ISOLATION AND IDENTIFICATION OF MAJOR URINARY METABOLITES OF RIFABUTIN IN RATS AND HUMANS

ILYA UTKitN, TATIANA KOUItAKoVA, TONI THOMPSON, CHARLES COTTRELL, EUGENIA IAItSIMItRKAIA, JOHN BARRY, PAUL VOItROS, AND NICHOLAs GERBER

Department of Pharmacology, College of Medicine (U.T., T.K., E.I., N.G.) and Campus Chemical Instrument Center (C.C.), The Ohio State University; and Department of Chemistry (T.T., J.B., P.V.), Northeastern University

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ABSTRACT:

The antimycobacterial drug rifabutin is extensively metabolized in humans and laboratory animals. About 40% of the dose is excreted in urine as unchanged drug, and lipophilic (extractable with 1-chlorobutane) and polar metabolites. Polar metabolites accounted for 59.1 ± 2.5% and 88.8 ± 4.4% of radioactivity in urine collected over 96 hr after intravenous administration of 25 and 1 mg/kg of [14C]-rifabutin to Sprague-Dawley rats, respectively. After 48 hr, all urinary radioactivity consisted of polar metabolites. The most abundant polar metabolite, identified by electrospray ionization-MS, collision-induced dissociation-MS, and comparison of HPLC retention times with the synthetic standard, was N-isobutyl-4-hydroxy-piperidine. Lipophilic metabolites accounted for <20% of urinary radioactivity. Major lipophilic metabolites, 25-O-deacetyl-rifabutin, 27-O-demethyl-rifabutin, 31-hydroxy-rifabutin, 32-hydroxy-rifabutin, and 20-hydroxy-rifabutin were isolated from both human and rat urine by HPLC and identified by electrospray ionization-MS, collision-induced dissociation-MS, and NMR spectrometry. In addition, two metabolites formed by the oxidation of the N-isobutyl-piperidyl group of rifabutin were found in the urine of rats, but not humans.

Materials and Methods

Chemicals. Rifabutin, 25-O-deacetyl-rifabutin, and two analogs of rifabutin bearing the spiro-piperidyl group N-substituted with an isopropyl or heptyl chain were provided by Pharmacia (Dublin, OH). Rifampicin, rifampycin SV, and rifamycin B were purchased from Sigma Chemical Co. (St. Louis, MO). Starting material for the synthesis of [14C]-rifabutin, 3-amino-4-deoxy-4-amino-rifamycin S, was a gift from Dr. Bruno Sardi (Pharmacia, Milan, Italy). Radiolabeled rifabutin (59.3 mCi/mmol) was synthesized at New England Nuclear (Boston, MA) as described by Fontana et al. (16) and Marsili et al. (1). The purity of rifabutin and [14C]-rifabutin was >98% and >97%, respectively. N-isobutyl-4-hydroxy-piperidine was obtained by reduction of N-isobutyl-4-piperidone with sodium borohydride. N-isobutyl-4-piperidone was produced by acid hydrolysis of rifabutin at 60°C for 3 hr as described elsewhere (14).

Acetonitrile (HPLC grade), methanol, 1-chlorobutane, α-phosphoric acid, and liquid scintillation cocktail ScintiSafe Plus 50% were purchased from Fisher Scientific (Fair Lawn, NJ). Dowex 1, Dowex-50W, MRFA and apomyoglobin were purchased from Sigma. Charcoal was from MCB Manufacturing Chemists (Cincinnati, OH). Other chemicals were received from Sigma or J. T. Baker, Inc. (Phillipsburg, NJ).
HPLC Analysis. All samples were analyzed on a Hewlett-Packard HPLC system (models 1050 or 1090 M, Palo Alto, CA) equipped with a UV diode-array detector. Radioactivity was measured on-line with a Radiomatic A-500 series Flow Scintillation Analyzer (Packard, Meriden, CT) with a solid scintillation flow cell. Separation was achieved using an analytical Hypersil C18 column (250 × 4.6 mm, 5 µm; Phenomenex, Torrance, CA) at a mobile phase flow rate of 1.5 ml/min. The mobile phases A, B, C, and D contained acetonitrile and/or one of the following aqueous solutions: 3% TEA and 0.3% trifluoroacetic acid titrated with phosphoric acid to pH 2.0 (A), 0.1% TEA and phosphoric acid (pH 6.8–7.0) (B), 0.01% TEA and phosphoric acid (pH 7.0) (C), and 1.5% TEA and phosphoric acid (pH 6.0) (D).

