

Resisting Population PK – The Story of P-gp inhibition and Co-administered Chemotherapy

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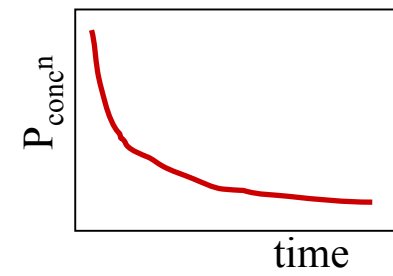
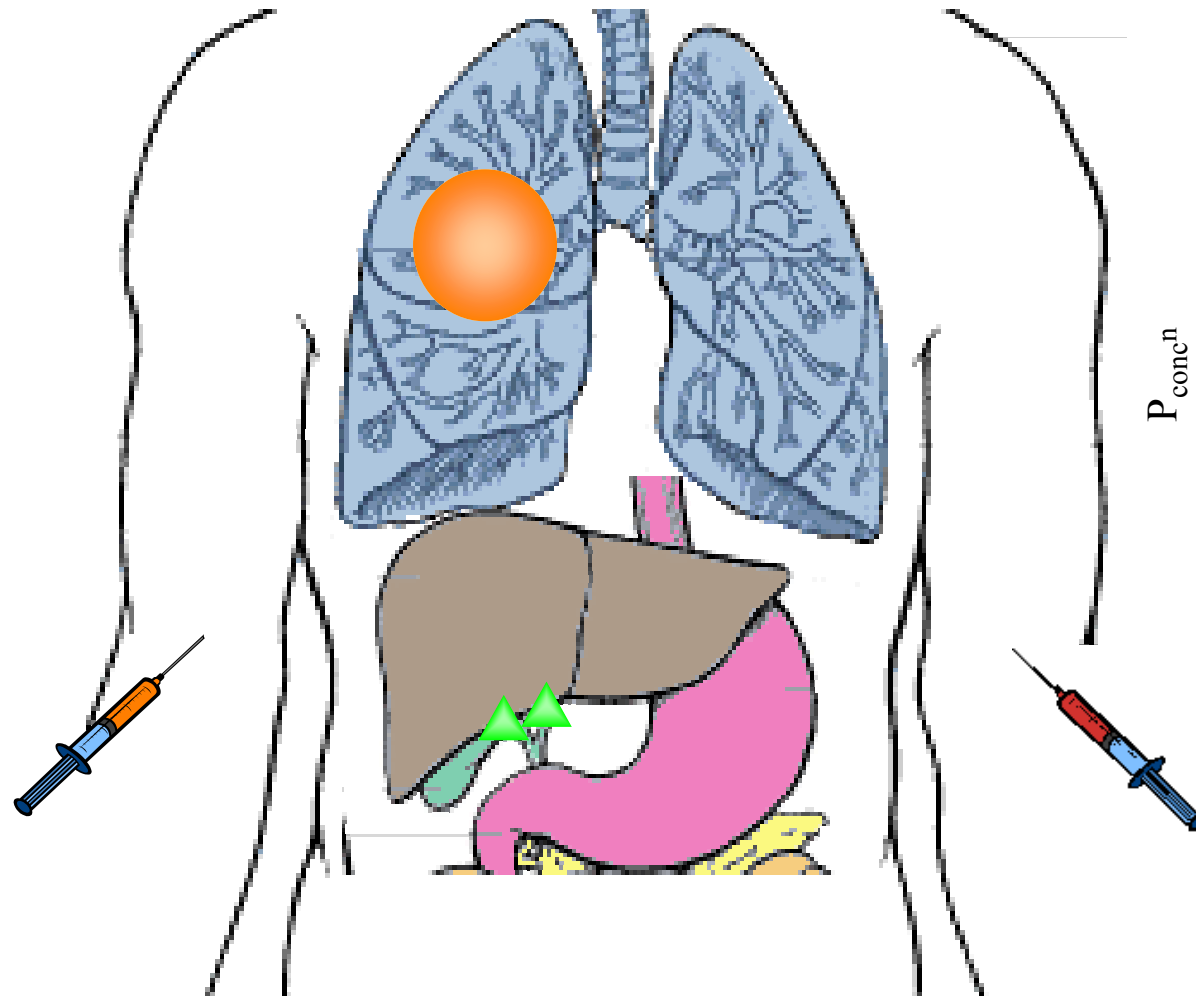
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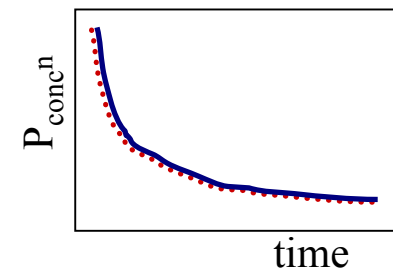
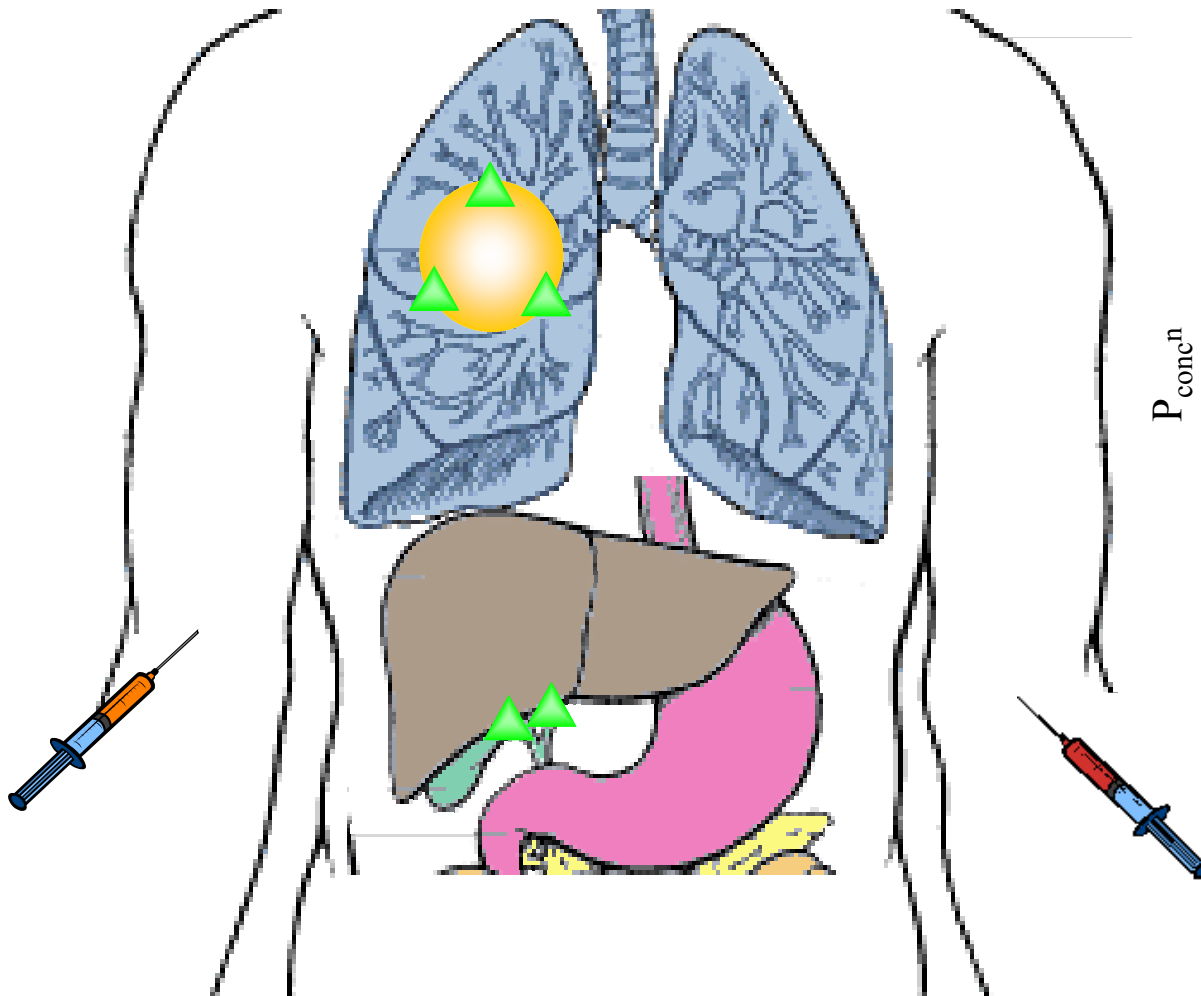
The Lilly logo is written in a red, cursive script font.

Answers That Matter.

