The Effect of Chronic Renal Failure on Drug Metabolism and Transport

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Abstract

Background—Chronic renal failure (CRF) has been shown to significantly reduce the nonrenal clearance and alter bioavailability of drugs predominantly metabolized by the liver and intestine.

Objectives—The purpose of this article is to review all significant animal and clinical studies dealing with the effect of CRF on drug metabolism and transport.

Methods—The National Library of Medicine PubMed was utilized with the search terms ‘chronic renal failure, cytochrome P450, liver metabolism, efflux drug transport and uptake transport’ including relevant articles back to 1969.

Results—Animal studies in CRF have shown a major downregulation (40-85%) of hepatic and intestinal cytochrome P450 (CYP) metabolism. High levels of parathyroid hormone, cytokines, and uremic toxins have been shown to reduce CYP activity. Phase II reactions and drug transporters such as P-glycoprotein (Pgp) and organic anion transporting polypeptide (OATP) are also affected.

Conclusion—CRF alters intestinal, renal, and hepatic drug metabolism and transport producing a clinically significant impact on drug disposition and increasing the risk for adverse drug reactions.

Keywords
chronic renal failure; cytochrome P450; CYP; drug transport; efflux transporters; hepatic clearance; P-glycoprotein; uremic toxins

1. Introduction

It is a basic principle of clinical pharmacokinetics that drugs predominantly cleared by the kidney require dose adjustment in acute and chronic renal failure (ARF and CRF). It is widely assumed that it is unnecessary to dose adjust drugs cleared exclusively by hepatic metabolism and transport in the CRF patient population. However, accumulated data from animal and human studies do not support this assumption [1-6]. In the past 5 years, a number of reviews have appeared in the literature, which detail the effects of CRF on drug metabolism and transport including hepatic, renal, and intestinal processes [7-11]. This review was undertaken to illustrate various mechanisms by which CRF can alter drug metabolism and transport and to provide examples of actual or potential clinical relevance. Citations were derived from PubMed in years 1990-2008 using the search terms, ‘liver metabolism and transport, chronic renal failure’.
renal failure, intestinal transport and cytochrome P450 metabolism, including other key articles dating back to 1969. Recently, exciting new studies including intestinal, hepatic, and renal drug transporters have begun to elucidate the mechanisms of this phenomenon. The major focus of this review is to summarize these recent studies.

2. Background Pharmacokinetics

The following is a short review of basic pharmacokinetic principles focusing on hepatic clearance[12]. Drug absorption, nonrenal clearance, and volume of distribution of drugs are in fact altered by CRF via changes in hepatic clearance, intestinal absorption and first pass metabolism, hepatic, renal, and intestinal transport, plasma protein binding, and tissue binding. Perturbations of plasma protein binding of drugs occur with CRF which usually leads to an increase in the free unbound fraction (fu). Conformational changes in albumin resulting in altered albumin binding affinity or competition with uremic toxins are thought to be the mechanism. These effects may be partially reversed by dialysis.[13-28]. CYP activity in the kidney is estimated to be 20% of liver activity per gram of tissue. Conjugation reactions such as glucuronidation and acetylation are also impaired by CRF [28].

Hepatic clearance (CLH) is a function of hepatic blood flow (QH) and hepatic extraction ratio (E_H) (Eq 1) [1,29].

\[
\text{CL}_H = Q_H E_H
\]

Eq(1)

The relationship between hepatic blood flow, intrinsic hepatic clearance, and protein binding are expressed in the well stirred model of total hepatic clearance (CL_HTOT) developed by Wilkinson and Shand (Eq 2) [29].

\[
\text{CL}_\text{HTOT} = \frac{Q_H \text{CL}_\text{INT} f_u}{Q_H + \text{CL}_\text{INT} f_u}
\]

Eq(2)

where: \(f_u\) = unbound fraction; \(\text{CL}_\text{INT}\) = intrinsic hepatic clearance

Thus, intrinsic hepatic clearance and free unbound fraction of the drug are the determinants of total hepatic clearance. Intrinsic hepatic clearance relates the rate of metabolism at steady-state to the free unbound fraction in blood. If the drug is highly extracted we assume \(\text{CL}_\text{INT} f_u >> Q_H\), then \(\text{CL}_\text{HTOT} = Q_H\) and total hepatic clearance is blood flow limited. If the drug is a low extraction agent then we assume \(Q_H >> \text{CL}_\text{INT} f_u\) then \(\text{CL}_\text{HTOT} = \text{CL}_\text{INT} f_u\) and would be highly influenced by changes in plasma protein binding and intrinsic hepatic clearance as simulated in empirically derived models [30]. Hepatic blood plasma flow has been shown to be unaffected by CRF in the absence of congestive heart failure and chronic liver disease [31].

