



PRECLINICAL PHARMACOKINETIC REPORT

Developmental Biology and Solid Tumor Program

P-PKSR Study 186814 - 1936417

STUDY TITLE:

SCREENING PLASMA AND TUMOR PHARMACOKINETICS (SPTPK) OF GEFITINIB IN FEMALE ATHYMIC NUDE MICE AFTER A SINGLE ORAL DOSE

SHORT TITLE: Gefitinib Screening Plasma and Tumor PK (SPTPK)

TEST ARTICLE: Gefitinib

SECTION: Nonclinical Pharmacokinetics (Non-GLP)

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Gefitinib Screening Plasma and Tumor PK (SPTPK)

Quality Statement

This non-GLP study was conducted using sound scientific principles and established techniques in accordance with the relevant guidelines and standard operating procedures (SOPs) of the Preclinical Pharmacokinetic Shared Resource (P-PKSR) and St. Jude Children's Research Hospital (SJCRH), Memphis, TN, USA. This report accurately reflects the data obtained during the course of this study.

These results represent part of an early phase preclinical pharmacology program. This study has been conducted to provide preliminary insights into the pharmacokinetic (PK) properties of the compound(s) in the indicated preclinical model(s). This study and its results are not intended to provide a comprehensive PK evaluation of the compound(s). The applied bioanalytical method was validated/qualified to support this specific study and discovery-style sample analyses.

Substantial study-to-study and inter-animal variability in preclinical PK exists. Such variability depends upon the in vivo scientists' experience, variations in compound purity and formulation, animal strains, sex and age, and other situational fixed effects (i.e. husbandry conditions, chow constituents, presence or absence of disease, concomitant drugs). As such, the actual PK, plasma or tissue compound concentrations, or equivalent dose in other studies or preclinical models may vary significantly from that reported herein.

Gefitinib Screening Plasma and Tumor PK (SPTPK)

1.0 METHODS

1.1 In Vivo Pharmacokinetic (PK) STUDY

The plasma and tumor pharmacokinetic (PK) profile of gefitinib was evaluated in female Athymic Nude mice (Charles River) approximately 12 weeks in age, bearing rhabdomyosarcoma (RMS) MAST39 tumors. The test article gefitinib (SJ-000520503-12, Santa Cruz Biotechnology, SC-202166, J2309) was suspended in 0.5% Hydroxypropyl Methylcellulose (HPMC) / 0.1% Tween 80 / 0.5% lactic acid in ultrapure water, at 1.5 mg/mL for a 15 mg/kg free base equivalent dose as a 10 mL/kg oral gavage. Terminal samples were collected over a 16 hour post-dose period by cardiac puncture using a 1 mL syringe, and the blood placed in a Sarstedt Microvette K3EDTA 500 μ L tube, and immediately processed to plasma. The carcass was then perfused with PBS, the tumor extracted, rinsed, and placed in a microcentrifuge tube. All samples were immediately stored on dry ice and transferred to -80 °C until analysis.

1.2 Bioanalysis

Tumor samples were weighed in 15 mL Lysing matrix D (MP Biomedicals, Santa Ana, CA), diluted with a 1:5 volume of ultrapure water, and homogenized using a FastPrep-24 system (MP Biomedicals, Santa Ana, CA) for five cycles of 1 min vibration at 6.5 M/S speed, with 5 min in ice bath between each cycle to prevent over-heating. The homogenates were then stored at -80 °C until analysis.

Plasma and tumor samples were analyzed for gefitinib (SJ000520503, Santa_Cruz, Lot # J2309, purity 99%) using a qualified liquid chromatography – tandem mass spectrometry (LC-MS/MS) assay. Plasma and tumor homogenate calibrators and quality controls were spiked with solutions, corrected for salt content and purity as necessary, prepared in methanol. Plasma samples, 25 μ L each, were protein precipitated with 150 μ L of 30 ng/mL afatinib dimaleate (SJ000855122, OCHEM, CAT# 140A737, Lot # 150318G1, purity 99%) in acetonitrile as an internal standard (IS). A 3 μ L aliquot of the extracted supernatant was injected onto a AB Sciex ExionLC high performance liquid chromatography system via a AB Sciex ExionLC autosampler. The LC separation was performed using a Waters BEH C18 (2.5 μ m, 75 mm x 2.1 mm) column maintained at 40 °C with gradient elution at a flow rate of 0.35 mL/min. The binary mobile phase consisted of water-formic acid (100:0.1 v/v) in reservoir A and acetonitrile-formic acid (100:0.1 v/v) in reservoir B. The initial mobile phase consisted of 5% B for 2.0 min with a linear increase to 95% B in 1.5 min. The column was then rinsed for 1.5 min at 95% B and returned to initial mobile phase conditions in 0.5 mins, then equilibrated the column for another 2.0 min for a total run time of 7.5 min. Under these conditions, the analyte and IS eluted at 4.05 and 4.06 min, respectively. For tumor samples analysis, the run time was shortened to 6.5 minutes by employing the initial mobile phase conditions of 5%B for 1.0 min with a linear increase to 98%B in 1.5 min and holding 98%B for another 1.5 min for column rinse and returned to initial conditions in 0.5 min, then equilibrated the column for another 2.0 mins. Gefitinib and afatinib were eluted at 3.04 and 3.05 min respectively under these gradient conditions.

