# APPLICATIONS



Quantitation of Human Plasma and Breast Milk Thiamine Monophosphate (TMP) and Thiamine by Applying Impact<sup>™</sup> Protein Precipitation Plate Technology with Gemini<sup>®</sup> 3µm NX-C18 HPLC Columns

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Thiamine monophosphate (TMP) and thiamine were extracted from human plasma and human breast milk by performing a rapid protein precipitation using Impact Protein Precipitation Plates followed by HPLC analysis using a Gemini 3µm NX-C18 100 x 3.0 mm HPLC column with fluorescence detection. Impact technology offers easy, fast protein removal while providing maximized recovery of the target analytes. The Gemini 3µm NX-C18 HPLC column produced excellent chromatographic resolution, sensitivity, and high peak capacities.

### Introduction

Vitamin B1, also known as thiamine, is mainly present in human body fluids and tissue as thiamine diphosphate (TDP), thiamine monophosphate (TMP) and free thiamine. TDP, the most abundant thiamine derivative, is well described as a cofactor of several important enzymes, whereas TMP and thiamine are thought to be simple intermediates for which no specific role has currently been defined<sup>1</sup>.

### **Materials and Methods**

### **Protein Precipitation**

- 1. Place the Impact plate onto a suitable 96-well sample manifold
- Dispense 100 μL of human plasma or breast milk into each well of the Impact plate
- 3. Add 300 µL of methanol to each well of the Impact plate
- 4. Mix 3 times by aspirating with a pipette tip
- 5. Apply vacuum to filter the sample and collect the purified filtrate in a collection plate

Transfer 50 µL to an autosampler vial (or allow it to remain in the collection plate). Add 50 µL of water and 50 µL of derivatizing reagent (15% sodium hydroxide solution plus 200 µL of 30 mM K<sub>3</sub>Fe(CN)<sub>6</sub>). Cover the vial or collection plate with a lid or sealing mat, respectively. Vortex for 15 seconds. Put the vial or collection plate into an autosampler. The sample is now ready to be injected onto the HPLC-FLD.

### **HPLC Conditions**

An Agilent<sup>®</sup> 1100 HPLC system (Agilent Technologies, Inc., Santa Clara, CA, USA) was used with a Shimadzu<sup>®</sup> RF-20A Prominence<sup>®</sup> Fluorescence Detector (Shimadzu, Japan) for LC/FLD analysis.

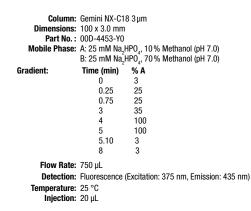
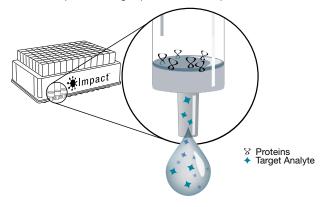


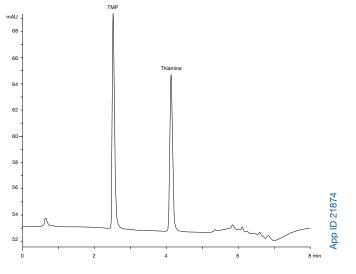
Figure 1

Protein Precipitation Using Impact Protein Precipitation Plates





200 nmol/L of TMP and thiamine in water filtered by an Impact Protein Precipitation Plate

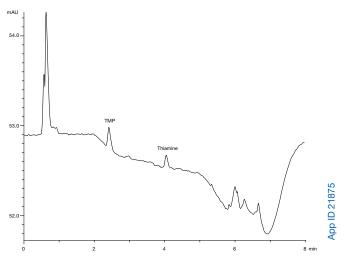




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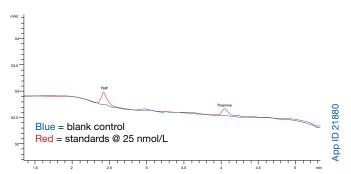
Figure 3a.