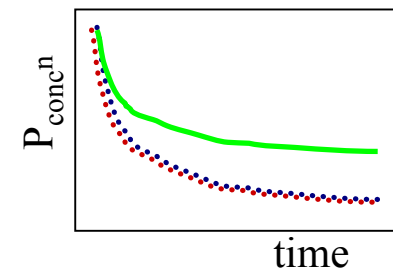
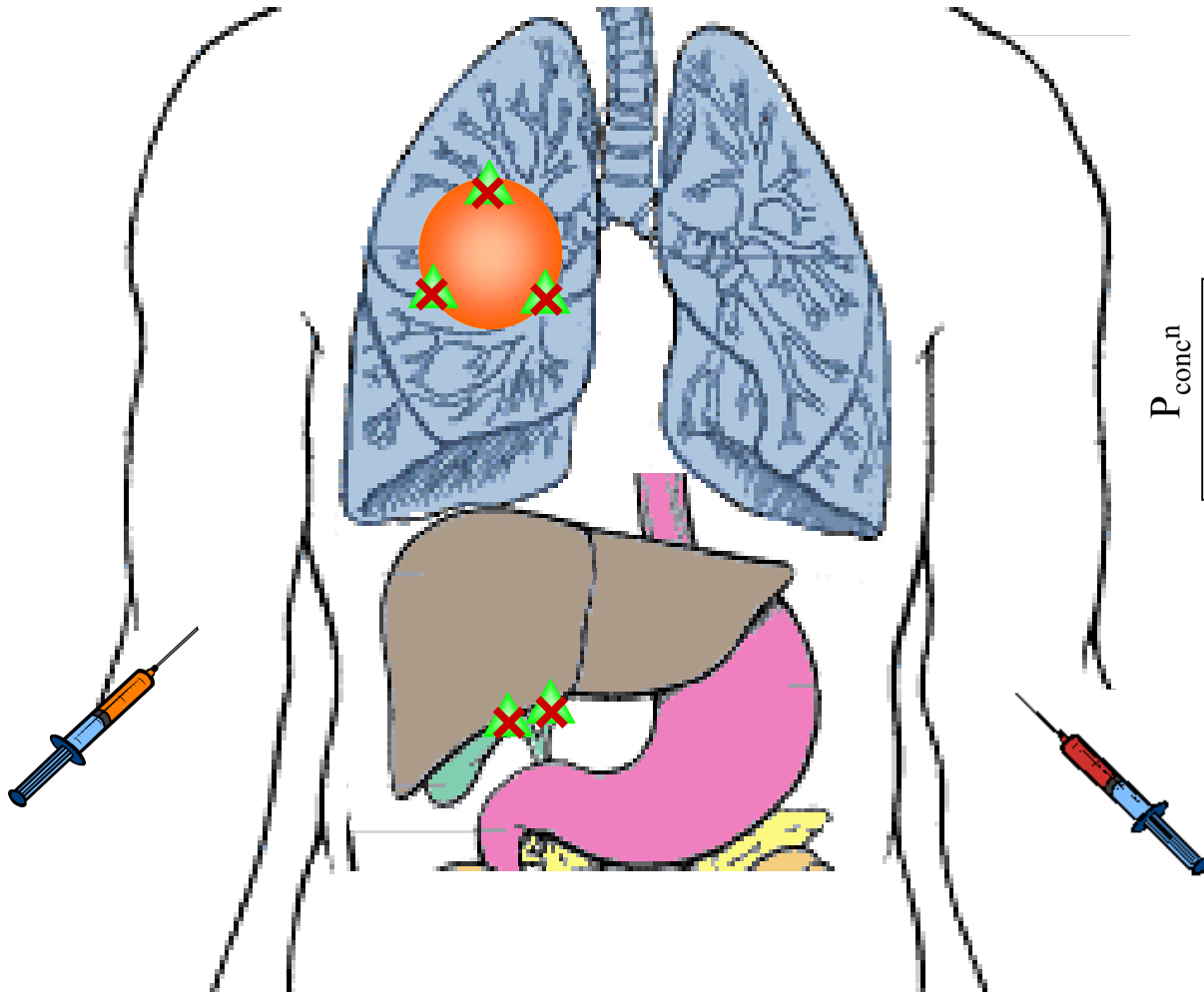
▲ P-gp



▲ P-gp



▲ P-gp



Agents Targeting P-gp Mediated Drug Resistance - The Early Generations

The drugs:

Verapamil

Quinidine

Quinine

Tamoxifen

Cyclosporin A

PSC-833

GF-120918

Reasons for failing:

Did not inhibit P-gp

Toxicity in its own right

PK interaction

PK Interactions with P-gp inhibitors & co-administered chemotherapy

In theory this depends on:

- Relative contribution of each transporter to drug efflux.
- The distribution of the transporter in normal tissue vs tumour
- Specificity of the inhibitor/drug

Modulator	Pgp	MRP1	MRP2	BCRP
PSC 833	+	-	+	-
VX-710	+	+	-	-
GF120918	+	-	-	+
Zosuquidar	+	-	-	-

Duration of inhibition ??

The Pharmacokinetic Effect of PSC833 on Doxorubicin (50 mg/m²) & Doxorubicinol

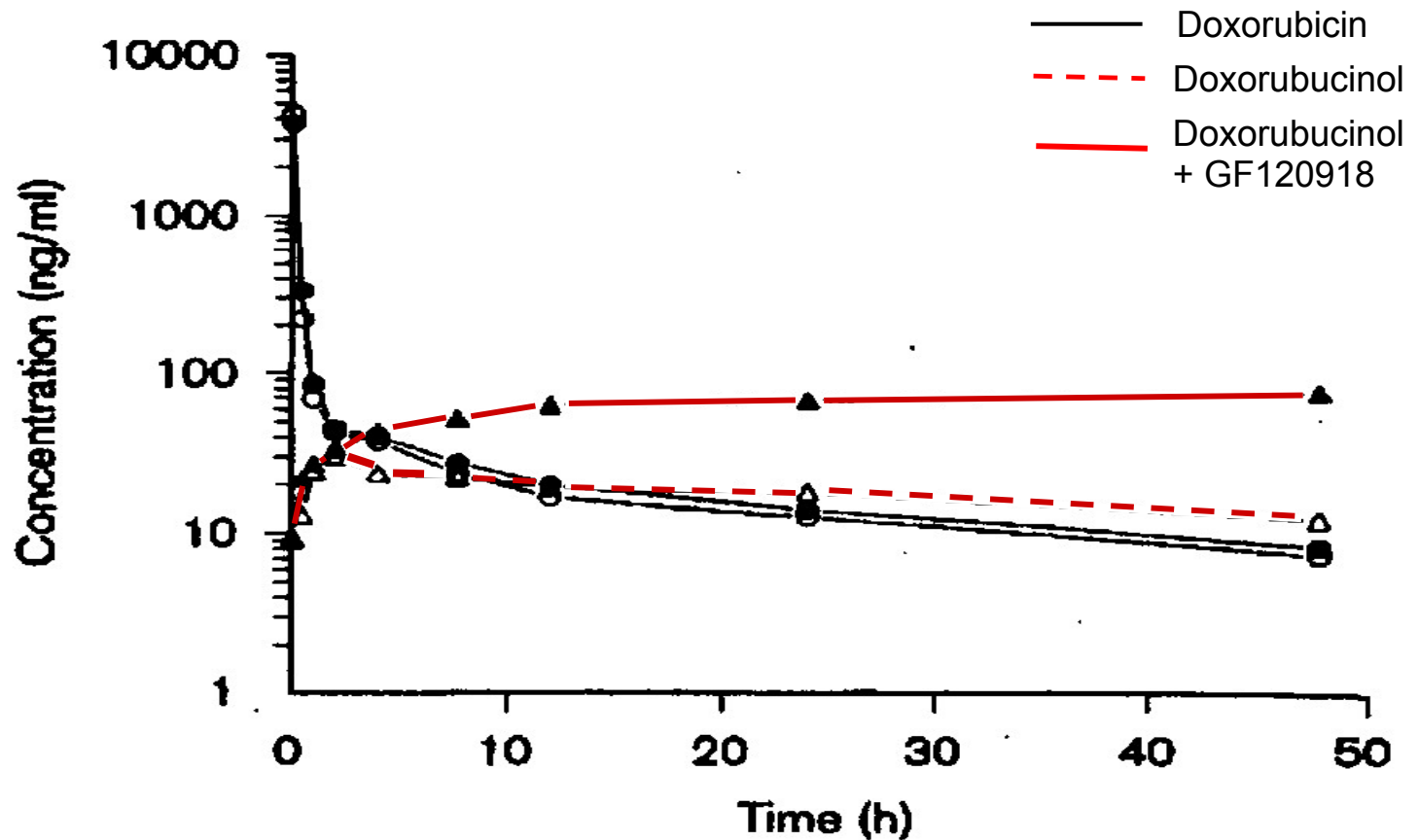
	% Change (based on arithmetic mean)
Doxorubicin PK	
CL	- 30
Doxorubicinol PK	
AUC _(0 - ∞)	+1063
C _{max}	+101

(Giaccone et al., Clinical Cancer Research. 3, 2005-2015, 1997)