Bioavailability is determined by the extent of intestinal absorption and intestinal and hepatic metabolism and transport. For a highly extracted drug with low systemic bioavailability (F), the bioavailability can be estimated by the following equation (Eq. 3) [1,29].

\[
F = \frac{Q_H}{Q_H + \text{CL}_\text{INT} f_u}
\]

Eq(3)
For highly extracted drugs, $\text{CL}_{\text{INT}} \, \text{fu} \gg Q_H$ and $F = \frac{Q_H}{\text{CL}_{\text{INT}} \, \text{fu}}$. Since CRF reduces $\text{CL}_{\text{INT}}$, bioavailability can be significantly increased. Reduced plasma protein binding due to CRF would be expected to increase the fu reducing bioavailability. The relative magnitude of changes in fu and $\text{CL}_{\text{INT}}$ will determine the net effect on F. Oral clearance and apparent oral clearance is influenced by F and defined by the following equation (Eqs 4 and 5) [29].

\[ \text{CL}_\text{oral} = \frac{\text{Dose}}{AUC} \quad \text{Eq(4)} \]

\[ \text{Apparent \, CL}_\text{oral} = \frac{\text{CL}}{F} \quad \text{Eq(5)} \]

The disposition of drugs by the liver can be altered by CRF in both high and low extraction states. For low extraction drugs, reduced $\text{CL}_{\text{INT}}$ leads to an increase in both free and total steady-state drug concentrations. Although the total hepatic clearance of high extraction drugs is blood flow limited, reductions in $\text{CL}_{\text{INT}}$ and/or increases in fu can alter F.

Sun has proposed that the relative contribution of drug transport and metabolism to drug disposition can be predicted based on the physiochemical properties such as the drug solubility and permeability criteria of the Biopharmaceuticals Classification System (BCS) [10,32]. In drugs with high solubility and high permeability, Class I, enzymes effects predominate. In Class II, high permeability low solubility, both enzymes and transporters are important. Class III, high solubility and low permeability, transporter effects predominate.

### 3. Animal Studies: Effect of Renal Failure on CYP Metabolism

Animal models of acute and chronic renal failure since the late 1960s have shown reduced CYP activity with retained susceptibility to induction [33-36]. An early ARF study in rats showed increased bioavailability of propranolol, a highly extracted drug [37]. The same investigators then showed reduced hepatic extraction of propranolol in isolated perfused liver when perfused with uremic serum. Both control and ARF livers showed a 50% reduction in $\text{CL}_{\text{INT}}$ when perfused with uremic serum [38]. Livers from ARF rats showed no reduction hepatic extraction when perfused with normal serum. This suggested that CYP metabolism of propranolol in uremic rats is not downregulated but that a rapidly acting inhibitory factor(s) exists in uremic serum, which directly effects CYP activity.

Later developments in molecular biology allowed the sequencing of the cytochrome P450 (CYP) gene superfamily. Probe drugs have been validated which are substrates of specific CYP isozymes which can be radioactively labeled or unlabeled. The intestinal and hepatic CYP metabolism in data from animal models of CRF are shown in Table 1A and 1B, respectively.

In CRF model in rats using subtotal nephrectomy there was reduced protein expression of CYP2C6, CYP2C11, CYP3A2 and CYP activity using trimethadione as non specific probe [39]. An in vivo study in rats using breath tests with the radiolabeled probe drugs aminopyrene (CYP2C11), erythromycin (CYP3A2), and caffeine (CYP1A2) showed a 35% reduction in CYP2C11 and CYP3A2 activities and no difference in CYP1A2[40]. These results correlated with protein expression. A later study by the same investigators confirmed these findings of reduced mRNA and protein expression and activity with no change in CYP2C6, 2D, 1A2,2E1 [41]. Protein expression of CYP2C11, CYP3A1, and CYP3A2 were reduced by greater than 40% and CYP1A2 remained unchanged as shown in previous studies. The downregulation of CYP3A1 and CYP3A2 was reversed by the inducing agents phenobarbital and dexamethasone. Another study in which various rat models of acute renal failure using 5/6 nephrectomy, bilateral ureteral ligation, intraperitoneal injection of cisplatin, and intramuscular injection of...
glycerol on intestinal and hepatic CYP showed differing effects on CYP metabolism depending on the model [42]. The hepatic CYP3A activity was decreased by 60% in partially nephrectomized and glycerol injected rats. The CYP2C activity approximately was reduced by 50% in cisplatin treated rats. Rats with bilateral ureteral ligation had no reduction CYP activity. Intestinal CYP3A was slightly increased in glycerol induced acute renal failure with no changes in the other models of renal failure. A possible explanation for the varied results from the different models of CRF may be due to differing amount of residual renal function. Another possible mechanism is varying degrees of inflammation and cytokine response in different models of renal failure. Partial nephrectomy and cisplatin models renal failure may have greater cytokine response then bilateral ureteral ligation and therefore more downregulation of CYP.