2.5~nmol/L of TMP and thiamine in water filtered by an Impact^ $^{\rm m}$  Protein Precipitation Plate



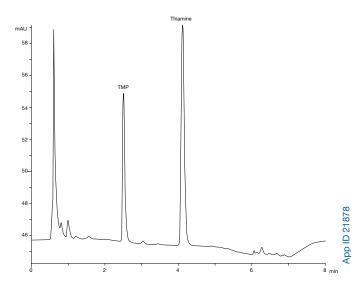
### Figure 3b.

Comparison of TMP and thiamine at 25 nmol/L vs. blank control (overlay and expanded chromatograms for human plasma filtered by an Impact Protein Precipitation Plate)

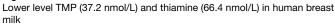


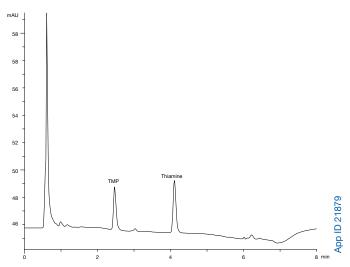
### Figure 4.

Higher level TMP (93.6 nmol/L) and thiamine (240.8 nmol/L) in human breast milk



### Figure 5.





#### Figure 6.

Standard curve of TMP filtered by an Impact Protein Precipitation Plate at a concentration range of 0 to 200 nmol/L

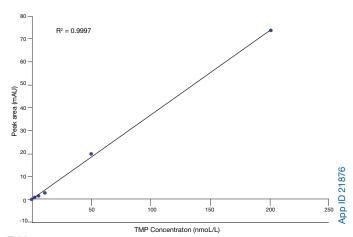


Table 1.

Standard curve (6-point) of TMP

TMP Standard Curve (6-point)	
Concentration (nmol/L)	Peak Area (mAU)
0	0
2.5	0.8
5	1.5
10	3.1
50	19.4
200	73.6

### Table 2.

Recovery of TMP and thiamine from human plasma after cleanup with an Impact Protein Precipitation Plate

Recovery for Human Plasma TMP			Recovery for Human Plasma Thiamine				
Added TMP (nmol/L	Observed (nmol/L)	Recovery (%)	Added Thiamine (nmol/L)	Observed (nmol/L)	Recovery (%)		
0	3.0		0	5.0			
5	7.0	87.5	5	10.8	108.0		
10	12.8	98.5	10	18.4	122.7		
	Mean	93.0		Mean	115.4		

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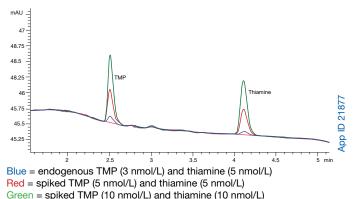
### Table 3

Accuracy studies for TMP and thiamine from human plasma after cleanup with an Impact<sup>™</sup> Protein Precipitation Plate

	TMP Sp	iked Plasma	Controls	Thiamine Spiked Plasma Controls			
Control No.	Expected (nmol/L)	Results (nmol/L)	Accuracy (%)	Expected (nmol/L)	Results (nmol/L)	Accuracy (%)	
Control 1	40	38.7	96.8	40	34.2	85.5	
Control 2	20	20.0	99.8	20	19.9	99.4	
Control 3	10	10.7	107.0	10	8.2	81.8	
		Mean	101.2		Mean	88.9	

#### Figure 7.

Overlay of endogenous and spiked TMP and thiamine in plasma filtered by Impact



### Table 4.

Retention time reproducibility studies

	Retention Time (RT)				
Inj. No.	TMP (min)	Thiamine (min)			
100	2.487	4.032			
150	2.534	4.133			
180	2.528	4.143			
240	2.491	4.122			
400	2.411	4.045			
420	2.452	4.061			
490	2.44	4.04			
Mean	2.4818	4.0822			
STEDV	0.0393	0.0483			
%CV	1.58	1.18			