Animals. Male Sprague-Dawley rats (Charles River Laboratories, North Wilmington, MA) weighing between 223 and 463 g, acclimated in a 12-hr light/dark cycle constant-temperature (24°C) vivarium with free access to standard chow and water for 1 week, were prepared with the iv catheters as described previously (15).

Dosing and Sample Collection. For metabolic profiles, [14C]rifabutin was administered to rats (three in each group) iv as a single dose of 25 mg/kg (specific activity: 2.7 mCi/mmol) or 1 mg/kg (specific activity: 59.3 mCi/mmol). Dosing solutions of [14C]rifabutin (10 and 0.4 mg/ml) were prepared in saline containing 10 mM HCl. Urine and feces were collected in containers immersed in ice before dosing and at 0–24, 24–48, 48–72, and 72–96 hr, and stored at −80°C. For isolation and structural analysis of metabolites, urine was collected from 20 rats over 0–24 hr after an iv dose of 25 mg/kg of rifabutin. Analysis of Urine. Before the analysis, urine was thawed, filtered through a 0.2-µm filter, and injected (200 µl) directly into the HPLC. Rifabutin and metabolites were separated as described previously (15) using a linear gradient of 27–42% acetonitrile in aqueous solution A over 45 min (fig. 1A). Separation of individual polar metabolites designated as PM1 and PM2 from the peak labeled P in fig. 1 was achieved by using 100% aqueous solution B as a mobile phase (fig. 2). Total radioactivity in urine was measured in liquid scintillation counting (100 µCi) using scintillation cocktail ScintiSafe Plus 50% (5 ml).

Isolation and Properties of Polar Metabolites. Chemical stability of polar metabolites was studied by incubating urine with equal volumes of 1 N HCl or 1 N KOH at 37°C for 22 hr, followed by HPLC analysis using 100% aqueous solution B as a mobile phase. PM2 was stable under these conditions while <25% of PM1 disappeared.

The following procedure was used to isolate the major polar metabolite PM2 for structural analysis. The urine (320 ml) was gently shaken twice in a separatory flask with 320 ml of 1-chlorobutane in the presence of 320 µl of isomyl alcohol. After separation of phases, the organic layer containing rifabutin and lipophilic metabolites was removed and dried under nitrogen. The aqueous phase containing polar metabolites was mixed with 5 volumes of methanol and kept overnight at 4°C to precipitate salts. After centrifugation (1200g for 10 min), the supernatant (1.8 liters) was filtered through a 0.2-µm filter and percolated through a layer of Dowex-50W (100 g) equilibrated at pH 2.2. The resin was washed with 300 ml of acidified distilled water (pH 2.2), and the metabolites were eluted with 2 liters of 2 N KOH. Activated charcoal (60 g) was shaken with the eluate over 10 min, separated by centrifugation (1200g for 10 min), and washed twice with 600 ml of 0.1 N KOH. The metabolites were eluted from the charcoal with 600 ml of a mixture consisting of methanol and trifluoroacetic acid (300:1, v/v) by shaking for 10 min, and then separated by centrifugation (1200g for 10 min). The supernatant was filtered through a 0.2-µm filter and concentrated to a volume of ~3 ml using a vacuum evaporator. The recovery of polar metabolites after precipitation of salts with methanol and solid-phase extraction with Dowex-50W, and charcoal was estimated to be ~20%. These steps permitted removal of contaminants that would have interfered with further purification by HPLC. The polar metabolites had insignificant absorbance at 210 nm, indicating the absence of chromophoric groups. For further purification of PM2 by HPLC, 150 µl of the concentrate was supplemented with a partially purified radioactive tracer [14C]PM2 (~16 nCi) obtained by a similar procedure (precipitation of salts with methanol and solid-phase extraction) from urine of a rat given [14C]rifabutin. Portions of the mixture were injected repeatedly into the HPLC and components separated using aqueous mobile phase B. Peaks of PM2 (RT ~15.1 min) were collected, combined, and dried under nitrogen. The residue was resuspended in 220 µl of mobile phase, of which portions were repeatedly injected into the HPLC to purify PM2 using a linear gradient of 0–80% acetonitrile in aqueous solution C. Fractions containing PM2 (RT ~12.2 min) were combined and dried under nitrogen.