When normal rat hepatocytes were incubated for 24 h with uremic serum from patients with advanced CRF the protein and mRNA expression of CYP2C6, 2C11, 3A1, and 3A2 were reduced by 35% and the N-Demethylation of erythromycin was also reduced by 35% [43]. Follow-up studies by the same group with human uremic serum demonstrated an inhibition of the mRNA and protein expression of CYP1A2, 2C11, 2D1/2D3, 3A2, 4A1/A3 mRNA in rat hepatocytes by more than 45% [44]. CYP3A and CYP1A activities were reduced by 51% and 59%, respectively [44]. The degree of inhibition was not affected by dialysis but reversed completely after renal transplantation [44]. The degree of inhibition of CYP2D was variable. The serum fraction of 10-15kDa contained the inhibitory activity [44]. Cytokines and PTH are in that molecular weight range and cytokines have been shown to inhibit CYP expression [45, 46]. The serum PTH in these patients correlated (R² = 0.79) with the degree of CYP downregulation in rat hepatocytes incubated with that same serum [44]. Subsequent rat hepatocyte incubation studies have shown that exogenous PTH downregulates CYP3A2 and CYP2C11 protein expression by approximately 60% and this effect of PTH can be reversed by adsorption of serum using anti PTH antibodies [47]. In contrast, parathyroidectomized rats with CRF do not exhibit downregulation of hepatic CYP. In a rat model, sera from CRF rats was shown to specifically inhibit intestinal CYP specifically CYP1A1 and CYP3A2 by 43% and 71%, respectively [48]. In an ARF model of ischemia-reperfusion, renal CYP2D6 and CYP2D9 were downregulated in the kidney [49]. These studies confirm the presence of circulating inhibitory factors in uremic serum which downregulate or directly inhibit hepatic and intestinal CYP activity and identify PTH and cytokines as putative inhibitory factors.

4. Clinical Investigations: Effect of Chronic Renal Failure on Drug Disposition

Several reviews of the effect of chronic renal failure on drug disposition exist in the literature [1-11]. One of the first clinical studies to address the issue of reduced drug metabolism in renal failure used sulfinpyrazone as an intravenous probe drug [50]. Sulfinpyrazone is acetylated, glucuronidated, and metabolized partially by CYP2C9. A summary of clinical data showing significant alterations in nonrenal clearance in CRF for a number of drugs is shown in (Table 2) [7, 51-61]. The majority of these studies involve ESRD patients. These alterations include both substantial reductions in nonrenal clearance ranging from 30 to 90% [1-11].

The effect of CRF manifests as increase in F-for high extraction drugs as shown in Table 3 [7, 62-65]. The propranolol F rises 3-fold as CRF develops but is partially reversed when the patient starts dialysis [65]. The same investigators also found that the propranolol F was greater just prior to dialysis (43%) compared to one day post dialysis (34%). These findings are in agreement with the animal data discussed earlier. Another investigator was not able to reproduce these findings in stable renal failure patients with creatinine clearance of approximately 15 ml/min [66]. These patients may have had enough residual renal function to prevent the inhibition of hepatic metabolism. The systemic clearance of nicardipine is also reduced by 50% and F is increased by 90% with chronic oral dosing in patients with moderate-
to severe chronic kidney disease (average GFR 39 ml/min) not yet on dialysis [67]. These change in the pharmacokinetics of nicardapine are reversed after starting hemodialysis, implicating a dialyzable inhibitory factor [67]. The apparent discrepancy between the results of Michaud [44] and Ahmed [67] in terms of the reversibility of downregulation of CYP3A4 with dialysis may lie in differences in study design. The study by Ahmed is an in vivo clinical pharmacokinetic study of the CYP3A4 substrate nicardapine and Michaud is an ex vivo study in which uremic serum from CRF patients on dialysis incubated with rat hepatocytes expressing the rat homolog CYP3A2 [44,67]. CYP3A2 responses to human uremic serum may differ from that of CYP3A4. There may be some uremic factors which inhibit CYP3A2 but not CYP3A4. Also, there is no information in either paper on what type of dialyzers were used, the blood flow rate, or dialysis duration, all of which could impact on the clearance of uremic inhibitory factors.

