Investigation of Clinical Absorption, Distribution, Metabolism, and Excretion and Pharmacokinetics of the HIV-1 Maturation Inhibitor GSK3640254 Using an Intravenous Microtracer Combined with EnteroTracker for Biliary Sampling

Bo Wen, Ying Zhang, Graeme C. Young, David Kenworthy, Adrian Pereira, Jill Pirhalla, Janine Doyle, Bethany Jordan, Joyce Zhan, and Mark Johnson

GSK, Collegeville, Pennsylvania (B.W., Y.Z., J.P., J.D., J.Z.); GSK, Ware, United Kingdom (G.C.Y., D.K.); GSK, Stevenage, United Kingdom (A.P., B.J.); and ViV Healthcare, Durham, North Carolina (M.J.)

Received May 23, 2022; accepted August 17, 2022

ABSTRACT

GSK3640254 is a next-generation maturation inhibitor in development for HIV-1 treatment, with pharmacokinetics (PK) supporting once-daily oral dosing in human. This open-label, nonrandomized, two-period clinical mass balance and excretion study was used to investigate the excretion balance, PK, and metabolism of GSK3640254. Five healthy men received a single intravenous microtracer of 100 μg [14C]GSK3640254 with a concomitant oral non-radiolabeled 200-mg tablet followed by an oral 85-mg dose of [14C]GSK3640254 14 days later. Complementary methods, including intravenous microtracing and accelerator mass spectrometry, allowed characterization of several parameters, including fraction absorbed, fraction escaping gut metabolism, hepatic extraction ratio, and renal clearance. Intravenous PK of GSK3640254 was characterized by low plasma clearance (1.04 l/h), moderate terminal phase half-life (21.7 hours), and low volume of distribution at steady state (28.7 L). Orally dosed GSK3640254 was absorbed (fraction absorbed, 0.26), with a high fraction escaping gut metabolism (0.898) and a low hepatic extraction ratio (0.00544), all consistent with low in vitro intrinsic clearance in liver microsomes and hepatocytes. No major metabolites in human plasma required further qualification in animal studies. Both unchanged parent GSK3640254 and its oxidative and conjugative metabolites were excreted into bile, with GSK3640254 likely subject to further metabolism through enterohepatic recirculation. Renal elimination of GSK3640254 as the parent drug or its metabolites was negligible, with >94% of total recovery of oral dose and >99% of the recovered radioactivity in feces. Altogether, the data suggest that systemically available GSK3640254 was slowly eliminated almost entirely by hepatobiliary secretion, primarily as conjugative and oxidative metabolites.

SIGNIFICANCE STATEMENT

The combination of an intravenous 14C microtracer with duodenal bile sampling using EnteroTracker in a human absorption, distribution, metabolism, and excretion study enabled derivation of absorption and first-pass parameters, including fraction absorbed, proportion escaping first-pass extraction through the gut wall and liver, hepatic extraction, and other conventional clinical pharmacokinetic parameters. This approach identified hepatic metabolism and biliary excretion as a major elimination pathway for absorbed drug, which would be overlooked based solely on analyses of plasma, urine, and fecal matrices.

INTRODUCTION

GSK3640254((R)-1-fluoromethyl-4-(17-((2-(1,1-dioxidothiomorpholino)ethyl)amino)-28-norlupa-2,20(29)-dienyl)-cyclohex-3-ene-1-carboxylic acid methanesulfonate) is a next-generation HIV-1 maturation inhibitor exhibiting panenotypic coverage of HIV-1 subtypes and polymorphic variants (Dicker et al., 2022) and is currently in phase IIb clinical development for HIV-1 treatment. Maturation is one of the last steps of the HIV-1 cycle wherein the mature structural Gag proteins are generated by viral protease-mediated cleavage (Wang et al., 2015). Maturation inhibitors specifically interfere with the protease-mediated processing of Gag, resulting in the release of HIV-1 particles that are immature and noninfectious.

Human absorption, distribution, metabolism, and excretion (referred to as ADME) studies quantitatively and comprehensively evaluate a drug’s overall disposition and are required for new drug approval (Coppola et al., 2019). Data obtained from human ADME studies provide
valuable information for confirming whether the toxicology species are relevant for assessing a drug’s safety and for developing a cohesive clinical pharmacology strategy, including investigation of potential drug-drug interactions and special patient populations (e.g., patients with hepatic or renal impairment).

