

## SUPPORTING INFORMATION

### Bromotyrosine-Derived Metabolites from a Marine Sponge Inhibit

#### *Pseudomonas aeruginosa* Biofilms

**Tam M. T. Tran**<sup>1</sup>, **Russell S. Addison**<sup>2</sup>, **Rohan A. Davis**<sup>3,4</sup>, **Bernd H. A. Rehm**<sup>1,5,\*</sup>

<sup>1</sup> Centre for Cell Factories and Biopolymers, Griffith Institute for Drug Discovery, Griffith University, Nathan, QLD 4111, Australia

<sup>2</sup> Preclinical ADME/PK, Griffith Institute for Drug Discovery, Griffith University, Nathan, QLD 4111, Australia

<sup>3</sup> NatureBank, Griffith Institute for Drug Discovery, Griffith University, Nathan, QLD 4111, Australia

<sup>4</sup> School of Environment and Science, Griffith University, Nathan, QLD 4111, Australia

<sup>5</sup> Menzies Health Institute Queensland, Griffith University, Gold Coast, QLD 4222, Australia

\* correspondence: [b.rehm@griffith.edu.au](mailto:b.rehm@griffith.edu.au)

**Table S1.** The degradation of ianthelliformisamine A in mouse liver microsomes.

**Table S2.** Metabolic stability parameters for ianthelliformisamine A in mouse liver microsomes.

**Table S3.** The degradation of ianthelliformisamine B in mouse and human liver microsomes.

**Table S4.** Metabolic stability parameters for ianthelliformisamine B in mouse and human liver microsomes.

**Table S5.** The degradation of ianthelliformisamine C in mouse and human liver microsomes.

**Figure S1.** Metabolic stability parameters for ianthelliformisamine A in mouse liver microsomes.

**Figure S2.** Metabolic stability of ianthelliformisamine B in mouse and human liver microsomes.

**Figure S3.** Metabolic stability of ianthelliformisamine C in mouse and human liver microsomes.

**Table S1.** The results for the degradation of ianthelliformisamine A (**1**) normalized to t = 0) in mouse liver microsomes

Time (min)	% Remaining	
	Mouse	
	Mean	SD
0	100.00	0.00
5	93.32	14.29
10	96.74	27.23
20	66.26	19.36
30	57.28	20.01
60	53.42	17.07

**Table S2.** Metabolic stability parameters for ianthelliformisamine A (**1**) based on NADPH-dependent degradation profiles in mouse liver microsomes.

Parameter	Units	Result
$k^1$	min <sup>-1</sup>	0.0218 ± 0.0125
Half life <sup>1</sup>	min	38.2 ± 17.2
$Cl_{int, in vitro}$	μL/min/mg protein	43.6
$Cl_{int}$	mL/min/kg	112.5
$Cl_{blood}$	mL/min/kg	58.1
Hepatic Extraction ( $E_H$ ) <sup>2</sup>	---	0.48

<sup>1</sup> mean ± SD, n = 3

<sup>2</sup> based on mean data for k

**Table S3.** The results for the degradation of ianthelliformisamine B (**2**) (normalized to t = 0) in mouse and human liver microsomes.

Time (min)	% Remaining			
	Mouse		Human	
	Mean	SD	Mean	SD
0	100.00	0	100.00	0
5	153.98	14.66	112.15	16.61
10	160.70	65.50	98.42	9.88
20	138.91	39.12	80.61	3.04
30	119.60	41.51	68.41	7.28
60	122.66	35.50	76.42	16.88

**Table S4.** Metabolic stability parameters for ianthelliformisamine B (**2**) based on NADPH-dependent degradation profiles in mouse and human liver microsomes.

