

Mechanistic population pharmacokinetics of total and unbound paclitaxel for a new nanodroplet formulation versus Taxol in cancer patients

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Abstract

Purpose Our objectives were (1) to compare the disposition and in vivo release of paclitaxel between a tocopherol-based Cremophor-free formulation (Tocosol Paclitaxel®) and Cremophor® EL-formulated paclitaxel (Taxol®) in human subjects, and (2) to develop a

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mechanistic model for unbound and total paclitaxel pharmacokinetics.

Methods A total of 35 patients (average \pm SD age: 59 ± 13 years) with advanced non-hematological malignancies were studied in a randomized two-way crossover trial. Patients received 175 mg/m^2 paclitaxel as 15 min (Tocosol Paclitaxel) or 3 h (Taxol) intravenous infusion in each study period. Paclitaxel concentrations were determined by LC–MS/MS in plasma ultrafiltrate and whole blood. NONMEM VI was used for population pharmacokinetics.

Results A linear disposition model with three compartments for unbound paclitaxel and a one-compartment model for Cremophor were applied. Total clearance of unbound paclitaxel was 845 L/h (variability: 25% CV). The prolonged release with Tocosol Paclitaxel was explained by the limited solubility of unbound paclitaxel of 405 ng/mL (estimated) in plasma. The 15 min Tocosol Paclitaxel infusion yielded a mean time to 90% cumulative input of $1.14 \pm 0.16 \text{ h}$. Tocosol Paclitaxel was estimated to release 9.8% of the dose directly into the deep peripheral compartment. The model accounted for the presence of drug-containing nanodroplets in blood.

Conclusions Population pharmacokinetic analysis indicated linear disposition and a potentially higher bioavailability of unbound paclitaxel following Tocosol Paclitaxel administration due to direct release at the target site. The prolonged release of Tocosol Paclitaxel supports 15 min paclitaxel infusions. This mechanistic model may be important for development of prolonged release formulations that distribute in and from the systemic circulation.

Keywords Paclitaxel · Ultrafiltration · Non-hematological malignancies · Mechanistic modeling · Population pharmacokinetics · NONMEM · Solubility limited disposition model

Introduction

Paclitaxel is a clinically important antitumor agent with activity against a wide array of cancers [27]. Due to the low aqueous solubility of paclitaxel, the most commonly used paclitaxel formulation (Taxol) contains Cremophor EL (polyoxyethylated castor oil) and ethanol (1:1, w/w) to solubilize paclitaxel. Cremophor is associated with hypersensitivity reactions due to complement activation [8, 42, 44]. Other studies [2, 47] suggest that Cremophor is involved in peripheral sensory neuropathy. Cremophor affects the pharmacokinetics (PK) of paclitaxel by entrapment of paclitaxel inside Cremophor micelles which are major “binding” sites for paclitaxel in vitro [34], in animals [32, 33], and in humans [14, 16]. Other binding sites for paclitaxel are albumin and α_1 -acid glycoprotein [22], red blood cells [34], and platelets [46].

Hepatic metabolism and biliary excretion are the primary routes of paclitaxel elimination [23, 43]. Non-linear PK of total paclitaxel in blood was associated with saturable elimination [30], saturable distribution [19], and binding to Cremophor [14, 16, 34, 40–42]. This non-linearity seems to be less pronounced if the duration of infusion is longer than 6 h [4, 45]. More recent studies measured unbound paclitaxel concentrations and found linear disposition of unbound paclitaxel [9, 15, 16].

For PK comparison of a Cremophor-containing formulation and a Cremophor-free formulation, it is important to measure both unbound and total paclitaxel concentrations. We studied a new tocopherol-based, Cremophor-free paclitaxel formulation (Tocosol Paclitaxel) in comparison to Taxol. Two key advantages of Tocosol Paclitaxel are the absence of Cremophor-related toxicity and the ability to administer Tocosol Paclitaxel as a 15-min infusion (compared to 3 or 24 h infusion for Taxol). This rapid infusion is possible, since paclitaxel is contained within nanodroplets that release paclitaxel over time. Tocosol Paclitaxel nanodroplets have a diameter of 40–80 nm [5, 6], are stable and neutrally charged. These nanodroplets have the potential to distribute into tumor tissue and release a fraction of the paclitaxel dose directly at the target site, since pore sizes of many tumor tissues are larger than the tight endothelial junctions of normal tissue vasculature [28]. Direct release of drug from drug-containing nanodroplets that distribute to the target site would result in a greater unbound area under the concentration time curve (AUC) at the target site compared to the AUC in plasma. This may reduce the toxicity of paclitaxel at sites not accessible to Tocosol Paclitaxel nanodroplets.

Population PK methodology was used to consider direct paclitaxel release to a peripheral compartment. Assessing the presence of paclitaxel containing nanodroplets in blood

samples requires a modeling approach, as unreleased paclitaxel in nanodroplets contributes to the observed concentrations. Additionally, it seems likely that some drug may be released from nanodroplets during the centrifugation of blood samples. The possible contribution of these processes was examined for the data collected using a modeling approach.

Our first objective was to compare the PK of paclitaxel in plasma ultrafiltrate and in whole blood between Tocosol Paclitaxel and Taxol by population modeling. Secondly, we sought to characterize the rate and extent of paclitaxel release from the Tocosol Paclitaxel formulation accounting for the presence of drug-containing nanodroplets in blood. We were especially interested in quantifying the amount of paclitaxel that was released directly into a peripheral compartment, since direct release at the target site would increase the availability at the site of action.

Materials and methods

Study design and patients

The study was a randomized, two-period, open label, multicenter (6 clinical sites), crossover study in 36 adult patients. The inclusion criteria were: (1) histological diagnosis of advanced non-hematological malignancy for which there was no curative therapy and for which treatment with a single agent taxane was appropriate according to the opinion of the treating physician, (2) no more than two prior cytotoxic chemotherapy regimens for metastatic disease (not including adjuvant or neoadjuvant chemotherapy for primary disease), (3) adequate hematological, renal and liver function, (4) peripheral neuropathy grade 0–1 per NCI-CTC, (5) Eastern Cooperative Oncology Group (ECOG) [26] performance status of 0–2, (6) absolute neutrophil count above 2,000 cells/mm³, and (7) platelet count above 100,000/mm³. The study was approved by all relevant Ethics Committees and Institutional Review Boards and was conducted following the revised version of the Declaration of Helsinki. Written informed consent forms were signed by each patient prior to inclusion in the study.

Drug administration

Each patient received a dose of 175 mg/m² paclitaxel (about 300 mg for a patient with 1.73 m² body surface area, BSA) as Tocosol Paclitaxel in study period I followed by the same dose of Taxol in study period II or vice versa. The washout period was 3 weeks. Paclitaxel was given as a constant rate infusion over approximately 15 min for Tocosol Paclitaxel and 3 h for Taxol.

Paclitaxel was infused intravenously with a controlled-rate infusion pump. The infusion rate was about 2 mL/min for Tocosol Paclitaxel. A paclitaxel concentration of 10 mg/mL yielded a total infused volume of about 30 mL for a patient with 1.73 m² BSA. Syringes, infusion pumps and intravenous (IV) tubing sets were provided by Sonus Pharmaceuticals. Patients were monitored closely and asked about any symptoms or adverse effects during the 15 min infusion. In case any symptoms occurred, the infusion was interrupted and resumed within 30 min, if resuming was deemed safe by the responsible physician. The exact infusion times and volumes were recorded and used for PK modeling. According to the clinical protocol, all patients were treated with a fixed regimen of antihistamines (H₁ and H₂ blockade) and glucocorticoids 30–60 min prior to drug administration in study phases I and II. This included diphenhydramine 50 mg IV, cetirizine 10 mg PO, or clemastine 2 mg IV as well as ranitidine 50 mg IV and dexamethasone 20 mg IV. Granisetron 10 µg/kg IV at 30–60 min before dosing was an optional premedication.