Intravenous (i.v.) pharmacokinetics (PK) are important for the calculation of primary PK parameters, such as bioavailability, clearance, and volume of distribution, which are necessary for a complete understanding of a drug’s absorption, distribution, and elimination (Denton et al., 2013; Harrell et al., 2019). In conventional intravenous PK studies, healthy participants typically receive separate single therapeutically relevant intravenous and extravascular doses in a two-way crossover design; however, the therapeutic intravenous dose would require an intravenous toxicology package to support its use in clinical studies. To bridge the “translational gap” between preclinical research and clinical development, exploratory clinical studies are nowadays encouraged to include microdosing to evaluate subtherapeutic exposures of drug candidates in first-in-human studies. Regulatory guidance approves ultralow dose (≤100 µg) i.v. administration in humans, which allows for a substantially reduced toxicology package (https://database.ich.org/sites/default/files/M3_R2_Guideline.pdf). In the microdosing approach, a 14C-isotope may be incorporated into a drug, which allows for the measurement of systemic exposure using technologies such as accelerator mass spectrometry (AMS) (Harrell et al., 2019). Using a so-called intravenous “microtracer” design, microdosing studies may include concomitant administration of the 14C-labeled microdose with a therapeutically relevant, extravascular dose to define intravenous PK (Lappin, 2016). Herein, we use an approach that intravenously administers a 14C-microtracer dose of GSK3640254 concomitantly with the therapeutically relevant oral tablet dose of GSK3640254 (2 × 100-mg tablets) that is under clinical development.

Biliary disposition is an integral part of the human ADME profile. Metabolites can be abundantly formed in the liver but may not circulate in plasma (Loi et al., 2013). In addition, fecal metabolites do not necessarily represent hepatic metabolism due to degradation and/or metabolism by gut microbiota (Wilson and Nicholson, 2017). Similar to applications of Entero-test devices, which are no longer commercially available, the EnteroTracker device (EnteroTrack LLC, Aurora, CO) can be used to noninvasively collect duodenal bile samples and investigate human biliary disposition in humans (Guiney et al., 2011). To evaluate biliary secretion of a parent drug and its metabolites, duodenal bile sampling is preferred to stool sampling for many reasons, including providing an improved bile-specific metabolic profile without the potential contribution of gut microbiota to metabolism (Bloomer et al., 2013; Harrell et al., 2019). Herein, we report the first-time application of EnteroTracker devices in a human ADME study in which duodenal bile samples were noninvasively collected, enabling characterization of biliary disposition of GSK3640254 and its metabolites and evaluation of the potential of enterohepatic recirculation.

Human ADME studies that include an intravenous dose can be used to estimate the absolute bioavailability (F) and oral fraction absorbed (F_oral) (Raje et al., 2018; de Vries et al., 2019; Harrell et al., 2019; Podoll et al., 2019; John et al., 2020). In preclinical species, GSK3640254 had low to moderate oral bioavailability dosed either as a spray dried dispersion or mesylate salt suspension (unpublished observations). Despite low turnover in liver microsomes or hepatocytes, absorbed drug-related material was primarily eliminated by metabolism followed by biliary secretion (~13% of the administered dose) in bile duct–cannulated rats dosed orally with [14C]GSK3640254 mesylate salt (unpublished observations). Investigation of both the intravenous and oral PK of GSK3640254 and that of drug-related material in humans enables contributions of F_oral from the gastrointestinal tract and first-pass hepatic extraction to systemic exposure to be quantified. The study discussed herein aimed to fully characterize and understand the human ADME and PK of [14C]GSK3640254 after both intravenous and oral administration.

**Study Objectives and Design**

This was an open-label, single-center, nonrandomized, two-period, single-sequence crossover study evaluating the PK, mass balance, excretion, and metabolism of GSK3640254 in healthy men (ClinicalTrials.gov identifier: NCT04507321). This study consisted of a screening visit and two on-treatment periods separated by a ≥13-day washout period (Fig. 1). In treatment period 1, participants received a single oral tablet 200-mg dose of GSK3640254 with a moderate-fat meal and a 1-hour i.v. microtracer infusion of 100 μg [14C]GSK3640254 mesylate salt [~3.7 kilobecquerels (kBq); 100 nanocuries (nCi)] 5 hours after the oral dose to coincide with the anticipated maximal systemic exposures from the oral tablet administration. In treatment period 2, participants received an oral suspension of 85 mg [14C]GSK3640254 [~3.15 megabecquerels (MBq); 85 microcuries (µCi)] with a moderate-fat meal. Participants were fasted overnight for ≥8 hours before study treatment initiation in both treatment periods. EnteroTracker capsules were supplied by EnteroTracker LLC (Aurora, CO) to collect duodenal bile samples. The chemical structure of GSK3640254 has been previously published (Dicker et al., 2022). The structure of [14C]GSK3640254 indicating the position of radiolabel incorporation is shown in Fig. 2.

The primary objectives were to assess the rate and extent of excretion and total recovery of radioactivity in urine and feces in treatment periods 1 and 2 to determine systemic concentrations of parent GSK3640254 and total drug-related radioactivity in treatment periods 1 and 2 and to calculate the absolute oral bioavailability of GSK3640254 when given with food. Secondary objectives were to evaluate the safety and tolerability of GSK3640254 after single intravenous or
Eligible participants were healthy men aged from 30 to 50 years with a body weight of \( \geq 50 \) kg, a body mass index between 19 and 31 kg/m\(^2\), and a history of regular bowel movements. Participants had no recent or chronic history of diarrhea, no history of drug abuse, no regular alcohol or tobacco use in the previous 6 months, no clinically relevant disease, and no exposure to significant radiation in the previous 3 years. Participants with use of over-the-counter or prescription medications, including analgesics, herbal medications, or grapefruit or Seville orange juices, within 14 days before study treatment until study completion were excluded. Full eligibility criteria can be found on ClinicalTrials.gov (NCT04507321; https://clinicaltrials.gov/ct2/show/NCT04507321).

