

Intra- and Interindividual Variabilities of Valacyclovir Oral Bioavailability and Effect of Coadministration of an hPEPT1 Inhibitor

Dana D. Phan,¹ Peter Chin-Hong,² Emil T. Lin,¹ Pascale Anderle,^{1†}
Wolfgang Sadee,^{1‡} and B. Joseph Guglielmo^{1*}

Schools of Pharmacy¹ and Medicine,² University of California, San Francisco, California

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Variability in valacyclovir bioavailability and the potential for cephalexin-valacyclovir interaction were evaluated. The intraindividual acyclovir area under the concentration-time curve (AUC) varied minimally, whereas interindividual differences were substantial. Coadministration of the human peptide transporter 1 (hPEPT1) substrates valacyclovir and cephalexin minimally reduced the acyclovir AUC. These results suggest a stable valacyclovir absorption phenotype, significant interindividual variability, and minimal interaction between these hPEPT1 substrates.

Human peptide transporter 1 (hPEPT1) is associated with the absorption of peptidomimetic drugs such as valacyclovir and cephalexin (2, 3, 10, 11). Animal and in vitro models have documented a 50% reduction in the hPEPT1 uptake of valacyclovir in the presence of equimolar cephalexin (4, 6–8). While valacyclovir is a substrate of hPEPT2 and organic anion transporter 1 (OAT1), uptake by hPEPT1 is thought to be the primary mechanism of valacyclovir absorption (1, 4–8). Since 99% of the absorbed valacyclovir is hydrolyzed to acyclovir within 3 h of oral administration (6–7), valacyclovir reaches hPEPT2 and OAT1 in the form of acyclovir, which is a substrate of neither hPEPT2 nor OAT1 (12). Considering its rapid metabolism to acyclovir, the oral bioavailability of valacyclovir is a reflection of the acyclovir area under the concentration-time curve (AUC).

The intra- and interindividual variabilities of the oral bioavailability of valacyclovir have not been well studied. We evaluated the variability of valacyclovir absorption, as measured by the acyclovir AUC, and the impact of cephalexin on the acyclovir AUC.

This study was conducted at the University of California at San Francisco (UCSF) General Clinical Research Center (GCRC). All study subjects gave informed consent. The protocol was approved by the Committee on Human Research.

Volunteers were excluded if they had (i) diabetes, cardiovascular disease, or renal or hepatic disease; (ii) a recent history of drug abuse, alcoholism, or nicotine dependence; (iii) a history of intolerance to acyclovir or its analogues, cephalosporins, or penicillins; (iv) participated in other studies during the preceding month; or (v) taken any medication or dietary supplement other than oral contraceptives, vitamins, or minerals within the preceding 2 weeks; and (vi) female volunteers

were excluded if they were pregnant, lactating, or sexually active without using adequate contraceptive measures. All female volunteers were required to provide urine for a urine dipstick pregnancy test.

With a random-number generator, subjects were randomly assigned to a single oral dose of (i) 500 mg of valacyclovir at both visits 1 and 2 (control group; $n = 6$) (ii), 500 mg of valacyclovir at visit 1 and 500 mg of valacyclovir plus 500 mg of cephalexin at visit 2 (treatment group A; $n = 5$), or (iii) 500 mg of valacyclovir plus 500 mg of cephalexin at visit 1 and 500 mg of valacyclovir at visit 2 (treatment group B; $n = 5$).

Subjects were admitted to the GCRC for two admissions, separated by at least 7 days. During each admission, subjects were allowed to continue their existing medication regimen. Subjects were required to abstain from alcohol-, caffeine-, or xanthine-containing products within 24 h prior to and during each admission, fast the night before each admission, and abstain from fluid intake within 2 h prior to and after the administration of each valacyclovir dose.

Blood samples were collected in 5-ml sodium heparinized tubes before dosing and then at 0.5, 1, 2, 4, 8, and 12 h following the administration of each valacyclovir dose. Plasma was separated via centrifugation at $1,300 \times g$ for 5 min and then frozen in 2 aliquots at -20°C . After overnight storage at -20°C , plasma samples were stored at -80°C until assayed.