Isolation of Lipophilic Metabolites. The 1-chlorobutanol extract of rat urine was dried, the residue dissolved in mobile phase (40% acetonitrile and 60% aqueous solution A), and portions repeatedly injected into the HPLC. Metabolites were eluted in a linear gradient of 27–42% acetonitrile in aqueous solution A over 45 min. P, peak of polar metabolites; R, rifabutin; 1, U1 (25-O-deacetyl-rifabutin); 2, U2 (31-hydroxy-rifabutin + 32-hydroxy-rifabutin); 3, U3 (27-O-demethyl-rifabutin); 4, U4 (20-hydroxy-rifabutin); 5, U5 (a mixture of two metabolites of rifabutin resulting from hydroxylation in the N-isobutyl-piperidyl group).
A dose of 25 mg/kg was injected (90 mC). The CID spectrum was based on comparison of CID spectra of rifabutin and 25-O-dideacetyl-rifabutin. The assignment of diagnostic fragments was based on comparison of CID spectra of rifabutin and 25-O-dideacetyl-rifabutin, rifampin, rifamycin SV, rifamycin B, and two analogs of rifabutin bearing the spiro-piperidyl group. The structures of rifabutin and lipophilic metabolites were determined using one-dimensional $^1$H NMR and/or two-dimensional $^1$H-$^1$H COSY and inverse $^1$H-$^1$C HMQC.

**Results**

**Rat Urinary Metabolites.** Polar Metabolites. After administration of 25 and 1 mg/kg of [14C]rifabutin as an iv bolus, 40.1 ± 4.3% and 34.1 ± 1.8% of the dose, respectively, were excreted in urine during 96 hr (15). Polar metabolites accounted for 59.1 ± 2.5% and 88.8 ± 4.4% of the urinary radioactivity after the 25 and 1 mg/kg dose, respectively. The proportion of urinary polar metabolites was higher during 24–48 hr than 0–24 hr (fig. 1), and by 48–72 hr all radioactivity in urine consisted of polar metabolites.

ESI-MS of the major polar metabolite PM2 showed [M + H]$^+$ at m/z 158. The CID spectrum of the m/z 158 ion is consistent with an assignment of a structure for the PM2 metabolite corresponding to hydroxylation of the piperidine ring, possibly forming N-isobutyl-4-hydroxy-piperidine (fig. 3). The HPLC RT of PM2 matched that of synthetic N-isobutyl-4-hydroxy-piperidine that confirmed this structural assignment.

**Lipophilic Metabolites.** Radiochemical and spectrophotometric detection (275 nm) of rifabutin and lipophilic metabolites yielded similar HPLC profiles, suggesting that rifabutin and lipophilic metabolites had similar extinction coefficients. ESI-MS of rifabutin and the metabolites U1, U2A, U2B, U3, U4, U5A, and U5B produced a protonated molecular ion [M + H]$^+$ as well as sodium [M + Na]$^+$ and potassium [M + K]$^+$ adducts. The fragmentation of rifabutin
could be described in terms of bond cleavage that yielded ions of the type A–G as indicated in fig. 4. Shift of the mass of any of these structurally diagnostic ions (table 1) in combination with NMR data (tables 2 and 3) provided a reasonable rationale for assigning the structure of new metabolites.

**Metabolite U1.** The $^1$H NMR spectrum of U1 showed the absence of the $\text{H}_{36}$ signal (s at 2.01 ppm in the spectrum of parent drug), thus indicating the loss of the acetyl group. The HPLC RT, ESI and CID mass spectra of this metabolite matched those of the authentic standard of $\text{O}$-deacetyl-rifabutin. Thus, metabolite U1 was identified as 27-$\text{O}$-deacetyl-rifabutin.

**Metabolite U3.** ESI-MS of metabolite U3 showed $[\text{M} + \text{H}]^+$ at $m/z$ 833, indicating removal of a methyl group. Otherwise, diagnostic daughter ions in the CID spectrum of U3 matched those of rifabutin (fig. 5). Furthermore, the signal of proton $\text{H}_{37}$ observed in the $^1$H NMR spectrum of rifabutin (s at 3.06 ppm) was missing in the spectrum of U3. On this basis, U3 was identified as 27-$\text{O}$-demethyl-rifabutin.