**Sample Collection and Processing**

Blood samples were collected in dipotassium ethylenediaminetetraacetic acid tubes through 8 days after dosing in treatment periods 1 and 2, and plasma was separated via centrifugation. Urine and fecal samples were collected through 8 days after dosing in treatment periods 1 and 2, which was extended up to 15 days after dosing in participants for whom excretion took longer than anticipated in treatment period 2. Duodenal bile was collected using a noninvasive bile string EnteroTracker device (Guiney et al., 2011). Briefly, participants pulled the first 10 to 20 cm of nylon out from the EnteroTracker capsule by the protruding loop and held the loop outside of the mouth while the capsule was swallowed with up to 350 ml of water. The string was then securely attached to the participant’s face with adhesive tape over the loop. In treatment period 1, the bile string was swallowed 2 hours after the oral dose and 3 hours before the intravenous infusion started, a duration recommended to allow transit of the string to the duodenum. Participants fasted from insertion of the bile string until the completion of the intravenous infusion, at which point a food cue (small, standard, high-fat meal) was given to stimulate gallbladder emptying. The EnteroTracker was removed 1.5 hours after the intravenous infusion stopped (\( \sim 7.5 \) hours after the oral dose) to capture the bile samples. In treatment period 2, the EnteroTracker was swallowed 2 hours after the oral dose. At 5.5 hours after the oral dose, a food cue (small, standard, high-fat meal) was given to stimulate gallbladder emptying. The collection bile string was removed 1.5 hours after the food cue (\( \sim 7 \) hours after the oral dose) to capture the duodenal bile samples expelled from the gallbladder. Samples of urine, feces, and bile strings were stored frozen before shipment for analysis.

**Sample Analysis**

**Mass Balance and Excretion.** Total radioactivity excreted in urine and fecal samples was determined using liquid scintillation counting (LSC) or AMS in treatment period 1 (intravenous; Pharmaron, Inc., Germantown, MD) and using LSC in treatment period 2 (oral; Labcorp Drug Development, Harrogate, UK; summarized in Supplemental Table 1). Accelerator mass spectrometry was used to analyze samples with radioactivity levels that were too low to be detected...
using LSC. Because LSC and AMS have been individually qualified in the literature (Garner et al., 2000; Keck et al., 2010), cross-validation between techniques was not performed for this study. Procedures for urine and fecal sample preparation and measurement of radioactivity by LSC and AMS have been previously described, and key methodology is summarized here (Harrell et al., 2019). Efficiency correlation curves for LSC were confirmed using $^{14}$C quenched standards. For AMS, instrument standards and process standards for graphitization were analyzed. Radioactivity in urine samples was measured in triplicate by LSC using a Packard TriCarb liquid scintillation counter (Canberra Packard, Pangbourne, Berkshire, UK). Fecal samples were combusted (Sample Oxidizer; PerkinElmer, Pangbourne, Berkshire, UK), absorbed in Carbo-Sorb (PerkinElmer), and combined with PermFluor E+ scintillation fluid (PerkinElmer). Oxidation efficiency was determined by combustion of quality control standards and had to be 95% to 105% at the beginning and end of each run to be acceptable. Values were not corrected for combustion efficiency. Lower limits of quantification (LLQs) were established as two times the mean of background disintegrations, with mean LLQ values of 0.01 ng GSK3640254 Eq/ml in urine and 186 ng Eq/g in fecal samples.

After undergoing combustion (oxidation) and graphitization (reduction), total radioactivity levels in urine and fecal samples were determined by AMS using a Single Stage AMS 250KV system (National Electrostatics Corporation, Middleton, WI) (Young et al., 2008). Mean LLQ values were 8.99 pg GSK3640254 Eq/ml in urine and 84.6 pg Eq/g in fecal samples.

**PK Assessments.** Plasma samples were analyzed for GSK3640254 concentrations by PPD (Middletown, WI) using a validated analytical method based on protein precipitation followed by liquid chromatography (LC) with tandem mass spectrometry (LC-MS/MS) as previously described (summarized in Supplemental Table 1) (Pene Dumitrescu et al., 2021). The lower limit of quantification was 3 ng/ml using a 50-μl aliquot of plasma, and the higher limit of quantification was 1000 ng/ml. Plasma total radioactivity was analyzed by Pharmaron, Inc. using AMS analysis, with lower limits of quantification of 48.4 pg GSK3640254 Eq/ml in treatment period 1 and 43.2 pg Eq/ml in treatment period 2. Plasma $^{14}$C[GSK3640254 concentrations were analyzed by Pharmaron, Inc. using a validated analytical method based on protein precipitation followed by LC + AMS analysis. Using a 500-μl aliquot of plasma, the lower limit of quantification was 60.9 pg/ml, and the higher limit of quantification was 1830 pg/ml.