Various phenotyping protocols have been used to assess CYP function in CRF. The cumulative recovery of 4-hydroxymephenytoin, a probe of CYP2C19, was reduced by 25% in subjects with $\text{Cl}_{\text{cr}}$ of less than 50ml/min [68]. The formation clearance of 6-hydroxyclozoxazone, a probe for CYP2E1, was not impacted by CRF [68]. The CYP2D6 probe drug debrisoquine recovery ratio was unchanged by CRF. In contrast another study using the CYP2D6 probe drug sparteine showed a 49% reduction in activity in CRF that correlated with the degree of renal insufficiency [69]. Our group showed a 50% reduction in CYP2C9 activity in ESRD using the S/R warfarin plasma ratio as specific probe [70]. S-warfarin is metabolized exclusively by CYP2C9, while R-warfarin is metabolized by multiple pathways. The erythromycin breath test (EBT) was used to measure CYP3A4 activity in ESRD and showed a 28% reduction [71]. Induction of CYP3A4 by rifampin was shown to overcome these effects [71].

The major hepatic conjugation reactions glucuronidation and acetylation are significantly reduced in CRF. The plasma area under the plasma concentration time curve (AUC) of zidovudine which is cleared predominantly by glucuronidation via glucuronosyltransferase (UGT2B7) was doubled in the CRF group [72]. Morphine glucuronidation, also via UGT2B7, is also significantly impaired in CRF [73]. N-acetyl-transferase (NAT-2) has been shown to exhibit genetic polymorphisms with a prevalence of rapid or slow acetylator phenotype of 50% in the African-American and Caucasian populations [74]. There were significant reductions is $\text{CL}_{\text{NR}}$ of isoniazid in CRF patients that were also rapid acetylators. The reduction $\text{CL}_{\text{NR}}$ isoniazid was more pronounced in slow acetylators with CRF. Both of these effects were reversed by transplantation [74,75]. These data suggest that with high extraction drugs that also exhibit polymorphic metabolism, poor metabolizers may be at greater risk for adverse effects of CRF on drug metabolism. These results are consistent with a recent rat study by Simard which showed a reduction in mRNA and protein expression of both Nat1 and Nat2 by 30% and reduction Nat2 activity by 50% [76]. Parathyroidectomy prevented the downregulation Nat expression and inhibition of Nat2 activity [76]. Addition of exogenous PTH to rat hepatocytes reduced Nat2 expression and activity suggesting that PTH may be a circulating inhibitory factor effecting NAT-2 activity in humans.

5. The Effect of Chronic Renal Failure on Drug Transport

Drug transport and metabolism are intimately connected. Drug uptake and efflux from intestinal, renal, and hepatic cells controls the size of the intracellular pool of drug and availability of substrate for CYP and other drug metabolizing enzymes. Drug uptake across the sinusoidal membrane of hepatocytes mediated by transporters such OATP may be rate limiting on hepatic elimination of drugs [77]. The Pgp and the multi-drug resistance associated protein (MRP2) are efflux transporters in the canalicular membrane of hepatocytes and luminal membranes of the renal proximal tubules and intestinal enterocytes and are responsible for extrusion of drugs[10]. Pgp is responsible for the efflux of neutral and cationic drugs many
of which are also substrates of CYP3A4 [10]. In fact the transcription factor PXR acts as xenobiotic sensor and controls the joint upregulation of Pgp and CYP3A4 [78]. MRP-2 transports anionic glucuronide, sulfate, and glutathione-conjugated compounds [10].

