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1 **The ATP-binding cassette proteins ABCC1 and ABCB1 as modulators of**
2 **glucocorticoid action**

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9
10 **Abstract**

11 Responses to hormones that act through nuclear receptors are controlled by modulating hormone
12 concentrations not only in the circulation but also within target tissues. The role of enzymes that
13 amplify or reduce local hormone concentrations has become well-established for glucocorticoid and
14 other lipophilic hormones; moreover, transmembrane transporters have proven critical in
15 determining tissue responses to thyroid hormones. However, there has been less consideration of the
16 role of transmembrane transport for steroid hormones. ATP-binding cassette (ABC) proteins were first
17 shown to influence the accumulation of glucocorticoids in cells almost three decades ago, but
18 observations over the past ten years suggest that differential transport propensities of both
19 exogenous and endogenous glucocorticoids by ABCB1 and ABCC1 transporters provides a mechanism
20 whereby different tissues are preferentially sensitive to different steroids. This Review summarises
21 this evidence and the new insights provided for the physiology and pharmacology of glucocorticoid
22 action, including new approaches to glucocorticoid replacement.

23

24

25 **Introduction**

26 Glucocorticoid hormones are vital for life; they confer diverse effects on multiple processes
27 and systems. The adverse consequences of glucocorticoid excess are well demonstrated by the frank
28 hypercortisolism of Cushing syndrome, but even subtle cortisol dysregulation has implications,
29 contributing to cardiovascular disease, for example.¹ Over the past 30 years it has become clear that
30 the concentration of glucocorticoid in the blood does not necessarily reflect that within tissues, as
31 enzymes (such as 11 β -hydroxysteroid dehydrogenase, which catalyses the interconversion of inert
32 cortisone and active cortisol) and delivery mechanisms of corticosteroid binding protein can confer
33 additional control over the absolute tissue levels.^{2,3}

34 As lipophilic molecules, glucocorticoids can diffuse across cell membranes to interact with
35 intracellular targets; however, they can also undergo active transmembrane transport. This process
36 was first described for the ABCB1 transporter (of the ATP-Binding Cassette [ABC] protein family),
37 which exports cortisol and a variety of synthetic glucocorticoids from 'sanctuary sites' including the
38 brain.^{4,5} Intriguingly, corticosterone is not readily exported by ABCB1, but we have discovered that the
39 ABCC1 transporter, found in tissues including adipose, exports corticosterone but not cortisol.⁶

40 In this Review, we will explore the implications of this tissue-specific glucocorticoid transport
41 in the central control of the hypothalamic–pituitary–adrenal (HPA) axis, adipose tissue metabolism
42 and pregnancy. We will also consider whether the steroid specificity of ABCB1 and ABCC1 transport
43 offers insights into the different roles of corticosterone and cortisol in humans and a potential
44 opportunity for developing glucocorticoid therapies that are better targeted than those currently
45 available to maximise efficacy and minimise toxicity.

46

47 **[H1] The movement of lipophilic hormones**

48 The 'free hormone hypothesis' determines that unbound lipophilic hormones move passively
49 down a concentration gradient⁷ and, indeed, steroids are taken up freely by cell types such as
50 keratinocytes without the relevant membrane transporters.⁸ Differences in tissue concentrations
51 were previously attributed to differences in physicochemical properties, such as lipophilicity, until the
52 discovery of the existence of specific thyroid hormone transporters challenged these traditional
53 assumptions. In the case of triiodothyronine (T3), which is highly lipophilic owing to the iodinated
54 aromatic ring, the level of hormone available to receptors not only depends on hormone synthesis
55 and peripheral enzymatic conversion, but also on transport into and out of cells, notably by the
56 monocarboxylate 8 (MCT8) transporter.⁹ The uptake of T3 into neurons is critically impaired in the
57 absence of MCT8, as occurs in the X-linked Allan-Herndon-Dudley syndrome of neurodevelopmental
58 anomalies associated with abnormal thyroid function.¹⁰

59 The cellular uptake of glucocorticoids by membrane transporters has been demonstrated in
60 *Drosophila melanogaster*, in which loss of the Ecdysone Importer (Eci) membrane transporter
61 produces a phenotype that is identical to that resulting from the loss of ecdysone or the ecdysone
62 receptor.¹¹ Organic anion transporting polypeptide transporters mediate the uptake of glucocorticoids
63 in rat liver *ex vivo*; however, this uptake has not been reproduced in humans.^{12,13} Furthermore, a
64 saturable glucocorticoid uptake mechanism across the blood–brain barrier (BBB) and blood–
65 cerebrospinal fluid barrier that was reported in mice was only discernible at supraphysiological
66 concentrations, and so might not be physiologically relevant.¹⁴

67 Our increasing understanding of the importance of transporters for thyroid hormone function
68 sets a biological precedent for a similar scenario for other lipophilic hormones; however, although the
69 active cellular import of glucocorticoids in humans has not been shown, there is mounting evidence
70 supporting the facilitated export of glucocorticoids from cells, particularly by two members of the ABC
71 transporter family.

72

73 [H1] The ABC protein family

74 As members of one of the most highly conserved protein superfamilies, ABC proteins shuttle
75 toxins, xenobiotics and signalling molecules across eukaryotic and prokaryotic cell membranes. These
76 proteins are classified into seven subfamilies according to their structural similarity and sequence
77 homology, and have been actively researched for decades, particularly in relation to multidrug
78 resistance. The evolution and relevance of this transporter superfamily in the context of cancer drug
79 efflux has been well reviewed;^{15,16} however, of the over 50 human ABC proteins that have been
80 identified, only ABCB1 and ABCC1 have recognised roles in glucocorticoid transport.¹⁷

81 The typical ABC transporter is a homodimer characterized by two transmembrane domains
82 (TMDs) and two cytoplasmic nucleotide-binding domains (NBDs) (FIG. 1).¹⁸ Each TMD domain contains
83 between six and ten transmembrane α -helices, depending on the specific transporter, and is involved
84 in substrate recognition. The cytoplasmic NBDs contain conserved motifs for ATP binding and
85 hydrolysis, including the ABC signature motif (or C-loop motif), Walker A motif (P-loop) and Walker B
86 motif.¹⁷ Together, these dimeric NBDs act to hydrolyse ATP and provide energy to drive transport
87 against concentration gradients.