ESI-MS of U2A, U2B, U4, U5A, and U5B showed a $[\text{M} + \text{H}]^+$ at $m/z$ 863, indicating the presence of an additional oxygen atom in these metabolites compared with rifabutin.

**Metabolites U2A and U2B.** The daughter ions of U2A and U2B at $m/z$ 112 and 423 corresponding to the unchanged fragments G and F (table 1, fig. 4), respectively, suggest no modification on that part of the molecule for either metabolite. For U2A, the diagnostic fragments at $m/z$ 561 (545 + 16), 589 (573 + 16), and 648 (632 + 16) pointed to carbons $\text{C}_{31}$, $\text{C}_{20}$, $\text{C}_{19}$, $\text{C}_{18}$, $\text{C}_{17}$, $\text{C}_{16}$, and $\text{C}_{30}$ as possible oxidation sites (table 1, fig. 4). In an HMRC experiment, the disappearance of the $\text{C}_{31}$ signal at 17.4 ppm and correlated $\text{H}_{31}$ signal at 0.85 ppm were accompanied by the appearance of a carbon signal at 63 ppm correlated with a proton signal at 3.54–3.60 ppm (tables 2 and 3).

Daughter ions of U2B at $m/z$ 545, 573, and 648 (632 + 16) implicated atoms $\text{C}_{32}$, $\text{C}_{32}$, and $\text{C}_{33}$ as possible oxidation sites (table 1, fig. 4). The signal corresponding to $\text{H}_{32}$ of parent drug (d at 1.04 ppm) disappeared, and a proton signal (3.80 and 3.98 ppm) appeared that correlated with $\text{H}_{22}$ at 1.78 ppm in $^1$H-$^1$H COSY and with $\text{C}_{32}$ at 61.9 ppm in $^1$H-$^1$C HMRC experiments (tables 2 and 3). The downfield shifts and splitting of $\text{H}_{31}$ and $\text{H}_{32}$ signals (table 2) in $^1$H NMR spectra of U2A and U2B, respectively, suggest hydroxylation of the corresponding methyl groups. Hydroxylation caused the methylene protons to become nonequivalent (appearing as an AB of ABX pattern), because they are adjacent to a chiral center. Consequently, splitting of the proton signals occurred. Downfield shifts of the corresponding $\text{C}_{31}$ and $\text{C}_{32}$ signals (table 3) confirm that hydroxylation occurred in the aforementioned positions. Thus, the metabolites U2A and U2B are 31-hydroxy-rifabutin and 32-hydroxy-rifabutin, respectively.

**Metabolite U4.** Diagnostic fragments observed in the CID spectrum of U4 at $m/z$ 423 and $m/z$ 561 (545 + 16) indicated that an oxidation site was located in the ansa chain of rifabutin between carbons $\text{C}_{15}$ and $\text{C}_{31}$ (table 1, fig. 4). In the $^1$H NMR spectrum of U4, the $\text{H}_{20}$ signal disappeared; the signals of protons $\text{H}_{19}$ and $\text{H}_{31}$ became a doublet and a singlet, respectively; and there were no changes in the chemical shifts and multiplicities of $\text{H}_{17}$ and $\text{H}_{18}$ (table 2). In an $^1$H-$^1$H COSY experiment, signals of coupling interactions between protons $\text{H}_{20}$ and $\text{H}_{19}$, $\text{H}_{20}$ and $\text{H}_{18}$, and $\text{H}_{20}$ and $\text{H}_{17}$ disappeared (fig. 6). Also, no signal was observed for $\text{C}_{30}$ in an HMRC experiment, confirming that there is no attached proton, and $\text{C}_{31}$ shifted downfield.

![Chemical structure, labeling position (*), and diagnostic fragments produced by CID-MS of rifabutin.](image)