The appropriate dose of Taxol was prepared as an IV admixture in 0.9% saline or 5% dextrose in water to yield a final concentration of approximately 1 mg of paclitaxel per mL of dosing solution (total volume: about 300 mL for a patient with 1.73 m² to be infused within 3 h). Taxol was infused at a constant rate of about 1.67 mL/min via a standard infusion set with a 0.22 µm in-line filter. Only glass, polypropylene and polyolefin containers were used, as Taxol solutions have been shown to extract DEHP plasticizer from PVC containers according to the Taxol package insert. Patients were carefully monitored and asked about symptoms and adverse effects during the infusion.

Blood sampling

In each study period, 4 mL of blood was taken at 18 specified time points. For Tocosol Paclitaxel, samples were taken immediately before start of the infusion, at 5, 10, and 15 min post-start and 5, 15, 30, and 60 min after end of Tocosol Paclitaxel infusion as well as at 2, 3, 5, 8, 14, 24, 48, 72, 96, and 120 h. For Taxol, blood sampling times were immediately before start of infusion and 1, 2, and 3 h post start, 5, 15, 30, and 45 min post end of infusion as well as at 2, 4, 6, 9, 14, 24, 48, 72, 96, and 120 h. If the infusion was interrupted, additional blood samples were taken immediately after such interruption and immediately before resuming infusion.

Blood was collected into EDTA tubes and samples were kept in a temperature controlled water bath at 37°C until processing. Two aliquots each of 0.5 mL whole blood were prepared from each blood sample. The remaining 3 mL blood was centrifuged at 10,000g for 15 min at 37°C to prepare two plasma aliquots from each sample. Blood and

plasma samples were frozen immediately and stored at –20°C prior to analysis.

Drug analysis

Preparation of plasma ultrafiltrate specimens from patient blood samples began within 20 min of drawing each blood sample and was completed without delay. Whole blood and plasma ultrafiltrate samples were frozen at the site, and shipped frozen to the analytical lab, where the samples were extracted by methyl tert-butyl ether and analyzed for paclitaxel concentrations using validated methods. The assay was validated from 0.1 to 50 ng/mL for plasma ultrafiltrate using an extraction volume of 80 µL and from 5 to 5,000 ng/mL for whole blood using an extraction volume of 100 µL. The method used liquid chromatography coupled to tandem mass spectrometry (LC–MS/MS), which included a Perkin Elmer Series 200 Micro LC pump (Perkin Elmer, Waltham, MA), a CTC-PAL auto sampler (LEAP Technologies, LLC, Carrboro, NC), and an API 3000 or 4000 mass spectrometer with a TurboIon Spray[®] source (Applied Biosystems, Foster City, CA). Chromatographic separation was performed on a Phenomenex Curosil-PFP column (3 µm, 50 × 2 mm, Phenomenex Inc, Torrance, CA) at a flow rate of 0.5 mL/min using a linear gradient elution of water and acetonitrile (0–2.0 min: 30–70% acetonitrile; 2.0–2.5 min: 70–90% acetonitrile; 2.5–2.7 min: 90% acetonitrile; and 2.7–2.9 min: 90–30% acetonitrile). Detection of analytes was by positive-ion mode using multiple reaction monitoring scan mode, with ion transitions *m/z* 854 → 286 for paclitaxel and *m/z* 860 → 292 for ¹³C₆-paclitaxel (internal standard). The lower limit of quantification was 0.1 ng/mL in plasma ultrafiltrate and 5 ng/mL in whole blood.

Correlation coefficients for regression of calibration curves were at least 0.984 for plasma ultrafiltrate and at least 0.988 for whole blood. For back-calculated standard concentrations, mean accuracy (% bias) ranged from –1.7 to 2.2% for ultrafiltrate and from –3.4 to 4.8% for whole blood; the precision (% coefficient of variation, %CV), ranged from 5.4 to 8.4% for ultrafiltrate and from 6.1 to 7.8% for whole blood. For quality control samples, the accuracy ranged from –20.3 to 9.8% for ultrafiltrate and from –9.9 to 8.5% for whole blood; the precision ranged from 8.0 to 15.9% for ultrafiltrate and from 8.4 to 14.3% for whole blood.

Laboratory testing

A complete blood count, chemistry panel, prothrombin time, partial thromboplastin time and pregnancy test (if applicable) were performed within 1 week before study drug administration in study period 1 and within 1 day before administration in study period 2. Additionally, a complete blood count including absolute neutrophil count

Table 1 Population pharmacokinetic parameter estimates and confidence intervals (CI)

Parameter	Symbol	Unit	Estimate (90% CI)	Variability (90% CI) ^a	ΔObj ^d
Disposition of paclitaxel					
Total clearance ^b	CL	L/h	845 (783–910)	25% (15–37%)	
Volume of central compartment ^b	V1	L	610 (430–718)	27% (0.3–40%)	
Volume of shallow peripheral compartment ^b	V2	L	1480 (1260–1710)	19% (3.8–27%)	
Volume of deep peripheral compartment ^b	V3	L	6830 (6080–7561)	24% (16–29%)	
Intercompartmental clearance to V2 ^b	CL _{d2}	L/h	724 (631–918)	23% (0.2–33%)	
Intercompartmental clearance to V3 ^b	CL _{d3}	L/h	317 (274–371)	23% (14–32%)	
Release from Tocol Paclitaxel nanodroplets					
Relative extent of bioavailability for Tocol Paclitaxel	F_{Tocol}		0.940 (0.882–0.997)	(F_{Taxol} fixed to 1)	9.9 ^d
Fraction of dose directly entering the deep peripheral compartment from nanodroplets	Fr3		0.098 (0.0443–0.153)		32
Maximum rate of release from nanodroplets	V_{max}	mg/h	356 (301–424)		
Amount of drug in nanodroplets for that the release rate is half-maximal	AN ₅₀	mg	50.0 (27.7–78.8)	47% (34–64%)	233 ^d
Solubility of unbound paclitaxel in plasma	C_{Sol}	ng/mL	405 (368–452)		155 ^d
Volume of distribution of nanodroplets ^b	V_{Nano}	L	6.66 (6.03–7.35)	21% (16–24%)	29 ^d
Half-life of release during sample handling if no saturation ^e	$t_{1/2}(\text{ka})$	min	26.8 ^e (19.8–36.2)	84% (56–120%)	936 ^d
Disposition of Cremophor					
Total clearance of Cremophor ^b	CL _{CrEL}	L/h	0.583 (0.257–0.970)	203% (96–405%)	491
Volume of distribution of Cremophor ^b	V_{CrEL}	L	6.01 (fixed)		
Paclitaxel binding					
Linear binding to plasma proteins and red blood cells	B_{P}		21.0 (19.0–22.6)	13% (4.1–17%)	51
Maximum binding capacity divided by K_{d}	$B_{\text{Sat,P}}/K_{\text{d}}$		8.01 (3.72–13.0)	103% (48–200%)	51
Dissociation constant	K_{d}	ng/mL	4.38 (3.02–6.99)		
Linear binding of paclitaxel to Cremophor ^c	B_{CrEL}	L/g	6.61 (5.65–7.93)		491
Unbound concentration					
Proportional error		%	30.2 (27.0–32.9)		
Additive error		ng/mL	0.110 (0.0802–0.143)		
Total concentration					
Proportional error		%	17.3 (15.6–18.7)		
Additive error		ng/mL	2.25 (1.73–2.79)		

^a Estimates are apparent coefficients of variation for the between-subject variability^b Per 1.73 m² body surface area^c Cremophor dose was 0.527 g purified Cremophor EL per 6 mg paclitaxel according to the Taxol package insert^d Increase in objective function after estimation of the simplified model, if the parameter mean was fixed to zero (exceptions: F_{Tocol} fixed to 1.0, V_{Nano} fixed to 5 L representing plasma volume, k_{a} fixed to 0/h, and C_{Sol} and AN₅₀ fixed to very large values). The final model was used as reference. An increase in the objective function by 3.84 for one degree of freedom (df) or by 5.99 for two df is associated with an α of 0.05 according to a chi-square distribution (likelihood ratio test)^e The release from nanodroplets during sample handling was assumed to be saturable due to the solubility of paclitaxel in plasma. The model predicted that a small fraction (on average about 1%) of unreleased paclitaxel in nanodroplets contributed to the observed concentration in plasma ultrafiltrate (see also Figs. 1, 5)

that is released directly into the DPC is Fr3. The remainder fraction of dose ($1 - \text{Fr3}$) is released simultaneously into the central compartment. Models with a direct release into the shallow peripheral compartment were also considered.