Analyte and IS were detected with tandem mass spectrometry using a SCIEX Triple Quad™ 3500 LC-MS/MS system in the positive ESI mode and the following mass transitions were monitored: gefitinib 447.2 \rightarrow 128, afatinib 486.3 \rightarrow 371.1. The method qualification and bioanalytical runs all passed acceptance criteria for non-GLP assay performance. A linear model ($1/X^2$ weighting) fit the plasma and homogenate calibrators across the 1 to 100 ng/mL range, with a correlation coefficient (R) of ≥ 0.9972 . The lower limit of quantitation (LLOQ), defined as a peak area signal-to-noise ratio of 5 or greater verses a matrix blank with IS, was 1 ng/mL for both plasma and tumor homogenates. Sample dilution integrity was confirmed. The intra-run precision and accuracy was $\leq 12.43\%$ CV and 91% to 105%, respectively.

Gefitinib Screening Plasma and Tumor PK (SPTPK)

1.3 Pharmacokinetic (PK) Analysis

Plasma and tumor concentration-time (Ct) data for gefitinib were grouped by matrix and nominal time point. Manual imputation of data below the lower limit of quantitation (BLOQ) was as follows: IF at any time point $\geq 2/3$ of the Ct results were above the LLOQ, the BLOQ data were replaced with a value of $1/2$ LLOQ, ELSE the entire time point's data were treated as missing. Summary statistics were calculated, and the arithmetic mean Ct values were subjected to noncompartmental analysis (NCA) using Phoenix WinNonlin 8.1 (Certara USA, Inc., Princeton, NJ). The extravascular model was applied, and area under the Ct curve (AUC) values were estimated using the "linear up log down" method. The terminal phase was defined as at least three time points at the end of the Ct profile, and the elimination rate constant (Kel) was estimated using an unweighted log-linear regression of the terminal phase. The terminal elimination half-life (T_{1/2}) was estimated as $0.693/\text{Kel}$, and the AUC from time 0 to infinity (AUC_{inf}) was estimated as the AUC to the last time point (AUC_{last}) + C_{last} (predicted)/Kel. Other parameters estimated included observed maximum concentration (C_{max}), time of C_{max} (T_{max}), concentration at the last observed time point (C_{last}), time of C_{last} (T_{last}), apparent clearance (CL/F = Dose/AUC_{inf}), and apparent terminal volume of distribution (V_z/F). The apparent plasma-to-tumor partition coefficient (K_{p,inf}) was estimated as the ratio of the AUC_{inf} in tissue to AUC_{inf} plasma, whereas K_{p,last} was similarly estimated using AUC_{last} values.

A clinically relevant dose (CRD) for mice was estimated from unbound plasma PK and exposure. The CRD was defined as the mouse dose achieving a predicted mean steady state unbound plasma AUC (AUC_u) similar to humans at the single agent maximum tolerated dose (MTD), recommended Phase II dose (RP2D), or FDA-approved dose. Dose proportional, linear, and time-invariant PK across species was assumed. Human and mouse plasma protein binding were assumed similar when data were not available. This is similar to the clinical relevance approach proposed by Spilker [1] which uses unbound plasma average steady state concentrations. Some latitude in dose rounding was permitted in the CRD recommendation, and an unbound exposure within 2-fold of the clinical target was considered acceptable. Additional considerations influenced the final recommended mouse dose, including mouse dosing regimens prevalent in the literature and the tolerability of the compound in mice.