The LC-MS/MS and LC + AMS analytical methods included quality control samples containing GSK3640254 and [1$^{4}$C]GSK3640254 at 3 different concentrations, which were stored and analyzed all samples against calibration standards, which were separately prepared. For analyses to be acceptable, ≤33% of quality control results and ≤50% of results from each concentration level could deviate from the nominal concentration by >15% for LC-MS/MS and >20% for LC + AMS. Applicable analytical runs met all predefined run acceptance criteria.

**Quantification and Characterization of Metabolites.** Procedures for the quantification and characterization of metabolites have been previously described, and key methodology is summarized here (Harrell et al., 2019). The metabolic profile of plasma extracts from treatment period 2 (oral), homogenized fecal extracts from treatment period 1 (intravenous), and bile string extracts from both treatment periods were characterized by Pharmaron, Inc. using LC + AMS (summarized in Supplemental Table 1). Plasma samples from individual participants in treatment period 2 were pooled using an area-under-the-curve (AUC) approach (Hamilton et al., 1981; Hop et al., 1998) from time 0 to 24 and 0 to 96 hours and then further pooled among participants by combining volumes in proportion to the time interval between individual samples to prepare a single plasma pool. The 0- to 24-hour AUC pools were used to allow a direct comparison with those of nonclinical species, and plasma toxicokinetic and $^{14}$C-metabolite profiles were determined over a 0- to 24-hour time period. Consequently, the 0- to 96-hour AUC pools were analyzed to capture the majority (>90%) of $^{14}$C radioactivity in human plasma samples. Homogenized fecal samples were pooled based on the ratio of the total weight of the sample excreted at each time point such that the pool represented ≥95% of the total excreted $^{14}$C across all participants. An equal weight of sample from each individual pool was obtained to prepare a combined fecal pool. Plasma and fecal samples were extracted with a 50:50 mixture of methanol to acetonitrile (volume/volume, 3–4 volumes) two times, dried under a stream of nitrogen gas, and reconstituted in a 10:10:80 mixture of dimethyl sulfoxide to acetonitrile to water (volume/volume/volume). Bile strings were removed from the freezer and thawed at room temperature for the shortest time that was practically possible. Each string was immediately placed inside the barrel of a hypodermic plastic syringe, and duodenal bile was extracted individually with high-performance LC (HPLC)-grade acetonitrile and then HPLC-grade water. The organic extracts from treatment period 1 (intravenous bile) were dried and then reconstituted with 250 μl of 10-mM ammonium formate (pH = 4) in a 9:1 mixture of LC-grade water to HPLC-grade acetonitrile containing 0.1% formic acid (pH = 4; volume/volume), whereas the combined extracts from treatment period 2 (oral bile) were diluted 100-fold with 10 mM ammonium formate (pH = 4). Extracts were pooled from individual participants, and extracts containing radioactivity above LSC background were pooled across participants.

The metabolic profile of fecal samples from treatment period 2 (oral) was characterized by Q2 Solutions (Morrisville, NC) using HPLC-MS/MS. Representative fecal samples containing >1% of the administered dose were pooled by total weight ratio to obtain a pool containing >95% of the radioactivity excreted in each participant. Fecal samples were extracted with a 50:50 mixture of methanol to acetonitrile, centrifuged evaporated under nitrogen, and reconstituted in a 1:1:8 (volume/volume/volume) mixture of dimethyl sulfoxide to acetonitrile to water. The amount of radioactivity in the extracts was determined by LSC before radio-HPLC analysis. Radio-HPLC data were captured offline by collecting fractions into 96-well Scintiplates (PerkinElmer, Waltham, MA). Each HPLC eluate was evaporated to dryness under centrifugal vacuum, the dried plates were sealed, and a microplate scintillation counter with 12 independent photomultiplier tubes was used to count each well; no background was subtracted.

Data analyses from AMS and scintillation counting data from offline analyses were converted into a form compatible with Laura software (LabLogic Systems, Inc., Tampa, FL) using its “LSC import” function to reconstruct AMS and radio-HPLC radiochromatograms. The area in each radioactive peak was expressed as a percentage of the total number of counts detected.

**Pharmacokinetics and Statistical Analyses**

No formal sample size calculation was performed. A sample size of four to six participants was deemed appropriate to achieve the objectives of the study (Penner et al., 2009). This study used WinNonlin software version 6.3 or higher (Certara, Princeton, NJ) to noncompartmentally analyze the parent drug after oral tablet or suspension administration and intravenous administration (plasma GSK3640254 and [1$^{4}$C]GSK3640254, respectively) as well as total radioactivity concentration-time data based on actual recorded sampling times for derivation of PK parameters. Clearance, volume of distribution at steady state ($V_{ss}$), and oral bioavailability for each oral formulation (tablet or suspension) were calculated for GSK3640254 in plasma. Pharmacokinetic parameters for GSK3640254, [1$^{4}$C]GSK3640254, and total radioactivity after intravenous infusion and oral administration were summarized descriptively. Absorption and first-pass PK parameters for GSK3640254, including fraction of drug escaping first-pass hepatic clearance ($F_{L}$), hepatic extraction ratio ($E_{L}$), fraction of drug escaping gut metabolism ($F_{G}$), and $F_{abs}$ were calculated as previously described (Harrell et al., 2019).