Observation model for Tocol Paclitaxel

The following output equation was used to describe the total paclitaxel concentration (C_{t}) in whole blood for Tocol Paclitaxel:

$$C_t = \frac{A_{\text{Nano}}}{V_{\text{Nano}}} + \frac{A_1 \cdot \text{BF}_{\text{TOC}}}{V_1} \quad (5)$$

$$\text{BF}_{\text{TOC}} = \left(1 + B_P + \frac{B_{\text{Sat,P}}}{K_d + C_1} \right). \quad (6)$$

The binding factor (BF_{TOC}) for Tocosol Paclitaxel is the ratio of total blood to unbound plasma concentration and B_P describes the linear binding and $B_{\text{Sat,P}}$ (maximum binding capacity) and K_d (dissociation constant) the saturable binding of paclitaxel to proteins and blood cells. It was assumed that the volume of distribution for nanodroplets includes plasma and that elimination of nanodroplets during the release phase was negligible. We assumed that paclitaxel is released from drug-containing Tocosol nanodroplets during processing of blood samples and assumed such time averaged 30 min. This choice only affects the estimate for half-life of drug release during centrifugation. The output equation for the drug concentration in plasma ultrafiltrate is derived in Appendix for Tocosol Paclitaxel.

Input and disposition models for Taxol

The drug input for Taxol was specified as a time-delimited zero-order input of paclitaxel into the central paclitaxel compartment and of Cremophor into the (central) Cremophor compartment (Fig. 1). For Taxol, the same disposition models as for Tocosol Paclitaxel were considered and the linear model for disposition of unbound paclitaxel as shown above for Tocosol Paclitaxel except for the saturable release from Tocosol nanodroplets was used as the final model.

Disposition of Cremophor was assessed with one-, two-, or three-compartment models with first-order elimination. As Cremophor concentrations were not measured, the volume of distribution for Cremophor was fixed to a literature estimate [38] of 6.01 L. As Cremophor concentrations were not measured in this study, the choice of this volume only affects the estimate for the Cremophor clearance and the parameter(s) for binding of paclitaxel to Cremophor but no other model parameter. Additionally, we incorporated the model from van den Bongard et al. [38] to describe the disposition of Cremophor.

Observation model for Taxol

The following output equation was used to describe the total concentration of paclitaxel in whole blood based on the modeled Cremophor concentration (C_{CrEL} , equation for Cremophor disposition not shown):

$$C_t = \frac{A_1 \cdot \text{BF}_{\text{Taxol}}}{V_1} \quad (7)$$

$$\text{BF}_{\text{Taxol}} = \left(1 + B_P + \frac{B_{\text{Sat,P}}}{K_d + C_1} + B_{\text{CrEL}} \cdot C_{\text{CrEL}} \right). \quad (8)$$

The binding factor for Taxol (BF_{Taxol}) contained one term describing the linear binding of paclitaxel to Cremophor (B_{CrEL}) in addition to the same terms as the binding factor for Tocosol Paclitaxel. It was assumed that the (central) volume of distribution for Cremophor includes the plasma volume. For Taxol, the paclitaxel concentration in plasma ultrafiltrate was the paclitaxel concentration in the central compartment.

Size model

The BSA was calculated by the Mosteller formula [24] and used to describe body size and to scale all clearance and volume terms linearly by BSA. The fractional change ($F_{\text{Size},i}$) in clearance and volume of distribution for the i th subject (with BSA_i) standardized to a BSA_{STD} of 1.73 m² is: $F_{\text{Size},i} = \text{BSA}_i / \text{BSA}_{\text{STD}}$.

Parameter variability model for PK

The between-subject variability (BSV) of PK parameters was estimated by assuming a log-normal distribution. The η_{BSV_i} is the log scale difference of the individual PK parameter estimate from its population mean for the i th subject. It is assumed that η_{BSV} is a normally distributed random variable with mean zero and standard deviation BSV. The BSV was estimated as variance. We report the %CV of a normal distribution on log-scale calculated by the formula $\%CV = [\exp(\text{variance}) - 1]^{0.5}$. The individual PK parameters were calculated as $\text{CL}_i = \text{CL}_{\text{POP}} \cdot F_{\text{Size},i} \cdot \exp(\eta_{\text{BSVCL},i})$.

The CL_{POP} is the group estimate of clearance for a patient of standard body size (BSA_{STD}), CL_i is the individual clearance for the i th subject, and $\eta_{\text{BSVCL},i}$ is the individual log-scale difference for clearance in the i th subject from the population mean. Similar formulas apply to the other PK parameters. Individual disposition parameter estimates were assumed to be the same for both formulations. Between-occasion variability of PK parameters was assumed to be small and was not included in the final model.

Residual unidentified variability

The residual unidentified variability was described by a combined additive and proportional error model for paclitaxel concentrations in plasma ultrafiltrate and whole blood.

Estimation

NONMEM[®] version VI level 1.1 (NONMEM Project Group, University of California, San Francisco, CA) [1] was used for population PK analysis. The first-order conditional estimation (FOCE) method with the interaction

estimation option in NONMEM was used for all analyses. Confidence intervals for model parameters were obtained by non-parametric bootstrapping with 35 randomly selected patients in each of the 500 bootstrap replicates.

Model qualification

The models were compared by their predictive performance, the objective function in NONMEM, individual model fits, observed versus predicted plots, and standard diagnostic plots. Predictive performance was assessed by visual predictive checks based on at least 6,000 virtual patients.

Monte Carlo simulations

Monte Carlo simulations were run in absence of residual error for 1,000 virtual patients to assess the paclitaxel release profile of Tocosol Paclitaxel and the disposition of Tocosol Paclitaxel and Taxol and calculate the median and 80% non-parametric prediction interval.

Non-compartmental analysis (NCA), statistics, and software

WinNonlinTM Professional (version 5.0.1, Pharsight Corp., Mountain View, CA) was used for NCA. The ANOVA and equivalence statistics with factors accounting for sequence, subject nested within sequence, period and treatment effects was obtained with SAS (version 9.1.3 for Windows, SAS Institute, Cary, NC). Maple[®] 10.05 (Maplesoft, Waterloo, ON, Canada) was used to explicitly solve the differential equations for the output equation.

Results

Patients

One patient dropped out after having received 48 mg Tocosol Paclitaxel during the first 2 min of the infusion. This patient was excluded from all PK analyses, as no post-dose blood samples were collected. The remaining 35 patients had heights of 167 ± 9.7 cm, total body weights of 68.5 ± 17.3 kg, BSA of 1.77 ± 0.23 m², and ages of 58.6 ± 12.5 years (mean \pm SD). Six patients were African Americans (5 females/1 male) and 29 patients were Caucasians (15 females/14 males). The ECOG performance status was 0 for 4 patients, 1 for 19 patients, and 2 for 12 patients. Type of primary cancer was breast in 9 patients, gastrointestinal in 6 patients, head and neck in 5 patients, lung in 3 patients, and 'other' in 12 patients.