88 Several models have been proposed to explain the relationship between ATP hydrolysis and
89 TMD-mediated transport,¹⁹ with most purporting that energy from ATP hydrolysis enables the TMDs
90 to switch between inward- and outward-facing configurations (FIG. 1A). Individual ABC transporters
91 are unidirectional: in eukaryotic cells, they are almost exclusively exporters, but both importers (of
92 nutrients) and exporters (of toxins and cell wall substrates) exist in bacteria.²⁰ Consistent with this
93 export function in eukaryotes, ABC transporters are typically found on the apical cell membrane at
94 **luminal surfaces** to limit xenobiotic exposure.¹⁷ Substrates range from ions to large proteins and there
95 is a high degree of overlap between transporters, although the molecular basis for this overlap
96 remains poorly documented.

97

[H1] ABCB1 and ABCC1 are steroid exporters**[H2] ABCB1 and steroid export**

Initially named P-glycoprotein (P-gp) and later multidrug resistance protein 1 (MDR1), ABCB1 has been extensively studied as the archetypal multidrug transporter, exporting a broad array of xenobiotics including antineoplastics, antimicrobials and antidepressants from cells (reviewed in ^{15,21,22}). In humans, the *ABCB1* gene, located on chromosome 7q21.12, encodes a protein of 1280 amino acids (141.5 kDa) in size with 12 membrane-spanning α -helices distributed among two TMDs.²³ The polyspecificity of ABC transporters is often purported to result from the plasticity of the drug-binding pocket, both in terms of side chain and backbone arrangements. Numerous attempts have been made over the years to determine the 3D structure of ABC proteins in an effort to understand their transport mechanisms and their substrate specificity; however, their size and hydrophobicity pose significant challenges.²⁴ Advances in the use of cryo-electron microscopy have enabled structural insights into substrate binding.²⁵⁻²⁷ Reconstitution of the structure of human ABCB1 in complex with chemotherapeutic drugs has revealed the drug-binding cavity to be globular in shape, with interactions contributed by all 12 membrane-spanning α -helices (FIG. 1B).²⁵ Substrate-induced structural changes in NBD2 are thought to confer changes in ATPase activity, which determines transport action.

A putative steroid-binding site has been identified in human ABCB1, but this is based upon a homology model of only the NBDs²⁸ and is not definitive. However, physiological data do support selective ABCB1-mediated transport of steroids. In the 1960s, murine fibroblasts were observed exporting steroids in an energy- and temperature-dependent manner, consistent with active transport.²⁹ Cortisol export was later (in 1992) demonstrated in a porcine renal tubular cell line (LLC-PK1) overexpressing human *ABCB1*.³⁰ Since then, several endogenous and synthetic steroids have been confirmed as ABCB1 substrates. Depending on the presence of hydroxyl groups at positions 11 and 17, steroids were stratified into three categories.³¹ ABCB1-mediated efflux was highest for

124 steroids with both hydroxyl groups (including dexamethasone, cortisol and prednisolone), lowest for
125 those with neither (deoxycorticosterone and progesterone), and intermediate in those with one
126 hydroxyl group (including corticosterone and aldosterone). A-ring planarity and 6 α - and 16 α - methyl
127 substitution were reported to enhance transport when compared to passive diffusion in the LLC-PK1
128 line, in keeping with the presence of a critical hydrophobic pocket in the steroid-binding region.³²
129 Methylprednisolone is the glucocorticoid most effectively exported by ABCB1, followed by
130 prednisolone, betamethasone, prednisone, dexamethasone, cortisol and cortisone.³¹⁻³³ Aldosterone
131 appears to be weakly transported, and there is no evidence that sex steroids or 11-
132 deoxycorticosterone undergo ABCB1-mediated export,³¹ although progesterone does bind avidly to
133 ABCB1 with an inhibitory effect.³⁴ Corticosterone — the predominant glucocorticoid in rats and mice
134 — was initially shown to be an ABCB1 substrate on the basis of efflux from murine macrophage-like
135 cells,³⁵ and subsequent in vitro work in murine adrenocortical cells has demonstrated that
136 pharmacological ABCB1 inhibition blocked the ability of these cells to secrete corticosterone.³⁶
137 However, this is in contrast to previous in vitro work showing that corticosterone was not exported in
138 the murine LMCAT fibroblast line.^{31,37-39} Studies of the human transporter have not shown
139 corticosterone to be transported by ABCB1, so affinity might be species specific.^{4,40} Importantly,
140 studies in murine thymoma cells overexpressing *Abcb1* in which corticosterone and cortisol transport
141 was compared showed a lower efflux of corticosterone compared with cortisol,³¹ indicating an overall
142 preference of this transporter for cortisol.

143

144 [H2] ABCC1 and steroid export

145 First identified and cloned as multidrug resistance-associated protein 1 (MRP1), ABCC1 was
146 also discovered in multidrug resistance studies where high levels of expression are poor prognostic
147 indicators in certain malignancies.⁴¹⁻⁴³ Since then, ABCC1 has been shown to efflux a diverse range of
148 conjugated xenobiotics and physiological organic anions.⁴⁴ Like ABCB1, ABCC1 demonstrates a

149 polarized distribution in epithelial cells, but is located on the basolateral rather than apical
150 membrane.⁴⁵

151 ABCC1 is encoded by the human *ABCC1* gene on the short arm of chromosome 16 (16p13.11).
152 Strikingly, ABCC1 and ABCB1 share only 23% sequence identity, and differ substantially in their
153 structural and physiological functions. To date, the structure of only bovine ABCC1 has been
154 determined by cryo-electron microscopy.⁴⁵ The 190 kDa ABCC1 protein has 17 transmembrane α -
155 helices distributed among three TMDs (TMD0, TMD1 and TMD2) rather than the two TMDs observed
156 in ABCB1 (FIG 1C).⁴⁵

157 The binding site between TMDs 1 and 2 is 'bipartite': it has a positively charged 'P pocket',
158 which forms hydrogen bonds with glutathione residues, and a second 'H pocket', which interacts with
159 hydrophobic moieties. This bipartite binding domain explains why glutathione coupling facilitates the
160 transport of a wide range of compounds.⁴⁵

161 ABCC1 substrates tend to be organic anions, whereas those for ABCB1 tend to be weak
162 cations;⁴⁵ and ABCC1 uniquely exhibits affinity for phase II hepatic metabolites (endogenous and
163 xenobiotic compounds conjugated with glutathione, glucuronide and sulphate to facilitate excretion).
164 There are differences in substrate preference between human and other mammalian isoforms — for
165 example, the glucuronide conjugate of 17 β -oestradiol is a substrate only in humans.⁴⁶ It has also been
166 shown in vitro, both in virally transfected mouse fibroblast LMCAT cells and subsequently in human
167 adipocytes, that ABCC1 can export corticosterone and 11-deoxycorticosterone, but not cortisol,
168 prednisolone or dexamethasone.^{6,39}