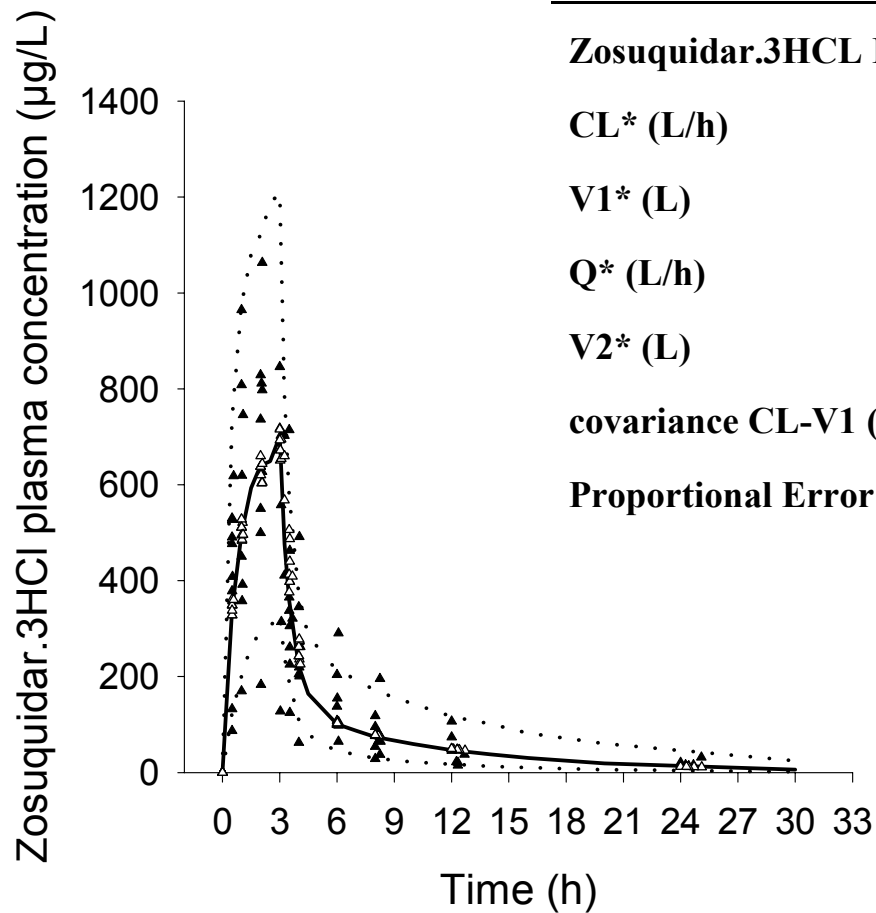
PSC 833 administered orally 2.5 to 25 mg/kg/d x 5 d

Impact on Doxorubicin- Doxorubicinol PK by GF120918

50mg/m² of Dox with or without
400mg BID of GF120918



Zosuquidar.3HCL PK Model



Zosuquidar.3HCL PK parameters

CL* (L/h)

Mean (SEE%)

127 (9.45)

IIV (SEE %)

35.1 (56.0)

V1* (L)

127 (16.5)

62.0 (51.4)

Q* (L/h)

79.9 (11.9)

NE

V2* (L)

412 (7.0)

11.3 (49.4)

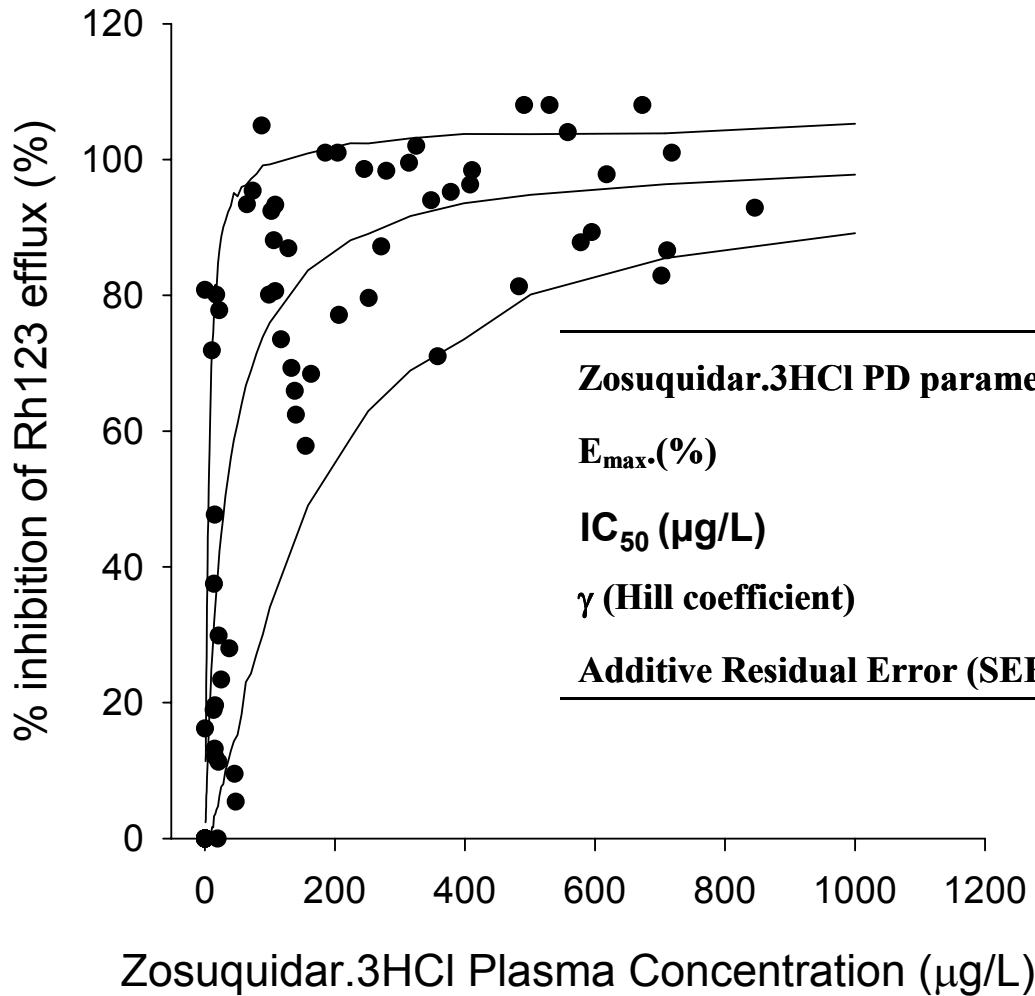
covariance CL-V1 (%)

43.5 (62.4)

Proportional Error (SE %)

27.0 % (9.3)