**Fig. 4.**

### TABLE 1

<table>
<thead>
<tr>
<th>Compound</th>
<th>$[\text{M} + \text{H}]^+$</th>
<th>A</th>
<th>B</th>
<th>C</th>
<th>D</th>
<th>E</th>
<th>F</th>
</tr>
</thead>
<tbody>
<tr>
<td>Rifabutin</td>
<td>847</td>
<td>815 (100)</td>
<td>755 (17)</td>
<td>632 (0.8)</td>
<td>573 (0.4)</td>
<td>545 (0.2)</td>
<td>423 (1.5)</td>
</tr>
<tr>
<td>U1</td>
<td>805</td>
<td>773 (100)</td>
<td>755 (1.4)</td>
<td>632 (0.06)</td>
<td>573 (0.02)</td>
<td>545 (0.01)</td>
<td>423 (0.6)</td>
</tr>
<tr>
<td>U2A</td>
<td>863</td>
<td>831 (100)</td>
<td>771 (16)</td>
<td>648 (1.2)</td>
<td>589 (0.2)</td>
<td>561 (0.06)</td>
<td>423 (1.0)</td>
</tr>
<tr>
<td>U2B</td>
<td>863</td>
<td>831 (40)</td>
<td>771 (7.9)</td>
<td>648 (0.3)</td>
<td>573 (2.2)</td>
<td>545 (0.9)</td>
<td>423 (13)</td>
</tr>
<tr>
<td>U3</td>
<td>833</td>
<td>815 (100)</td>
<td>755 (19)</td>
<td>632 (4.4)</td>
<td>573 (2.4)</td>
<td>545 (6.5)</td>
<td>423 (7.2)</td>
</tr>
<tr>
<td>U4</td>
<td>863</td>
<td>831 (23)</td>
<td>ND*</td>
<td>ND</td>
<td>561 (0.2)</td>
<td>423 (0.7)</td>
<td>112 (8.6)</td>
</tr>
<tr>
<td>U5A</td>
<td>863</td>
<td>831 (100)</td>
<td>771 (15)</td>
<td>648 (1.4)</td>
<td>589 (0.4)</td>
<td>561 (0.6)</td>
<td>439 (0.4)</td>
</tr>
<tr>
<td>U5B</td>
<td>863</td>
<td>831 (100)</td>
<td>771 (21)</td>
<td>648 (1.8)</td>
<td>589 (0.7)</td>
<td>NA*</td>
<td>439 (1.3)</td>
</tr>
</tbody>
</table>

CID spectra of U1 and U3 were obtained using argon with a collision energy of $-81$ and $-61$ eV, respectively. For the other metabolites, collision energy was set at $-51$ eV.

*ND, not detected.

*NA, no data available.*
to 30 ppm (table 3). Thus, it is concluded that hydroxylation occurred at carbon C\text{20}, and metabolite U\text{4} is assigned the structure of 20-hydroxy-rifabutin.

Metabolites U\text{5A} and U\text{5B}. There was a 16 mass units increase in the mass of all characteristic fragments observed in the CID spectra of U\text{5A} and U\text{5B}, compared with the parent drug (table 1), thus indicating that these metabolites contained an additional oxygen atom in the N\text{-isobutyl-piperidyl moiety (fig. 4). The lack of authentic standards and difficulties in interpretation of the CID and NMR spectra did not permit a more complete structural characterization of these metabolites.

Human Urinary Metabolites. Figure 7\text{A} shows a typical HPLC profile of lipophilic metabolites of rifabutin extracted with 1-chlorobutane from 0- to 48-hr human urine. Individual peaks were collected and rechromatographed in a gradient of 30 –50% acetonitrile in aqueous solution D (see Materials and Methods) for 1 hr. This demonstrated that HU\text{1}, HU\text{3}, and HU\text{4} consisted of one major metabolite each, whereas HU\text{2} was composed of two compounds designated as HU\text{2A} and HU\text{2B}. Chemical identity of the metabolites HU\text{1}, HU\text{2A}, HU\text{2B}, HU\text{3}, and HU\text{4} with the corresponding rat metabolites U\text{1}, U\text{2A}, U\text{2B}, U\text{3}, and U\text{4} was established by comparison of HPLC RTs, NMR, and/or mass spectra. Thus, the major lipophilic metabolites were identified as 25-O-deacetyl-rifabutin, 31-hydroxy-rifabutin, 32-hydroxy-rifabutin, 27-O-demethyl-rifabutin, and 20-hydroxy-rifabutin.

ESI-MS of metabolite HU\text{0} showed a \([M + H]^{+}\) at \(m/\text{z} 821\) suggesting the loss of the acetyl group and the presence of an additional oxygen atom. The loss of the acetyl group was confirmed by the absence of the H\text{36} signal in the \(^{1}H\) NMR spectrum of HU\text{0}. The unchanged fragment F at \(m/\text{z} 423\) in the CID spectrum of HU\text{0} indicated that oxidation occurred in the ansa chain. However, the amount of HU\text{0} was insufficient for complete characterization by NMR and MS.