The effects of CRF on drug transport have been studied only recently. A more complete review of the effects of CRF and ARF on intestinal and hepatic drug transport can be found in Sun [10] and are shown in Table 1A and 1B, respectively. In one study by Laouari, CRF in 80% nephrectomized rats showed a 70-200% increase in protein expression and mRNA for MRP2 in both kidney and liver with no change in Pgp in either organ [79]. These results are in contrast to study by Naud that showed rats with chronic renal failure developed an increase in protein and mRNA expression of hepatic Pgp of 25% and 40%, respectively [80]. CRF increased hepatic MRP2 mRNA expression by 40% and did not alter hepatic MRP protein expression [80]. Oatp2 showed a decrease in protein expression 35% with no change in mRNA expression of the uptake transporter. Isolated rat hepatocytes were incubated with serum from uremic rats and this reproduced the in vivo protein expression data showing an increase in Pgp, decrease in Oatp2 and no change MRP2. The isolated hepatocyte data differed from the in vivo data in that it did not show change in mRNA levels of any of the transporters [80]. Naud speculated that the degree of CRF in their rat model was greater in their study compared to Laouari and that may have explained the different expression of Pgp and MRP2. Another study in rat model of glycerol-induced, acute renal failure showed a 2.5-fold increase in protein expression of renal Pgp but no change in liver and brain [81]. However, there was a suppression of Pgp activity in all three organs using rhodamine 123 as probe substrate. CRF also reduced the protein expression of rat renal organic cation transporter (OCT2) by 35% and inhibited transport activity, which was reversed by testosterone [82].

There is also data in regard to intestinal transport. In a study by Naud, CRF rats showed reduced intestinal Pgp transport function and protein expression without a change in mRNA[83]. Veau observed a decrease in intestinal Pgp function with no change protein expression or mRNA levels [84]. Rat enterocytes incubated with rat uremic serum reproduced the reduction in MRP2 and Pgp protein expression and reduction Pgp activity, which implicated a role circulating uremic factors [83]. Decrease in protein expression without a change in mRNA suggests that post-translational mechanisms such as protein degradation and not downregulation of transcription are responsible for the loss in drug transporter function. The discrepancies in the findings were attributed to technical differences in isolation of rat enterocytes. Another possible mechanism is circulating uremic factors acting as competitive inhibitors of transport function. These findings could explain the increased bioavailability of drugs in CRF.

Recent studies have shown the effect of known uremic toxins on uptake transport and metabolism of drugs. In the basolateral proximal tubular membrane of the kidney are organic anion transporter (OAT1 and 3) OCT. The sinusoidal membrane of hepatocytes contains the uptake transporter OATP and OCT. The uremic toxin, such as 3-carboxy-4-methyl-5-propyl-2-furan-propanoic acid (CMPF) has been shown to directly inhibit the uptake of erythromycin by Oatp2 in freshly isolated rat hepatocytes [85]. Another uremic toxin, indoxyl sulfate, inhibits the enzymatic breakdown of erythromycin [85]. Use of the oral adsorbent AST-120, which binds uremic toxins such as indoxyl sulfate in rats with CRF showed prevention of downregulation of OAT1 in the kidney[86]. Uptake of renal uremic toxins have been shown to be mediated rOat1/hOAT1 and rOat3/hOAT3 on the basolateral membrane of the proximal tubules [87,88]. Both OAT1 and OAT3 contribute equally to the renal uptake indoxyl sulfate, OAT3 to CMPF uptake, and OAT1 to indolacetate and hippurate uptake [88].
6. Effect of Altered Drug Transport on Interpretation of Pharmacokinetic Data in CRF

There is a complex interaction between drug transport and metabolism and CRF may simultaneously perturb both processes leading to difficulties in interpretation pharmacokinetic data. As shown by Frassetto, the erythromycin breath test may be confounded by CRF induced changes in the drug transporters OATP (Organic Anion Transporter Protein) and Pgp which control the intracellular pool of erythromycin in the hepatocytes [89]. Nolin showed an increase in EBT post hemodialysis compared to predialysis indicating a dialyzable uremic factor may be inhibiting erythromycin metabolism [90]. The confounding role of transport effects on EBT are illustrated in this important clinical study by Nolin [91]. This group showed no change in midazolam IV and oral clearance in ESRD patients using midazolam as specific CYP3A4 probe. An earlier study by Vinik also showed no change unbound clearance of IV midazolam in CRF patients consistent with the later findings [92]. This indicates intestinal and hepatic CYP3A4 are not altered in the well dialyzed ESRD patient [91]. The OATP and Pgp probe drug fexofenadine was given to these same patients and there was 63% reduction in the oral clearance. Since fexofenadine is not a substrate of CYP3A4 and largely excreted unchanged in the bile these results indicate reduced hepatic Pgp and/or OATP activity [91]. This suggests the mechanism for reduced EBT is inhibition of OATP mediated uptake of erythromycin leading to reduced intracellular pool of erythromycin. Theoretically, increased hepatic Pgp mediated efflux of erythromycin would also decrease the intracellular pool of erythromycin and reduce the EBT but that would not be be consistent with the reduced fexofenadine oral clearance observed by Nolin [91]. Another level of complexity is added with intestinal Pgp and OATP which can also effect oral clearance by altering F. Intestinal Pgp mediates the efflux of drugs into the intestinal lumen reducing drug absorption. It is inhibited by CRF which would tend to increase F and therefore decrease the apparent oral clearance (CL/F) of fexofenadine, Eq (5). Intestinal OATP facilitates uptake of drug and inhibition of OATP by CRF would reduce F and therefore increase the apparent oral clearance of fexofenadine, Eq (5).