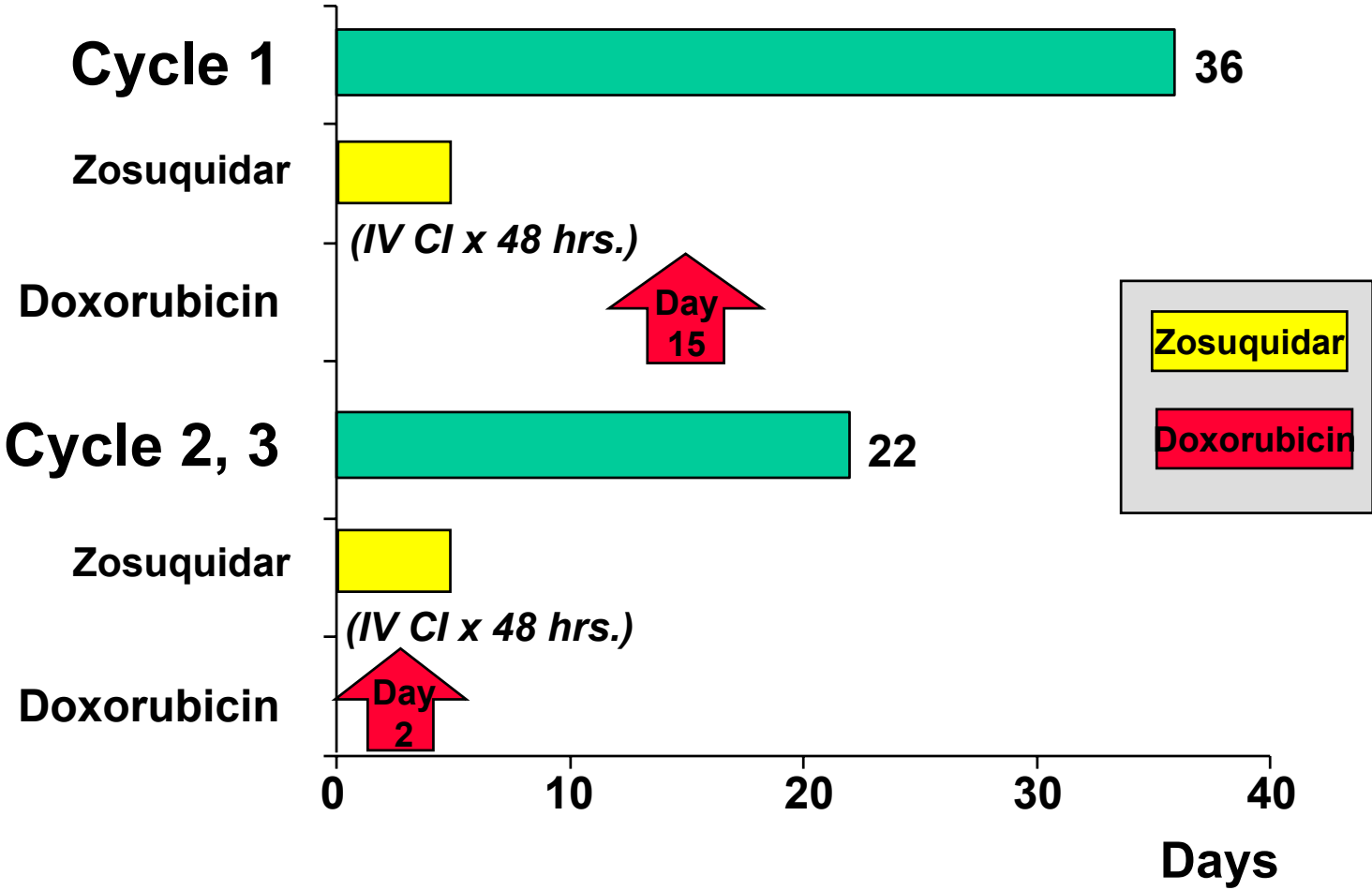
Zosuquidar.3HCL PK/PD Model



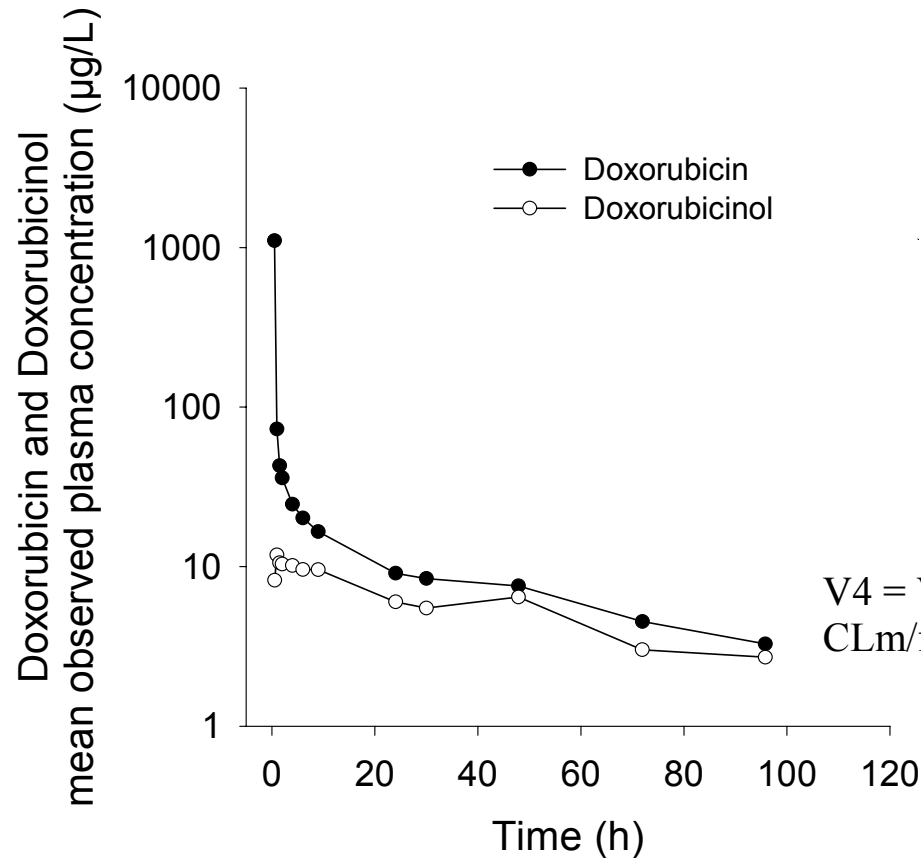
Zosuquidar.3HCl PD parameters	Mean (SEE%)	IIV (SEE %)
E_{max} (%)	100 Fixed	
IC_{50} (µg/L)	31.7 (32.2)	103 (30.0)
γ (Hill coefficient)	1.31 (11.7)	
Additive Residual Error (SEE%)**	13.5 % (57.9)	

Estimated $IC_{90} = 169 \mu\text{g/L}$

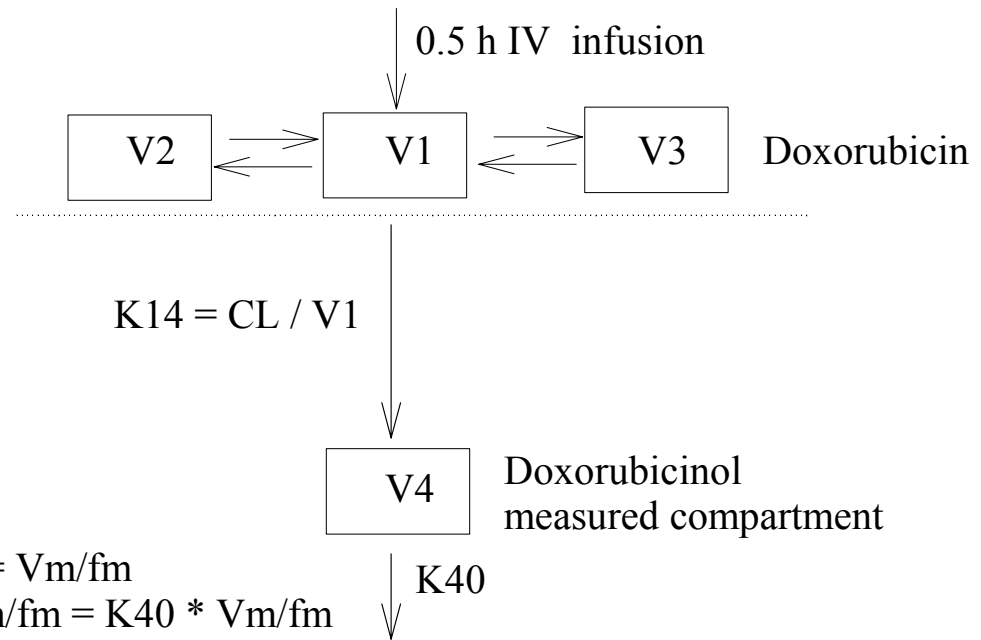
Clinical Study of Intravenous Zosuquidar in Combination with Doxorubicin



Dox-Doxol PK Model in Absence of Zosuquidar

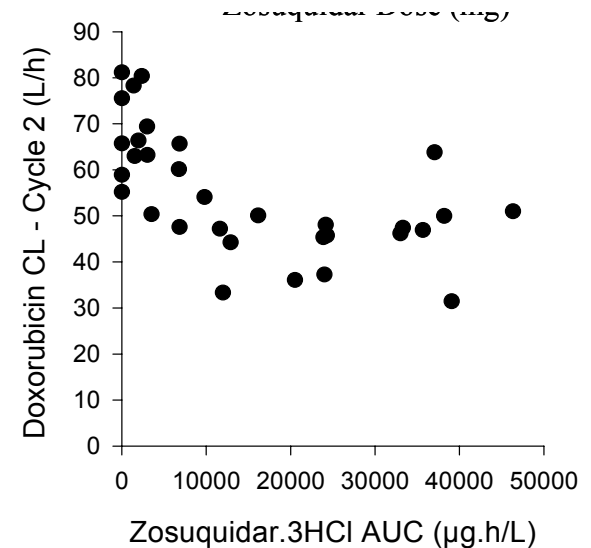
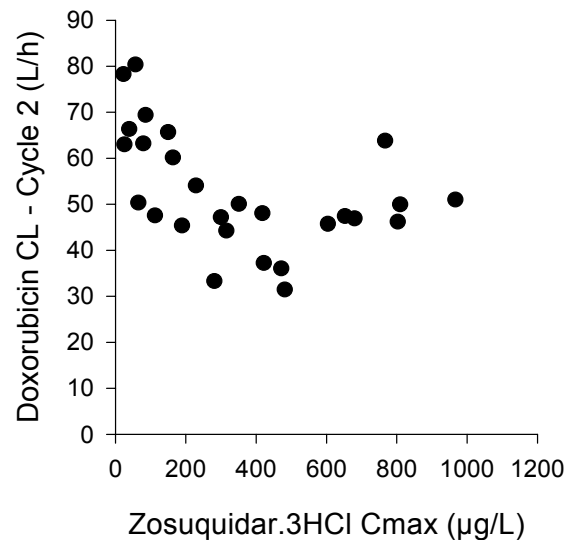
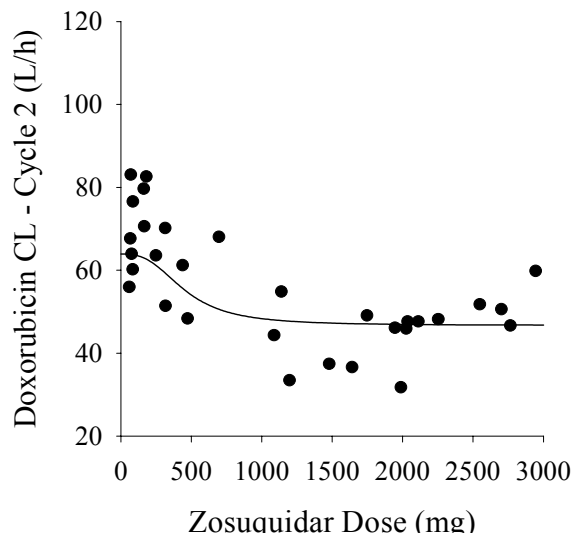
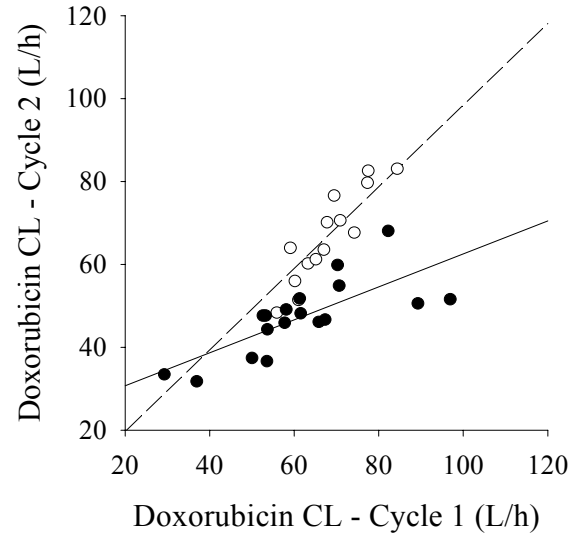