Discussion

Our results showed that radiolabeled polar metabolites PM\text{1} and PM\text{2} (figs. 1 and 2) accounted for most of the radioactivity excreted in rat urine after iv administration of [\(^{14}\text{C}\)]rifabutin. The retention of radiolabel indicated that these polar metabolites were derived from the N-isobutyl-piperidyl group of rifabutin (fig. 4) and, possibly, its lipophilic metabolites. PM\text{2} was identified as N-isobutyl-4-hydroxy-piperidine. Similarity of physiochemical properties of PM\text{1} and PM\text{2} (chemical stability, binding to a strong cation exchanger and activated charcoal, and the absence of chromophoric groups) suggested that the more polar metabolite PM\text{1} is a hydroxy derivative of PM\text{2}. The

<table>
<thead>
<tr>
<th>Atom</th>
<th>Rifabutin</th>
<th>U\text{2A}</th>
<th>U\text{2B}</th>
<th>U\text{4}</th>
</tr>
</thead>
<tbody>
<tr>
<td>H\text{17}</td>
<td>6.25 d ((J = 10.3))</td>
<td>6.30 d ((J = 10.2))</td>
<td>6.28 d ((J = 9.9))</td>
<td>6.22 d ((J = 10.4))</td>
</tr>
<tr>
<td>H\text{18}</td>
<td>6.38 dd ((J = 10.3, 15.8))</td>
<td>6.58 dd ((J = 10.2, 15.6))</td>
<td>6.35 dd ((J = 9.9, 15.5))</td>
<td>6.38 dd ((J = 10.4, 15.6))</td>
</tr>
<tr>
<td>H\text{19}</td>
<td>6.02 dd ((J = 15.8, 6.7))</td>
<td>5.98 dd ((J = 15.6, 6.6))</td>
<td>6.15 dd ((J = 15.5, 6.8))</td>
<td>5.99 d ((J = 15.6))</td>
</tr>
<tr>
<td>H\text{20}</td>
<td>2.37 m</td>
<td>2.50 m</td>
<td>2.44 m</td>
<td>ND\text{a}</td>
</tr>
<tr>
<td>H\text{21}</td>
<td>3.65 m</td>
<td>4.07 m</td>
<td>3.75 m</td>
<td>3.83 m</td>
</tr>
<tr>
<td>H\text{22}</td>
<td>1.74 m</td>
<td>1.90 m</td>
<td>1.78 m</td>
<td>1.90 m</td>
</tr>
<tr>
<td>H\text{23}</td>
<td>2.04 s</td>
<td>2.09 s</td>
<td>2.07 s</td>
<td>2.09 s</td>
</tr>
<tr>
<td>H\text{24}\text{a}</td>
<td>0.85 d ((J = 6.9))</td>
<td>3.60, 3.54 AB of ABX</td>
<td>0.96 d ((J = 6.6))</td>
<td>1.26 s</td>
</tr>
<tr>
<td>H\text{24}\text{b}</td>
<td>1.04 d ((J = 6.9))</td>
<td>1.06 d ((J = 6.6))</td>
<td>3.98, 3.80 AB of ABX</td>
<td>1.15 d ((J = 6.6))</td>
</tr>
</tbody>
</table>

\(\text{dd, doublet of doublets; } J\) coupling constant.

\(\text{a ND, not detected because H}_{36}\) has been replaced by a hydroxy group.

\(\text{b These hydroxymethylene protons have become nonequivalent because they are adjacent to a chiral center. } J_{\text{AB}} = -11\text{ Hz for both U2A and U2B.}\)
metabolite PM1 could not be identified conclusively because of the lack of essential synthetic standards.

The proportion of different urinary metabolites of rifabutin in the rat was dose-dependent. The fraction of the dose excreted in urine as polar metabolites was lower at a higher (25 mg/kg) dose than at a lower (1 mg/kg) dose apparently due to saturation of enzymatic and/or transport systems involved in the formation and excretion of these metabolites. This is in agreement with our previous findings (15) that first-pass metabolism of rifabutin was more significant at the lower dose (bioavailability after po administration of 25 mg/kg and 1 mg/kg of rifabutin was >90% and 44%, respectively).