2.0 RESULTS

The gefitinib Ct data demonstrated high variability between and within mice, with coefficients of variation ranging from 6.15% to 171%. The tumor Ct data was extremely variable, with CVs averaging 120%. Most of the variability in plasma concentrations was seen during the putative absorption phase. The absorption rate of gefitinib was rapid, with the T_{max} occurring at 0.5 hours post-dose for plasma, whereas the tumor appearance rate was moderate with the T_{max} occurring at 1 hour. After C_{max}, plasma and tumor concentrations diminished in a bi-exponential manner. The apparent plasma and tumor terminal half-life of gefitinib were 6.02 and 4.79 hours respectively. The apparent plasma clearance (CL/F) of gefitinib was moderate at 41 mL/min/kg, or approximately 45% of murine hepatic blood flow. The apparent plasma terminal volume of distribution (V_z/F) for gefitinib was also high at 21.4 L/kg, in excess of total body water. The plasma to tumor partitioning of gefitinib was moderate, with a K_{p,inf} of 0.304, and K_{p,last} of 0.329. Compared with our previous plasma PK study using 15 mg/kg in a non-acidified suspension (RPT.186810-1936320), the current plasma C_{max} and AUC_{inf} values were 4- and 3-fold higher, respectively.

In a pediatric Phase 1 trial in combination with irinotecan, the MTD of gefitinib was 150 mg/m² PO QD [2]. The single agent MTD of gefitinib has been estimated at 400 mg/m² PO QD in children, with the total plasma AUCs at steady state being 12900 and 29300 hr-ng/mL, respectively for 150 and 400 mg/m² [3]. The fraction unbound in plasma (F_{u,p}) for gefitinib has been estimated as 0.034 and 0.051 in humans and mice, respectively [4]. Using the PK findings from our current study, gefitinib CRDs calculated by unbound AUCs between humans and mice range from 20 to 50 mg/kg PO QD.

Gefitinib Screening Plasma and Tumor PK (SPTPK)

3.0 REFERENCES

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Gefitinib Screening Plasma and Tumor PK (SPTPK)

4.0 TABLES, LISTINGS, AND FIGURES (TLFS)

Figure 4.1: Mean (SD) Ct Profiles by Analyte and Group

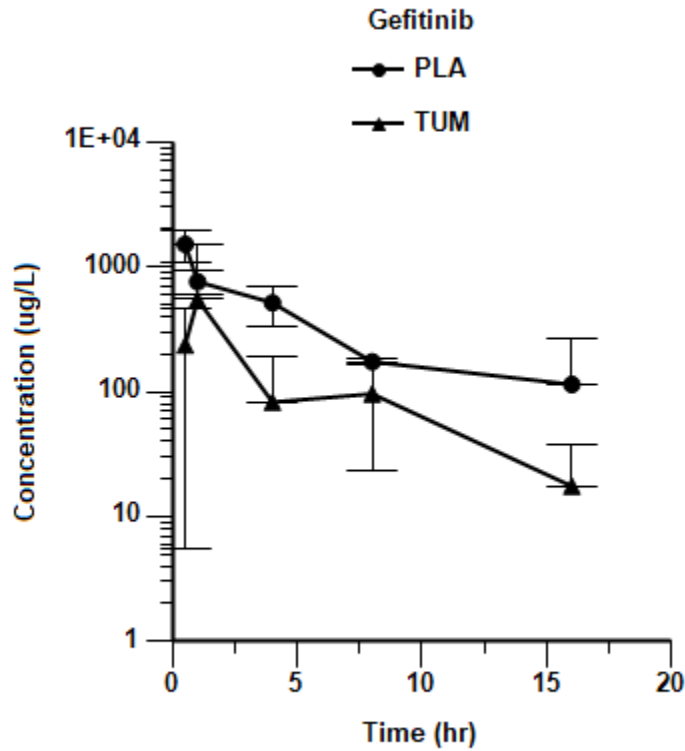


Table 4.1: NCA Parameter Estimates by Analyte and Group

		Analyte	
		Gefitinib	
		Group	
		PLA	TUM
Parameter	Unit	Value	
Cmax	ug/L	1520	548
Tmax	hr	0.500	1.00
AUClast	hr*ug/L	5210	1720
AUCinf	hr*ug/L	6090	1860
Kel	1/hr	0.115	0.141
T1/2	hr	6.02	4.91
CL/F	L/hr/kg	2.46	8.05
Vz/F	L/kg	21.4	57.1
Clast	ug/L	114	17.5

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		Analyte	
		Gefitinib	
		Group	
		PLA	TUM
Parameter	Unit	Value	
Tlast	hr	16.0	16.0
Kp_last			0.330
Kp_inf			0.306