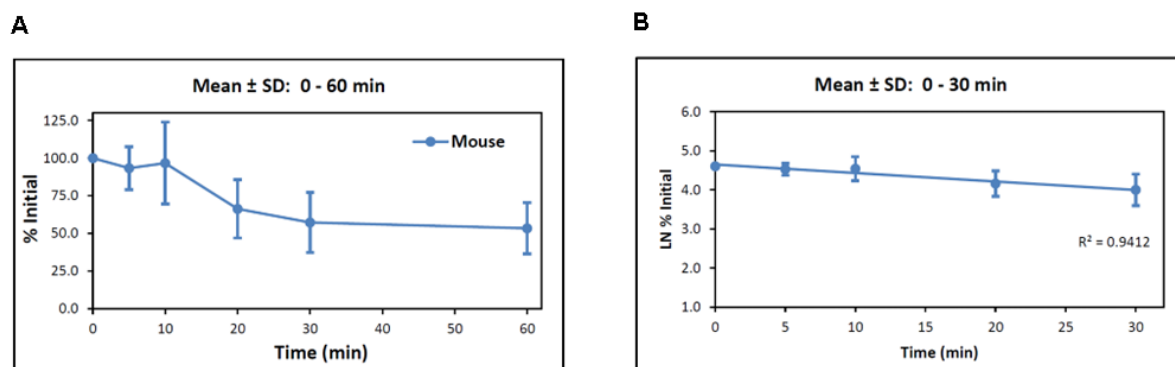
Parameter	Units	Mouse	Human
$k^1$	min <sup>-1</sup>	0.0165 ± 0.0030	0.0195 ± 0.0011
Half life <sup>1</sup>	min	42.8 ± 7.6	35.7 ± 2.1
$Cl_{int, in vitro}$	μL/min/mg protein	33.0	39.0
$Cl_{int}$	mL/min/kg	85.1	32.1
$Cl_{blood}$	mL/min/kg	49.8	12.6
Hepatic Extraction ( $E_H$ ) <sup>2</sup>	---	0.42	0.61

