and/or disorganizes several 317 а actin-cytoskeletal components, including actin itself, actin-associated proteins, 318 and membrane-cytoskeletal cross-linked proteins (Keenan and Kelleher 1998; 319 Larsson 2006). 320

The kind of canalicular protein that is internalized under oxidative-stress con-321 ditions and the signalling molecule involved seem to depend on the pro-oxidant 322 agent employed and on the magnitude of the oxidative damage. Low concentrations 323 of the oxidizing compound, ethacrynic acid, do not translocate cPKC but "novel" 324 PKC isoforms (nPKC). Under these conditions, Mrp2 but not Bsep is internalized, 325 by a mechanism probably involving Ca²⁺-dependent activation of inducible nitric 326 oxide (NO) synthase (iNOS), followed by NO-mediated cGMP increase and further 327 cGMP-activated nPKC (Sekine et al. 2006). However, higher ethacrynic acid doses, 328 capable of activating cPKC isoforms as well, induce internalization of both Bsep 329 and Mrp2 (Sekine et al. 2006). 330

As summarized in Fig. 140.2, a picture is emerging on the effect of acutely 331 induced OS on the hepatobiliary function. Under mild ROS challenge not affecting 332 hepatocellular viability, Ca²⁺ elevations induce cPKC and/or nPKC activation. This 333 brings on a number of alterations in both function and localization of structures 334 relevant to bile formation, such as actin cytoskeleton, canalicular transporters, and 335 tight-junctional components. This impairs, in turn, the biliary secretion and the 336 further retention of solutes that provide osmotic driving force for bile formation. 337 Other factors such as GSSG-induced *cis*-inhibition of bile-salt transport, reduced 338 biliary excretion of glutathione due to intracellular glutathione depletion, and 339 hepatocellular death may become contributing factors, depending on the magnitude 340 341 of the pro-oxidant condition.

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Fig. 140.2 Effect of the model oxidizing compound *tert*-buthylhydroperoxyde (*t*BOOH) on the canalicular transport of bile salts via Bsep and of oxidized glutathione (GSSG) via Mrp2. In normal cells, the pericanalicular localization of F-actin allows for the normal localization of the canalicular transporters in their membrane domain and the proper barrier function of the tight-junctional structures. The acute exposure to *t*BOOH induces mobilization of Ca^{2+} across the plasma and from the calciosome membranes and the further activation of Ca^{2+} -dependent, "classical" PKC isoforms (cPKC). This activation leads to relocalization of F-actin to the cell body, which in turn induces blebbing, Bsep/Mrp2 internalization, and tight-junctional disorganization; these two latter events explain the impairment of bile-salt (BS) and GSSG biliary excretion and their further canalicular retention. Cytosolic Ca^{2+} elevations may also activate inducible nitric oxide (NO) synthase (iNOS), which leads to NO-mediated activation of guanylate cyclase (GC) and further cyclic guanosine monophosphate (cGMP)-mediated activation of "novel" PKC isoforms (nPKC); nPKC activation internalizes selectively Mrp2

342 The Antioxidant Adaptive Hepatic Response and Bile Secretion

Hepatocytes develop an adaptive response against ROS when the oxidative insult is 344 sustained (Fig. 140.3). This response involves induction of antioxidant enzymes 345 346 such as catalase (Sen et al. 2005) and manganese superoxide dismutase (Kwak et al. 2001), as well as increments of glutathione synthesis via induction of γ -glutamyl-347 cysteinyl synthetase (Yamane et al. 1998). In addition, hepatic adaptation integrates 348 the distinctive metabolizing and secretory capacity of the organ to reinforce these 349 antioxidant mechanisms. In this context, it is crucial the OS-mediated induction of 350 the phase-II-detoxifying enzymes glutathione-S-transferase (GST) (Kohle and 351 Bock 2007) and UDP-glucuronosyltransferase (UGT) (Kwak et al. 2001). 352