Concentrations of acyclovir in plasma were determined by a validated liquid chromatography-tandem mass spectrometry method developed at the UCSF Drug Study Unit (E. T. Lin, unpublished data). The standard curve was linear over a concentration range of 50 to 6,000 ng/ml ($r^2, \geq 0.996$). The lower limit of quantitation was 50 ng/ml, and the coefficients of variation were less than 15%. The assay was reproducible and accurate (coefficient of variation, $\leq 10\%$) and without interference from the matrix and anticoagulant. Although the samples were analyzed within 1 month of collection, acyclovir was stable in frozen human plasma at -80°C for at least 3 months. Acyclovir was stable in plasma for up to five freeze-and-thaw cycles.

The above-mentioned liquid chromatography-tandem mass

* Corresponding author. Mailing address: Department of Clinical Pharmacy, Box 0622, C-144, University of California at San Francisco, San Francisco, CA 94143. Phone: (415) 476-1927. Fax: (415) 476-6632. E-mail: bjpg@itsa.ucsf.edu.

† Present address: ISREC, 1066 Epalinges, Switzerland.

‡ Present address: Ohio State University, Columbus, OH 43210.

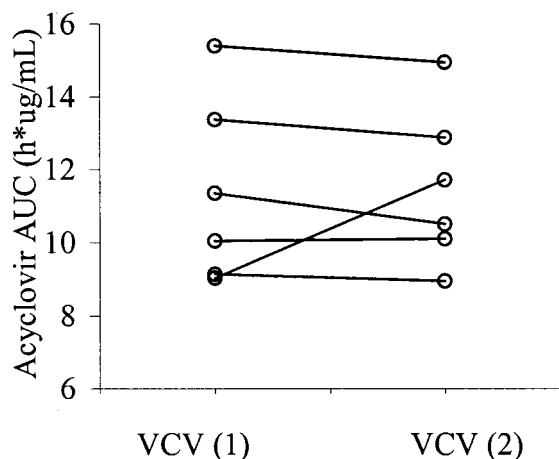


FIG. 1. Acyclovir $AUC_{0 \rightarrow \infty}$ in the absence of cephalixin. VCV (1) is valacyclovir during PK visit 1, and VCV (2) is valacyclovir during PK visit 2. The interindividual standard deviations for VCV (1) and VCV (2) were 2.5 and 2.2, respectively. The intraindividual standard deviation was 0.56 or about four times less than the interindividual variability. There was no difference in the acyclovir $AUC_{0 \rightarrow \infty}$ between the periods ($P = 0.82$).

spectrometry method was modified to monitor the conversion of valacyclovir to acyclovir.

WinNonLin 3.1 was used to perform noncompartmental pharmacokinetic analysis. Acyclovir concentration-versus-time curves were integrated and extrapolated by the linear/log trapezoidal rule and the $1/\text{concentration}$ weighted method to yield the AUC from zero to infinity ($AUC_{0 \rightarrow \infty}$). λ_z was calculated by selecting the terminal portion of the curve and subsequently used to extrapolate the AUC from the last measurable concentration to infinity ($AUC_{\text{last} \rightarrow \infty}$).

Eleven previously identified single-nucleotide polymorphisms (SNPs) in and surrounding the hPEPT1-encoding gene were selected for sequencing on the basis of their allelic vari-