Of those 35 patients, 7 patients (2 patients for Taxol and 5 patients for Tocosol Paclitaxel) had their paclitaxel infusion interrupted and resumed within 30 min. Three patients (1 patient for Tocosol Paclitaxel and 2 patients for Taxol) dropped out before receiving the dose in study period II. As those patients contributed data from study period I, these three patients were included in the population PK analysis. However, these three patients and one patient with insufficient data for NCA in study period I were excluded from NCA and ANOVA statistics.

NCA

Figure 2 shows the median plasma concentrations for paclitaxel in plasma ultrafiltrate and in whole blood. Ultrafiltrate concentrations for Tocosol Paclitaxel revealed the prolonged release of paclitaxel from nanodroplets. Total concentrations showed a sharp peak at 0.25 h for Tocosol Paclitaxel. There was no apparent prolonged release for Taxol.

The following data are geometric means (%CV). Area under the curve from time zero to infinity was 603 ng h/mL (22%) for Tocosol Paclitaxel and 360 ng h/mL (23%) for Taxol in plasma ultrafiltrate and was 29.9×10^3 ng h/mL (27%) for Tocosol Paclitaxel and 14.3×10^3 ng h/mL (27%) for Taxol. Total clearance was 511 L/h (28%) for Tocosol Paclitaxel and 852 L/h (29%) for Taxol in plasma ultrafiltrate and was 10.3 L/h (33%) for Tocosol Paclitaxel and 21.4 L/h (32%) for Taxol in whole blood. Terminal half-life was 26.0 h (26%) for Tocosol Paclitaxel and 25.2 h (28%) for Taxol in plasma ultrafiltrate and was 21.5 h (26%) for Tocosol Paclitaxel and 22.9 h (37%) for Taxol in whole blood. The ratio of AUC for Tocosol Paclitaxel divided by Taxol was 1.67 (1.55–1.80) for ultrafiltrate concentrations and 2.08 (1.97–2.20) for whole blood concentrations [point estimate (90% confidence interval) from ANOVA].

Population PK model

Visual predictive checks and the log-likelihood significantly favored a three- versus two-compartment disposition model. After accounting for the time-dependent binding of paclitaxel to Cremophor due to elimination of Cremophor and for the unreleased drug in nanodroplets in blood samples, models with saturable elimination or saturable distribution neither improved the objective function nor the predictive performance compared to models with first-order disposition of unbound paclitaxel.

Linear and saturable binding to plasma proteins and red blood cells (combined) and linear and saturable binding to Cremophor were considered. The final model included linear and saturable binding to plasma proteins and red