The proportion of polar metabolites was higher in urine collected at later times; unchanged drug and lipophilic metabolites were not detected in 48- to 96-hr urine. Large amounts of polar metabolites in urine and an increase in their percentage at later times were observed previously after po administration of [14C]rifabutin to humans and rats (11) and after iv administration to rats (12). The predominant urinary excretion of radioactivity as polar metabolites indicates that the cleavage of the N-isobutyl-piperidyl group plays a major role in renal elimination of rifabutin after iv and po administration of the drug.

The mechanism of PM2 formation in vivo is unclear. We hypothesize that it is a two-step process that includes oxidative N-dealkylation of rifabutin by cytochrome P450 to form N-isobutyl-4-piperidone and subsequent reduction of N-isobutyl-4-piperidone to N-isobutyl-4-hydroxy-piperidine by alcohol dehydrogenase or carbonyl reductase.

It has been shown that polar metabolite(s) of rifabutin are present in human urine (11) and also in incubations of the drug with human liver and enterocyte microsomes (17). However, the identity of these metabolites remains to be established conclusively.

In previous work (11), two rat urinary metabolites were identified as 25-O-deacetyl-rifabutin and 31-hydroxy-rifabutin by comparison of their HPLC RTs with those of authentic standards. In addition to these metabolites, we found in rat urine 32-hydroxy-rifabutin, 27-O-demethyl-rifabutin, 20-hydroxy-rifabutin, and two products that resulted from oxidation of the N-isobutyl-piperidyl group.

Five rifabutin metabolites were isolated from human urine and identified in this study as 25-O-deacetyl-rifabutin, 31-hydroxy-rifabutin, 32-hydroxy-rifabutin, 20-hydroxy-rifabutin, and 27-O-demethyl-rifabutin. The first three have been described previously (13, 14),
Materials and Methods

Aqueous phase A over 45 min as described in 1-chlorobutane and separated by HPLC in a gradient of 27–42% acetonitrile in aqueous phase B over 45 min as described in Materials and Methods.

Urinary metabolites of rifabutin and unchanged drug were extracted with 1-chlorobutane and separated by HPLC in a gradient of 27–42% acetonitrile in aqueous phase A over 45 min as described in Materials and Methods. About 5% of the dose is excreted in urine as rifabutin (11).

Values are means ± SD for 10 subjects.

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Urinary metabolites of rifabutin and unchanged drug were extracted with 1-chlorobutane and separated by HPLC in a gradient of 27–42% acetonitrile in aqueous phase A over 45 min as described in Materials and Methods. Previously reported 30-hydroxy-rifabutin and 25-hydroxy-rifabutin were not found in this study. Although several common major lipophilic metabolites of rifabutin have been identified in human and rat urine, there is a difference in the metabolism of the drug in these species. The metabolites USA and USB excreted in rat urine were not found in significant amounts in human urine.

whereas 20-hydroxy-rifabutin and 27-O-demethyl-rifabutin are newly identified. 20-Hydroxy-rifabutin and 25-O-deacetyl-rifabutin seem to be the major urinary lipophilic metabolites of rifabutin in humans (fig. 7, table 4). Previous reported 30-hydroxy-rifabutin and 25-O-deacetyl-rifabutin-N-oxide (13, 14) were not found in this study, possibly because of their low quantities in human urine. Although several common major lipophilic metabolites of rifabutin have been identified in human and rat urine, there is a difference in the metabolism of the drug in these species. The metabolites USA and USB excreted in rat urine were not found in significant amounts in human urine.

MAJOR URINARY METABOLITES OF RIFABUTIN

TABLE 4

Relative amounts* of rifabutin metabolites as a % of unchanged drug in human urine (0–48 hr) after a single dose of 300 mg of rifabutin

<table>
<thead>
<tr>
<th>Rifabutin</th>
<th>HU1</th>
<th>HU2</th>
<th>HU3</th>
<th>HU4</th>
</tr>
</thead>
<tbody>
<tr>
<td>100 ± 29</td>
<td>29 ± 16</td>
<td>9.4 ± 3.3</td>
<td>6.0 ± 2.0</td>
<td>36 ± 10</td>
</tr>
</tbody>
</table>

Urinary metabolites of rifabutin and unchanged drug were extracted with 1-chlorobutane and separated by HPLC in a gradient of 27–42% acetonitrile in aqueous phase A over 45 min as described in Materials and Methods.

* Values are means ± SD for 10 subjects.

** About 5% of the dose is excreted in urine as rifabutin (11).

References