7. Regulatory Issues

The FDA is currently revising its guidance for pharmacokinetic studies in chronic renal failure. A concept paper is currently available on line at the FDA website at http://www.fda.gov/ohrms/dockets/ac/08/briefing/2008-4351b1-01-FDA.pdf. The impact of the 1998 guidance is reviewed by Huang which showed an increase in the renal impairment studies and dosage adjustment recommendations based on PK changes from 1996-1997 to 2003-2007 [93]. There is a recognition by the FDA of the effect of CRF on drug metabolism and transport. The previous guidance had recommended PK studies in renal patients for drugs with a low therapeutic index (TI) which are cleared predominantly by nonrenal mechanisms. The current concept paper recommends at least one abbreviated PK study in ESRD patients. Given that the current data suggests that circulating uremic toxins which inhibit drug transport and metabolism can be cleared partially by dialysis, advanced chronic kidney disease GFR < 20 not yet on dialysis may be the most seriously effected and should be included in an abbreviated PK study.

8. Conclusions

There is abundant evidence that the nonrenal clearance, protein binding, and volume of distribution of drugs are altered in chronic renal failure, increasing the risk of adverse drug reactions. The majority of the data are in patients with ESRD. Data spanning four decades have shown reduction in nonrenal clearance of drugs extracted predominantly by the liver. In the past 10 years mechanisms for this reduction of nonrenal clearance of drugs have been elucidated. The pharmacologic and physiological effects of CRF on intestinal and hepatic drug
metabolism and transport are summarized in Figure 1. Downregulation of hepatic and intestinal CYP by circulating uremic factors have been shown in animal models particularly CYP2C and CYP3A isozymes. Human studies have shown a reduction in CYP2C9 and CYP2C19 activity using serum warfarin S/R ratio and mephenytoin. CYP3A4 activity was not reduced in ESRD patients using midazolam as a probe substrate but clearance of the transport probe fexofenadine was diminished indicating reduced in OATP mediated hepatic uptake or reduced Pgp efflux. EBT in ESRD is confounded by transport effects probably due to reduced OATP mediated hepatic uptake of erythromycin. More recently animal models of CRF have shown effects on hepatic and intestinal Oatp2, MRP2, and Pgp that can have significant impact on drug disposition and metabolism. The effect of altered transport effects on warfarin metabolism have not been studied. Sun has proposed a scheme for predicting the relative importance of transport effects vs enzyme effects based on the permeability and solubility in the biopharmaceutical drug classification [10].

9. Expert Opinion

It is clear that the effects of CRF on drug metabolism and transport are real and are clinically significant, based on many years of pharmacokinetic studies showing major alterations in nonrenal clearance in patients with CRF. Mechanisms point to circulating inhibitory factors some of which are dialyzable. Patients with advanced CRF Stage IV (GFR 15-29) and Stage V (0-15 ml/min) may behave differently depending whether or not they are on dialysis since these factors may be partially removed by dialysis. Estimating the magnitude of these nonrenal effects is difficult without empirical studies. Caution should be exercised in dosing the drugs in Tables 2 and 3, which show significant reductions in nonrenal clearance and increased bioavailability. Careful titration from the lowest dose should be attempted. Education of practitioners about the existence of this phenomenon is important.