**Results**

**Study Population**

Of the 17 individuals screened, five participants were enrolled and completed the study. One protocol deviation was reported for a participant who met exclusion criteria (past or intended use of over-the-counter or prescription medications); this participant was re-screened after meeting eligibility criteria and enrolled. All participants were men, with a mean age of 37 years and a mean body mass index of 24.8 kg/m² (Supplemental Table 2).

**Mass Balance and Excretion**

The mean recovery of radioactivity over time after intravenous and oral administration of [1$^{4}$C]GSK3640254 is shown in Fig. 3. After intravenous administration of [1$^{4}$C]GSK3640254, the majority of radioactivity was excreted within 96 hours (53% of dose) and a total of 76% of the administered dose was recovered within 7 days, with 74% of the
dose being recovered in feces and <2% in the urine. Normalized for recovery, fecal excretion was predominant, accounting for 97.5% of the total recovered radioactivity, with ~2.5% in the urine (Fig. 3A). After oral administration of [14C]GSK3640254, 75% of the administered dose was recovered within 96 hours and a total of >94% of the administered dose was recovered within 7 days. Similarly, excretion was predominately in the feces, accounting for >99% of the total recovered radioactivity (Fig. 3B), with <0.6% of the administered dose recovered in the urine.

Pharmacokinetic Results

After oral administration of GSK3640254 (2 × 100-mg tablets) in treatment period 1, the maximum observed concentration (Cmax) was reached with a median time of Cmax (Tmax) of 7 hours followed by a slow decline in concentrations, with a geometric mean terminal phase half-life (t1/2) of 24.1 hours (Fig. 4A; Table 1). The geometric mean F of GSK3640254 after oral tablet administration was 21.4% [90% confidence interval (CI), 19.4% to 23.5%; Table 2] based on dose-normalized AUC from time 0 extrapolated to infinity (AUC0–∞) relative to the intravenous microtracer dose.

After intravenous infusion administration of [14C]GSK3640254 in treatment period 1, similar concentration-time profiles were demonstrated for both [14C]GSK3640254 and total radioactivity, with Cmax occurring at the end of the 1-hour infusion (median Tmax = 0.983 hours; Fig. 4B; Table 1). These data demonstrated that GSK3640254 is the predominant contributor of total radioactivity in plasma, with a minimal contribution of metabolite(s) to the drug-related material in systemic circulation. Plasma concentrations of [14C]GSK3640254 steadily declined from Cmax, with a geometric mean t1/2 of 21.7 hours. The geometric mean plasma clearance of [14C]GSK3640254 was low (0.0438 and 0.180, respectively). After oral suspension administration, it is estimated that ~26% of GSK3640254 is absorbed (Fabs) across the gastrointestinal tract and only ~0.5% of the absorbed drug is cleared (Eh) before reaching the systemic circulation. Minimal loss of GSK3640254 was likely due to high fractions of drug escaping metabolism in the gut wall and liver, with an Fg value of 89.8% and an Fh value of 99.5%, respectively.

Metabolism

A putative metabolic scheme for GSK3640254 is shown in Fig. 2. Accelerator mass spectrometry analysis of time-resolved pooled plasma extracts (0–24 hours and 0–96 hours) from oral-dosed participants (treatment period 2) showed that GSK3640254 was the predominant circulating...
Fig. 4. Mean (S.D.) plasma concentrations of (A) parent GSK3640254 after oral tablet (2 × 100 mg) and i.v. infusion (100 μg) administration and (B and C) parent GSK3640254 vs. total radioactivity after (B) i.v. infusion (100 μg) and (C) oral suspension (85 mg) administration.
After intravenous infusion in treatment period 1, unchanged GSK3640254 was the predominant radioactive component in pooled fecal samples, accounting for 44% of total radioactivity and 32% of the dose; other notable radioactive components included M4 (oxidation; 7% of radioactivity), M8 (N-dealkylation; 3.6% of radioactivity), and M9 (N-dealkylation) plus M28 (oxidation; labeled as “Unknown” and accounted for 9.0% of combined radioactivity; Fig. 6A). All other radioactive components accounted for <4% of the total radioactivity. In pooled bile extracts after intravenous infusion, M1 (formed by acyl glucuronidation) was the major radioactive component, representing 41% of total radioactivity; other radioactive components included unchanged GSK3640254 (15% of radioactivity), M4 (oxidation; 11% of radioactivity), and M8 (d-oxidation; 6% of radioactivity; Fig. 6B). All other radioactive components individually accounted for <4% of the total radioactivity.

