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UNIVERSITY OF CALIFORNIA SAN DIEGO

Novel Fluorescent Guanine Analogs as Guanine Deaminase Substrates and Inhibitors

A thesis submitted in partial satisfaction of the requirements for the degree

Master of Science

in

Chemistry

by

You Wu

Committee in charge:

Professor Yitzhak Tor, Chair  
Professor Seth Cohen  
Professor Emmanuel Theodorakis

2018

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2018

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## ABSTRACT OF THE THESIS

Novel Fluorescent Guanine Analogs as Guanine deaminase substrates/inhibitors

by

You Wu

Master of Science in Chemistry

University of California San Diego, 2018

Professor Yitzhak Tor, Chair

Two fluorescent guanine analogues (<sup>th</sup>G, <sup>tz</sup>G) derived from thieno[3,4-d]-pyrimidine and isothiazole[4,3-d]-pyrimidine, respectively were synthesized and analyzed for their isomorphism and isofunction as the natural occurring guanine. Photophysical analysis demonstrated the promoted properties, such as visible emission, high quantum yield, sensitivity to polarity and pH, while the enzymatic conversion illustrated the highly isofunctional of <sup>tz</sup>G as guanine through its rapid reactivity with guanine deaminase (GDA) as the native substrate, which makes <sup>tz</sup>G be a good surrogate to help monitor the activity of GDA and assist the research for this significant enzyme in neuronal recovery.

## 1. Introduction

GDA is the abbreviation of guanine deaminase which can convert guanine to xanthine through a hydrolytic deamination.<sup>[1- 3]</sup> It is an important metalloenzyme due to its correlations with neuronal recovery.<sup>[4]</sup> First, the conversion from guanine to xanthine is necessary for forming the dendritic spines and consequently benefits the proper brain development.<sup>[5, 6]</sup> Furthermore, GDA performs one of the steps of purine salvage pathway, which finally produces uric acid, an important antioxidant correlated with neuronal recovery and help reduce the damage from traumatic brain injury, stroke, and spinal cord injury.<sup>[3, 7, 8]</sup> At the circuit level, overexpression of GDA plays significant roles in shaping neuronal circuits and regulating the synaptic plasticity, confers resistance to neuronal death after traumatic brain injury, which makes it be one of the most significant current treatments for neurological disease.<sup>[9, 10, 11]</sup> These relations all support the significance of GDA in neurology. In addition, it is also reported that GDA is correlated with liver function while a high level of GDA activity is observed during liver diseases and transplant rejection.<sup>[12]</sup>

These significant correlations make GDA become a potentially attractive drug target for the treatment of cognitive disorder, liver diseases and purine metabolism deficiency. As a result, it facilitates the research of GDA and also the identification of its novel activator and inhibitors. However, the effective and accurate methods used to monitor the activity of GDA has not been extensively investigated.<sup>[13]</sup>

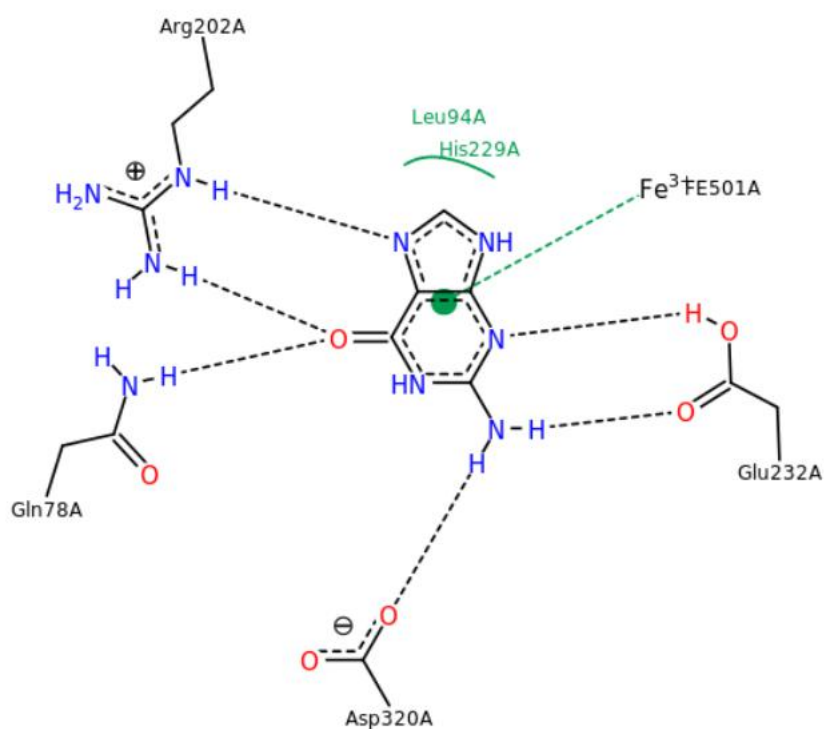
Absorbance technique is currently used for detecting the activity of GDA, but there

are a number of disadvantages for this method. First, the absorbance spectra may not be accurate enough due to the absorbance of solvent and protein, which usually absorb around 260 nm. Second, the low concentration of substrate in the enzymatic reaction results in the noise in the absorbance spectra and makes it more inaccurate. Therefore, a fluorescence technique, which is better than absorbance, was introduced here. The detection of fluorescence is much more sensitive than absorbance with less noise and it would not be influenced by solvent and non-emissive protein.