<sup>1</sup> mean ± SD, n = 3

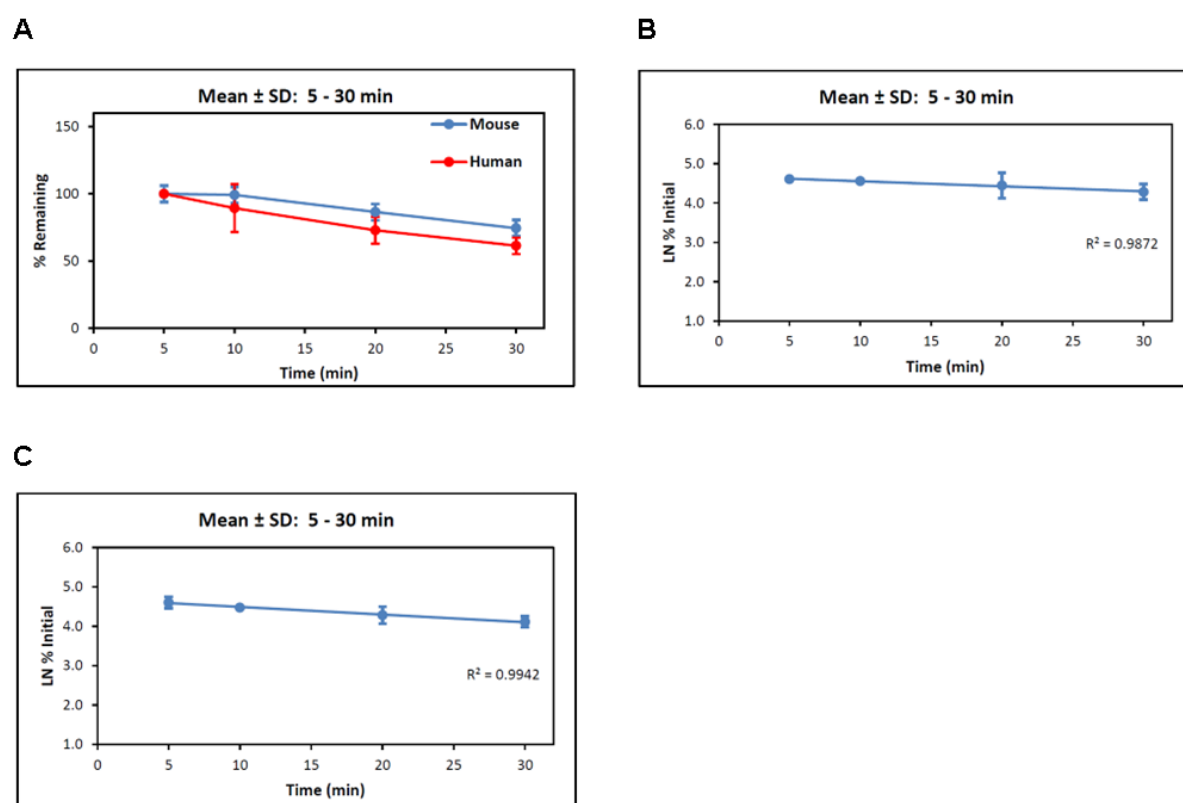
<sup>2</sup> based on mean data for k

**Table S5.** The results for the degradation of ianthelliformisamine C (**3**) normalized to t = 0) in mouse and human liver microsomes.

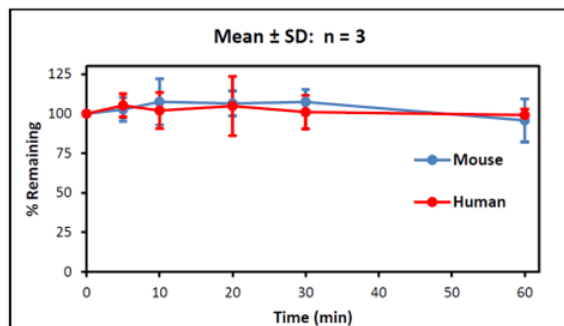
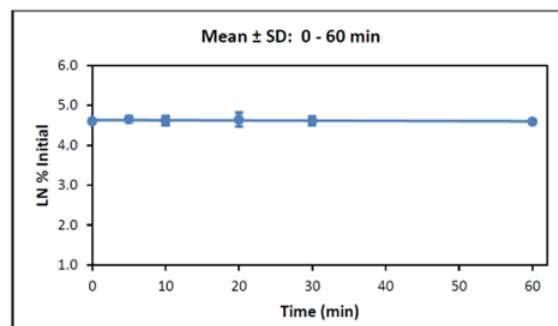
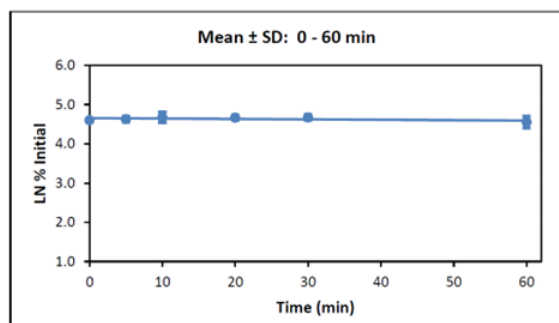
Time (min)	% Remaining			
	Mouse		Human	
	Mean	SD	Mean	SD
0	100.00	0	100.00	0
5	102.69	7.394	105.22	7.357
10	107.54	14.67	102.09	11.38
20	106.48	7.882	104.88	18.772
30	107.42	7.931	101.07	10.566
60	95.80	13.574	99.19	3.752



**Figure S1.** (A) Metabolic stability parameters for ianthelliformisamine A (**1**) based on NADPH-dependent degradation profiles in mouse liver microsomes. (B) Metabolic stability of **1** (mean  $\pm$  SD;  $n = 3$ ; 0 – 30 min) in mouse liver microsomes: LN-transformed data.



**Figure S2.** (A) Metabolic stability of ianthelliformisamine B (**2**) (mean  $\pm$  SD;  $n = 3$ ) in mouse and human liver microsomes. (B) Metabolic stability of **2** (mean  $\pm$  SD;  $n = 3$ ) in mouse liver microsomes: LN-transformed data (normalized to  $t = 5$  min). (C) Metabolic stability of **2** (mean  $\pm$  SD;  $n = 3$ ) in human liver microsomes: LN-transformed data (normalized to  $t = 5$  min).