After oral administration in treatment period 2, the principal radioactive component in feces was unchanged GSK3640254, representing 84% of total radioactivity and 79% of the dose (Fig. 7A). Other notable radioactive components included M4 (oxidation; 3% of radioactivity) and M9 coeluting with M28 (5% of radioactivity; formed by N-dealkylation and oxidation, respectively). The remaining lesser radioactive components included M5 (oxidation), M7 (hydration), M8 (d-oxidation), M13 (hydration and oxidation), M14 (oxidation and N-dealkylation), M18 (oxidation), M27 (oxidation), and M29 (oxidation and N-dealkylation), each <2% of radioactivity (Supplemental Table 4). Unchanged GSK3640254 also accounted for the majority of total radioactivity in pooled bile extracts after oral administration (94%), with other radioactive components each accounting for <1% of total radioactivity (Fig. 7B); however, comparison of biliary metabolite profiles from intravenous- and oral-dosed participants suggested that GSK3640254 in bile samples from treatment period 2 was composed of primarily the unabsorbed GSK3640254. With the high
radioactive dose in treatment period 2, it is likely that any potential contamination from the unabsorbed radioactive oral dose had significant impact on the metabolic profiles from oral-dosed participants.

**Safety**
No deaths, serious adverse effects (AEs), or AEs leading to study withdrawal were reported. Overall, 1 (20%) participant reported three

**TABLE 2**

<table>
<thead>
<tr>
<th>Dose-Normalized PK Parameter</th>
<th>Dose-Normalized Adjusted Geometric Mean</th>
<th>Ratio (% CVw)</th>
<th>90% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Absolute Bioavailability</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$AUC_{0-24h}$/dose (h-ng/ml per mg)</td>
<td>Test (oral tablet) 206</td>
<td>0.214 (7.2)</td>
<td>0.194–0.235</td>
</tr>
<tr>
<td></td>
<td>Reference (intravenous) 966</td>
<td></td>
<td></td>
</tr>
<tr>
<td>$AUC_{0-96h}$/dose (h-ng/ml per mg)</td>
<td>Test (oral tablet) 204</td>
<td>0.219 (7.8)</td>
<td>0.198–0.243</td>
</tr>
<tr>
<td></td>
<td>Reference (intravenous) 933</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Relative Bioavailability</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$AUC_{0-24h}$/dose (h-ng/ml per mg)</td>
<td>Test (oral suspension) 224</td>
<td>1.090 (8.2)</td>
<td>0.973–1.212</td>
</tr>
<tr>
<td></td>
<td>Reference (oral tablet) 206</td>
<td></td>
<td></td>
</tr>
<tr>
<td>$AUC_{0-96h}$/dose (h-ng/ml per mg)</td>
<td>Test (oral suspension) 222</td>
<td>1.090 (8.4)</td>
<td>0.969–1.216</td>
</tr>
<tr>
<td></td>
<td>Reference (oral tablet) 204</td>
<td></td>
<td></td>
</tr>
<tr>
<td>$C_{max}$/dose (ng/ml per mg)</td>
<td>Test (oral suspension) 7.39</td>
<td>1.140 (12.1)</td>
<td>0.992–1.318</td>
</tr>
<tr>
<td></td>
<td>Reference (oral tablet) 6.46</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

CVw, within-participant coefficient of variation.

Fig. 5. Reconstructed AMS radiochromatograms of (A) pooled human AUC$_{0-24h}$ plasma and (B) pooled human AUC$_{0-96h}$ plasma after administration of 85 mg $[^{14}C]$GSK3640254 as an oral suspension. AUC$_{0-24h}$, area under the concentration-time curve from time 0 to 24 hours; AUC$_{0-96h}$, area under the concentration-time curve from time 0 to 96 hours.
AEs during the study. This participant reported AEs of asthenia and nausea that were mild in intensity during treatment period 1 and an additional AE of asthenia that was moderate in intensity during treatment period 2. All AEs were considered to be drug related and resolved within 2 days of onset. No clinically significant findings were reported for chemistry or hematology values, electrocardiographic parameters, or vital signs.

Discussion

A combination of a 14C intravenous microtracer and noninvasive EnteroTracker bile sampling techniques was used to fully characterize the ADME and PK of GSK3640254 in healthy men, with comprehensive information obtained for mass balance and excretion data, GSK3640254 and total radioactivity in plasma, and metabolite profiles in multiple matrices. This approach allowed for the derivation of absorption and first-pass parameters that are not typically reported in conventional clinical PK studies, including parameters such as Fabs, Fg, Eh, and geometric mean renal clearance. Incorporating a 14C-isotope into the parent GSK3640254 molecule enabled human metabolism to be fully evaluated in both plasma and excreta, allowing for the assessment of exposure to circulating metabolites as well as the identification of elimination pathways. To ensure pharmacological relevance, the intravenous microtracer dose was concomitantly administered with a therapeutically relevant GSK3640254 oral dose (2 × 100-mg tablets), which is currently under clinical development. The first-time application of the noninvasive bile sampling via EnteroTracker devices enabled the assessment of biliary disposition of GSK3640254 and its metabolites. Most importantly, hepatic metabolism and biliary excretion were identified as a major elimination pathway for absorbed GSK3640254, which would otherwise be overlooked based solely on analysis of plasma, urine, and fecal matrices.