From a regulatory standpoint, these findings would argue for the need of pharmacokinetic studies to evaluate nonrenal clearance in CRF patients with varying degrees of renal impairment particularly when the drug under development is a substrate of CYP2C9, CYP3A4, and NAT-2. Multiple animal as well as human studies have consistently shown that CYP2C9, CYP2C19, CYP3A4 and NAT-2 and the rat homologs are suppressed by CRF and that these effects are clinically significant. Animal studies by Simard showed downregulation of hepatic Nat1 and Nat2 with reduction in Nat2 activity by 50% which appeared to mediated at least partially by PTH [77]. The recent work by Nolin suggests the previous EBT results were confounded by transport effects and that CYP3A4 activity is not reduced in well dialyzed hemodialysis patients when midazolam is used as a probe [86]. Patients with advanced CRF not yet on dialysis may behave differently and studies in this population are needed. There is less animal and human evidence concerning transport by OATP, MRP2, or Pgp but preliminary studies suggest that CRF may perturb the activity of these transporters and significantly alter drug disposition. The FDA concept paper recommends PK studies for all drugs in ESRD. The scheme proposed by Sun [10] based on permeability and solubility may be useful in directing clinical studies based on whether transport or enzyme effects predominate. If the drug under development is BCS Class III where transport effects predominate and the drug is transported by MRP2, Pgp, and OATP then full pharmacokinetic (PK) studies are required in all stages of CKD particularly if the drug has low TI [87]. If the drug is Class I, where enzyme effects predominate, and the drug is a substrate of CYP2C9, CYP2C19, and NAT-2 then full PK studies should be done in all stages of CKD especially stage IV-V and the drug has low TI.
Bibliography


### Abbreviations

<table>
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<tr>
<th>Abbreviation</th>
<th>Description</th>
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<tbody>
<tr>
<td>ARF</td>
<td>Acute Renal Failure</td>
</tr>
<tr>
<td>AUC</td>
<td>area under the concentration time curve</td>
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<tr>
<td>BCS</td>
<td>Biopharmaceutical Classification System</td>
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<tr>
<td>CKD</td>
<td>Chronic Kidney Disease</td>
</tr>
<tr>
<td>CL</td>
<td>clearance</td>
</tr>
<tr>
<td>CL&lt;sub&gt;H&lt;/sub&gt;</td>
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<td>CL&lt;sub&gt;NR&lt;/sub&gt;</td>
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<td>CMPF</td>
<td>3-carboxy-4-methyl-5-propyl-2-furan-propanoic acid</td>
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<td>CRF</td>
<td>Chronic Renal Failure</td>
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<td>CYP</td>
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<td>E</td>
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<td>ESRD</td>
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*Expert Opin Drug Metab Toxicol. Author manuscript; available in PMC 2009 September 16.*
end stage renal disease

F
bioavailability

Fu
unbound fraction

MRP-2
multi-drug resistance protein-2

NAT-2
N-acetyltransferase 2

OAT
organic anion transporter

OATP
organic anion transporting polypeptide

OCT
organic cation transporter

Pgp
P-glycoprotein

PK
pharmacokinetics

PTH
parathyroid hormone

Q
blood flow rate

QH
hepatic blood flow

TI
therapeutic index

TPMT
thiopurine methyl transferase

V_D
volume of distribution
Figure 1.
Summary of Physiological and Pharmacologic Effects of CRF on Drug Metabolism and Transport.
### TABLE 1A Animal Studies of CRF and Intestinal Metabolism and Transport

<table>
<thead>
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<th>INTESTINE</th>
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<tr>
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<td>↓</td>
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<tr>
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<td>↔ Naud [83]</td>
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<td>CYP3A2</td>
<td>↓70% Leblond [48]</td>
<td>↓25% Leblond [48]</td>
<td>↑</td>
</tr>
</tbody>
</table>