GST induction enhances the coordinated inactivation, via GSH conjugation, of DNA hydroperoxides and lipid hydroperoxides formed as secondary metabolites during OS (Ketterer and Meyer 1989). GST also catalyzes GSH conjugation of highly reactive, toxic α,β-unsaturated lipid aldehydes, 4-hydroxy *trans*-2-nonenal



Fig. 140.3 Main hepatocellular adaptive changes induced by sustained OS in the expression of enzymes, and transporters involved in: (*i*) glutathione (GSH) synthesis via γ -glutamylcysteine synthetase (γ -GCS) and GSH synthetase (GS), (*ii*) GSH conjugation of lipid peroxides or their aldehydic derivatives (LP) via GSH *S*-transferase (GST), and (*iii*) plasmatic and biliary extrusion of conjugated LP and oxidized glutathione (GSSG) via both Mrp1 and Mrp2, respectively. This adaptive response in mainly governed by the redox-sensitive transcription factor Nrf2, which escapes from its cytosolic repressor Keap1 and translocates to the nucleus under OS conditions. Once in there, it binds to the *antioxidant response element* (ARE) and activates ARE-dependent gene transcription, including that of γ -GCS, GST and Mrp2; Mrp1 activation is instead independent on Nrf2. The effect of Nrf2 on γ -GCS is reinforced by the activation of the transcription factor AP-1, via a proximal AP-1 element (PAP-1E). For further details, see section "The Antioxidant Adaptive Hepatic Response and Bile Secretion"

(HNE) being the most abundant (Renes et al. 2000). These lipid peroxides combine
spontaneously with cysteine, histidine, and lysine residues of proteins, which
modifies protein function and leads eventually to cellular toxicity.

UDP-glucuronosyltransferase induction improves glucuronidation of pro-360 oxidant toxicants, such as benzo(a)pyrene (Byczkowski and Gessner 1987), penta-361 chlorophenol (Umemura et al. 2006), acetaminophen (Clement and Williams 2005), 362 aliphatic alcohols (Ebner and Burchell 1993), and manadione (Liu et al. 1993). 363 These phase-II products are then extruded from the cell via the hepatocellular efflux 364 pumps MRP1, MRP2, MRP3, and MRP4 (ABCC4) and breast cancer resistance 365 protein (BCRP, ABCG2), all of which are also upregulated by ROS (Adachi et al. 366 2007; Aleksunes et al. 2008; Vollrath et al. 2006). Glutathione conjugates are 367 substrates of MRP2 and MRP1 (Geier et al. 2007). Since these transporters also 368 transfer GSSG, MRP1/2-mediated GSSG extrusion helps to maintain low 369

intracellular GSSG levels, when GSSG reduction back to GSH via GSSG reductase 370 becomes rate limiting. Unlike MRP1, the basolateral carriers MRP3, MRP4, and 371 BCRP transport glucuro- and sulfoconjugates and bile salts (Geier et al. 2007). 372 MRP1, MRP3, and MRP4 are normally expressed at very low levels in 373 the basolateral membrane of the hepatocytes. Upregulation of basolateral extrusion 374 pumps during a sustained oxidant insult is expected to shift the transfer of substrates 375 normally excreted into bile towards blood, to permit urinary excretion. As 376 an untoward effect of this adaptive response, this phenomenon might decrease the 377 biliary excretion of bile salts, which would contribute to the cholestatic 378 phenomenon. 379

All these adaptive mechanisms are transcriptional in nature and involve the activation of a number of redox-sensitive transcription factors, such as Nrf2, NF- κ B, and AP-1. The transcription factor activated depends on the magnitude of the oxidant insult. Low OS induces Nrf2, whereas higher levels trigger an inflammatory response through the activation of NF- κ B and AP-1 (Halliwell and Gutteridge 1999).