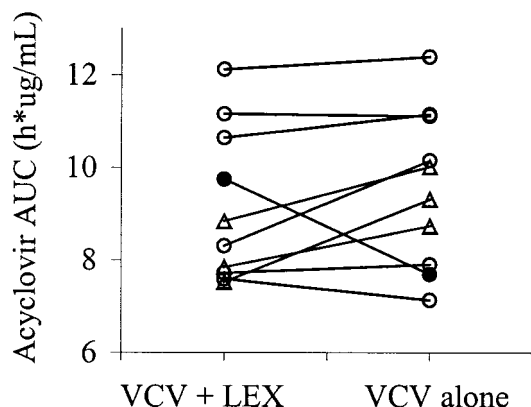


FIG. 2. Acyclovir $AUC_{0 \rightarrow \infty}$ in the presence and absence of cephalixin. VCV + LEX is concomitant administration of valacyclovir and cephalixin, and VCV alone is single treatment with only valacyclovir. The 7.1% reduction in oral bioavailability was determined without the outlier (\bullet). Concomitant cephalixin reduced the acyclovir $AUC_{0 \rightarrow \infty}$ from 9.8 ± 1.7 to $9.1 \pm 1.8 \text{ h} \cdot \mu\text{g}/\text{ml}$ ($P = 0.034$). The $AUC_{0 \rightarrow \infty}$ was reduced by 20% in subjects represented by the symbol Δ .

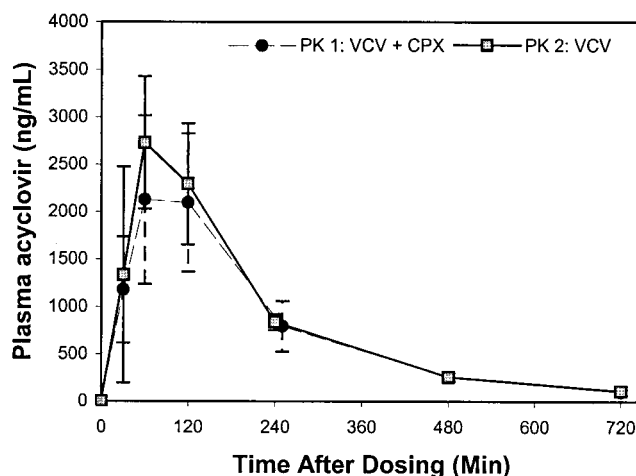


FIG. 3. Mean plasma concentration-versus-time profiles of acyclovir in treatment group A ($n = 5$; 500 mg of valacyclovir [VCV] plus 500 mg of cephalixin [CPX] at PK 1 and 500 mg of VCV at PK 2). Standard deviations are shown as error bars.

ation frequency ($>1\%$) or phenotype (P. Anderle and W. Sadee, unpublished data).

Genotyping of hPEPT1 was performed as previously described (9). Briefly, primers for exons and adjoining intronic regions (ca. 50 bp) for hPEPT1 (National Center for Biotechnology Information reference sequence, NM_005073) were designed by using the Virtual Genome center website at <http://alces.med.umn.edu/VGC.html> and ordered from Operon (Alameda, Calif.). A collection of 247 ethnically identified genomic DNA samples was obtained from the Coriell Institute of Medicine and used to screen for hPEPT1 variants. PCR was performed with TaqGold on the GeneAmp 9700 thermocycler from PE. Samples were pooled 3 deep so that 96 samples were ready for high-performance liquid chromatography (dHPLC).

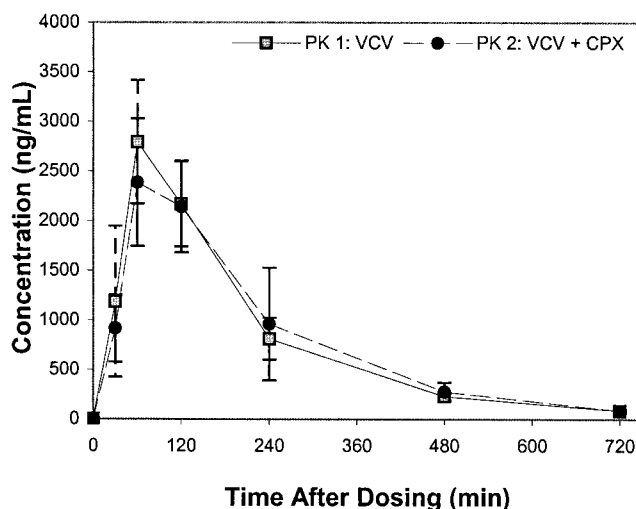


FIG. 4. Mean plasma concentration-versus-time profiles of acyclovir in treatment group B ($n = 5$; 500 mg of valacyclovir [VCV] at PK 1 and 500 mg of VCV plus 500 mg of cephalixin [CPX] at PK 2). Standard deviations are shown as error bars.