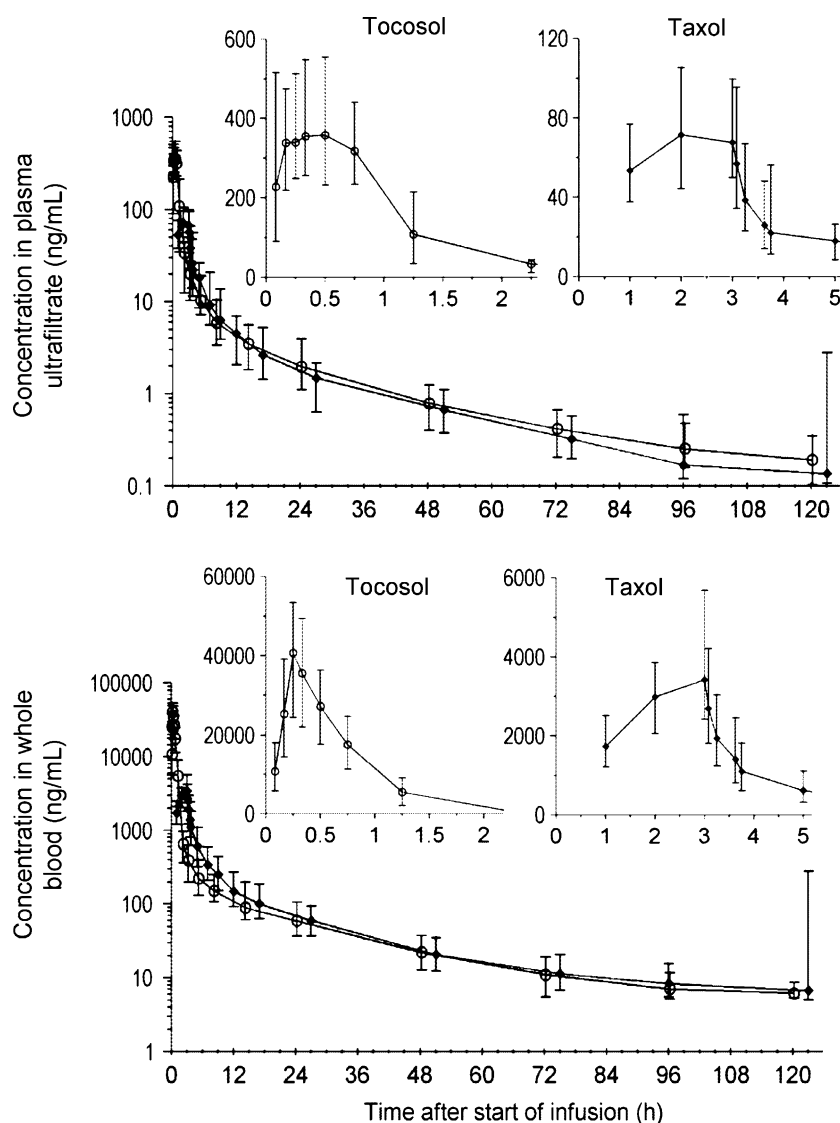


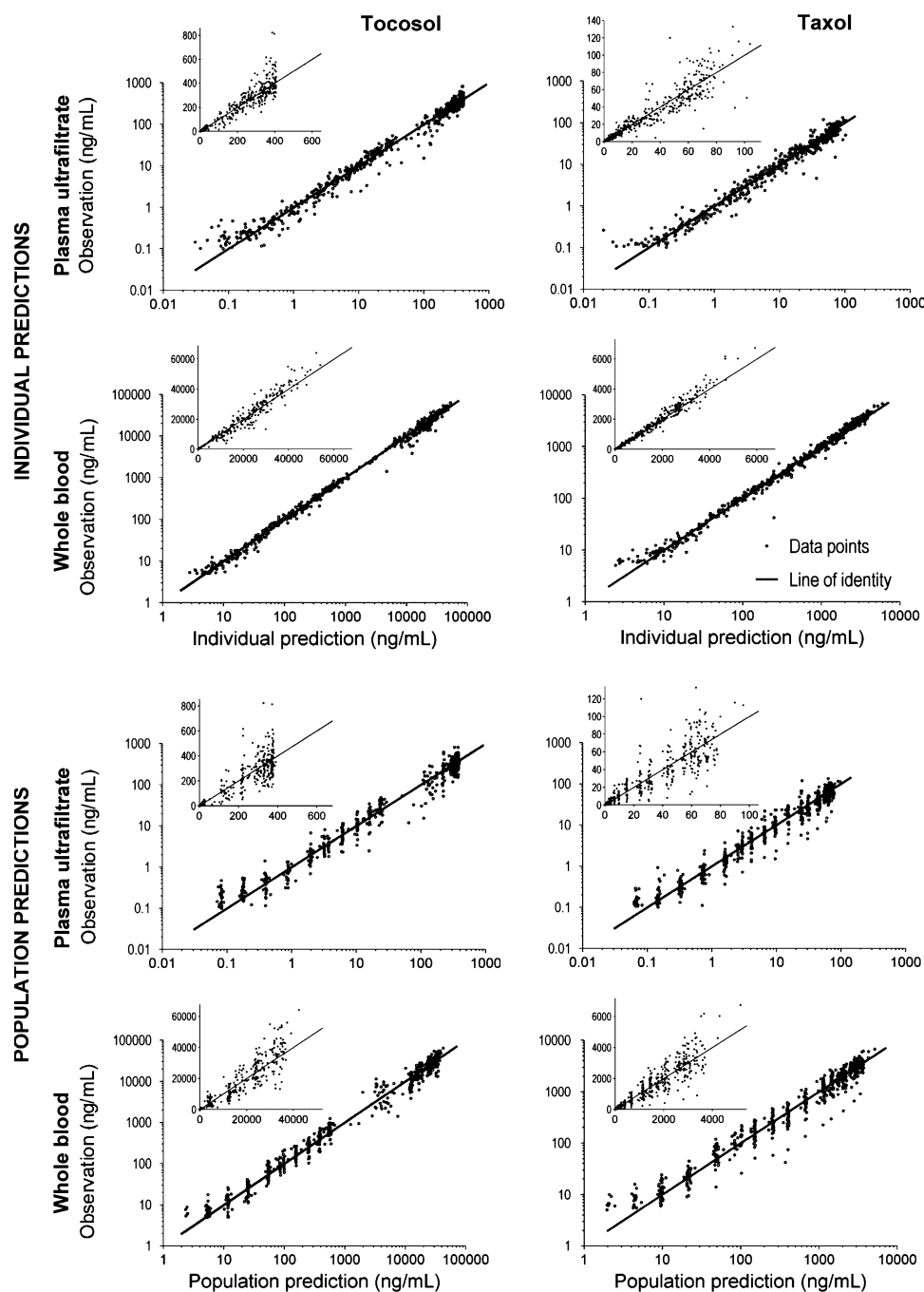
Fig. 2 Median (10–90% percentile) paclitaxel concentrations versus time after administration of Tocosol Paclitaxel (left, open circles) and Taxol (right, filled diamonds). Inserts show early time profiles

blood cells and linear binding to Cremophor (Fig. 1). Use of a one-compartment PK model for Cremophor significantly improved the model fits and log-likelihood compared to models with a constant Cremophor concentration. When multi-compartment models for Cremophor with first-order elimination were estimated, the objective function improved by 41 for the two-compartment Cremophor model and by 49 for the three-compartment Cremophor model both relative to the one-compartment model. However, the individual curve fits, the population predicted versus observed and individual predicted versus observed plots were very similar for one-, two- and three-compartment models for Cremophor. As we did not measure Cremophor concentrations, the one-compartment model was chosen following the rule of parsimony. Use of

the population PK model for Cremophor from van den Bongard et al. [38] yielded a considerably worse objective function (increase by 118).

Drug input for Taxol was described adequately by zero-order input into the central compartment. For Tocosol Paclitaxel, a mixed-order release from the nanodroplet to the central compartment (Fig. 1) improved the predictive performance and the log-likelihood significantly compared to models with first-order release. The output equations accounted for drug entrapped in Cremophor micelles (Taxol only) and unreleased drug in nanodroplets (Tocosol Paclitaxel only). The volume of distribution of the nanodroplets was estimated to be 6.66 L (21% CV). The objective function improved significantly, when release of paclitaxel during sample handling was included (Fig. 1).

Fig. 3 Observations versus individual predictions (*top rows*) and observations versus population predictions (*bottom rows*) for paclitaxel concentrations in plasma ultrafiltrate and whole blood on logarithmic scale (*inserts* show the same data on linear scale). The assumed solubility limit of unbound paclitaxel that was estimated as 405 ng/mL causes the boundary in the *top left panel* (see Fig. 1; Table 1)



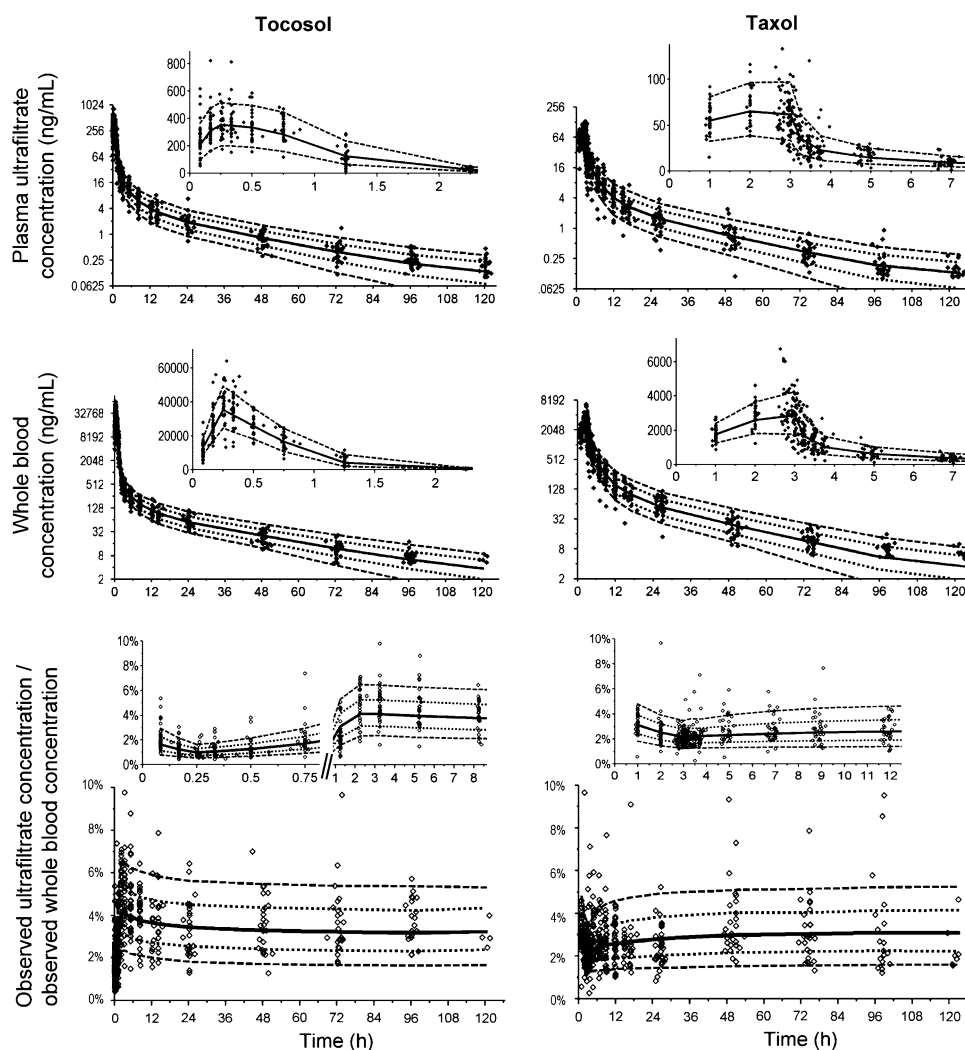
Without this feature, model fits were biased and models could not predict the complex time course for the ratio of concentrations in plasma ultrafiltrate and whole blood.

The final population PK parameter estimates are shown in Table 1. Plots of observed versus individual predicted concentrations (Fig. 3, top) indicated unbiased and precise fits for plasma ultrafiltrate and whole blood for both treatments. There was no bias for observed concentrations versus population predictions (Fig. 3, bottom). As these predicted versus observed plots indicated precise and unbiased fits without including between occasion

variability in the model, between occasion variability was probably rather small and was not included in the final model. Visual predictive checks of the final population PK model revealed excellent predictive performance for Tocosol Paclitaxel and Taxol, as the central tendency and variability of the observed concentrations were well captured (Fig. 4). The model also captured the time course of the ratio of observed concentrations in plasma ultrafiltrate and whole blood.