169 Whilst ABCB1 is thought to transport substrates partitioning through the bilipid cell
170 membrane (the 'hydrophobic vacuum'),⁴⁷ ABCC1 is only open to substrates within the cytoplasm.⁴⁵

171

172 **[H1] ABCB1 and ABCC1 expression in tissues**

173 The mRNA expression profiles of human ABCB1 and ABCC1 in various tissues are summarised
174 in FIG. 2. ABCB1 is highly expressed (both at mRNA and protein level) in the adrenal gland, but also
175 found at absorptive surfaces (for example, of the intestines), protective barriers (for example, testis,
176 BBB and placenta) and in secretory tissues (for example, biliary canaliculi and renal tubule).²³ ABCC1
177 is widely expressed in almost all cell types, with highest levels in the thymus, parathyroid glands and
178 skeletal muscle. It seems to be poorly expressed in the liver⁴⁸ and nervous system but, notably, is
179 found in greater quantities than ABCB1 in adipose tissue and skeletal muscle.^{23,49,50}

180 A model for the consequences of this tissue-specific transporter expression on the
181 intracellular concentrations of different glucocorticoids is outlined in FIG. 3. Combining in vitro studies
182 from three different laboratories, glucocorticoids can be separated into three groups depending on
183 their relative propensity to be exported by ABCB1 and ABCC1.^{6,31,39} According to this model, the
184 intracellular concentrations of cortisol will be lower in tissues that predominantly express ABCB1
185 (including the central HPA axis negative feedback sites behind the blood-brain barrier), and those of
186 corticosterone will be lower in tissues that predominantly express ABCC1, such as adipose tissue. We
187 can use experimental data from animal and human studies to show how this may modulate the
188 physiology of the HPA axis, influence lipogenesis within adipocytes and alter glucocorticoid transfer
189 across the placenta.

190

191 **[H2] ABCB1 and ABCC1 and the HPA axis**

192 **[H3] *Insights from murine models.***

193 Central control of the HPA axis depends on feedback from circulating glucocorticoids to the
194 hypothalamus and pituitary, but to reach the brain the glucocorticoids must traverse the tightly
195 packed endothelium of the BBB, where ABCB1 is found.⁵¹ Murine models have been used extensively
196 to assess ABCB1-dependent modulation of steroid concentrations within tissues, including the brain.

197 Importantly, rodents have two ABCB1 isoforms: ABCB1A (also known as MDR1A or MDR3) and ABCB1B
198 (also known as MDR1B or MDR1),^{52,53} which broadly share the characteristics of the human protein.⁵³
199 Indeed, *Abcb1a*-knockout mice accumulate 87 times more of the ABCB1 substrate ivermectin in brain
200 than do wild-type animals,⁵⁴ while ABCB1 inhibition with tariquidar increases cerebral retention of
201 labelled verapamil on PET imaging and demonstrates the role of ABCB1 at the human BBB.⁵⁵

202 *Abcb1a*-knockout mice exhibit enhanced retention of cortisol and dexamethasone in the
203 brain.^{4,5,54,56} As seen in vitro, results for corticosterone export in vivo are varied, perhaps reflecting
204 redundancy between the murine isoforms. One study reported no difference in the levels of infused
205 radiolabelled corticosterone in the brains of adrenalectomised *Abcb1a*-knockout compared to wild-
206 type mice.⁴ However, the *Abcb1ab*-double knockout mouse retained an excess of cortisol and
207 corticosterone in the brain;⁵⁷ this retention was greater for cortisol than for corticosterone, suggesting
208 that, overall, ABCB1 activity in mice favours cortisol over corticosterone transport, as was also found
209 in vitro. However, another group reported the opposite effect: a retention of both glucocorticoids in
210 *Abcb1a*-knockout mice, and cortisol retention alone in the *Abcb1ab* double knockout mice.⁵⁸⁻⁶⁰ The
211 authors highlight methodological differences between the studies which might limit comparisons: for
212 instance, in one study isotope radioactivity rather than intact steroid concentration was measured,
213 and the use of labelled corticosterone in adrenally intact animals might have resulted in isotope
214 dilution by endogenously secreted corticosterone.

215 From these findings we might predict that the HPA axis would be relatively suppressed by the
216 accumulation of glucocorticoids in the brain if ABCB1 activity is reduced. Indeed, *Abcb1a*-knockout
217 mice do show evidence of HPA axis suppression, with lower basal and stress-stimulated levels of
218 corticosterone, adrenocorticotrophic hormone (ACTH) and corticotrophin-releasing hormone than
219 control animals, with the effect localised to the hypothalamic level.⁶¹ Furthermore, mice treated with
220 the ABCB1 inhibitor tariquidar show an attenuated corticosterone response to stressful stimuli.⁶²

221

222 **[H3] Insights from dogs and humans.**

223 The ABCB1 protein is well conserved in larger, cortisol-dominant species, with a notable exception
224 being in Collie-derived dogs. Like *Abcb1a*-knockout mice,⁵⁴ these animals are exquisitely sensitive to
225 ivermectin owing to a 4-base pair deletion mutation (termed *Mdr1-1Δ*) for which 40–50% of this breed
226 are homozygous.^{63,64} This mutation results in a severely truncated protein (<10% of normal length),
227 which is predicted to be non-functional. Anecdotally, Collie dogs are reported to recover relatively
228 slowly from illness,⁶⁵ and animals with the MDR1^{-/-} genotype showed chronic suppression of the HPA
229 axis, with lower basal cortisol levels and greater ACTH suppression in response to dexamethasone
230 than their wild-type counterparts. It has been hypothesised that enhanced brain retention of cortisol
231 (the dominant canine glucocorticoid) leads to this HPA axis suppression, and predisposes the animals
232 to a form of relative corticosteroid insufficiency.⁶⁵ This hypothesis has been supported by a
233 metabolomics study demonstrating lower urinary cortisol metabolites in MDR1^{-/-} dogs than controls
234 [reaching significance for allotetrahydrocortisol (11.2 ± 3.4 ng/L versus 20.7 ± 14.9 ng/L, $P=0.006$) and
235 β -cortol (105.5 ± 63.3 ng/L versus 221.0 ± 225.5 ng/L, $P=0.025$)].⁶⁶

236 In a human study, the corticosterone:cortisol ratio in brain autopsy specimens was five times
237 greater than the corresponding ratio in plasma in age- and sex-matched healthy controls.⁴ Similarly,
238 the ratio of corticosterone to cortisol in live subjects is 5-6 times higher in cerebrospinal fluid than in
239 plasma.⁶⁷ Many drugs, including verapamil and cyclosporin A, inhibit ABCB1, but their experimental
240 use to test ABCB1 physiology in humans is hampered by toxicity at levels that are too low to carry out
241 meaningful studies of ABCB1 inhibition.⁶⁸