### **Results and Discussion**

Traditionally, a protein precipitation step is used for fast cleanup of plasma or breast milk samples. Protein precipitation is normally performed using a centrifuge tube or a 96-well collection plate; however this process requires that supernatant be collected while being careful not to disrupt pelleted protein in the bottom of the tube or collection plate. This step was greatly simplified by using Impact Protein Precipitation Plates. The Impact plate allows for the analysis of 96 samples at once, eliminates the transfer steps that are commonly associated with protein precipitation, and can also be automated. Protein precipitation was performed within the wells of the Impact plate and sample was not allowed to pass through the filter of the plate until vacuum was applied. This ensured that the precipitated protein was left within the wells of the Impact plate while protein free sample was allowed to pass through the filter and into a collection plate (**Figure 1**).

After the protein precipitation step, the plasma and breast milk samples were derivatized and analyzed by HPLC-FLD using a Gemini<sup>®</sup>  $3\mu$ m NX-C18 HPLC column. The Gemini  $3\mu$ m NX-C18 HPLC column contains a unique silica-organic layer that is grafted onto the base silica which mechanically strengthens the particle while providing excellent efficiencies. Efficiency and resolution were necessary in this analysis because the separation of TMP and thiamine (**Figures 2**, **3a**, and **3b**) was crucial in order to accurately quantify each compound in plasma (**Figure 7**) and breast milk (**Figures 4** and **5**).

The reproducibility of our analysis was determined by producing a standard curve of TMP at a concentration range of 0 to 200 nmol/L, resulting in a correlation coefficient of  $R^2 = 0.9997$  (**Figure 6**). Thiamine was also subjected to a linearity curve at a concentration range of 0 to 200 nmol/L, resulting in a correlation coefficient of  $R^2 = 0.9993$  (not shown). Even at low levels of detection, our method proved to be reproducible for both TMP and thiamine.

Compared with a water blank, the lower level TMP and thiamine standards (2.5 nmol/L) can be clearly distinguished (**Figure 3b**), showing that our extraction and HPLC method are extremely sensitive. Endogenous levels of TMP and thiamine were also studied in plasma (**Figure 7**), which showed that 3 nmol/L and 5 nmol/L were present, respectively.

Resulting recoveries of both target compounds averaged 93% for human plasma TMP and 115% for thiamine (**Table 2**). Accuracy studies resulted in an average accuracy of 101.2 for TMP and 88.9 for thiamine (**Table 3**), suggesting that our method not only provided acceptable recoveries but was also accurate and reproducible.

Retention times (RT) were stable by comparing several retention times between injection number 100 and 490, resulting in % CV's of 1.58 % for TMP and 1.18 % for thiamine (**Table 4**).

### Conclusion

Protein precipitation has always been a popular sample preparation method however the process can be improved upon. Using Impact Protein Precipitation Plates, TMP and thiamine were cleaned up from plasma and breast milk providing benefits such as minimal method development and processing time as well as ease of use. The resulting cleanup method can also be automated, allowing laboratories to save time by increasing productivity while improving reproducibility and reducing the risk of human error. Preparation by automated protein precipitation is also a versatile sample preparation technique as the resulting extract can be analyzed by several different methods including HPLC/UV, LC/MS/MS, and HPLC-FLD. Using HPLC-FLD, our separation method on the Gemini 3µm NX-C18 HPLC column provided excellent resolution of TMP and thiamine and was sensitive enough to detect down to low levels.

### References

1. Wolfgan Stuetz, Verena Ilona Carrara, Rose McGready, Sue Jean Lee, Hans Konrad Biesalski and Francois Henry Noste. PLOS ONE, 2012.

# PLICATIONS



3 µm Microb	ore, Minibore and	MidBore <sup>™</sup> Colu	mns (mm)					SecurityGu	ard™ Cartridges (mm)	
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