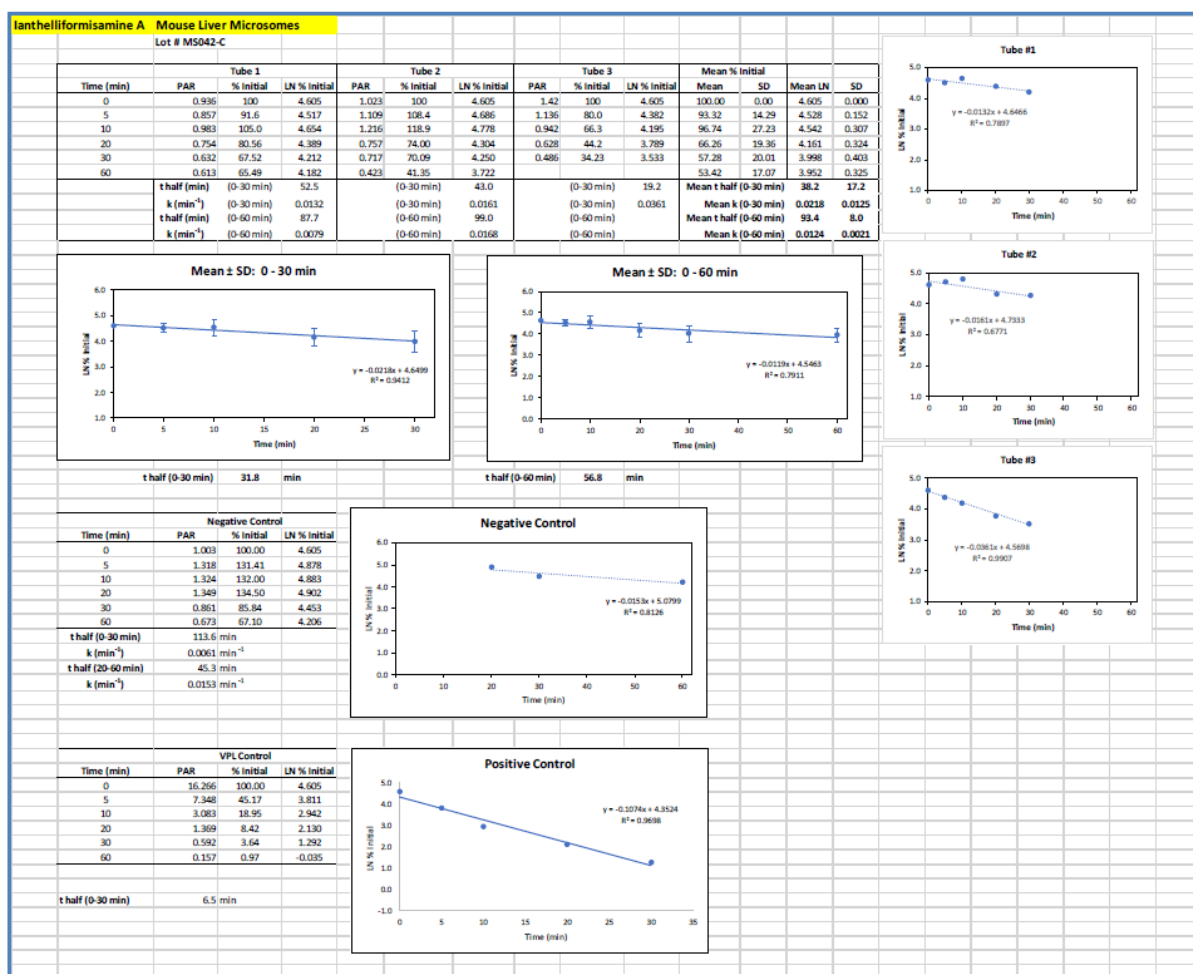
**A****B****C**

**Figure S3.** (A) Metabolic stability of ianthelliformisamine C (**3**) (mean  $\pm$  SD; n = 3) in mouse and human liver microsomes. (B) Metabolic stability of **3** (mean  $\pm$  SD; n = 3) in human liver microsomes: LN-transformed data. (C) Metabolic stability of **3** (mean  $\pm$  SD; n = 3) in mouse liver microsomes: LN-transformed data.

