## Pre-Clinical Pharmacokinetic Pharmacodynamic Modelling study of 4-Hydroxyisoleucine using Validated Ultra Performance Liquid Chromatography-Tandem Mass Spectroscopy

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## Supplementary data

Extracting solvent	Sample volume (µL)	Volume of extraction solvent added (mL)	Vortex time (min)	Centrifugation [speed (g), time (min)]	% Recovery	Remarks
Methanol	300	3	2	15000, 10	28-31	Inconsistent and poor recovery
Acetonitrile	300	3	2	15000, 10	40-42	Consistent recovery was noticed

Table S1.: Optimization of protein precipitation extraction method for sample preparation

Table S2.: Optimization parameters of liquid chromatography and mass spectroscopy

Sr.No.	Chromatographic	Parameters	Remarks		
	condition				
1.	Mode	ES-	Low intensity of peak		
		ES+	High Intensity of peak		
Based on	the results, ESI+ mode wa	as selected and used further f	or optimizing the chromatographic conditions		
2.	Mobile Phase	ACN: Water	No proper ionization of analytes which results		
	(50: 50)		in decreased peak intensity		
		ACN: 0.1% glacial	Splitting in the peaks was observed		
		acetic acid			
		ACN: 0.1% formic acid	Symmetrical peak with high peak intensity		
From these re	sults, it was conformed th	at the ACN: 0.1% formic aci	d was used as mobile phase and further optimized		
3.	Mobile phase	40:60	Retention time was more than 2 min		
	composition (ACN:	30: 70	Peak splitting was noticed		
	0.1% formic acid)	20:80	Symmetric peaks of analyte and IS was		
			observed with proper resolution		
Based on	the results, it was conclude	ed that ACN:0.1% formic aci	d with 20:80 was used as the mobile phase for		
	further m	ethod development and valid	lation of parameters		

Compound	Formula	Parent m/z	Cone voltage	Daughters	Collision	Ion mode
Name					energy	
4-Hydroxy	C <sub>6</sub> H <sub>13</sub> NO <sub>3</sub>	148.19	24	74.02	12	ES+
isoleucine			24	102.13	10	ES+
			24	84.16	18	ES+
			24	95.88	14	ES+
L-isoleucine	$C_6H_{13}NO_2$	132.17	24	86.19	12	ES+
			24	44.08	18	ES+
			24	69.04	20	ES+
			24	41.08	24	ES+

**Table S3.:** Intellistart MRM Optimization of analytes

Table S4.: Regression parameters of the calibration curve generated for each weighting factor.

Wi	b	a	r <sup>2</sup>
unweighted	9.65e-06	6.82E-04	0.9993
1/var	1.02e-05	6.86E-04	0.9990
$1/x^{2}$	1.02e-05	6.49E-04	0.9991
1/x	1.02e-05	6.17E-04	0.9988
$1/y^{2}$	1.05e-05	6.25E-04	0.9966
1/y	1.03e-05	6.16E-04	0.9291

Wi, weighing factor; b, slope; a, constant; r<sup>2</sup>, regression co-efficient

Based on the above results,  $1/x^2$  was selected for further analysis based on the regression coefficient value.<sup>1</sup>

## Reference

 H. Gu, G. Liu, J. Wang, A.-F. Aubry and M. E. Arnold, *Anal. Chem.*, 2014, 86, 8959– 8966.



Fig S1.: Splitting pattern for 4-hydroy isoleucine and l-isoleucine



Fig S2.: Statistical residuals plotted against concentrations (ng/mL) (a) 1/variance; (b)  $1/x^2$ ; (c) 1/x; (d)  $1/y^2$ ; (e) 1/y