

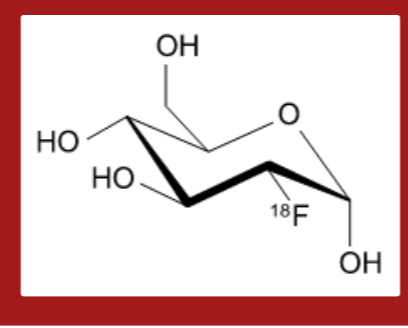


Background and Objectives

- Evaluation of tumor glucose metabolism has been suggested as an alternative to tumor size to assess early tumor response to therapy in cases where clinical benefit is observed but a change in tumor size is limited or delayed (e.g. cytostatic drugs) [1].
- Sunitinib (Sutent®) is a multi-targeted tyrosine kinase inhibitor with anti-angiogenic properties, approved for the treatment of imatinib-resistant or -intolerant gastro-intestinal stromal tumors (GIST).

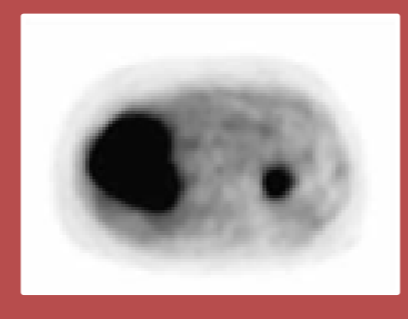
Radiotracer administration

- [¹⁸F]-FDG intravenous administration
- 60-minutes uptake period



Whole body PET/CT scanning

- Selection of up to 5 lesions evaluable by CT
- Measurement of radiotracer concentration in the region of interest (ROI)



Standardized uptake value (SUV) calculation in the ROI

- Q : tumor radiotracer concentration in ROI (MBq/l)
- Q_{inj} : injected activity (MBq)
- LBW: lean body weight (kg)

$$SUV = \frac{Q \cdot LBW}{Q_{inj}}$$

Fig. 1: FDG-PET scan procedure and SUV calculation

The objectives of the present study are:

- To investigate potential relationships between sunitinib exposure and the time-course of the maximum and mean SUV (SUV_{max} and SUV_{mean} [2]) of individual lesions in GIST patients treated with sunitinib
- To characterize both inter-individual (IIV) and inter-lesion variability (ILV) in SUV responses
- To investigate potential relationships between SUV_{max} and overall survival (OS)

Methods

Patients and Data

Sixty-six imatinib-resistant GIST patients treated with sunitinib were included in the analysis [3].

Table 1: Summary of the data

Dosing schedule (weeks on/off)	6-week cycles (4/2): 50 mg q.d. 4-week cycles (2/2): 25, 50, 75 mg q.d. 3-week cycles (2/1): 50 mg q.d.
Pharmacokinetics	Individual PK parameters for 44 patients [4] Typical values for 22 patients [4]
SUV for individual lesions	607 baseline and post-baseline SUV_{max} and SUV_{mean} data from 172 lesions Up to 5 lesions per patient Median duration of follow-up: 10 weeks Maximum duration of follow-up: 102 weeks

Individual lesion SUV model

- Structural model:** Indirect response (IDR) models with inhibition of the production (R_{in}) or stimulation of the loss (k_{out}) of response were investigated to describe the time-course of SUV of individual lesions. Linear, power and E_{max} drug-effect relationships driven by the daily dose or the daily AUC were considered during model building. Linear and non-linear disease progression models were tested.
- ILV model:** The ILV model for a parameter θ_{ij} for the i^{th} individual and the j^{th} lesion can be written as:

$$\theta_{ij} = \theta \cdot \exp(\eta_i + \kappa_{ij})$$
 where θ is the typical value in the population, η model between individual differences and κ model between lesion differences, with $\eta \sim N(0, \omega^2)$ and $\kappa_j \sim N(0, \pi_j^2)$. A common variance is assumed between lesions, i.e. $\pi_1^2 = \pi_2^2 = \dots = \pi_L^2$. Both IIV and ILV were evaluated in all model parameters.
- Residual error model:** Individual lesions' SUV assessed from the same FDG-PET scan were allowed to have different residual error values (ϵ), which arose from a multivariate normal distribution parameterized with a zero mean vector and a covariance matrix Σ .
 - ✓ The diagonal elements of Σ containing the variances for each ϵ were assumed to be the same.
 - ✓ The off-diagonal elements of Σ containing the covariances between ϵ were assumed to be the same, denoting the same correlation between the SUV of all lesions assessed simultaneously.
 - ✓ This was implemented in NONMEM through a Cholesky decomposition of the sigma matrix and the level-2 (L2) item.
- Model development was performed using SUV_{max} data. The final SUV_{max} structural model was applied to SUV_{mean} data.