## Supplementary Data

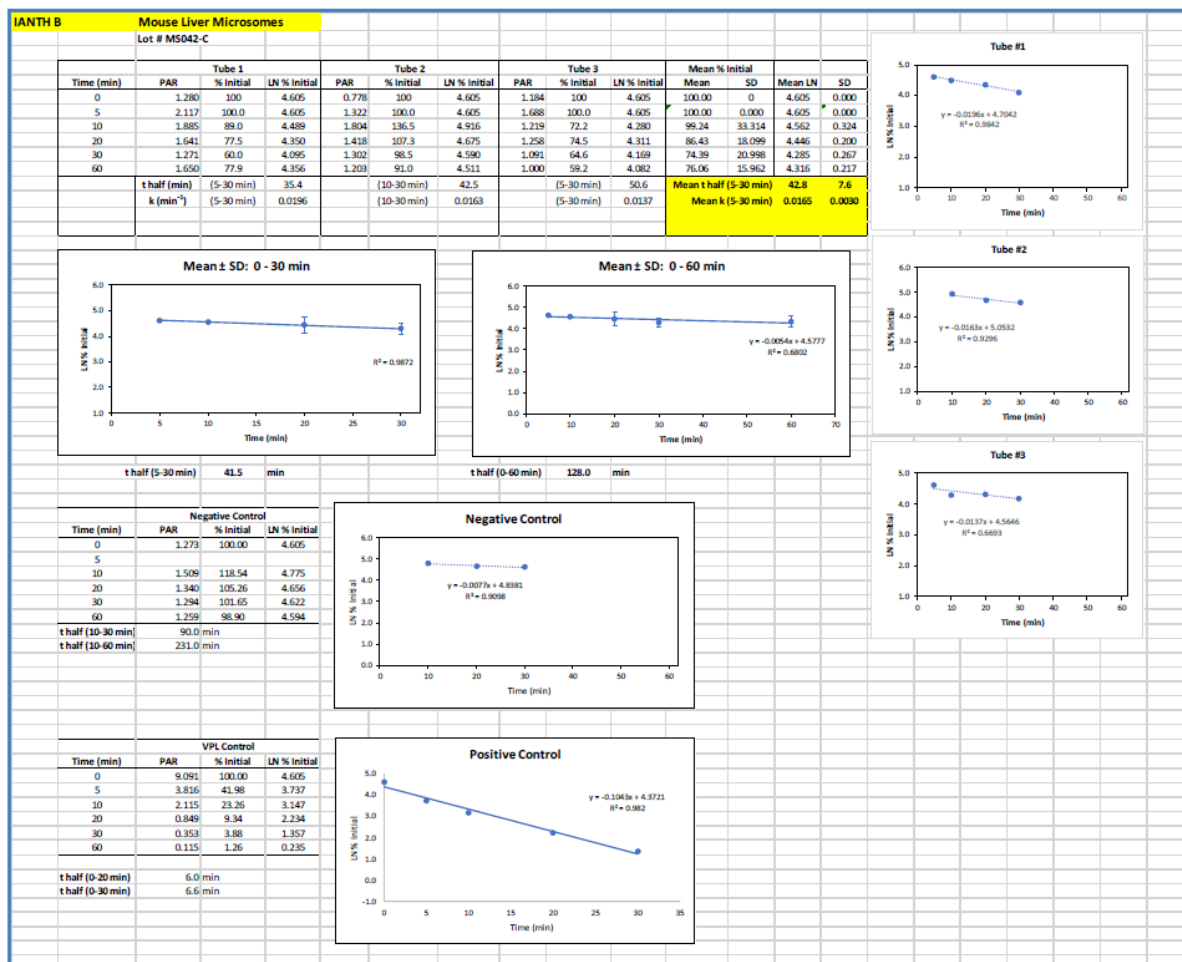
### Ianthelliformisamine A

- Mouse liver microsomes