Nrf2 is a key transcription factor of the hepatic adaptation to sustained OS. Its 386 induction has been linked to different oxidant agents such as the cancer 387 chemoprotective agent 3H-1,2-dimethiole-3-thione (Kwak et al. 2001), alcohol 388 (Gong and Cederbaum 2006), tert-butylhydroquinone (Adachi et al. 2007), acet-389 aminophen (Aleksunes et al. 2008), and bile salts (Tan et al. 2007). The action of 390 Nrf2 depends on its accumulation in the nucleus, where it interacts with the 391 antioxidant response element (ARE) (Nguyen et al. 2003). This is a cis-acting 392 enhancer sequence that contains the 5'-TGAC-3' tetranucleotide present in the 393 genes of enzymes associated with glutathione biosynthesis, redox proteins with 394 active sulfhydryl moieties, drug-metabolizing enzymes, and transporters. Nrf2 395 induces the transcription of γ -glutamylcysteine synthetase, the rate-limiting enzyme 396 responsible for glutathione synthesis (Kwak et al. 2001). The effect of Nrf2 on this 397 enzyme is reinforced by AP-1. A proximal AP-1 element (-263 to -269) has been 398 identified to be critical in mediating the effect of OS-induced increase in the 399 transcription of the human catalytic subunit of this enzyme (Rahman et al. 1996). 400 Nrf2 also induces GST and UDP-glucuronosyltransferase (Kohle and Bock 2007; 401 Kwak et al. 2001; Yueh and Tukey 2007), as well as the hepatocellular transporters 402 MRP2 (Vollrath et al. 2006), MRP3 (Aleksunes et al. 2008), MRP4 (Aleksunes et al. 403 2008), and BCRP (Adachi et al. 2007). Nrf2-induced coordinated GST and Mrp2 404 expression increases the biliary excretion of conjugated BSP and possibly other 405 glutathione-conjugated compounds, such as DNA and lipid hydroperoxides 406 (Reisman et al. 2009). Furthermore, Nrf2 constitutes a defense system against 407 oxidative stress generated in the liver by experimental models of both extrahepatic 408 (Okada et al. 2009) and intrahepatic (Tanaka et al. 2009) cholestasis. Finally, Nrf2 is 409 required for the upregulation of basolateral bile-salt efflux pumps that counteract the 410 deleterious effects of hepatocellular build up of bile salts in cholestasis, as part of the 411 adaptive response against this condition (Tanaka et al. 2009). 412

418 Impairment of Hepatobiliary Function Induced by Sustained OS

The adaptive, spontaneous mechanisms that take place in hepatocytes to minimize the deleterious effects of ROS are, however, not always sufficient to prevent hepatocellular oxidative damage. When ROS production is maintained with time at high levels, alterations occurs in the capability of the hepatocyte to produce bile and to secrete cholephilic compounds, mainly because of changes in the expression of transporters.

The Long-Evans Cinnamon rat, an animal model of Wilson's disease, is 421 a prototypical model of high, chronic hepatic OS. These rats have a genetic defect 422 in Atp7b gene, which is homologous to the human Wilson's disease gene, resulting 423 in inability to mobilize copper from the liver (Harada et al. 2000). Apart from 424 copper, these rats also have high hepatic iron levels (Kato et al. 1993). Chronic 425 copper/iron accumulation increases lipid peroxidation by 50 %, presumably due to 426 the capability of these metals to induce OS via both mitochondrial dysfunction 427 (Sternlieb et al. 1995) and Fenton-type, copper/iron-catalyzed Haber-Weiss reac-428 tion (Yamamoto et al. 2001). 429

These mutant rats have histological features of cholestasis (Du et al. 2004) and 430 exhibit a number of alterations in hepatic transporter expressions. They have 431 a reduced basal bile-salt biliary excretion due to a posttranscriptional impairment 432 in Bsep expression (Chiba et al. 2007; Levy et al. 2007). On the other hand, mRNA 433 levels of the bile-salt uptake systems Ntcp and Oatp (isoforms Oatp1a1 and 434 Oatp1a4) are decreased, although this has not been confirmed at the protein level 435 (Chiba et al. 2007). Apart from alterations in bile-salt hepatic handling, Long-Evans 436 Cinnamon rats have both hyperbilirubinemia (Du et al. 2004; Yamamoto et al. 2001) 437 and impairment in the excretion of the Mrp2 substrate BSP (Itagaki et al. 2004). If 438 these alterations involve changes in the expression of Mrp2 at a protein level is 439 unknown, but Mrp2 mRNA levels are normal (Chiba et al. 2007). Alternatively, the 440 above-mentioned transcriptional downregulation of Oatps, which also transport 441 non-bile-salt organic anions such as BSP and bilirubin, may be a contributing factor. 442