242

243 **[H3] ABCB1 and ABCC1 modulate the HPA axis.**

244 These results are all consistent with the hypothesis that ABCB1 at the BBB exports cortisol and thereby
245 modulates negative feedback of the HPA axis in cortisol-dominant species. The absence of ABCC1 in

246 the brain and BBB is consistent with corticosterone being retained to a greater extent than cortisol in
247 brain. One additional complexity, however, is that the pituitary gland (which expresses both
248 transporters)⁶⁹ lies outside the BBB but also contributes to the control of the HPA axis. We have
249 demonstrated that administration of probenecid, an inhibitor of ABCC1, induces greater tonic
250 negative feedback of the HPA axis in healthy subjects than placebo as judged by elevations in ACTH
251 and cortisol during combined mineralocorticoid and glucocorticoid receptor antagonism.⁷⁰ This finding
252 is consistent with ABCC1 also contributing to the export of corticosterone from the pituitary gland or
253 other central feedback areas, and warrants further investigation in animal models.

254

255 **[H2] ABCC1 transporters in adipose tissue**

256 **[H3] *Insights from mice and humans.***

257 In contrast with the BBB, where ABCB1 is more abundant than ABCC1, the reverse is true in adipose
258 tissue. Glucocorticoids within adipose tissue induce lipogenesis; in particular, they stimulate the
259 accumulation of lipids in visceral tissue and the production of adipokines.⁷¹ *Abcc1*-knockout mice
260 infused with corticosterone and cortisol showed an enhanced accumulation of corticosterone but not
261 cortisol in adipose tissue, accompanied by the upregulation of both glucocorticoid-responsive and
262 adipogenic genes.⁶

263 We have also demonstrated that human adipocytes preferentially accumulate cortisol over
264 corticosterone, and that this accumulation was reversed in vitro after treatment with the ABCC1
265 inhibitors probenecid or MK-571.⁶ It was also accompanied by activation of glucocorticoid-responsive
266 and adipogenic genes (*PER1*, *ADIPOQ*, *ATGL*, *HSL*) and resulted in the increased accumulation of fatty
267 acids in lipid droplets.⁶ Moreover, during infusion of cortisol or corticosterone in vivo in patients with
268 primary adrenal insufficiency, the induction of glucocorticoid-responsive gene expression (*PER1*, *LPL*)
269 in adipose tissue was greater in response to cortisol than to corticosterone (achieved at plasma

270 glucocorticoid levels which were equipotent for ACTH suppression).⁶ This suggests that corticosterone
271 could have a more favourable metabolic profile than cortisol in glucocorticoid replacement,
272 particularly when ACTH suppression is a target.

273

274 **[H2] ABCB1 and ABCC1 in the placenta**

275 As the interface between the mother and the fetus in pregnancy, the placenta functions both as a
276 nutritive source and a barrier, including to glucocorticoid transport. The fetus is unable to synthesize
277 cortisol until the third trimester, and therefore depends on maternal cortisol; however, although
278 maternal cortisol levels increase by several-fold during pregnancy, this increase is not transferred to
279 the fetus indiscriminately.⁷² In early pregnancy, excessive glucocorticoids are detrimental to the fetus,
280 so the placenta provides a glucocorticoid barrier,⁷³ but it confers a more facilitative role towards term
281 for fetal organ maturation.⁷⁴

282

283 **[H3] *The placental glucocorticoid barrier: 11 β -hydroxysteroid dehydrogenase 2.***

284 The enzyme 11 β -hydroxysteroid dehydrogenase 2 is viewed as the main component of the placental
285 glucocorticoid barrier, converting active cortisol to inactive cortisone.⁷⁵ However, the results of a study
286 in which 11 β -hydroxysteroid dehydrogenase 2 was inhibited during ex vivo perfusion of human
287 placentas collected on ice immediately after delivery suggested that the enzyme might contribute only
288 part of the glucocorticoid barrier, as cortisol transfer was restricted even at maximal inhibition of 11 β -
289 hydroxysteroid dehydrogenase 2.⁷⁶ The role of other mechanisms that are operating at the placental
290 barrier, such as transmembrane transport, therefore warrants further consideration.

291

292 **[H3] *The placental glucocorticoid barrier: ABCB1 and ABCC1.***

293 ABCB1 is located within syncytiotrophoblasts at the apical brush-border membrane, in direct contact
294 with maternal blood.⁷⁷ It is highly expressed in early pregnancy and decreases towards term,
295 consistent with the physiological role suggested above.⁷⁸ As occurs in other tissues, glucocorticoids
296 have been shown to upregulate the expression of *ABCB1* in the placenta in the first trimester, which
297 might enhance the barrier effect.⁷⁹ Data demonstrating low concentrations of ABCB1 substrates
298 (antiretrovirals, for example) in the fetal circulation both at birth and in the ex vivo perfused placenta
299 indicate that ABCB1-mediated export towards the maternal circulation is active in vivo.⁸⁰ ABCC1 is
300 located on the fetal-facing placental surface and has been identified in cytotrophoblasts,
301 syncytiotrophoblasts and the fetal endothelium.⁸¹ This localization might be consistent with a role in
302 transferring ABCC1 substrates such as folic acid to the fetus and, in contrast with *ABCB1*, *ABCC1* is
303 upregulated towards term.^{81,82} Studies of other ABCC1 substrates using the inhibitors probenecid and
304 MK-571 have not demonstrated a clear effect on cross-placental transfer, so cannot be extrapolated
305 to corticosterone transport.⁸³ It has been shown that the cortisol:corticosterone ratio is higher in the
306 maternal circulation (15:1) than in the umbilical vein (7:1) at term,⁸⁴ which might be accounted for by
307 fetal adrenal cortisol:corticosterone secretion rates or by the facilitated transport of maternal
308 corticosterone by ABCC1 into the fetal circulation.