$V4 = Vm/fm$
 $CLm/fm = K40 * Vm/fm$

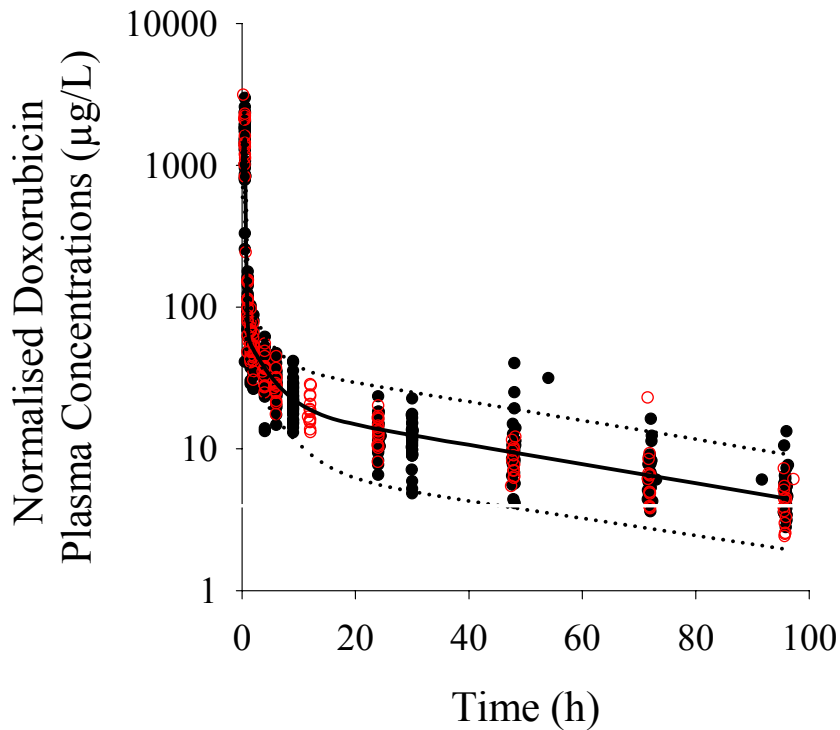


$CLm/fm =$ Apparent CL, fm is fraction of doxorubicin converted to doxorubicinol

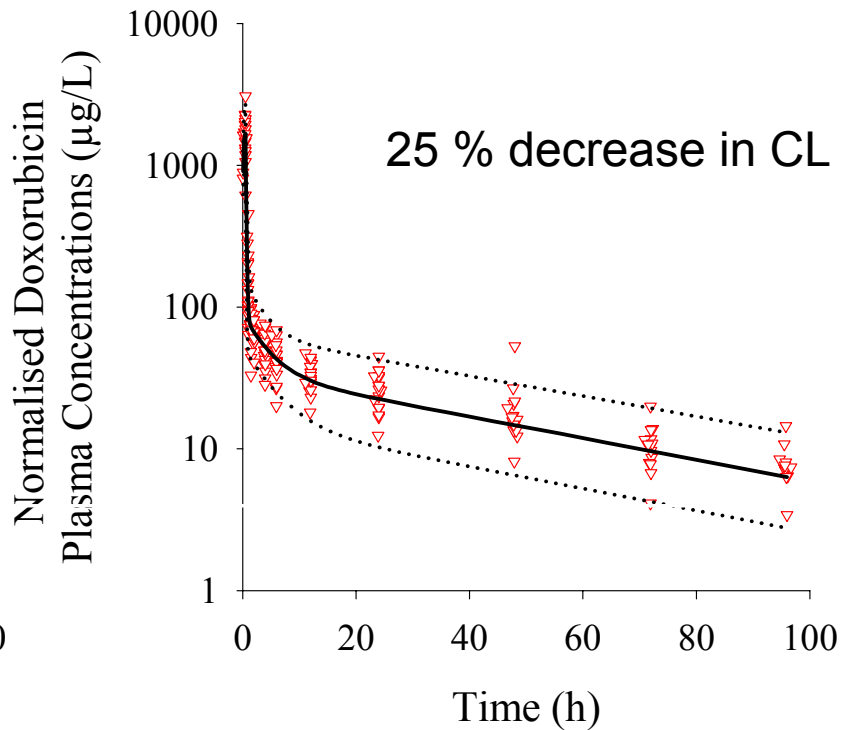
Relationship between Dox PK and Zosuquidar Dose & Exposure



Observed & Predicted Dox PK in the presence & absence of Zosuquidar

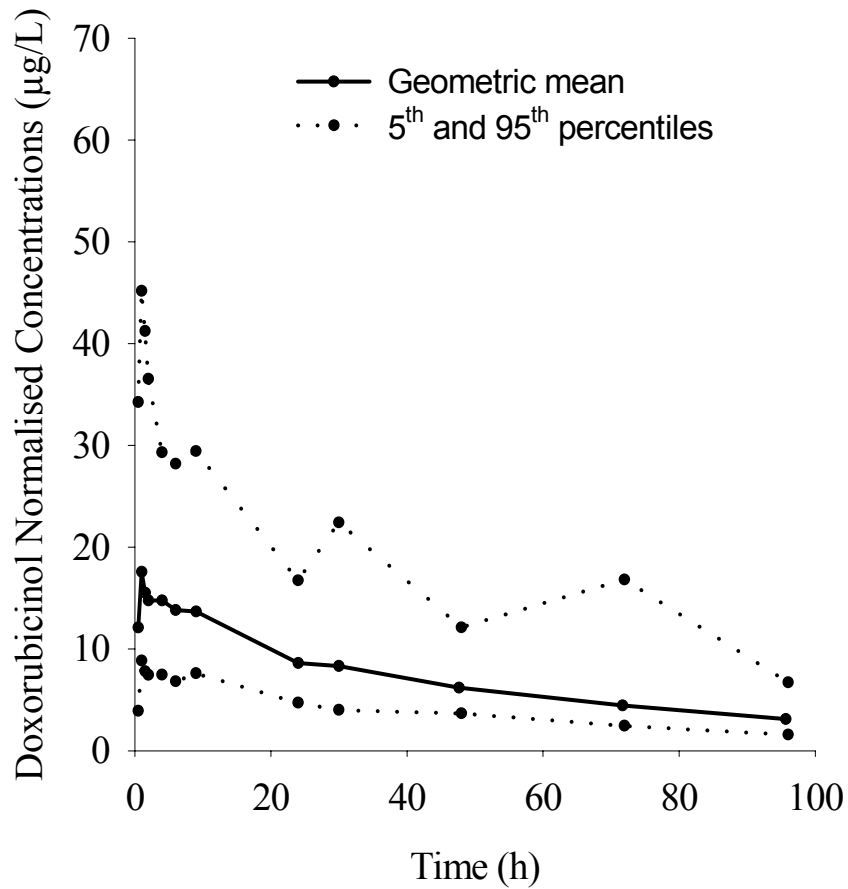


Zosuquidar Dose either 0 or <500mg

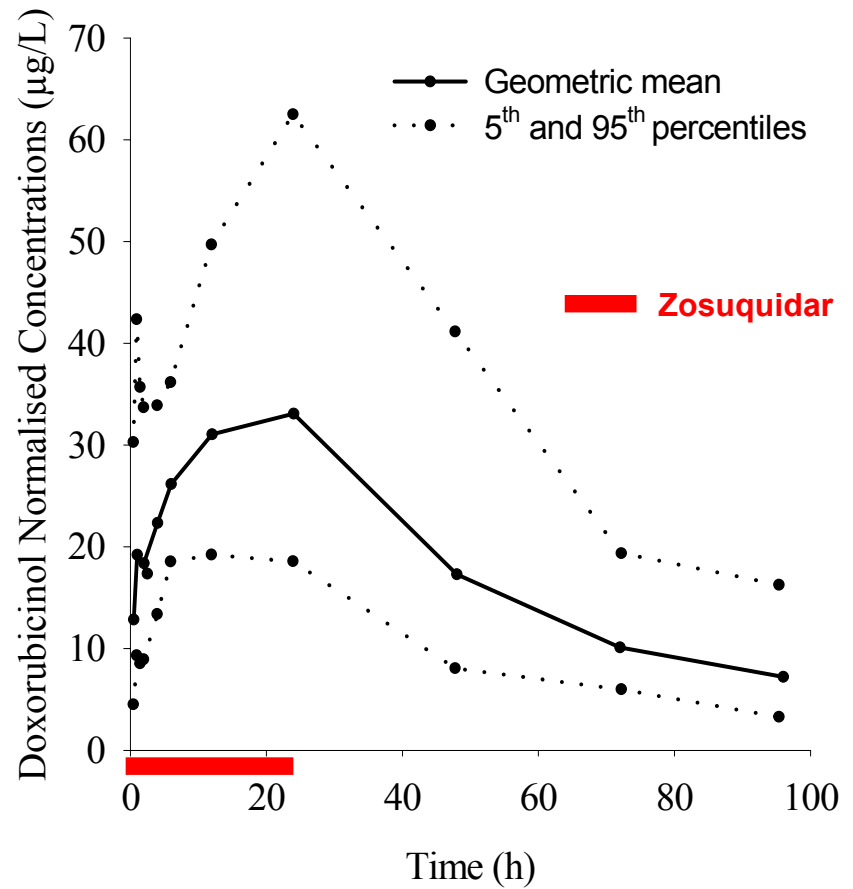


Zosuquidar Dose ≥ 500 mg

The Observed Doxorubicinol Concentration-Time Curves in the Presence and Absence of Zosuquidar