Au6

Another experimental model of metal-induced chronic OS is that afforded by 443 long-term aluminum (Al³⁺) exposure to rats (Gonzalez et al. 2007). When admin-444 istered intravenously for 1-2 weeks, Al³⁺ reduces bile flow, and this impairment 445 correlates directly with Al³⁺ hepatic content; this was associated with elevations of 446 serum bile salts, suggesting impaired hepatic handling of bile salts (Klein et al. 447 1988). An even more chronic exposure to Al^{3+} (3 months), which doubles the lipid-448 peroxidation levels, also reduced bile flow and the biliary output of bile salts 449 (Gonzalez et al. 2004). Compartmental analysis of the plasma decay of BSP 450 revealed that both sinusoidal uptake and canalicular excretion of the dye are 451 decreased, the latter phenomenon being associated with a decrease in Mrp2 protein 452 expression (Gonzalez et al. 2004). All these alterations were prevented by admin-453 istration of the antioxidant vitamin E, suggesting that OS was the main, if not the 454 only, mechanism (Gonzalez et al. 2007). 455



Fig. 140.4 Changes in the expression of transporters relevant to bile flow generation in hepatic injury induced by ischemia-reperfusion. An ischemic period of 60 min, followed by 1 day of reperfusion, decreases the mRNA levels of the basolateral transporters Ntcp and OATP, as well as the canalicular export pumps Mrp2 and Bsep (only confirmed at the protein level for Mrp2). It is unknown whether ischemia-reperfusion exerts these transcriptional effects via ROS-induced changes in degradation/function of RNA polymerase II (RNA Pol II) or, indirectly, by promoting the release of proinflammatory cytokines, which may both downregulate transcription factors (TF) that function as transactivators of hepatocellular transporters and reduce their DNA binding activity. Since downregulation of Mrp2 protein is more severe than that of its mRNA, an additional posttranscriptional mechanism is proposed, which involves OS-induced transporter internalization, followed by lysosomal degradation. For further details, see section "Impairment of Hepatobiliary Function Induced by Sustained OS"

Hepatic ischemia-reperfusion injury is another prototypical OS-mediated
hepatopathy associated with both cholestasis (Lee et al. 2000; Lemasters and
Thurman 1997) and changes in expression of hepatocellular transporters (Tanaka
et al. 2006, 2008) (Fig. 140.4).

The impact of this maneuver on both bile flow generation and the expression of 460 transporters depends on the duration of the ischemia. A 30-min ischemia decreases 461 both bile flow and biliary bile-salt output after 1 day of reperfusion, but no changes 462 occur in mRNA and protein levels of the main basolateral and canalicular bile-salt 463 transporters, Ntcp and Bsep, respectively (Accatino et al. 2003); in this case, 464 changes in localization/intrinsic activity of these transporters or impairment in the 465 466 expression of other bile-salt transporters, such as Oatps, may explain bile-saltsecretory failure. Similarly, Mrp2, mRNA, and protein expressions are unaffected, 467 in agreement with the absence of changes in the maximum-secretory rate of 468 469 the Mrp2 substrate ceftriaxone (Accatino et al. 2003). Unlike a 30-min ischemia, a 60-min ischemic period followed by 1 day of reperfusion decreases the mRNA
levels of the basolateral transporters Ntcp and Oatp (all isoforms), as well as those
of the canalicular export transporters, Mrp2 and Bsep (Tanaka et al. 2006); this has
been confirmed at the protein level only for Mrp2 (Tanaka et al. 2008).