309

310 [H1] Regulation and dysregulation

311 [H2] Regulation of ABCB1

312 The mechanisms underpinning the regulation of the expression of *ABCB1* have been reviewed
313 thoroughly elsewhere.⁸⁵⁻⁸⁷ The *ABCB1* promoter contains a number of areas of interest, including
314 binding sites for the tumour suppressor p53, heat shock proteins and adopted orphan receptors,
315 including the pregnane X receptor (PXR) and constitutive androstane receptor (CAR), which bind a
316 number of xenobiotic ligands.⁸⁸ Xenobiotics, inflammatory mediators and cellular stress (such as
317 irradiation, heat shock, hypoxia) typically upregulate *ABCB1* expression through common pathways

318 involving nuclear factor kappa B (NF- κ B) and Y-box binding protein.^{89,90} This upregulation appears to
319 be a protective response, and polymorphisms in NF- κ B are linked with increasing colon cancer risk,
320 potentially owing to enhanced cellular exposure to toxins.⁹¹

321 Glucocorticoids modulate the expression of ABCB1 mRNA and protein in rodents and humans.
322 This modulation has been demonstrated across multiple tissues using dexamethasone, prednisolone,
323 cortisol, methylprednisolone and some inhaled glucocorticoids.^{33,79,92-97} Although glucocorticoids
324 predominantly induce the expression of *ABCB1*, this effect might be specific to some species or cell
325 types, as there are also instances of *ABCB1* expression being downregulated.⁹⁸ This glucocorticoid
326 effect is inhibited in the presence of the glucocorticoid receptor blocker RU486, indicating that this
327 effect is at least partly mediated via the glucocorticoid receptor, but as no consensus glucocorticoid
328 response element has been found in the human *ABCB1* promoter, it is assumed to be an indirect
329 genomic effect. Dexamethasone-mediated upregulation of *ABCB1* in retinal pigment epithelium was
330 reported to be abolished when the PXR receptor was silenced, implying that PXR (which does contain
331 a consensus glucocorticoid response element) is either a co-regulator or a target of the glucocorticoid
332 receptor.^{97,99,100} This upregulation of expression raises concerns about increased drug efflux when
333 glucocorticoids are used in combination with other ABCB1 substrates (as often occurs in
334 chemotherapy protocols), and is theorised to be a cause of glucocorticoid resistance in conditions such
335 as asthma;³³ however, this effect has also been exploited clinically — for example, methylprednisolone
336 is used in the treatment of paraquat toxicity to increase excretion of the drug.¹⁰¹

337 However, the regulation of ABCB1 in inflammation is complex and potentially biphasic.
338 Evidence from rodent studies indicates that, in the very early stages of inflammation, ABCB1 is
339 functionally inhibited by lipopolysaccharides and inflammatory cytokines, despite mRNA levels
340 remaining constant, perhaps owing to ABCB1 being trafficked away from the cell membrane; later in
341 the evolution of inflammation, however, ABCB1 mRNA and protein levels are upregulated by the
342 cytokines tumour necrosis factor and endothelin 1 converging on the NF- κ B pathway.⁸⁹ Protein

343 turnover at the cell surface under normal conditions is relatively slow (the half-life of ABCB1 is
344 estimated at just over 24 hours)¹⁰² and there might be a role for post-translational and other
345 mechanisms in modulating this turnover. Taken together, this evidence suggests that in times of
346 increased physiological stress (for example, in response to illness or injury), ABCB1 can be upregulated
347 both by stress-activated glucocorticoids and by signals released by cellular damage. This upregulation
348 might result in positive feedback on cortisol production by further restricting glucocorticoid access to
349 sites of higher negative feedback.

350

351 **[H2] Regulation of ABCC1**

352 Most research on factors affecting the expression levels of *ABCC1* and its protein activity relates to
353 cancer biology and chemotherapeutics, whilst physiological regulation has been poorly studied to
354 date. Basal transcription of *ABCC1* is stimulated by the SP1 transcription factor¹⁰³ which is, in turn,
355 inhibited by the tumour suppressor protein p53.¹⁰⁴ It has not been clearly established whether PXR
356 affects *ABCC1* transcription^{105,106} and, although early mapping of the *ABCC1* promoter in a human
357 leukaemic cell line did reveal a putative glucocorticoid response element, dexamethasone has not
358 been shown to alter *ABCC1* expression in the human placenta or in lymphocytes.^{94,107-109} Furthermore,
359 we cannot clearly conclude whether *ABCC1* is affected by acute inflammation in the same way that
360 *ABCB1* is, as both unchanged and increased mRNA expression have been reported in response to
361 mediators such as lipopolysaccharide, tumour necrosis factor, IL-1 and IL-6.¹¹⁰⁻¹¹²

362 In vitro studies investigating the metabolic regulation of *ABCC1* have focused on endothelium,
363 and have demonstrated that expression of the transcript is downregulated in a hyperglycaemic
364 environment.¹¹³ Metformin, a drug commonly used in the treatment of type 2 diabetes mellitus, is
365 known to reduce *ABCC1* expression in a human hepatocellular carcinoma cell line through the AMP-
366 activated protein kinase–hypoxia-inducible factor 1 pathway.¹¹⁴

367 Whilst limited, overall this evidence suggests that *ABCC1* is regulated differently from *ABCB1*,
368 and is predominantly responsive to metabolic and immunomodulatory signals rather than to
369 mediators of acute stress or inflammation.

370

371 **[H2] Pathological dysregulation**

372 There have been few studies of variations in ABC transporter expression beyond the
373 descriptions in various cancers mentioned above. A transcriptomic analysis utilising single-cell RNA
374 sequencing showed upregulation of *ABCB1* in the adrenal cortex of patients with ACTH-dependent
375 Cushing disease.³⁶ This upregulation probably reflects the effects of glucocorticoids on *ABCB1*
376 expression, but might contribute to pathogenicity by further enhancing the export of cortisol from the
377 gland. Hypothesizing that steroid retention in adipocytes due to low levels of *ABCC1* could be a driving
378 mechanism for obesity, we actually found that *ABCC1* mRNA levels were upregulated in the adipose
379 tissue (subcutaneous and visceral) of individuals with obesity compared with lean individuals, which
380 may reduce glucocorticoid concentrations in adipocytes, although this reduction might only be true
381 for corticosterone.⁶

382

383 **[H2 Lessons from human genetics**

384 **[H3] Germline mutations in *ABCB1*.**

385 Human germline mutations in *ABCB1* are rare. To our knowledge, there are only two publications of
386 *ABCB1* mutations: twin girls with recurrent reversible toxic encephalopathy alongside febrile illness,
387 ¹¹⁵ and a 13-year old boy with ivermectin sensitivity.¹¹⁶ In both cases, the mutations were identified
388 by whole exome sequencing and show **compound heterozygosity**. The twin girls were found to have
389 a nonsense mutation (p.Pro1182X) combined with a splice variant (c.2786 + 1 G>T) and showed
390 markedly enhanced CNS retention of ¹¹C-verapamil on PET imaging in comparison with their parents.