Overall survival model

- A parametric time-to-event model was developed to describe the OS data.
 - ✓ Weibull and exponential distributions were evaluated to describe the baseline hazard for OS.
 - ✓ The following predictors were evaluated on OS:
 - Maximum individual lesion SUV_{max} at baseline (observed and predicted)
 - Summed SUV_{max} at baseline (observed and predicted)
 - Summed SUV_{max} time-course
 - Relative change from baseline in summed SUV_{max} over time
 - Maximum relative change in individual lesion SUV_{max} from baseline at week 1 and at week 2
 - Baseline observed tumor size (sum of longest diameters, SLD)
 - ✓ Censoring was described by an exponential model.

Estimation and simulations were performed using NONMEM 7.3.

References

- [1] Wahl et al. Nucl Med. 2009; 50 Suppl 1): p. 122S-50S.
[2] Vanderhoek et al. Nucl Med. 2013; 54(8): p.1188-94.
[3] Demetri et al. Clin Cancer Res. 2009; 15(18): p. 5902-9.
[4] Houk et al. Clin Cancer Res. 2009; 15(7): p.2497-506.

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Results

Individual lesion SUV model

- Log-transformed individual lesion SUV_{max} data were well described by an IDR model with stimulation of k_{out} through a linear drug effect driven by daily AUC:

$$\frac{dSUV}{dt} = R_{in} - k_{out} \cdot (1 + DRUG \cdot AUC_{daily}) \cdot SUV(t)$$

with $R_{in} = Base \cdot k_{out}$ and $SUV_{t=0} = Base$

- The estimated IIV was larger than the estimated ILV for both SUV_{max} baseline (Base) and the drug effect parameter (DRUG).
- The typical doubling time of SUV_{max} for return to baseline during off-treatment periods was ~2 weeks.
- The model predicts a typical decrease in SUV of 49% after 14 days of sunitinib treatment (50 mg q.d.).
- No significant disease progression was identified.
- VPCs illustrate the capability of the model to predict the drug effect on individual lesion SUV_{max} (Figure 2A), as well as the summed SUV_{max} of all lesions (Figure 2B) at each time point.
- In addition, the final structural model successfully described individual lesion SUV_{mean} data.

Table 2: Final parameter estimates for the individual lesion SUV_{max} and SUV_{mean} models

Parameter	Individual lesion SUV_{max}			Individual lesion SUV_{mean}		
	Typical value (RSE%)	IIV CV% (RSE%)	ILV CV% (RSE%)	Typical value (RSE%)	IIV CV% (RSE%)	ILV CV% (RSE%)
Base	7.73 (6)	33 (15)	23 (17)	5.81 (6)	29 (16)	25 (15)
k_{out} (week ⁻¹)	0.378 (28)	-	-	0.340 (33)	-	-
DRUG (AUC ⁻¹)	1.03 (18)	60 (33)	51 (21)	1.00 (19)	64 (28)	51 (23)
Residual error (%)	43.0 (16)	-	-	43.5 (15)	-	-

Base: individual lesion SUV baseline value; DRUG: SUV reduction rate constant induced by sunitinib; k_{out} : loss of response rate constant; RSE: relative standard error.