Allowing a small amount of the dose to be released from nanodroplets into the DPC improved the objective function

Fig. 4 Visual predictive check for paclitaxel concentrations in plasma ultrafiltrate and in whole blood based on 10,000 simulated, virtual patients. Data are median, 25–75% percentile (dotted line), and 10–90% percentile (dashed line). Ideally, 10% of the observations should fall outside the 10–90% percentile of simulated concentrations at each time point on each side



by 32.1 points ($P < 0.001$, likelihood ratio test). This fraction of dose was estimated to be 9.8%. Release from nanodroplets into the shallow peripheral compartment did not improve the objective function and the associated fraction of dose was estimated to be less than 1%. Therefore, direct release into the shallow peripheral compartment was not included in the final model.

The increase in the objective function after exclusion of one specific model component is shown in Table 1 with the final model as reference. After estimation of the simplified population PK models, the objective function favored the full model significantly ($P < 0.01$, likelihood ratio test) compared to each simplified model.

Monte Carlo simulations for a 15 min infusion of 300 mg Tocosol Paclitaxel showed that 90% of the dose was released into the central compartment or DPC after 1.14 ± 0.16 h (average \pm SD). The prolonged release profile for Tocosol Paclitaxel nanodroplets is shown in Fig. 5a. During the first hour, the amount of drug in

nanodroplets was the major fraction of the observed concentration in whole blood (Fig. 5b). These results were qualitatively consistent with an equilibrium dialysis study that showed a significantly slower in vitro release of paclitaxel for Tocosol Paclitaxel compared to Taxol (data on file at Sonus). The release from nanodroplets during sample handling was assumed to be saturable due to the solubility of paclitaxel in plasma. The model predicted that a small fraction (on average about 1%) of unreleased paclitaxel in nanodroplets contributed to the observed concentration in plasma ultrafiltrate. This small fraction causes plasma ultrafiltrate concentrations to be greater than unbound plasma concentrations during the first hour (Fig. 5c).

We simulated the unbound plasma concentrations which differ from ultrafiltrate concentrations in that unbound concentrations do not contain any drug released from nanodroplets during sample processing. A comparison of unbound plasma concentrations for Tocosol Paclitaxel and Taxol is shown in Fig. 5 for unbound concentrations in the

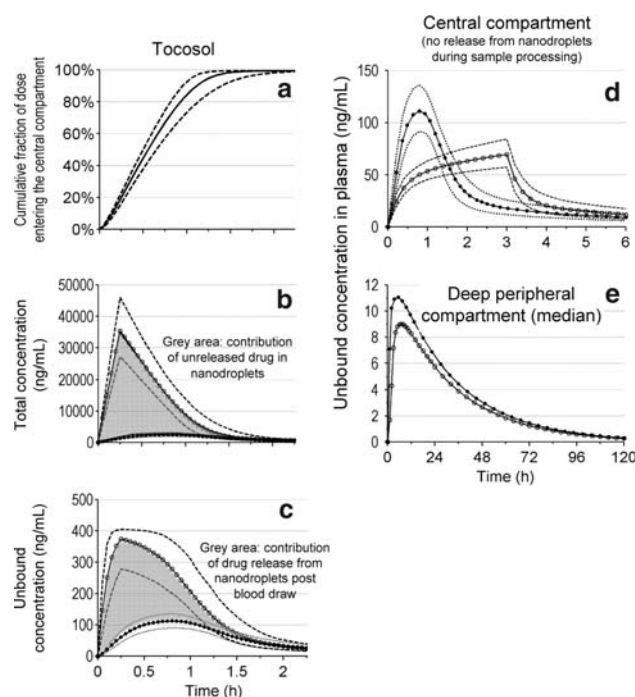


Fig. 5 Median (10–90% percentile) from 1,000 simulated patients with 1.73 m² BSA for a dose of 300 mg paclitaxel given as 15 min infusion (*Tocosol Paclitaxel*) or as 3 h infusion (*Taxol*) in absence of residual error; **a** cumulative fraction of dose reaching the central compartment, **b** observed total concentration (*open circle*) and total concentration without paclitaxel in nanodroplets (*filled circle*), **c** unbound concentration in plasma ultrafiltrate (*open circle*) and unbound plasma concentration without contribution from nanodroplets (*filled circle*), **d** and **e** unbound plasma concentration in central and deep peripheral compartment without release from nanodroplets sample processing (*Tocosol Paclitaxel*: *filled circle*, *Taxol*: *open circle*)

central compartment (d) and unbound concentrations in the DPC (e). Despite the substantially shorter duration of infusion (15 min for *Tocosol Paclitaxel* vs. 3 h for *Taxol*), the peak of the unbound concentrations was only about 60% higher for *Tocosol Paclitaxel* than for *Taxol* (Fig. 5d). The relative paclitaxel exposure in the central compartment for *Tocosol Paclitaxel* versus *Taxol* was 94% with a 90% confidence interval from 88 to 100%.

Discussion

A mechanistic population PK model is proposed that describes the release and disposition of paclitaxel in plasma ultrafiltrate and whole blood for a new tocopherol-based, Cremophor-free paclitaxel formulation and for *Taxol*. The model described simultaneously the release and disposition of unbound paclitaxel in plasma and binding to Cremophor, proteins, and red blood cells. The kinetics of nanodroplets and disposition of Cremophor were incorporated (Fig. 1). The final model had excellent predictive performance for

paclitaxel concentrations in whole blood and plasma ultrafiltrate (Fig. 4) and could predict the complex time course of the ratio of plasma ultrafiltrate to whole blood concentrations. The low plasma ultrafiltrate to whole blood concentration ratios of about 1–2% from 0 to 1 h for *Tocosol Paclitaxel* (Fig. 4, bottom left panel) were consistent with the assumption that blood samples contained unreleased paclitaxel in nanodroplets (Fig. 5b).

Accounting for the release of paclitaxel during sample processing removed a systematic bias in the model fits and improved the objective function and predictive performance significantly. Although nanodroplets are stable for several hours in plasma (data on file at Sonus Pharmaceuticals), centrifugation to separate plasma from blood cells or ultrafiltration may release a small fraction of drug from the nanodroplets. The model predicted that on average only about 1% of the total unreleased paclitaxel in nanodroplets contributed to the observed concentration in plasma ultrafiltrate. As Cremophor is considered a high-affinity “binding site” for paclitaxel and as Cremophor micelles persist for several hours after dilution to concentrations below the critical micelle concentration of 0.009% (w/v, at equilibrium) [21, 34], no net release of paclitaxel from Cremophor micelles following blood sample collection was assumed.

The point estimates (90% confidence intervals) from NCA and ANOVA for the relative extent of bioavailability were 1.67 (1.55–1.80) for paclitaxel in plasma ultrafiltrate and 2.08 (1.97–2.20) for paclitaxel in whole blood for *Tocosol Paclitaxel* compared to *Taxol*. The PK model accounted for the time-course of drug release from nanodroplets and for the presence of drug-containing nanodroplets in blood samples (Figs. 1, 5b, c). As these processes cannot be accounted for by standard non-compartmental techniques, the NCA seems likely to provide biased estimates for drug exposure of slow release formulations given intravenously. In these cases, a modeling approach seems more appropriate. The estimated relative extent of bioavailability in the central compartment (90% confidence interval) from population PK modeling was 94% (88–100%). This estimate seems reasonable from a modeling perspective for two formulations administered intravenously. However, this estimate probably does not reflect the bioavailability at the site of toxicity (see companion article).