391 Their symptoms were suspected to be caused by retention of inflammatory mediators within the brain
392 during acute illness, and it was shown in a mouse model by the authors that cytokines Tumour
393 Necrosis Factor, IL-1, IL-6 and Ccl-2 were retained in brain at 24 hours after lipopolysaccharide injection
394 in *Abcb1ab* knockout versus wild-type animals. The investigators estimated from studies in
395 lymphocytes that only ~10% of functional ABCB1 protein was expressed. In the other case, the
396 affected boy presented with severe neurological side effects after a single oral dose of ivermectin to
397 treat scabies and was found to have inherited a nonsense mutation in ABCB1 from each parent (c.2380
398 C>T and c.3053_3056delTTGA), both of which are predicted to result in the loss of the carboxy-
399 terminal nucleotide-binding domain. The boy and twin girls were otherwise healthy and growing
400 normally in each case.

401 **[H3] Germline mutations in ABCC1.**

402 Similarly, there is only one published mutation of *ABCC1* of clinical significance: a heterozygous
403 missense mutation (c.1769 A>G) identified as causing familial sensorineural deafness.¹¹⁷ *ABCC1* has
404 been found within the rodent cochlea, where it could be protective against neurotoxins.¹¹⁸ This
405 mutation is thought to disrupt hydrogen bonds, and thus stability between the helices of the
406 transmembrane domains in the proteins, but analysis of lymphoblastoid cell lines derived from
407 affected family members showed loss of around 40–45% of *ABCC1* mRNA expression when compared
408 with those unaffected, suggesting an additional impairment in mRNA stability. Extrusion of SNARF-1,
409 a known *ABCC1* substrate, from lymphoblastoid cells as a measure of transport activity was
410 subsequently shown to be slower.¹¹⁷

411 **[H3] Polymorphisms in ABCB1 and ABCC1.**

412 With nonsense and frameshift mutations being rare, there have been attempts to correlate common
413 polymorphisms with clinically relevant outcomes (reviewed in ¹¹⁹).

414 Three *ABCB1* variants are common in humans: c.2677 G>A/T, c.3435 C>T and c.1236 C>T. The c.3435
415 C>T allele is **synonymous**, but might affect mRNA stability;¹²⁰ c.1236 C>T is silent; and c.2677 G>A/T
416 results in an amino acid substitution (alanine to serine or threonine), which could potentially result in
417 substrate changes. There is marked variation in the frequency of these polymorphisms across different
418 races: for example, c.3435 C>T is much less common in African populations (~80% of people from
419 West Africa are homozygous for the C allele versus ~20% of individuals from western Europe).^{120,121}
420 However, it has not been convincingly demonstrated that these variants affect substrate transport,
421 for instance levels of the *ABCB1* substrate digoxin have been found to be increased, decreased and
422 unchanged in the plasma of individuals with these polymorphisms. Subsequent attempts to correlate
423 polymorphisms with response to chemotherapeutics, drug side effects, and resistance to anti-
424 retroviral and anti-epileptic therapies have been similarly inconclusive.¹²²⁻¹²⁴

425 Studies of the HPA axis in individuals with *ABCB1* variants have unfortunately been
426 inadequately powered. No differences were found in the levels of evening cortisol and ACTH in 30
427 Japanese men with C/C, C/T or T/T c.3435 genotypes (the variant associated with potentially reduced
428 transporter mRNA stability); however, another study, of 51 women, reported lower levels of cortisol
429 in the plasma, taken at 6 pm, of individuals with one or two copies of the T allele compared with C/C
430 controls; these lower levels reached significance only in the follicular menstrual phase so an
431 interaction with sex hormones is proposed.^{125,126} In one candidate gene study of over 5,000 Japanese
432 individuals, the c.2677 G>A/T variant was highly associated with increased body mass index, which
433 could potentially reflect increased HPA axis activity, whilst in a study of 154 individuals with
434 depression, the response of cortisol (but not ACTH) to corticotrophin-releasing hormone was lower in
435 c.2677 TT homozygotes than in the major allele (GG) or heterozygous (TG) groups, which was taken
436 to reflect reduced adrenal cortisol release.^{36,127} However, neither plasma cortisol levels nor body mass
437 index has been associated with any *ABCB1* polymorphisms in larger cohorts.

438 Genetic studies have also been undertaken in patients taking exogenous steroids. In a cohort
439 of 171 patients requiring long-term treatment with glucocorticoids for adrenal insufficiency, those
440 patients with the c.3435 TT genotype had lower bone density than CC or CT groups, suggesting greater
441 systemic steroid absorption or enhanced bone retention.¹²⁸ There have been attempts to correlate
442 glucocorticoid treatment outcomes in patients with rheumatoid arthritis, inflammatory bowel disease,
443 immune thrombocytopenic purpura and nephrotic syndrome with ABCB1 polymorphisms.¹²⁹⁻¹³² Most,
444 but not all, indicate a higher steroid response with the minor allele of the studied polymorphism, but
445 studies are limited by sample size and a failure to control for multiple testing.

446 Documented polymorphisms for *ABCC1* are mostly rare and non-coding, and have not been
447 assessed in the context of HPA axis activity or metabolism.¹³³ Three polymorphisms might predict the
448 outcome of acute myeloid leukaemia, but any corresponding effect of these polymorphisms on
449 transporter expression or function has so far not been established.¹³⁴

450

451 **[H1] Implications and future research**

452 The observations that two ABC transporters influence the retention of glucocorticoids in
453 tissues allow us to add membrane transporters to the list of factors that are involved in the
454 metabolism of glucocorticoids at the pre-receptor level (FIG. 4). These observations provide insights
455 into HPA axis physiology and how corticosterone and cortisol might carry out different functions in
456 species that produce both steroids. These findings also provide therapeutic opportunities for anti-
457 inflammatory and physiological replacement steroid therapies that might better target tissues
458 mediating efficacy while avoiding those mediating toxicity.

459

460 **[H2] Revised glucocorticoid physiology**

461 In rodents, the lack of steroid 17-hydroxylation necessitates that corticosterone is the sole
462 endogenous glucocorticoid.¹³⁵ In humans and other species in which both glucocorticoids circulate, it
463 is common to consider them interchangeable. Indeed, cortisol and corticosterone share similar
464 metabolic pathways (for example, susceptibility to metabolism by 11 β -hydroxysteroid dehydrogenase
465 enzymes) and affinities for the glucocorticoid and mineralocorticoid receptors.¹³⁶⁻¹³⁹ However,
466 corticosterone does exhibit differences to cortisol, including more rapid clearance from the
467 circulation, and a greater response to ACTH, such that the corticosterone:cortisol ratio rises under
468 stress.¹⁴⁰⁻¹⁴²