dHPLC was performed on Varian HPLC machines with Varian Helix columns. The sequence for each amplicon was submitted to the dHPLC melt program at Stanford University (<http://insertion.stanford.edu/melt.html>). This information was used as the basis for analysis. Typically, the highest recommended temperature was run along with at least one other temperature, depending on the complexity of the melt profile. The results were scored manually. If a well was scored positive, the three PCRs corresponding to that well were cleaned with 2 U of SAP and ExoI (from USB), sequenced with ABI BigDye v2, cleaned with 96-well gel filtration blocks from Edge Bio-Systems, and run on an ABI 3700 DNA analyzer.

Parametric, paired, two-tailed *t* tests were used to determine the statistical significance ($P = 0.05$) of intra- and interindividual variability, the period effect, and the effect of concomitant treatment with cephalexin on acyclovir $AUC_{0-\infty}$.

Sixteen healthy volunteers (nine females, seven males) with a mean age of 27 years (range, 22 to 39 years) enrolled in the study. Their mean height and weight were 167 cm (range, 152 to 182 cm) and 62 kg (range, 45 to 77 kg), respectively. All subjects were calculated to be within their ideal body weight range.

The acyclovir $AUC_{0-\infty}$ values for the control and treatment groups are shown in Fig. 1 and 2. While considerable interindividual variability was observed, no significant intraindividual variability in the AUC was demonstrated ($P = 0.82$) between study periods. Coadministration of cephalexin reduced the acyclovir $AUC_{0-\infty}$ by 7.1% or from 9.8 ± 1.7 to 9.1 ± 1.8 h · $\mu\text{g}/\text{ml}$ ($P = 0.034$). However, this AUC reduction was only observable after exclusion of an outlier who had an increased acyclovir AUC with concomitant cephalexin. In 2 of 10 subjects, oral bioavailability was reduced by 20% with concomitant administration of cephalexin. Figures 3 and 4 display acyclovir concentrations in plasma over time in patients receiving valacyclovir with or without cephalexin. While a reduction in the peak concentration of acyclovir in plasma and the AUC was observed with concomitant administration of cephalexin, no change in the time to the peak concentration of acyclovir in plasma or the terminal half-life took place.

Genotypic analysis did not reveal any contributions of the selected SNPs to the observed responses. In a separate study, none of the nonsynonymous SNPs assayed here had any effect on hPEPT1 transport activity for cephalexin (Anderle and Sadee, unpublished).

The experimental procedures and medications were generally well tolerated. Most subjects reported nonpainful erythema at the site of venous access.

While considerable interindividual variability was observed, the intraindividual oral bioavailability of valacyclovir was remarkably constant. As hPEPT1 is assumed to play a major role in valacyclovir absorption, the stability of hPEPT1 activity, as indicated by minimal intraindividual variability at two observation periods, is consistent with results from animal studies (9, 10). Interindividual variability, while significantly greater than intraindividual variability, did not exceed a twofold range, thereby, supporting the utility of hPEPT1 as a drug target. While a more substantial reduction in valacyclovir absorption (measured by the acyclovir AUC) was associated with concom-

itant administration of cephalexin in some patients, the reduction was less than the 50% decrease suggested by isolated hPEPT1 models (5–7). Furthermore, one patient actually experienced an increased acyclovir AUC with simultaneous administration of cephalexin. One explanation for the discrepancy between studies is that the isolated hPEPT1 models used much higher concentrations and differing ratios of the respective agents. The reduction in bioavailability may be more significant with increased valacyclovir doses, longer duration of therapy, and concomitant administration with a more potent hPEPT1 competitive substrate, such as cefadroxil (11). It is possible that additional subjects with increased genetic polymorphism would produce more substantial differences in bioavailability. The results also suggest that isolated models of hPEPT1 do not account for all of the factors affecting the pharmacokinetics of valacyclovir. It is also possible that cephalexin and valacyclovir are absorbed via other transporters and less dependent on hPEPT1-mediated transport than previously suggested.

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