

SUPPLEMENTARY TABLES

Supplementary Table 1. The microsomal stabilities of the compounds.

Compound	Human liver microsomes		Rat liver microsomes		Mouse liver microsomes	
	$t_{1/2}$ (min)	CL_{int} (mL/min/kg)	$t_{1/2}$ (min)	CL_{int} (mL/min/kg)	$t_{1/2}$ (min)	CL_{int} (mL/min/kg)
Bortezomib	21.4	58.3	12.4	200.5	10.7	508.7
NNU219	32.4	43	27.8	50	16.7	83

Note: The microsomal stabilities of bortezomib and NNU219 were determined in the presence of pooled human, rat or mouse liver microsomes. $T_{1/2}$ (min) and CL_{int} (mL/min/kg) were detected by LC/MS. Antipyrine and testosterone (5 μ M each) were used as positive and negative controls, respectively. Abbreviations: t = time; CL_{int} , intrinsic clearance.

Supplementary Table 2. Pharmacokinetic profiles of NNU546 in mice.

	$t_{1/2}$ (h)	CL (mL·h ⁻¹ ·kg ⁻¹)	V _z (mL·kg ⁻¹)	AUC _{0-t} (h·ng·mL ⁻¹)	MRT (h)	F(%)
i.v.	2.08±0.991	986±75.4	2914±1247	1890±73.8	1.55±0.133	---
i.g.	2.41±0.420	---	---	208±26.7	2.35±0.221	11.0±1.41

Note: Blood samples were collected at baseline and after intravenous or oral administration of 2 mg/kg of NNU546. Each time-point represents the average value of three animals. NNU546 concentration in blood and plasma samples was determined by liquid chromatography-tandem mass spectrometry (LC/MS/MS) in a non-good laboratory practice lab. Pharmacokinetic analysis of the blood and plasma concentration data was performed using WinNonlin version 5.2 (Pharsight Corp.). Kinetic parameters were estimated using a noncompartmental model with sparse sampling mode (model 201 for plasma and blood). Area under the concentration vs. time curve (AUC) was calculated using the linear trapezoidal rule. Abbreviations: t = time; CL, clearance; V_z, apparent volume of distribution; AUC_{0-t}, AUC (area under the curve) from 0 to t h; MRT, mean residence time; F%, oral bioavailability.