The total recovery of radioactivity achieved after oral suspension administration of 85 mg [14C]GSK3640254 (3.15 MBq radioactivity) with food was 94%, which is in line with expectations for an acceptable mass balance recovery as established by the European Medicines Agency (https://www.ema.europa.eu/en/investigation-drug-interactions). Although recovery of radioactivity was lower (76%) after intravenous infusion, it was only slightly below the 80% recovery threshold reported in a retrospective analysis of other mass balance clinical studies (Roffey et al., 2007). It is noteworthy to mention that the much lower radioactive microtracer dose (3.7 kBq) delivered by intravenous infusion did not permit recovery to be used for release criteria due to the slow data turnaround; thus, a fixed collection period of 7 days after dosing was established in the study protocol. Excretion of GSK3640254 predominately occurred through the feces, accounting for 97.5% and >99% of total recovered radioactivity after intravenous and oral administration, respectively. By contrast, urinary excretion was minimal after administration of [14C]GSK3640254 as an intravenous infusion (<2.5%) or oral suspension (<1%). Due to negligible radioactivity excreted in urine, no metabolite investigation was conducted in this matrix.

Intravenous administration of [14C]GSK3640254 resulted in plasma concentration-time profiles characterized by low clearance and low Vss. GSK3640254 exhibited blood clearance equal to 2% of human liver blood flow, and the Vss (28.7 L) was less than the previously reported value for total body water (42 L) (Davies and Morris, 1993); these results suggest that limited distribution of GSK3640254 occurred outside of the plasma. Regardless of formulations, oral administration of GSK3640254 demonstrated good dose proportionality between the 200-mg oral tablets and 85-mg oral suspension. After oral suspension

![Fig. 6. Reconstructed AMS radiochromatograms of (A) pooled human feces and (B) pooled human duodenal bile after administration of 100 μg [13C]GSK3640254 as an intravenous infusion.](image-url)
administration, it is estimated that ~26% of dose is absorbed ($F_{abs}$) across the gastrointestinal tract. Although the oral $F_{abs}$ was low, any absorbed GSK3640254 is likely subject to poor ‘first-pass’ metabolism by the gut wall ($F_g = 0.898$) and liver ($F_h = 0.005$; $E_h = 0.995$). Thus, it is postulated that the relatively low oral $F$ (21.4%) was primarily driven by the low oral absorption rates rather than metabolic clearance or ‘first-pass’ effects. Key factors driving oral $F$ can be better understood by dissecting oral absorption into specific parameters such as absorption rate and gut and hepatic extraction (Harrell et al., 2019). Determining the ‘first-pass’ drug burden to the gastrointestinal tract and liver may also improve the evaluation of potential drug-drug interactions using physiologically based PK models. In addition, the low $E_h$ and poor ‘first-pass’ effects of GSK3640254 in human are parallel with the observed low intrinsic clearance in vitro in incubations with human liver microsomes and hepatocytes (unpublished observations). Despite the low $F_{abs}$, the minimal ‘first-pass’ effect and low clearance ensures delivery of adequate exposures of GSK3640254 reaching the systemic circulation. The geometric mean elimination $t_{1/2}$ values were similar, ranging from 22 to 24 hours after administration as oral tablets, intravenous infusion, or oral suspension, suggesting that the rate of terminal elimination was not affected by the administration route (intravenous or oral). Based on geometric mean plasma $\text{[^{14}C]}$GSK3640254-to-total radioactivity ratios for $C_{\text{max}}$ and AUC, unchanged GSK3640254 was the predominant contributor of total radioactivity in plasma after both intravenous infusion (>90%) and oral administration (82% to 93%).

The predominant circulating drug-related component in plasma was unchanged GSK3640254 after oral dosing. Several minor radioactive peaks representing ~2% to 3% of sample radioactivity were composed of three metabolites formed by acyl-glucuronidation, oxidation, and N-dealkylation. Therefore, no individual metabolite exceeded the 10% threshold of the total drug-related exposure outlined by the International Conference on Harmonization and US Food and Drug Administration (https://database.ich.org/sites/default/files/M3_R2__Guideline.pdf; https://www.fda.gov/media/72279/download), above which further nonclinical evaluation may be justified. The metabolites identified in human plasma had chemical structures that were formed via common oxidative and conjugative routes and are considered innocuous and unlikely to be reactive. Therefore, no human GSK3640254 metabolites need further evaluation in nonclinical studies. In addition, the levels of circulating radioactivity were closely aligned with the levels of
circulating concentrations of GSK3640254, suggesting minimal levels of metabolites in plasma from the intravenous administration of $^{14}$C-GSK3640254. The acyl glucuronide M1 was a major component (41%) in human duodenal bile after intravenous administration but was not detected in fecal extracts (Fig. 6), suggesting a rapid hydrolysis of the acyl glucuronide metabolite by the gut microflora. This likely contributed to a potential enterohepatic recirculation of GSK3640254, which may explain small fluctuations (secondary uplifts) observed in the plasma PK profiles. Although only “snapshot” measurements, LC + AMS analysis of bile string extracts suggested that acyl glucuronidation (as M1) was a likely major route of elimination pathway of GSK3640254. Given that no or only trace levels of M1 were detected in other biologic matrices, including plasma and feces, it is of importance that duodenal bile sampling bridges the gap in understanding the major elimination pathways of GSK3640254 in human. Based on in vitro phenotyping experiments, several uridine diphosphateglucuronosyltransferase (UGT) isoforms, including UGT1A4 and UGT2B7, were involved in glucuronidation of GSK3640254, with less contribution from UGT1A9 (unpublished observations).