The mechanisms by which these transcriptional alterations occur are far from 474 being understood. First, we have to bear in mind that high levels of OS globally 475 inhibit gene transcription by inducing RNA polymerase II degradative 476 ubiquitination and decrease in histone H3 and H4 acetylation; histone acetylation 477 dissociates DNA from the histone complex, allowing transcription to proceed 478 (Berthiaume et al. 2006). However, at least part of these transcriptional alterations 479 may be due to the ROS-dependent hepatic inflammatory response, which leads to 480 release of cholestatic, proinflammatory cytokines (e.g. TNF- α and IL-1 β) with 481 capability to transcriptionally impair transporter expression (Geier et al. 2007). 482 This phenomenon involves downregulation of the ubiquitous heterodimerization 483 partner retinoid X receptor (RXR α), leading to impairing of the binding activity 484 of nuclear receptor heterodimers that requires $RXR\alpha$ for their transcriptional 485 activity or, in particular, for Ntcp and Mrp2, respectively, and to reduction of the 486 nuclear levels of the monomeric transcription factors hepatocyte nuclear factor- 1α 487 (HNF-1a) (Geier et al. 2007) and interferon regulatory factor 3 (IRF3) (Hisaeda 488 et al. 2004). 489

490 In ischemia-reperfusion injury, downregulation of Mrp2 protein is more profound than that of mRNA, suggesting additional posttranscriptional mechanisms 491 (Tanaka et al. 2008). Although the causes underlying the latter phenomenon are 492 presently unknown, the early OS-induced transporter internalization, sustained with 493 time, may lead to delivery of the endocytosed transporters to the lysosomal com-494 partment, followed by degradation, as was suggested to occur late in LPS-induced 495 cholestasis (Kubitz et al. 1999) and in obstructive cholestasis (Paulusma et al. 2000) 496 in rats, two cholestatic models exhibiting OS. Apart from alterations in transporter 497 expression/function, the tight-junctional barrier is impaired in ischemia-reperfusion 498 injury, as shown in 24- or 48-h-cold-stored isolated perfused rat livers subjected to 499 reperfusion (Almada et al. 2003). 500

Taken together, these models of long-lasting OS show consistently that cholestasis and/or impairment of the constitutive expression of transporters relevant to bile formation is a common feature in prolonged OS and that both transcriptional and posttranscriptional mechanisms are involved. Proinflammatory cytokines released by the inflammatory response to the oxidative liver damage may be key mediators.

508 Future Directions

Cholestasis is a common feature under OS conditions, even at OS levels far lower than those affecting hepatocellular viability. Despite considerable progresses have been made in the characterization of the effects of ROS on the biliary secretory machinery, the characterization of the molecular mechanisms underlying these

effects is in its infancy. A bridge needs to be built between early events and late 513 consequences of OS on bile secretion, in order to reconstruct the cascade of events 514 leading to the posttranscriptional changes in transporter protein expression 515 observed eventually during the sustained oxidative challenge. Also, we need to 516 distinguish direct ROS-mediated effects from secondary consequences of the oxi-517 dative injury (e.g., inflammatory response, accumulation of biliary solutes). In 518 addition, we must fully characterize the redox-sensitive signalling pathways 519 involved in these effects; a more complete picture should provide more selective 520 therapeutic strategies to interfere with ROS-mediated harmful pathways or to 521 enhance the protective ones. Studies on the impact of OS on transport function of 522 biliary epithelial cells are also eagerly awaited; cholangiocytes may contribute to 523 bile secretory failure, as they are the main target of cholangiopathies associated with 524 periductal inflammation and OS. And finally, it remains to be ascertained the actual 525 contribution of OS in both transcriptional and posttranscriptional changes in liver 526 transporter expression occurring in chronic cholestatic liver diseases in humans 527 (Geier et al. 2007). 528

Genomic and proteomic tools are accelerating the discovery of new ROS-529 responsive genes and the molecular targets of ROS action. These approaches are 530 expected to greatly help to achieve the goals above. Meanwhile, we hope this 531 preliminary information contributes to draw attention about the convenience of 532 limiting OS in hepatopathies with cholestatic features. Some short-scale clinical 533 studies using co-adjuvant, antioxidant therapies have shown encouraging results 534 (Vendemiale et al. 2002), but multicentric and long-term clinical trials are needed 535 to determine whether this strategy holds promise for the future. 536

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