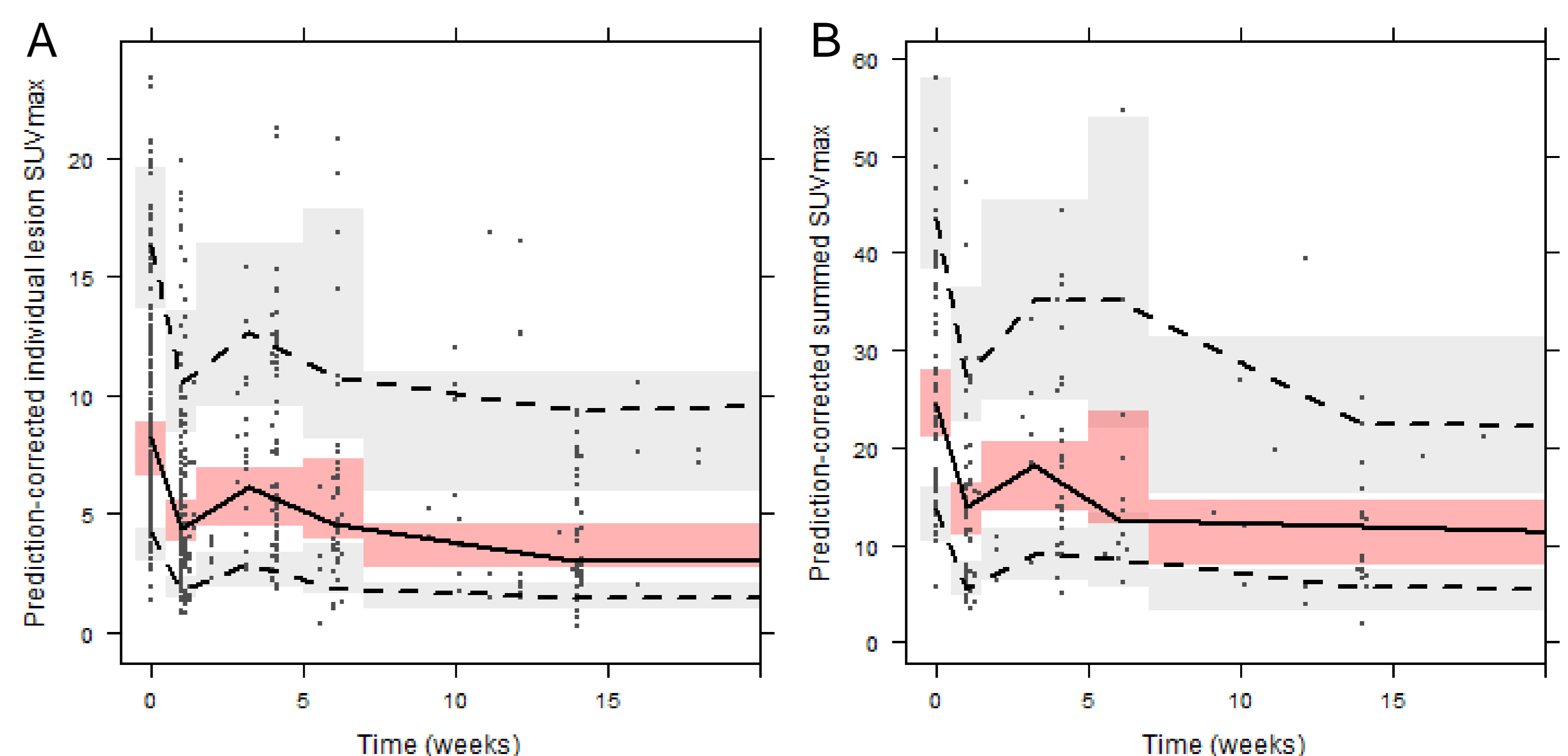


Fig. 2: Visual predictive checks of the final individual lesion SUV_{max} model.

(A) Individual lesion SUV_{max} (B) Summed SUV_{max} from all target lesions. Median (solid line), 10th and 90th percentiles (dashed lines) of the observed data are compared to the 95% confidence intervals (shaded areas) for the median, 10th and 90th percentiles of the simulated data (based on 1000 simulations).

Overall survival model

- The underlying hazard for OS was best described by an exponential distribution.
- The relative change from baseline in summed SUV_{max} over time was the most significant predictor for OS (dOFV = -25.2), followed by the maximum relative change in individual lesion SUV_{max} from baseline at week 1 (dOFV = -6.0). In both cases, the hazard decreases as the relative reduction in SUV increases.