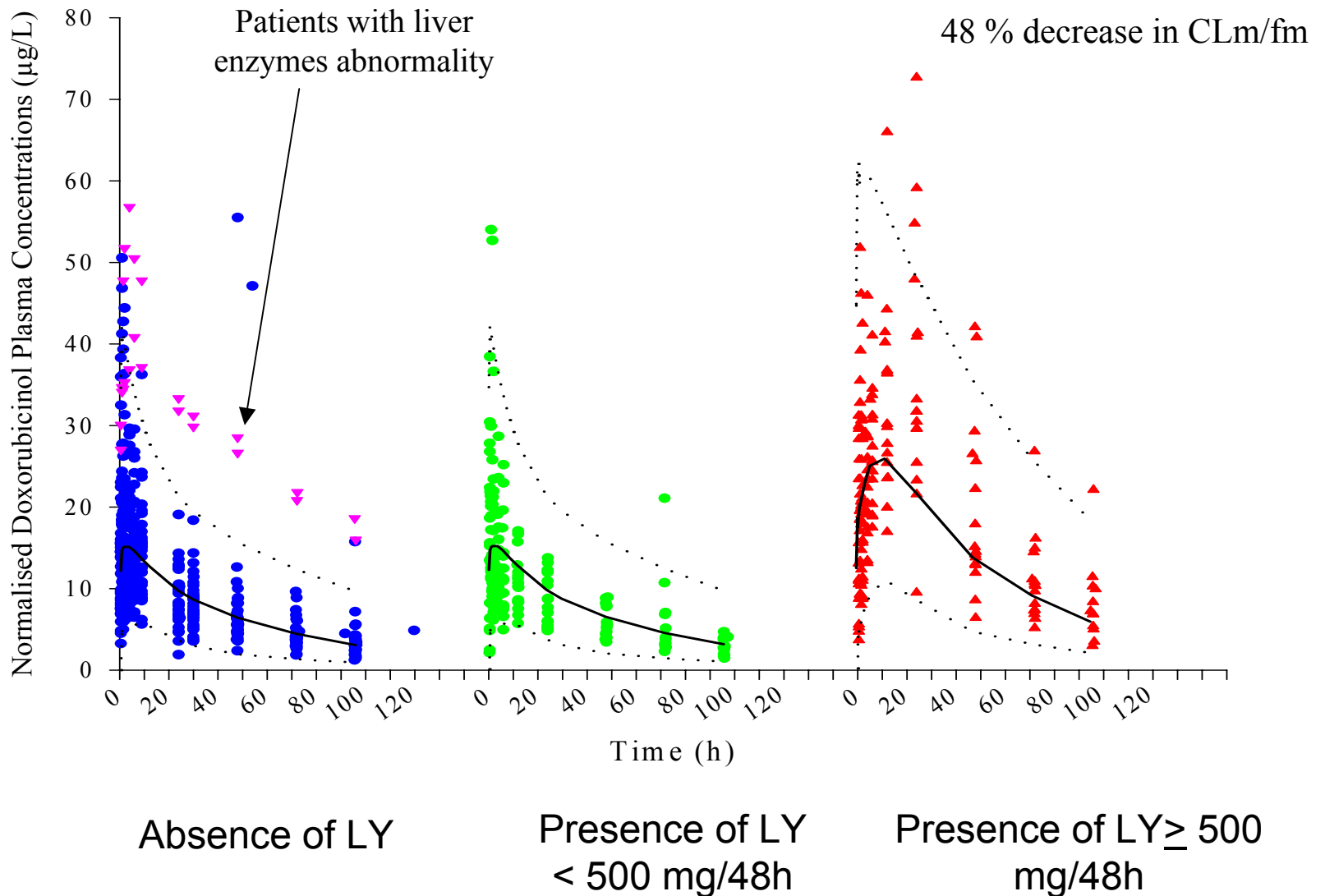


Dose either 0 or <500mg

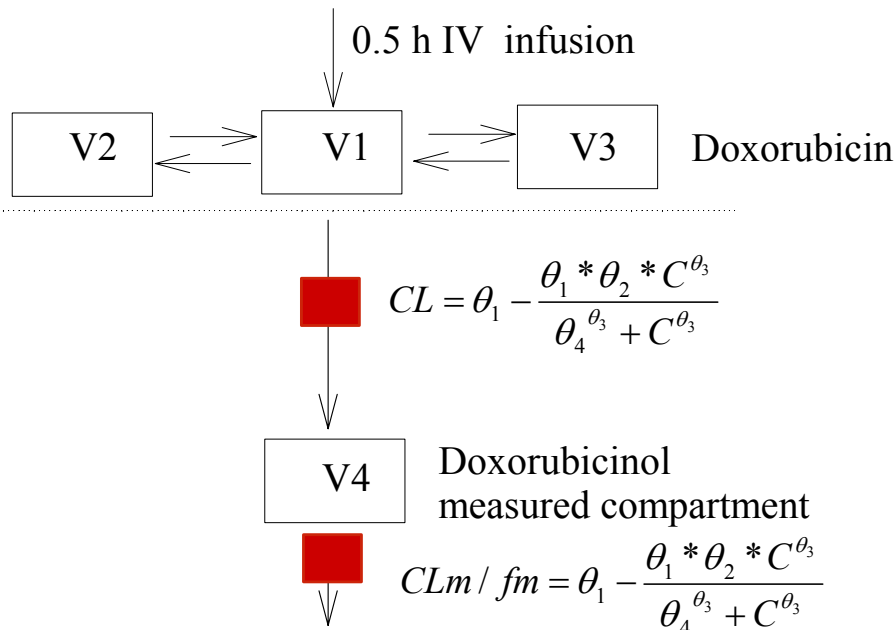


Dose ≥ 500 mg

Posterior Predictive Check of Dox-Doxol Model

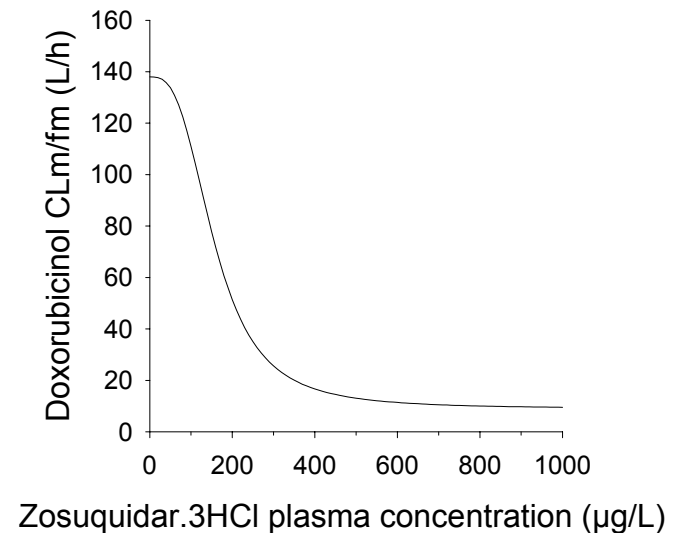
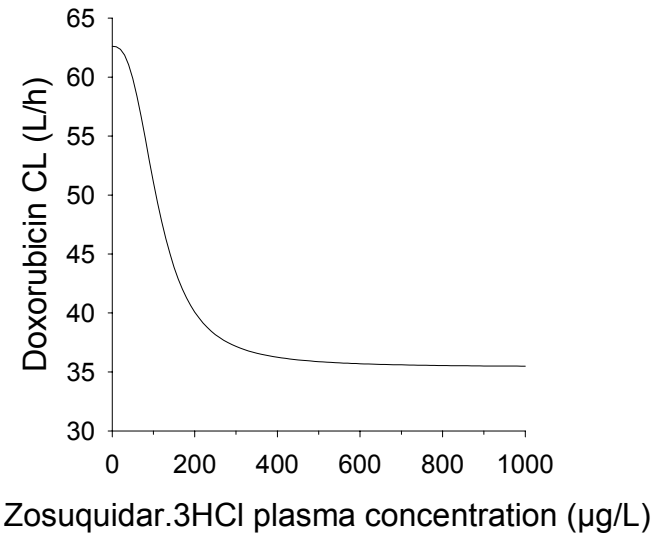