469 The findings outlined in this Review further demonstrate that cortisol and corticosterone are
470 not interchangeable with respect to glucocorticoid action. Specifically, in tissues where ABCB1 but not
471 ABCC1 is present, such as the brain, cortisol concentrations are constrained by export back into the
472 circulation and corticosterone can play a disproportionate role. Conversely, in tissues such as adipose
473 where ABCC1 but not ABCB1 is expressed, corticosterone is exported and the response to cortisol can
474 be disproportionate (FIG. 5). This observation raises the concept of a distinctive role for corticosterone
475 in mediating HPA axis negative feedback. In the stressed state, the ability to restrict the high levels of
476 circulating cortisol from accessing higher centres might prevent axis suppression and facilitate
477 recovery, as demonstrated by the *Mdr1-1 Δ* Collie dogs who lack this capacity.⁶⁵ It is recognised in
478 other species that the ratio of cortisol to corticosterone and the peak levels of circulating
479 glucocorticoids vary seasonally,¹⁴³ possibly in response to photoperiod length. If corticosterone is
480 more accessible to negative feedback sites, and less peripherally anabolic than cortisol (in terms of
481 effects on adipose tissue), then the energy-expending stress response might be restrained and access
482 to vital adipose energy stores when food is scarce might be improved. Conversely, with a slower
483 turnover than corticosterone in the circulation and adipose tissue in comparison with other tissues
484 such as brain and liver,¹⁴⁴ cortisol might provide the option for medium-term adjustments, in
485 comparison with the acute changes mediated by corticosterone.

486 Understanding the implications of the differential control and actions of cortisol and
487 corticosterone in glucocorticoid physiology will require a detailed dissection of the dynamics of ligand
488 availability for receptors within human target tissues in vivo. The increasing use of exome-wide
489 sequencing in clinical as well as research settings might well identify further individuals or families
490 with significant *ABCB1* and *ABCC1* mutations and offer new routes to addressing these key
491 physiological issues.

492

493 [H2] Novel glucocorticoid therapies

494 A major limitation of current glucocorticoid therapies is their narrow **therapeutic index**.
495 Despite extensive efforts, it has proved difficult to develop selective glucocorticoid receptor
496 modulators with pharmacodynamic interactions that discriminate between efficacious and toxic gene
497 transcription.¹⁴⁵ An alternative approach depends on the premise that efficacious and toxic effects are
498 often mediated in different tissues, suggesting that the therapeutic index could be improved by
499 modifying the pharmacokinetics of steroid drugs to ‘target’ them to the tissues where efficacy is
500 mediated while avoiding tissues where toxicity is mediated. Could this be achieved by using steroids
501 with different affinities for the *ABCB1* and *ABCC1* transporters?

502 When considering physiological replacement in patients with adrenal insufficiency, the
503 challenges of this narrow therapeutic index are well documented, with adverse outcomes including,
504 but not limited to, obesity, osteopenia and insulin resistance attributable to the steroid regime of
505 these patients.^{146,147} Such challenges are particularly evident in patients with congenital adrenal
506 hyperplasia (CAH), in whom doses of glucocorticoid that achieve adequate adrenal androgen
507 suppression are invariably associated with morbidity.¹⁴⁶ All glucocorticoids currently used to replace
508 cortisol (hydrocortisone, prednisolone, dexamethasone, and the active metabolites of pre-drugs
509 cortisone and prednisone) are substrates for *ABCB1* but not *ABCC1*. Although pharmacokinetic
510 adjustments, such as delayed release preparations, might confer some benefits,^{148,149} they cannot

511 overcome the closeness of the dose-response relationship between efficacy and toxicity, and the
512 prospect of choosing a glucocorticoid based on affinity for ABCC1 over ABCB1 is an intriguing
513 therapeutic prospect.

514 As one such glucocorticoid, corticosterone is not currently available in an oral form, but our
515 experimental work using intravenous corticosterone has provided proof-of-concept of the potential
516 advantages of corticosterone in avoiding harmful metabolic effects mediated in adipose tissue. As
517 described earlier, infused cortisol induced a greater response of glucocorticoid-responsive gene
518 expression compared with infused corticosterone in the adipose tissue of patients with Addison
519 disease.⁶ In a similar study, 14 individuals with CAH also underwent ramped cortisol and
520 corticosterone infusions; despite higher plasma levels of corticosterone being achieved, the amount
521 of insulin released was greater in response to cortisol than to corticosterone – a marker of
522 glucocorticoid effect on adipose to induce insulin resistance.¹⁵⁰

523 The potential for glucocorticoid therapies that avoid toxicity in metabolic tissues deserves
524 further investigation but would require the generation of an oral corticosterone preparation for
525 practical administration to patients.

526

527 **Conclusions**

528 We have collated evidence from cell, animal and human studies that the ATP-binding cassette
529 transporters ABCB1 and ABCC1 differentially export cortisol, corticosterone and synthetic
530 glucocorticoids from tissues and contribute to pre-receptor glucocorticoid regulation. Differing
531 transporter expression profiles in the brain, placenta and adipose confer different tissue sensitivities
532 to these steroids, which might be important for optimising the responsiveness of the HPA axis,
533 controlling fetal exposure to steroids throughout gestation, and optimising adipose fuel metabolism.
534 Although much is known about these transporters in the context of multidrug resistance, their

535 physiological roles and regulation remain largely unexplored. The prospect of developing steroid
536 therapies with transporter affinities that are tailored to give improved efficacy, without deleterious
537 peripheral toxicity, offers new avenues for exploration for the management of inflammatory and
538 endocrine diseases.

539

540 Box 1 - Multidrug Resistance

541

542 Multidrug resistance (MDR) is the ability of malignant cells to evade the actions of a broad range of
543 chemotherapeutic agents. Tumours which are initially very sensitive can become resistant to multiple
544 agents over the course of the disease, ultimately resulting in treatment failure and disease
545 progression. There are several potential reasons for this, but the key mechanism is increased drug
546 efflux out of malignant cells by membrane transporters, particularly those of the ABC family. Some
547 tumours have innately high levels of transporter expression, but others develop this after exposure to
548 chemotherapy.¹⁵¹

549

550 ABCB1 is the transporter most widely associated with MDR, particularly since alkylating agents,
551 anthracyclines and vinca alkaloid drugs are all substrates.²¹ ABCC1 and ABCG2 (aka Breast cancer
552 Resistance Protein) are also implicated in MDR.