An alternative reason for a higher AUC for unbound paclitaxel would be a saturation of clearance at high concentrations. Earlier studies [11, 18–20, 30] presumed non-linearity in PK of total paclitaxel concentrations by saturable elimination and saturable distribution. More recent studies [9, 14–16, 32, 34] explained the apparent non-linearity of total paclitaxel by an entrapment of paclitaxel within Cremophor micelles with Cremophor concentrations

changing over time and dependent on Cremophor dose. Several studies [9, 14–16] successfully described the PK of unbound paclitaxel by linear two- or three-compartment models. A short (15 min) Tocosol Paclitaxel infusion yielded linear paclitaxel elimination and linear distribution for observed concentrations in plasma ultrafiltrate up to about 500 ng/mL. Human liver microsome data for CYP2C8 metabolism showed a high average Michaelis–Menten constant of 9 μM (equivalent to 7.7×10^3 ng/mL) [15]. Linear elimination was also found for docetaxel formulated with polysorbate 80 that is degraded rapidly in plasma [39]. These studies support first-order elimination of paclitaxel.

Our estimated terminal half-life of unbound paclitaxel was in good agreement with the terminal half-life from Henningsson et al. [15] who used a comparable sampling schedule. The estimate for total clearance of paclitaxel in whole blood (Table 1) was within the reported range of clearances for total paclitaxel [7, 13, 17, 25]. Our unbound clearance of 845 L/h (25% CV for BSV) was higher than the range of average unbound clearances from 301 to 429 L/h in the literature [9, 14–16]. The BSV in unbound clearance was up to 44% with individual clearances ranging from 83.7 to 1,055 L/h [15]. Literature studies [9, 14–16] applied equilibrium dialysis with a tritiated paclitaxel tracer [3] to determine unbound concentrations, whereas paclitaxel in plasma ultrafiltrate was determined by LC–MS/MS in this study. This could have contributed to the differences in clearance of unbound paclitaxel. Further studies are required to resolve these differences.

As we did not measure Cremophor, the volume of distribution of Cremophor was fixed to the estimate from van den Bongard et al. [38]. Our estimate for Cremophor clearance of 0.583 L/h was at the upper end of the range of Cremophor clearances in literature [9, 10, 35, 41], if areas up to the last quantifiable concentration were used for clearance calculation. The median of the individual Cremophor half-lives was 10.6 h in this study. This half-life was estimated from binding of paclitaxel to Cremophor that changes over time due to elimination of Cremophor. Cremophor is a non-ionic surfactant consisting of a complex mixture of chemical compounds [31]. Therefore, this half-life reflects the net half-life of the Cremophor component(s) relevant for paclitaxel binding. Terminal half-lives up to about 100 h were reported for Cremophor [10, 35]. We found longer terminal half-lives, when we tested two- and three-compartment models for Cremophor. As these models did not improve any critical feature of the model and provided essentially indistinguishable curve fits, the simplest one-compartment model for Cremophor was chosen as the final model. Cremophor binding was probably less important at lower Cremophor concentrations that might have dropped below the critical micelle concentration for Cremophor [21].

We posited saturable paclitaxel release from nanodroplets into blood and saturable release of paclitaxel during sample processing based on the low aqueous solubility of paclitaxel. The estimate for paclitaxel solubility of 405 ng/mL (equivalent to 0.47 μM) for unbound paclitaxel in plasma is in good agreement with the observed paclitaxel solubility of 0.4 μM (range: 0.2–0.9 μM) at equilibrium [12, 29, 37]. It was encouraging to see that implementing a physicochemical mechanism improved the objective function, model fits, and predictive performance significantly. Figure 2 shows a “plateau” of concentrations in plasma ultrafiltrate for Tocosol Paclitaxel, whereas the associated concentrations in whole blood showed a pronounced peak. Saturable binding of paclitaxel would cause the opposite behavior. The final model predicted the time course of plasma ultrafiltrate and whole blood concentrations as well as their ratio (Fig. 4). In agreement with results of a phase I dose escalation study with Tocosol Paclitaxel [36], the model predicted a slightly more than proportional increase of whole blood AUC with dose. In addition, the model predicted an approximately proportional increase of plasma ultrafiltrate AUC and a proportional increase of unbound paclitaxel AUC in the central compartment for Tocosol Paclitaxel doses from 25 to 225 mg/m².

Prolonged release of paclitaxel from nanodroplets was found (Fig. 5a) that supports administering Tocosol Paclitaxel as a 15 min infusion. Additionally, the new Cremophor-free Tocosol Paclitaxel formulation does not carry the risk for Cremophor-related toxicity. A small fraction (9.8%) of Tocosol Paclitaxel dose was estimated to enter directly into the DPC. As models with a direct release of paclitaxel into the shallow peripheral compartment did not improve the objective function and as the associated fraction of dose was minimal (<1%), only release into the deep peripheral compartment was included in the final model. A direct release of a fraction of dose into the deep peripheral compartment increases the unbound AUC in this compartment much more compared to a direct release of the same fraction of dose into the shallow peripheral compartment, since the equilibration between the shallow peripheral and central compartment was much faster than the equilibration for the deep peripheral compartment. As described in the companion article, the direct release of drug into the deep peripheral compartment was one mechanism that explained differences in toxicity.

As Tocosol Paclitaxel nanodroplets have a diameter of 40–80 nm [5, 6] and as pore sizes of many tumor tissues are larger than tight endothelial junctions of normal tissue vasculature [28], this estimate seems reasonable (see companion article for a more detailed discussion). Nanodroplets were assumed to release a small fraction (9.8%) of the Tocosol Paclitaxel dose into the whole DPC (Fig. 5e). If only a part of the DPC is accessible to nanodroplet release,

paclitaxel concentrations at these sites will have greater unbound concentrations than shown in Fig. 5e. Tocol Paclitaxel might therefore show more toxicity at sites accessible to release from nanodroplets compared to Taxol at the same dose. More toxicity for Tocol Paclitaxel has been observed for neutrophil counts (see companion article).

In conclusion, the proposed mechanistic population PK model showed that disposition of unbound paclitaxel was linear. The prolonged release of paclitaxel from Tocol Paclitaxel nanodroplets was well described by a solubility-limited process. For a 15 min Tocol Paclitaxel intravenous infusion, the average (\pm SD) time to 90% cumulative drug input was predicted to be 1.14 ± 0.16 h. Predicted unbound peak paclitaxel plasma concentrations were about 60% (median) higher for a 15 min Tocol Paclitaxel infusion compared to a 3 h Taxol infusion, due to the prolonged release for Tocol Paclitaxel and limited solubility of paclitaxel. The prolonged paclitaxel release supports 15 min infusion of Tocol Paclitaxel. A small fraction of dose (9.8%) was estimated to be directly released into the DPC. The PK model proposed herein that is predicated on the limited solubility of paclitaxel may be important to optimize dosage regimens for nanodroplet formulations of drugs with a low aqueous solubility like paclitaxel.