Dox-Doxol Regimen Independent PK model in the presence of Zosuquidar

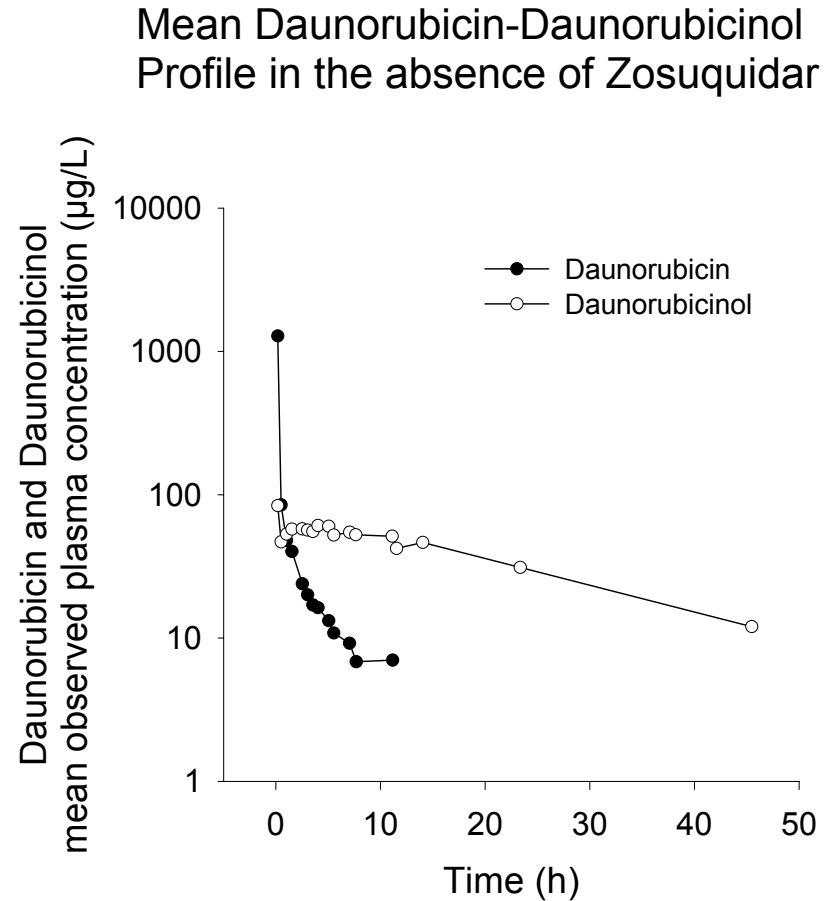
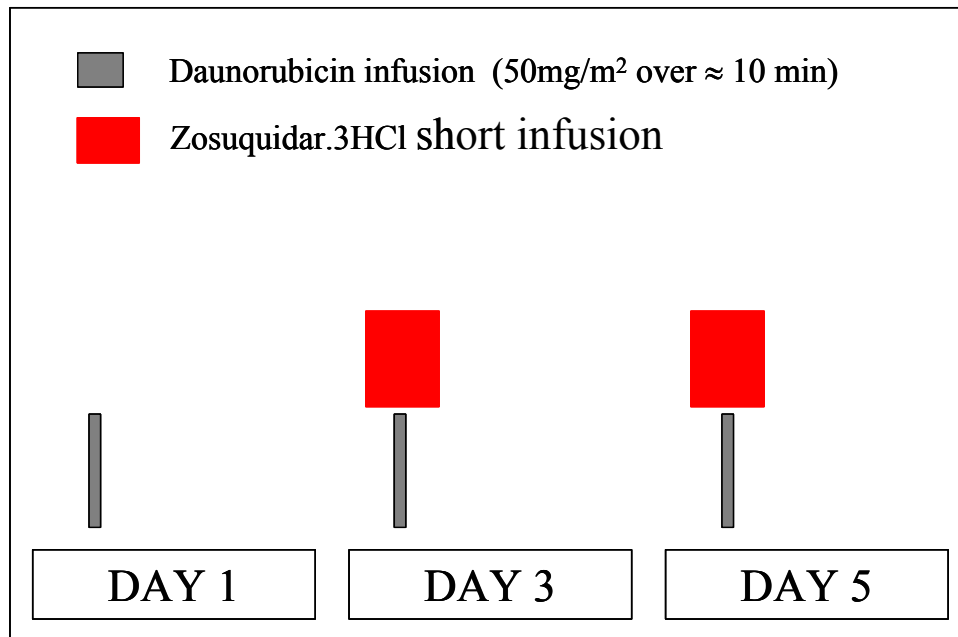


■ Pgp inhibition effect on PK
 C = Zosuquidar.3HCl predicted concentration

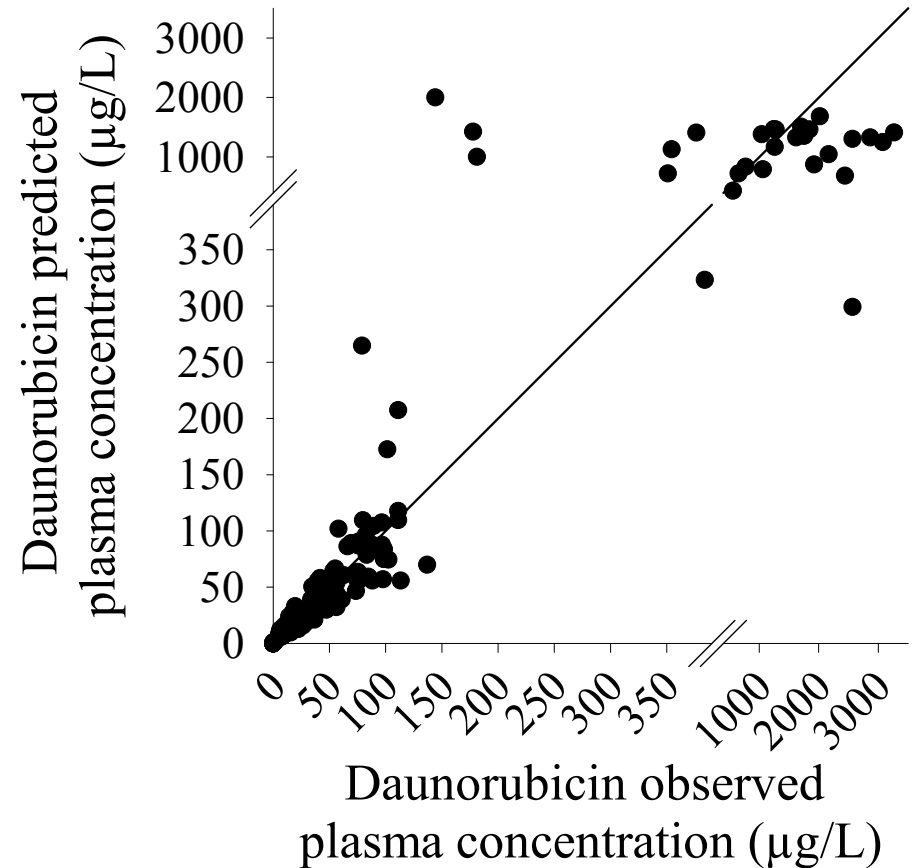
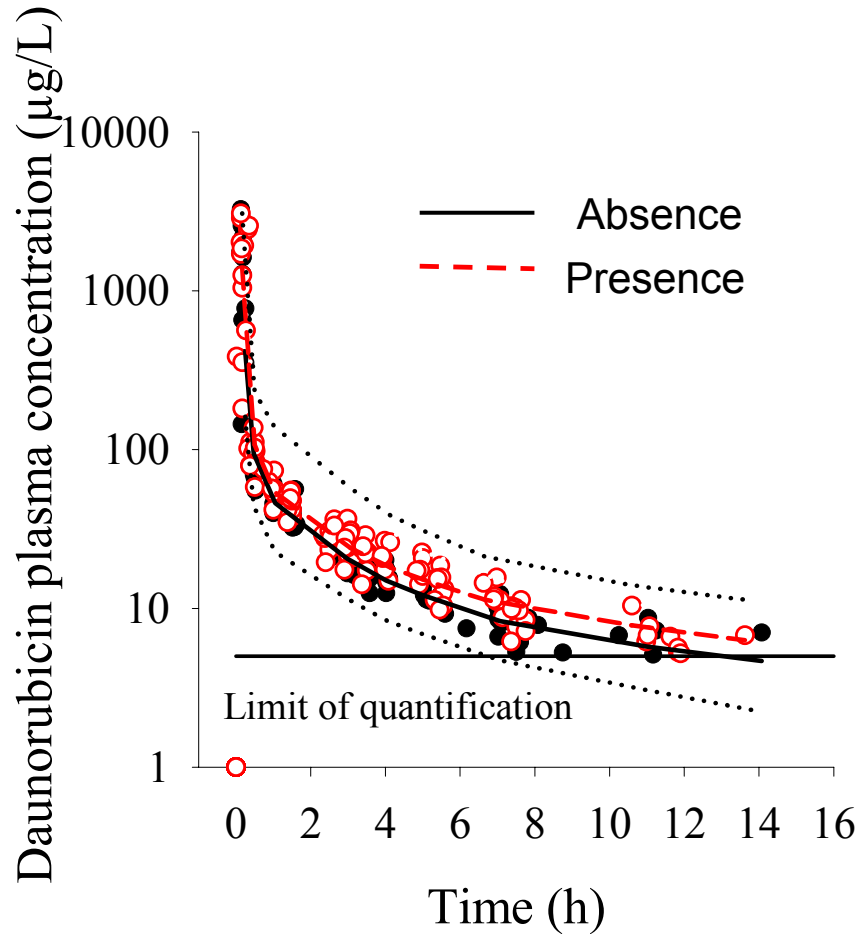
CLm/fm = Apparent CL, fm is fraction of doxorubicin converted to doxorubicinol