Table 4.2: Full Summary Statistics of Ct Data by Analyte and Group

		Analyte	
		Gefitinib	
		Group	
		PLA	TUM
Time (hr)		Concentration (ug/L)	
0.500	N	3	3
	Mean	1520	236
	SD	422	231
	Min	1030	47.1
	Median	1720	168
	Max	1800	493
	CV%	27.8	97.7
	Geometric Mean	1470	158
	CV% Geometric Mean	31.6	173
1.00	N	3	3
	Mean	761	548
	SD	169	937
	Min	565	3.00
	Median	853	9.64
	Max	864	1630
	CV%	22.3	171
	Geometric Mean	747	36.1
	CV% Geometric Mean	24.5	27400
4.00	N	3	3
	Mean	516	82.6
	SD	180	108
	Min	367	17.8
	Median	464	22.2
	Max	716	208
	CV%	35.0	131
	Geometric Mean	496	43.4
	CV% Geometric Mean	34.9	231
8.00	N	3	3
	Mean	174	96.2

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		Analyte	
		Gefitinib	
		Group	
		PLA	TUM
Time (hr)		Concentration (ug/L)	
	SD	10.7	73.0
	Min	161	12.9
	Median	179	126
	Max	180	149
	CV%	6.15	75.9
	Geometric Mean	174	62.4
	CV% Geometric Mean	6.28	235
16.0	N	3	3
	Mean	114	17.5
	SD	151	19.7
	Min	12.1	3.00
	Median	43.5	9.64
	Max	287	39.9
	CV%	132	112
	Geometric Mean	53.3	10.5
	CV% Geometric Mean	341	209

Table 4.3: Ct Data Listings by Subject, Analyte, Group, and Time

Subject	Analyte	Group	Time (hr)	Concentration (ug/L)
M1	Gefitinib	PLA	0.500	1800.50
M1	Gefitinib	TUM	0.500	168.29
M2	Gefitinib	PLA	0.500	1717.10
M2	Gefitinib	TUM	0.500	493.49
M3	Gefitinib	PLA	0.500	1031.60
M3	Gefitinib	TUM	0.500	47.09
M4	Gefitinib	PLA	1.00	853.42
M4	Gefitinib	TUM	1.00	9.64
M5	Gefitinib	PLA	1.00	863.59
M5	Gefitinib	TUM	1.00	1629.90
M6	Gefitinib	PLA	1.00	565.14
M6	Gefitinib	TUM	1.00	3.00
M7	Gefitinib	PLA	4.00	715.94
M7	Gefitinib	TUM	4.00	17.79
M8	Gefitinib	PLA	4.00	366.83
M8	Gefitinib	TUM	4.00	207.71
M9	Gefitinib	PLA	4.00	463.87
M9	Gefitinib	TUM	4.00	22.16

Gefitinib Screening Plasma and Tumor PK (SPTPK)

Subject	Analyte	Group	Time (hr)	Concentration (ug/L)
M10	Gefitinib	PLA	8.00	161.41
M10	Gefitinib	TUM	8.00	12.88
M11	Gefitinib	PLA	8.00	179.42
M11	Gefitinib	TUM	8.00	126.29
M12	Gefitinib	PLA	8.00	180.40
M12	Gefitinib	TUM	8.00	149.28
M13	Gefitinib	PLA	16.0	12.13
M13	Gefitinib	TUM	16.0	3.00
M14	Gefitinib	PLA	16.0	43.54
M14	Gefitinib	TUM	16.0	9.64
M15	Gefitinib	PLA	16.0	287.33
M15	Gefitinib	TUM	16.0	39.94

Table 4.4: Ct Summary (Mean, SD, N) by Analyte and Group

Analyte	Group	Time (hr)	Mean (ug/L)	SD (ug/L)	N
Gefitinib	PLA	0.500	1520	422	3
Gefitinib	PLA	1.00	761	169	3
Gefitinib	PLA	4.00	516	180	3
Gefitinib	PLA	8.00	174	10.7	3
Gefitinib	PLA	16.0	114	151	3
Gefitinib	TUM	0.500	236	231	3
Gefitinib	TUM	1.00	548	937	3
Gefitinib	TUM	4.00	82.6	108	3
Gefitinib	TUM	8.00	96.2	73.0	3
Gefitinib	TUM	16.0	17.5	19.7	3

5.0 ATTACHED FILES

- Attached File 5.1** Gefitinib Screening Plasma PK v2.0.docx – *Final in vivo study plan as executed*
- Attached File 5.2** Gefitinib Screening Plasma PK TLFs.docx – *Report TLFs as a Word document for manipulation, plotting, and further presentation*
- Attached File 5.3** Copy of 186814-1936417_GEF_SPTPK_2020-10-26.xlsx – *Study digital data collection form*
- Attached File 5.4** IMG_9491.jpg – *Photocopy of paper study data collection form*