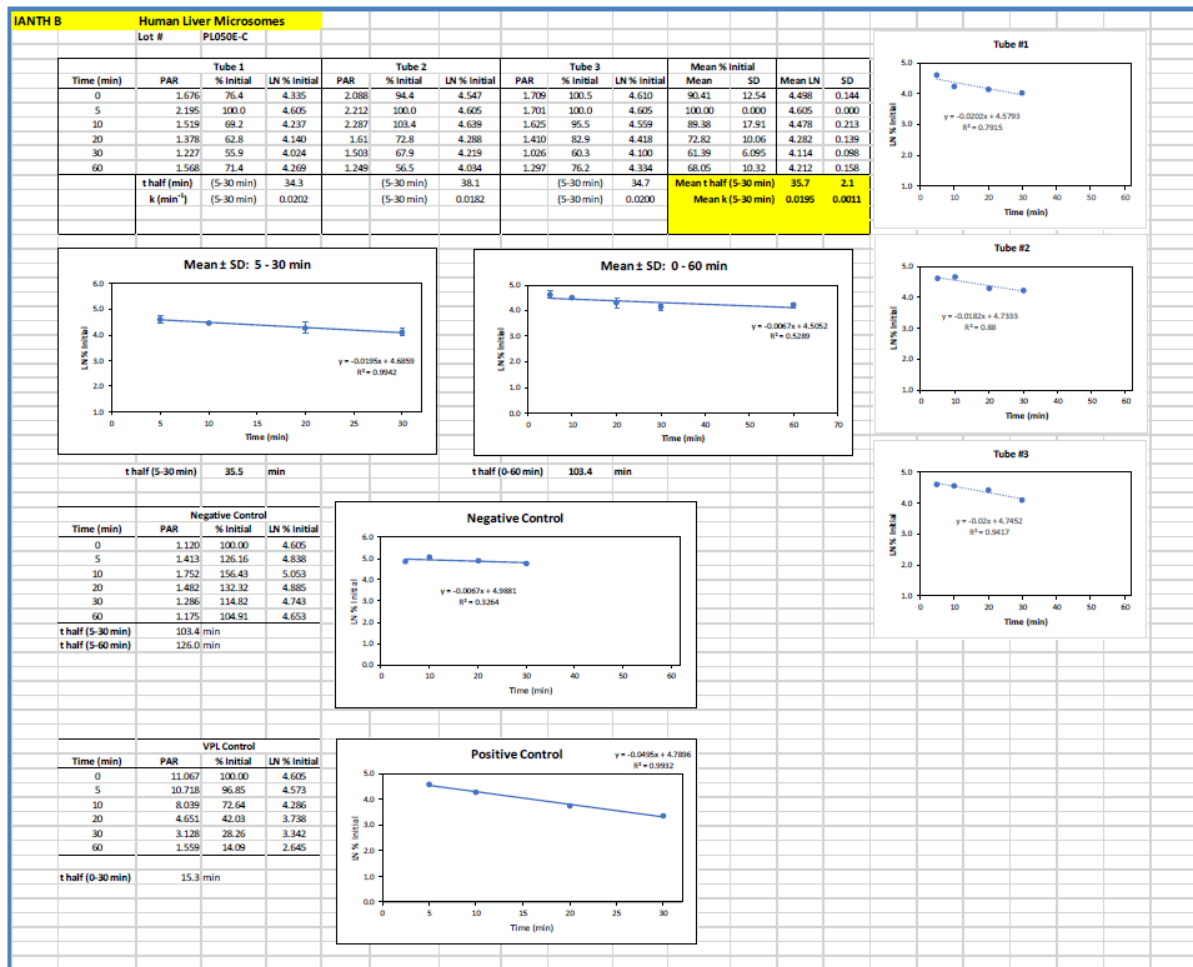
# Ianthelliformisamine B

- Mouse liver microsomes



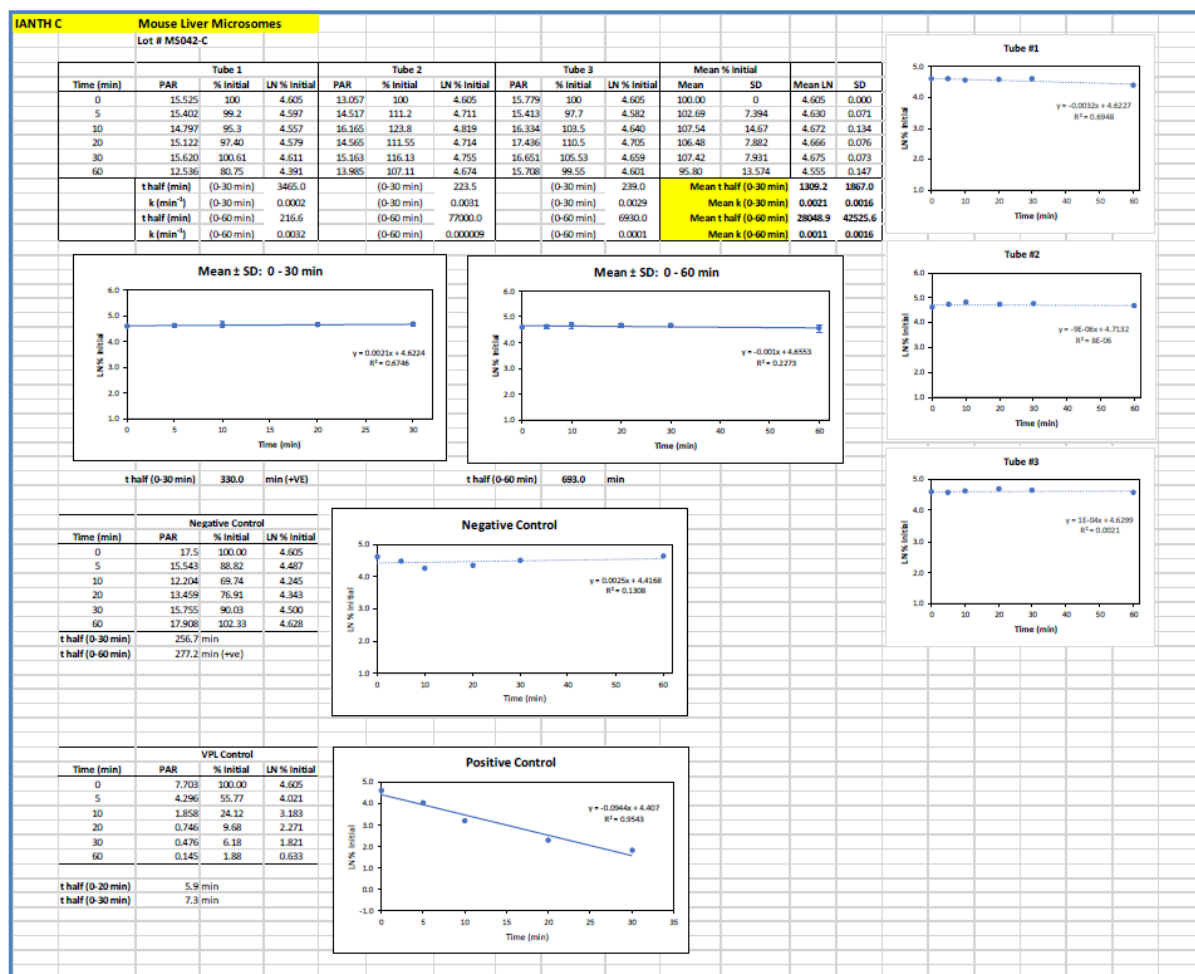


- Human liver microsomes



# Ianthelliformisamine C

- Mouse liver microsomes



- Human liver microsomes