553

554 As examples, survival rates from lung cancer, multiple myeloma and acute myeloid leukaemia have
555 been inversely associated with levels of ABCB1 expression.¹⁵²⁻¹⁵⁴ High levels of ABCC1 expression are
556 associated with poor outcomes in childhood neuroblastoma,¹⁵⁵ whilst over-expression of ABCG2 is a
557 negative prognostic factor in pancreatic ductal adenocarcinomas.¹⁵⁶

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1058 **Related links**

1059 The Human Protein Atlas: <https://www.proteinatlas.org/>

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1061 **Key points**

- 1062 • Humans have two circulating glucocorticoid hormones, cortisol and corticosterone, which
1063 diffuse into cells to become transcription factors when bound to their intracellular receptors.
- 1064 • The availability of glucocorticoids to interact with their receptors depends not only on their
1065 plasma concentration but also on their intracellular concentration, which is modulated by
1066 intracellular enzymes and by transmembrane transporters.
- 1067 • Glucocorticoids are susceptible to cellular export by membrane transporters from the ABC
1068 (ATP-binding cassette) transporter family: cortisol is a substrate for the ABCB1 transporter,
1069 and corticosterone for ABCC1.
- 1070 • Tissues expressing ABCB1 (such as the brain) might be relatively sensitive to corticosterone
1071 over cortisol; those expressing ABCC1, such as adipose, might be more sensitive to cortisol.
- 1072 • In future, therapeutic glucocorticoids could be selected on the basis of lower tendency to be
1073 exported from sites of efficacy and higher tendency for export from sites where harmful side
1074 effects occur.

1075 **Glossary terms:**

1076 sanctuary sites – areas within the body that are relatively protected from access by drugs (e.g. anti-
1077 cancer agents) and toxins

1078 α -helices – a form of secondary protein structure formed by hydrogen bonding between amine and
1079 carbonyl groups of amino acids 4 apart, and resulting in a stable rod shape

1080 luminal surfaces – the lining surfaces of body channels, such as the intestines or blood vessels

1081 polyspecificity – the capacity to bind multiple unrelated substrates

1082 glutathione coupling – conjugation with the tri-peptide glutathione

1083 phase II hepatic metabolites – conjugation of a substance to another molecule, such as glutathione
1084 or glucuronide, in the liver to make it more water soluble and thus facilitate excretion

1085 syncytiotrophoblasts – cells forming the outer layer of the placenta, and the major site of gas and
1086 nutrient exchange between mother and fetus

1087 cytotrophoblasts – the inner stem cell layer of the placenta villi – cellular precursors to
1088 syncytiotrophoblasts

1089 adopted orphan receptors – an orphan receptor is a receptor whose ligand has not been identified.
1090 It can later be termed an “adopted orphan receptor” when a ligand is discovered.

1091 compound heterozygosity – the presence of two different mutant alleles at a genetic locus

1092 lymphoblastoid cell lines – immortalised cells which are derived from, and closely resemble,
1093 peripheral blood lymphocytes

1094 synonymous – a silent genetic mutation where a change in DNA sequence does not result in a
1095 change in the amino acid sequence of the protein produced

1096 therapeutic index – the margin between the desirable and undesirable effects of a drug. The
1097 narrower the margin, the more likely it is that side effects will occur at a therapeutic dose.

1098 **Figure legends**

1099 **Figure 1: Action and structure of ABCB1 and ABCC1.** a| In general, most ABC transporters are
1100 comprised of two transmembrane domains (TMDs) and 2 nucleotide binding domains (NBDs). In the
1101 proposed model of action, binding of ATP dimerises the NBDs and induces a conformational change
1102 within the TMDs, resulting in the switch between 'inward' and 'outward' facing configurations.^{17,18}
1103 Subsequent hydrolysis of ATP returns the transporter to baseline status. b| Ribbon diagram of human
1104 ABCB1 (Protein Data Bank ID 6QEX) and c| ribbon diagram of bovine ABCC1 (Protein Data Bank ID
1105 5UJA). The amino (N)- and carboxyl (C)- terminal halves are coloured magenta and blue, respectively.
1106 NBD1 and NBD2 are coloured green and yellow, respectively, with the drug-binding pocket
1107 highlighted.

1108

1109 **Figure 2: Tissue-specific expression of ABCB1 and ABCC1.** Human expression of *ABCB1* and *ABCC1*, as
1110 derived from data from the Human Protein Atlas, is shown. Expression is normalised to an Nx
1111 (normalised expression) value based on outputs from the Human Protein Atlas, the genotype-tissue
1112 expression (GTEx) project and FANTOM5 transcriptomic analyses (data available online from The
1113 Human Protein Atlas).²³ Tissues are ranked in order of *ABCB1:ABCC1* ratio, such that those towards
1114 the top of the Y axis have greater *ABCB1* expression, and those at the bottom have higher *ABCC1*
1115 expression.

1116

1117 **Figure 3: Tissue ABC transporter expression determines glucocorticoid sensitivity.** The influence of
1118 ABCB1 and ABCC1 on the retention of common glucocorticoids within human target tissues according
1119 to transporter affinity is depicted. Steroids in red are predominantly substrates for ABCB1, those in
1120 dark blue are predominantly substrates for ABCC1, and those in light green are substrates for neither
1121 transporter. Passive diffusion is indicated by double-headed arrows.

1122

1123 **Figure 4: Intracellular glucocorticoid regulatory pathways.** After diffusing into cells (double-headed
1124 arrows), the glucocorticoids cortisol and corticosterone might (1) be exported by the membrane-
1125 bound ATP transporters ABCB1 and ABCC1; (2,3) might undergo enzymatic metabolism by 11 β -
1126 hydroxysteroid dehydrogenase (11 β -HSD), 5 α reductase or carbonyl reductase enzymes; or (4) might
1127 become incorporated in the intracellular steroid pool. These processes restrict access to the nuclear
1128 glucocorticoid and/or mineralocorticoid receptors (GR and MR), which mediate the cellular response
1129 (5).

1130

1131 **Figure 5: Modulation of the hypothalamic–pituitary–adrenal (HPA) axis by ABCB1 and ABCC1.**
1132 Glucocorticoids are secreted from the adrenal cortex upon stimulation by signals from the
1133 hypothalamus and pituitary . They act peripherally on sites throughout the body, and feed back to the
1134 hypothalamus, pituitary and higher centres to maintain homeostasis. ABCB1 present at the blood–
1135 brain barrier might act to restrict the access of cortisol to feedback sites. Conversely, ABCC1, which is
1136 found without ABCB1 in adipose and skeletal muscle, exports corticosterone but not cortisol. The
1137 activity of the adrenal enzyme CYP17 (17-hydroxylase) determines the ratio of secreted
1138 cortisol:corticosterone.

1139

1140 **Text for Table of Contents**

1141 This Review discusses the ATP-binding cassette (ABC) proteins ABCB1 and ABCC1 and their
1142 preferential cellular export of cortisol and corticosterone, respectively, as well as exploring the
1143 potential to select therapeutic glucocorticoids on the basis of their different tendencies for export to
1144 avoid harmful side effects.

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