### TABLE 1B Animal Studies of CRF and Hepatic CYP Metabolism and Transport

<table>
<thead>
<tr>
<th>LIVER</th>
<th>Protein</th>
<th>Activity</th>
<th>Effect</th>
</tr>
</thead>
<tbody>
<tr>
<td>TRANSPORTERS</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Pgp</td>
<td>↑25% Naud [80]</td>
<td>↑45% [80]</td>
<td>↑ Biliary Excretion</td>
</tr>
<tr>
<td>ABCB1</td>
<td>↔ Laouari [79]</td>
<td></td>
<td>?</td>
</tr>
<tr>
<td>MRP2</td>
<td>↔ Naud [80]</td>
<td>NA</td>
<td>NA</td>
</tr>
<tr>
<td>ABCC2</td>
<td>↑ 70-200% Laouari [79]</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Oatp2</td>
<td>↓40% Naud [80]</td>
<td>NA</td>
<td>↓ Biliary Excretion</td>
</tr>
<tr>
<td>SLCO</td>
<td></td>
<td></td>
<td>↓ metabolic CL</td>
</tr>
<tr>
<td>ENZYMES</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CYP1A1</td>
<td>↔</td>
<td>↔ Leblond [40,41]</td>
<td>↔ Hepatic CL</td>
</tr>
<tr>
<td>CYP2C11</td>
<td>↓40-45% Leblond [40]</td>
<td>↓35% Leblond [40,41]</td>
<td>↓ CL</td>
</tr>
<tr>
<td>Uchida [39]</td>
<td>Uchida [39]</td>
<td></td>
<td></td>
</tr>
<tr>
<td>CYP2D</td>
<td>↔ Leblond [41]</td>
<td>↔ Leblond [41]</td>
<td>↔ CL</td>
</tr>
<tr>
<td>INTESSTINE</td>
<td>Protein</td>
<td>Activity</td>
<td>Effect on F</td>
</tr>
<tr>
<td>------------</td>
<td>---------</td>
<td>----------</td>
<td>-------------</td>
</tr>
<tr>
<td>CYP3A1</td>
<td>↓75-85% Leblond [40,41]</td>
<td>↓35% Leblond [40,41]</td>
<td>↓ CL</td>
</tr>
<tr>
<td>CYP3A2</td>
<td>↓45-65% Leblond [40,41] Uchida [39]</td>
<td>↓35-50% Leblond [40,41]</td>
<td>↓ CL</td>
</tr>
<tr>
<td>Nat 1</td>
<td>↓33% Simard [77]</td>
<td>↓45% Simard [77]</td>
<td>↓ CL</td>
</tr>
<tr>
<td>Nat 2</td>
<td>↓50% Simard [77]</td>
<td>↓50% Simard [77]</td>
<td>↓ CL</td>
</tr>
</tbody>
</table>

? = limited data
TABLE 2
Effect of CRF on Nonrenal Clearance in Human Subjects

<table>
<thead>
<tr>
<th>Drug</th>
<th>% Change CLnr</th>
<th>Enzyme</th>
<th>Metabolism</th>
</tr>
</thead>
<tbody>
<tr>
<td>Captopril</td>
<td>-50</td>
<td>TPMT *</td>
<td>sulfoxidation</td>
</tr>
<tr>
<td>Morphine</td>
<td>-40</td>
<td>UGT2B7</td>
<td>glucuronidation</td>
</tr>
<tr>
<td>Procainamide</td>
<td>-60</td>
<td>NAT-2</td>
<td>acetylation</td>
</tr>
<tr>
<td>Imipenem</td>
<td>-58</td>
<td>dehydropeptidase</td>
<td></td>
</tr>
<tr>
<td>Nimodipine</td>
<td>-87</td>
<td>CYP3A4</td>
<td>dealkylation</td>
</tr>
<tr>
<td>Verapamil</td>
<td>-54</td>
<td>CYP3A4</td>
<td>demethylation</td>
</tr>
<tr>
<td>Metoclopramide</td>
<td>-66</td>
<td>CYP2D6</td>
<td>dealkylation, sulfation</td>
</tr>
<tr>
<td>Desmethyldiazepam</td>
<td>-63</td>
<td>CYP2C9</td>
<td>hydroxylation</td>
</tr>
<tr>
<td>Warfarin</td>
<td>-50</td>
<td>CYP2C9</td>
<td>hydroxylation</td>
</tr>
</tbody>
</table>

TPMT thiopurine methyl transferase
<table>
<thead>
<tr>
<th>Drug</th>
<th>Effect</th>
<th>Enzyme</th>
</tr>
</thead>
<tbody>
<tr>
<td>Propranolol</td>
<td>+300%</td>
<td>CYP2D6</td>
</tr>
<tr>
<td>Erythromycin</td>
<td>+100%</td>
<td>CYP3A4</td>
</tr>
<tr>
<td>Propoxyphene</td>
<td>+100%</td>
<td>CYP3A4</td>
</tr>
<tr>
<td>Dihydrocodeine</td>
<td>+70%</td>
<td>CYP2D6</td>
</tr>
<tr>
<td>Oxprenolol</td>
<td>+100%</td>
<td>CYP2D6</td>
</tr>
</tbody>
</table>

**TABLE 3**
Effect of CRF on Bioavailability in Human Subjects