It remains challenging to predict which metabolites formed in the liver will eventually circulate in plasma (Loi et al., 2013). Despite extensive hepatic metabolism observed in duodenal bile, unchanged parent GSK3640254 is the predominant circulating component in human plasma, with only trace levels of metabolites observed (Fig. 5). This is parallel with the low plasma ML (ratio of metabolite to total radioactivity) after intravenous (ML = 0.0438) and oral (ML = 0.18) administration (Table 3). Routes of metabolism included various oxidations, including acyl glucuronidation, N-dealkylation, mono- and dioxygenations, and combinations thereof. Apart from UGT-mediated conjugation reactions, cytochrome P450 3A4 was identified as the only cytochrome P450 isoform metabolizing GSK3640254 in incubations with human liver microsomes (unpublished observations). Overall, these data suggest that once absorbed, the systemically available GSK3640254 was eliminated primarily by hepatobiliary secretion in the form of metabolites. A high proportion of metabolic elimination is likely to result from acyl glucuronidation. Renal clearance is a negligible route of elimination.

In conclusion, this small, two-part study using concomitant administration of intravenous and oral doses, state-of-the-art analytical technology, and thorough evaluation of ADME and PK parameters has rigorously characterized GSK3640254 disposition in human. These findings subsequently helped the development of a cohesive strategy for clinical pharmacology studies, including drug-drug interactions and study of special patient populations.

### Acknowledgments

The authors would like to acknowledge the contributions of Mark Bush, Samit Joshi, and Martin Gartland for clinical study design and data interpretation, Jessica Shoultz and Yijun Yi (Q2 Solutions, Indianapolis, IN) for conducting the LC-MS/MS metabolite identification work, Stephen English (Pharmaron, Inc., Germantown, MD) for contributing to the AMS and LC + AMS analyses, and Lee Crossman (Labcorp Drug Development, Harrogate, UK) for contributing to the human excretion and mass balance analyses. Editorial assistance was provided under the direction of the authors by Megan Schmidt and Lauren Bragg, ELS, and MedThinkSci.com and was funded by ViiV Healthcare.

### Authorship Contributions

**Conducted experiments:** Pirilla, Doyle.

**Contributed new reagents or analytic tools:** Young, Kenworthy, Pereira.

**Performed data analysis:** Wen, Zhang, Pereira, Pirilla, Doyle, Zhan.

**Wrote or contributed to the writing of the manuscript:** Wen, Zhang, Young, Kenworthy, Pereira, Pirilla, Doyle, Joran, Zhan, Johnson.

### References


**TABLE 3** Summary of plasma metabolite load, hepatic clearance, hepatic extraction, and absorption parameters

<table>
<thead>
<tr>
<th>PK Parameter</th>
<th>Treatment</th>
<th>N</th>
<th>n</th>
<th>Geometric Mean (% CVb)</th>
<th>95% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>MLiv</td>
<td>Intra</td>
<td>5</td>
<td>5</td>
<td>0.0348 (363)</td>
<td>0.0058–0.3310</td>
</tr>
<tr>
<td>MLpo</td>
<td>Oral</td>
<td>5</td>
<td>5</td>
<td>0.180 (8.7)</td>
<td>0.162–0.2010</td>
</tr>
<tr>
<td>CLh,iv,plasma (l/h)</td>
<td>Intra</td>
<td>5</td>
<td>5</td>
<td>1.02 (20.0)</td>
<td>0.794–1.3000</td>
</tr>
<tr>
<td>CLh,iv,blood (l/h)</td>
<td>Intra</td>
<td>5</td>
<td>5</td>
<td>0.542 (20.0)</td>
<td>0.424–0.6930</td>
</tr>
<tr>
<td>EM</td>
<td>Intra</td>
<td>5</td>
<td>5</td>
<td>0.00544 (20.0)</td>
<td>0.000426–0.000969</td>
</tr>
<tr>
<td>Fh</td>
<td>Oral</td>
<td>5</td>
<td>5</td>
<td>0.898 (5.5)</td>
<td>0.839–0.9610</td>
</tr>
<tr>
<td>Fh</td>
<td>Oral</td>
<td>5</td>
<td>5</td>
<td>0.995 (0.1)</td>
<td>0.993–0.9960</td>
</tr>
<tr>
<td>Fabs</td>
<td>Oral</td>
<td>5</td>
<td>5</td>
<td>0.260 (8.9)</td>
<td>0.232–0.2900</td>
</tr>
</tbody>
</table>

MLiv/hepatic blood clearance; CLh,iv,plasma, hepatic plasma clearance; CVb, between-participant coefficient of variation; MLpo, plasma metabolite load; MLiv, oral metabolite load.