Design of Phase 1 Study with Daunorubicin + Zosuquidar short infusion

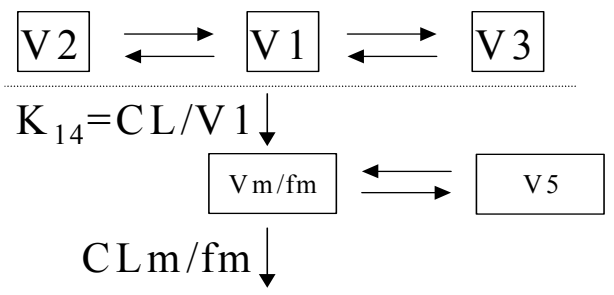


Impact on Daun PK (6 h or less Zosuquidar infusion) - Daun CL decreased by 10 %

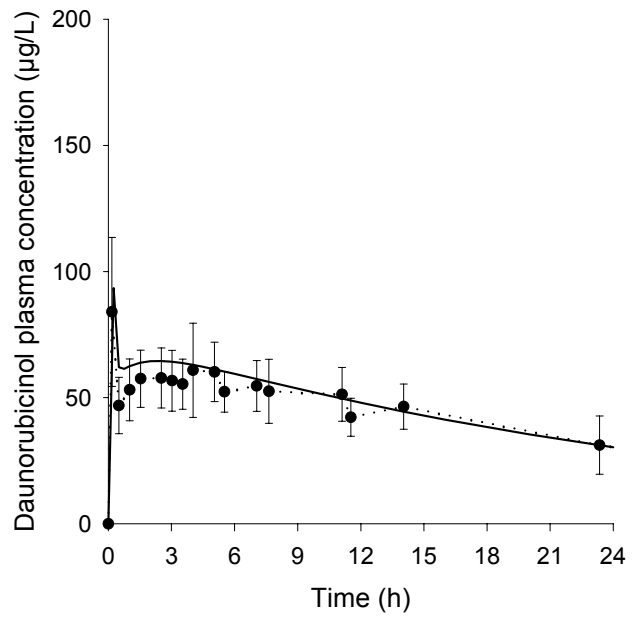


Impact on Daunol PK (55 % decrease in CL_m/f_m) in the Presence of Zosuquidar

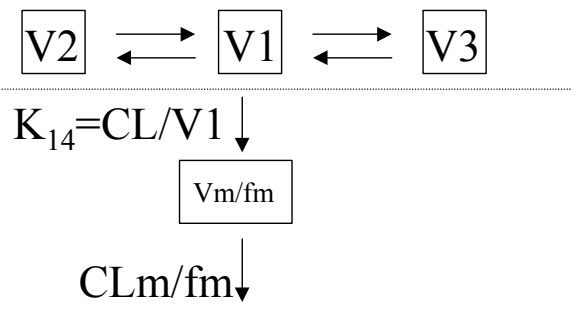
Daunorubicin



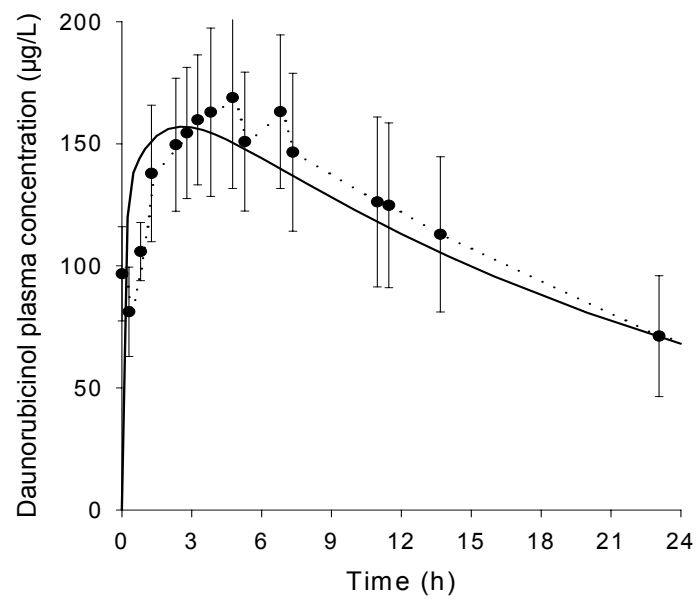
Daunorubicinol



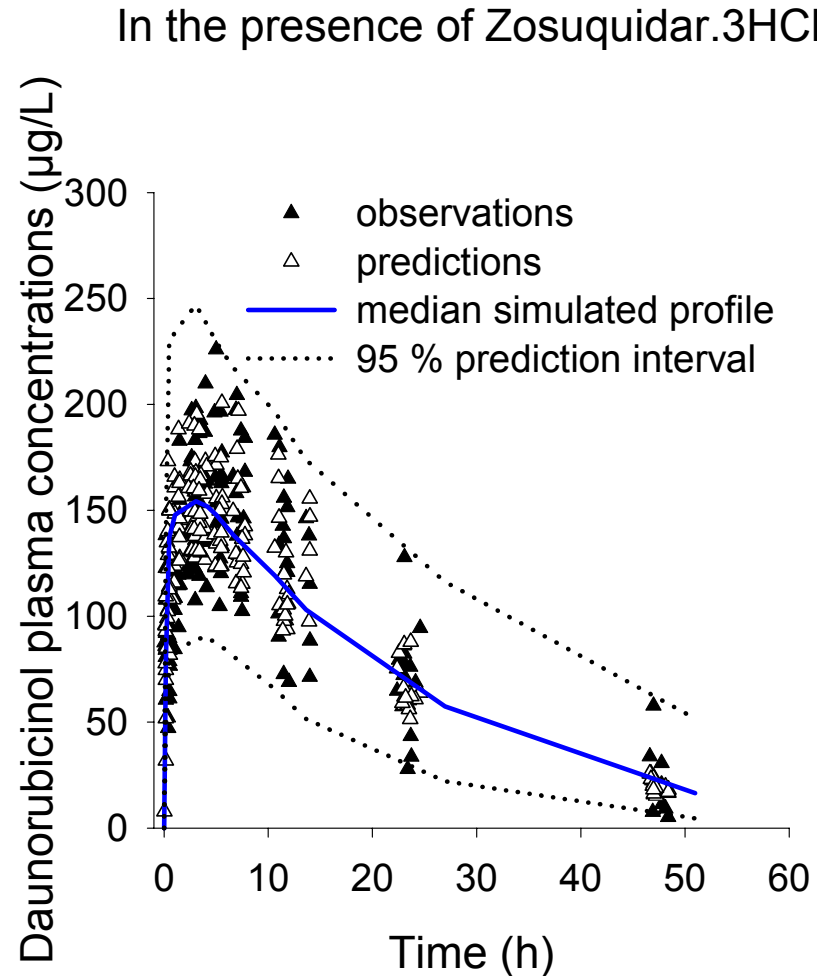
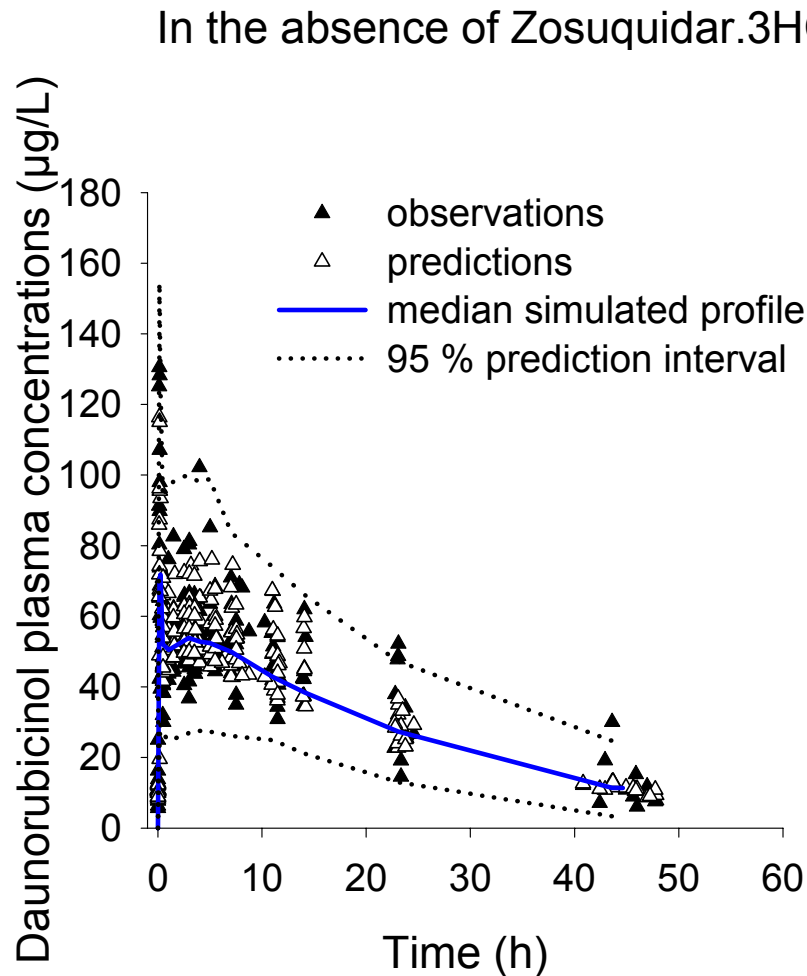
Daunorubicin



Daunorubicinol



Observed vs Simulated PK Profiles for Daunol (Median & 95 % Prediction Intervals)



Conclusions

- Lack of modelling of this type has precluded the efficient testing of P-gp inhibition in the clinic
- We developed a population PK model which allowed us to explore the balance between effect and toxicity. The model highlights the duration of inhibition as an important determinant of toxicity.
- Based on our understanding we developed a short IV infusion of Zosuquidar which is feasible and produces maximal P-gp inhibition, but reduced PK interactions.
- A randomized Phase 3 study with full dose chemotherapy in each arm is required to validate these hypotheses and the models from this work.
- This study is already on-going in 450 AML patients, application of the model has already allowed 390 patients enrolled without any obvious imbalance of toxicity, which hitherto has not been possible.
- We await the efficacy results from this trial with interest.