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Appendix

Mechanistic model for drug release from Tocol Paclitaxel nanodroplets

The following equations describe the amount of paclitaxel in nanodroplets (A_{Nano}) and in the unstirred water layer (A_{Layer}) (see also Fig. 1):

$$\frac{dA_{\text{Nano}}}{dt} = F_{\text{Tocol}} \cdot R_{\text{Input}} - ka_1 \cdot \left(1 - \frac{C_{\text{Layer}}}{C_{\text{Sol}}}\right) \cdot A_{\text{Nano}} \quad (\text{A1})$$

$$\frac{dA_{\text{Layer}}}{dt} = ka_1 \cdot \left(1 - \frac{C_{\text{Layer}}}{C_{\text{Sol}}}\right) \cdot A_{\text{Nano}} - ka_2 \cdot A_{\text{Layer}} \quad (\text{A2})$$

where C_{Sol} is the solubility of unbound paclitaxel in the volume of distribution for nanodroplets (V_{Nano}), C_{Layer} is

the paclitaxel concentration in the unstirred water layer, R_{Input} is the zero-order input into the nanodroplet compartment, and F_{Tocol} is the relative bioavailability of Tocol Paclitaxel compared to Taxol. The release of drug from nanodroplets into the unstirred water layer is described by the first-order rate constant ka_1 and the transfer of drug from the unstirred water layer to the central and peripheral compartment by the first-order rate constant ka_2 (Fig. 1). Initial conditions for A_{Nano} and A_{Layer} are zero.

If ka_2 is much larger than ka_1 , A_{Layer} will show a rapid initial rise and thereafter the differential dA_{Layer}/dt will be very small compared to dA_{Nano}/dt . In this situation, the quasi-stationary solution for A_{Layer} that is valid after the rapid initial rise can be calculated by setting the differential dA_{Layer}/dt to zero and solving for A_{Layer} . This yields:

$$A_{\text{Layer}} = \frac{ka_1 \cdot A_{\text{Nano}}}{ka_2 + \frac{ka_1 \cdot A_{\text{Nano}}}{C_{\text{Sol}} \cdot V_{\text{Layer}}}} \quad (\text{A3})$$

The V_{Layer} represents the volume of the unstirred water layer. Inserting this equation into Eq. A1 yields the following equation for the mixed-order (Michaelis–Menten) process from the nanodroplet to the central compartment:

$$\begin{aligned} \frac{dA_{\text{Nano}}}{dt} &= F_{\text{Tocol}} \cdot R_{\text{Input}} - \frac{ka_2 \cdot C_{\text{Sol}} \cdot V_{\text{Layer}} \cdot A_{\text{Nano}}}{\frac{ka_2 \cdot C_{\text{Sol}} \cdot V_{\text{Layer}}}{ka_1} + A_{\text{Nano}}} \\ &= F_{\text{Tocol}} \cdot R_{\text{Input}} - \frac{V_{\text{max}} \cdot A_{\text{Nano}}}{AN_{50} + A_{\text{Nano}}} \end{aligned} \quad (\text{A4})$$

The maximum rate of drug transfer is denoted as V_{max} and is equal to the product of ka_2 , C_{Sol} , and V_{Layer} . The amount of drug in the nanodroplet compartment (AN_{50}) that results in a transfer rate of 50% of V_{max} is V_{max}/ka_1 . Based on the final estimates, we confirmed that this quasi-stationary solution is an accurate approximation of the full set of differential equations.

The product of ka_2 and V_{Layer} is the clearance from the unstirred water layer (CL_{Layer}) which is identifiable, whereas the individual terms are not. Therefore, a mixed-order transfer from the nanodroplet to the central compartment can be explained mechanistically by solubility-limited transfer through an unstirred water layer.

Output equation for plasma ultrafiltrate for nanodroplet formulation

Blood samples drawn during the release phase of Tocol Paclitaxel nanodroplets contain nanodroplets with paclitaxel. This unreleased paclitaxel was assumed to be in part released during the processing of blood samples. Due to the

vigorous conditions during centrifugation, we assumed that the unstirred water layer is less important. The release of paclitaxel from nanodroplets into plasma was described by a first-order process that becomes saturated, if the unbound plasma concentration approaches the estimated solubility limit (Fig. 1). This yields the following equations for the amounts of drug in nanodroplets in a blood sample ($A_{\text{Nano},S}$) and in the blood sample excluding drug in nanodroplets ($A_{\text{Blood},S}$):

$$\frac{dA_{\text{Nano},S}}{dt} = -k_a \cdot A_{\text{Nano},S} \cdot \left(1 - \frac{A_{\text{Blood},S}}{V_{\text{Sample}} \cdot \text{BF}_{\text{TOC}} \cdot C_{\text{Sol}}}\right) \quad (\text{A5})$$

$$\frac{dA_{\text{Blood},S}}{dt} = k_a \cdot A_{\text{Nano},S} \cdot \left(1 - \frac{A_{\text{Blood},S}}{V_{\text{Sample}} \cdot \text{BF}_{\text{TOC}} \cdot C_{\text{Sol}}}\right). \quad (\text{A6})$$

The binding factor for Tocosol paclitaxel (BF_{TOC}) was assumed to be constant during sample handling, as the saturable component of protein binding was negligible for high unbound concentrations. The first-order release rate constant under sample handling conditions is denoted as k_a . The paclitaxel amount in sampled blood excluding drug in nanodroplets ($A_{\text{Blood},S}$) divided by the sample volume (V_{Sample}) and by the binding factor BF_{TOC} yields the unbound plasma concentration. The initial condition was $A_{\text{Nano}} \cdot V_{\text{Sample}} / V_{\text{Nano}}$ for $A_{\text{Nano},S}$ and the unbound concentration in central compartment (C1) multiplied by V_{Sample} and BF_{TOC} for $A_{\text{Blood},S}$.

These two differential equations were solved in Maple and the solution was used as the output equation for the paclitaxel concentration in plasma ultrafiltrate for Tocosol Paclitaxel [$E(t_{\text{Proc}})$, N1, and N2 are intermediary variables to simplify the equations]:

$$E(t_{\text{Proc}}) = \exp\left(\frac{k_a \cdot (A_{\text{Nano}} - V_{\text{Nano}} \cdot C_{\text{Sol}} \cdot \text{BF}_{\text{TOC}} + V_{\text{Nano}} \cdot \text{BF}_{\text{TOC}} \cdot C1) \cdot t_{\text{Proc}}}{V_{\text{Nano}} \cdot \text{BF}_{\text{TOC}} \cdot C_{\text{Sol}}}\right) \quad (\text{A7})$$

$$N1 = -C_{\text{Sol}} \cdot A_{\text{Nano}} + A_{\text{Nano}} \cdot C1 - V_{\text{Nano}} \cdot C_{\text{Sol}} \cdot \text{BF}_{\text{TOC}} \cdot C1 \quad (\text{A8})$$

$$N2 = V_{\text{Nano}} \cdot \text{BF}_{\text{TOC}} \cdot C1^2 + C_{\text{Sol}} \cdot E(t_{\text{Proc}}) \cdot A_{\text{Nano}} \quad (\text{A9})$$

$$A_{\text{blood},S}(t_{\text{Proc}}) = \frac{V_{\text{Sample}} \cdot \text{BF}_{\text{TOC}} \cdot [N1 + N2]}{-\text{BF}_{\text{TOC}} \cdot C_{\text{Sol}} \cdot V_{\text{Nano}} + V_{\text{Nano}} \cdot \text{BF}_{\text{TOC}} \cdot C1 + E(t_{\text{Proc}}) \cdot A_{\text{Nano}}}. \quad (\text{A10})$$

Dividing the amount of paclitaxel in the blood sample (excluding the unreleased amount in nanodroplets) at the end of sample processing (t_{Proc}) by the sample volume and the binding factor yields the output equation for the concentration in plasma ultrafiltrate for Tocosol Paclitaxel:

$$\text{Cu}(t_{\text{Proc}}) = \frac{A_{\text{blood},S}(t_{\text{Proc}})}{V_{\text{Sample}} \cdot \text{BF}_{\text{TOC}}}. \quad (\text{A11})$$

We assumed an average time of sample processing of 30 min according to the procedures described in the clinical protocol and estimated the half-life of the first-order release rate constant k_a . The sample volume (V_{Sample}) was arbitrarily set to 1 L, as this choice did not influence the results.

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