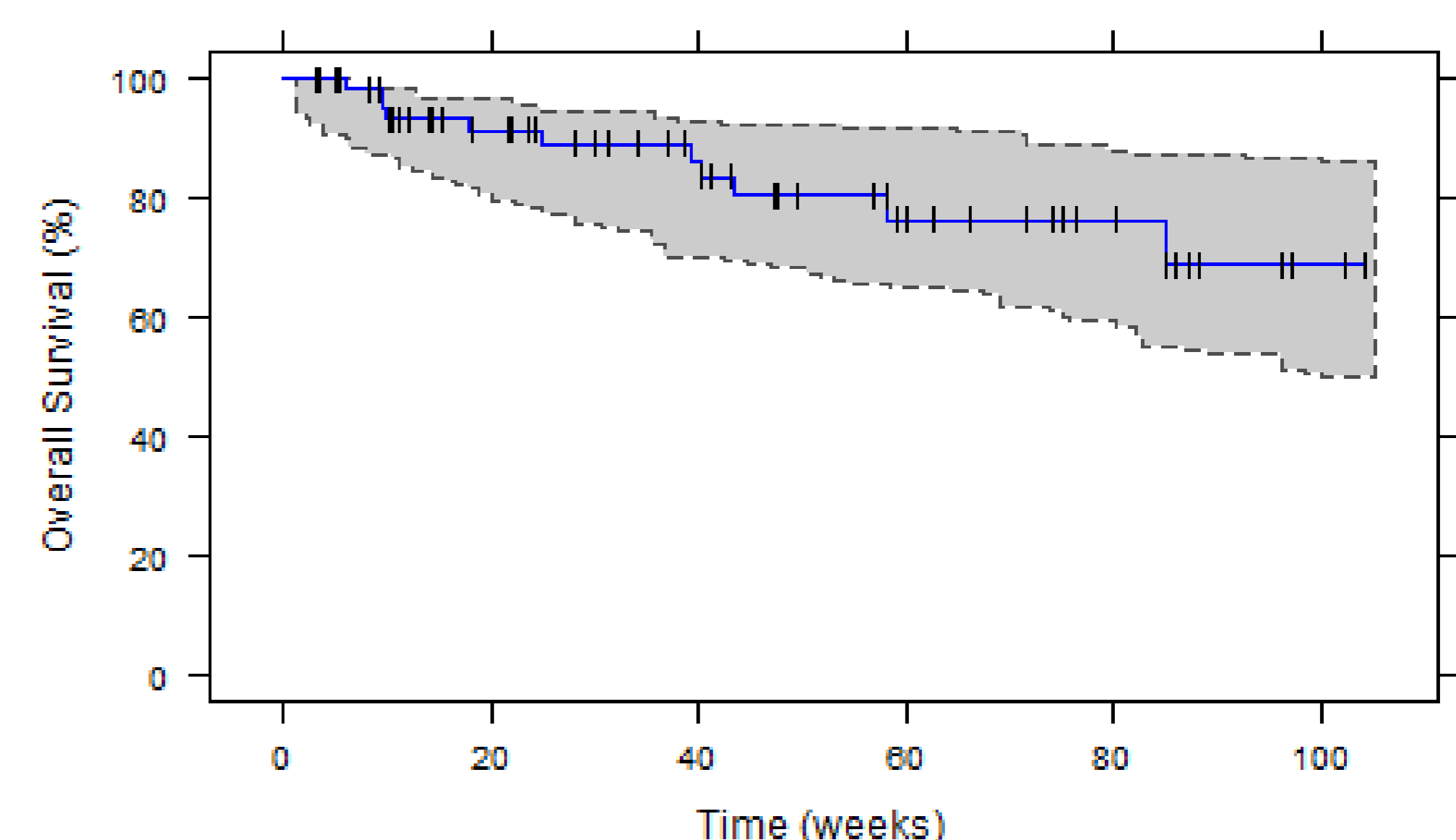


Fig. 3: Visual predictive check for the Kaplan-Meier survival curve. The observed Kaplan-Meier curve (solid line) is compared to the 90% confidence interval (shaded area) derived from model simulations (based on 200 samples) of the survival model including the relative change from baseline in the summed SUV_{max} across lesions. Vertical black lines represent censored observations

Discussion and Conclusions

- The developed model appropriately described the time-course of individual lesion SUV_{max} and SUV_{mean} in GIST patients treated with sunitinib. Significant IIV and ILV in SUV responses could be identified.
- Preliminary results showed that SUV-based metrics are related to OS, supporting that FDG-PET can be of interest to early predict clinical outcome during treatment.
- This methodology can be used to leverage data collected on individual lesions (e.g. tumor size, volume, metabolic activity) during oncologic clinical trials.