To monitor the activity of GDA by fluorescence technique, a fluorescent substrate of GDA needs to be generated, however, it's not easy to obtain good fluorescent surrogates. Generally, the structural resemblance, photophysical properties improvement and biochemical compatibility need to be considered together.<sup>[13]</sup> First, the analogues require a structural similarity to the natural counterparts, such as similar H-bonding pattern and comparable molecular size, which allows the analogues to bind to the enzyme with high specificity. Second, improved physicochemical properties than the native substrate, such as red-shifted absorbance, additional fluorescence properties and sufficient quantum yield, make the analog a more valuable alternative. Finally, the analog needs to have the same or at least similar biochemical activity as the natural substrate since the analogues are intended to be used as surrogates in the research of enzymatic reactions.

Structural compatibility plays an significant role in creating the guanine analogues.<sup>[1, 2]</sup> The size of guanine analogues should be similar as the native guanine since the overlarge substrate will result in steric clash and consequently prevent the closure of C-terminal loop, which has been reported to be necessary for the completion of the enzymatic reaction.<sup>[12]</sup>

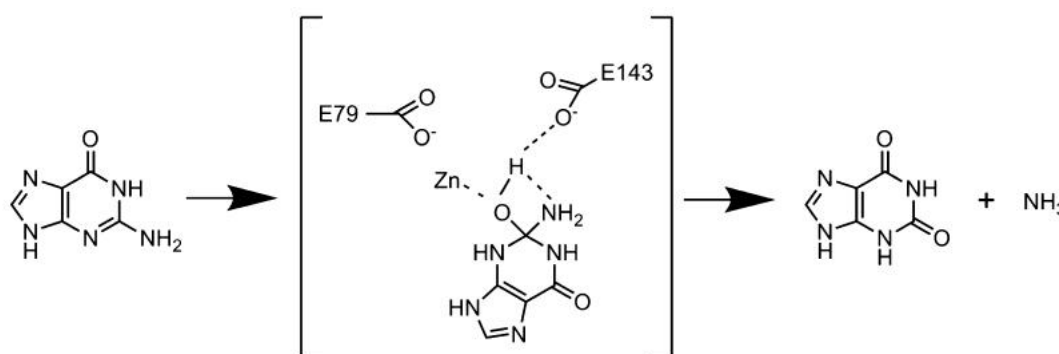
Secondly, it's reported that correct binding between substrate and GDA ensure the stabilization of the substrate at the appropriate location in the active site in preparation for the subsequent deamination process.<sup>[13, 15, 16, 17]</sup> Therefore, the guanine analogues should also have the same binding ability as native guanine with GDA (Figure 1).



**Figure 1.** Crystal Structure of Guanine Deaminase from *C. acetobutylicum* with bound guanine in the active site<sup>[17]</sup>

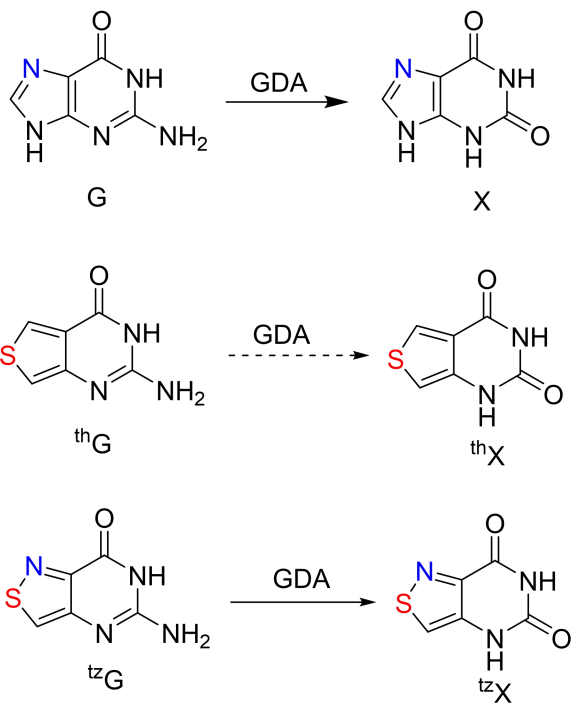
Moreover, the significance of structural similarity between guanine and the analogues can also be seen from the mechanism of the enzymatic reaction.<sup>[18]</sup> As a metalloenzyme, GDA converts the guanine to xanthine through a hydrolytic deamination while the zinc atom activates the catalytic water molecule that is coordinated to the carboxylate group and generated hydroxide nucleophile.<sup>[18]</sup> It then attacks the C2 atom adjacent to the amino group resulting in the formation of a tetrahedral intermediate.<sup>[13, 17, 18]</sup> The carboxylate groups act as

the proton shuttle assist the formation of the tetrahedral structure and the collapse of this intermediate, which promotes the completion of the deamination process (Figure 2).<sup>[13, 15]</sup> Thus, the target amino group of guanine analogues should be in close proximity to the Zn atom, activated water and the carboxylate groups, and the neighbor N should be retained to support the forming of tetrahedral structure, which all ensure the ability to react with GDA.



**Figure 2.** Reactions Catalyzed by Guanine Deaminase from *Nitrosomonas europaea* (NE0047) <sup>[13]</sup>

Based on the discussion above, two fluorescent guanine analogs, derived from thieno [3,4-d]-pyrimidine (<sup>th</sup>G) and isothiazole [4,3-d]-pyrimidine (<sup>tz</sup>G), respectively, have been synthesized (Figure 3).<sup>[19- 24]</sup> Both of these analogues have similar structure and size compared to the native guanine, which make them have the possibility to be the surrogate of native guanine and react with GDA.<sup>[25]</sup> However, the <sup>th</sup>G may not have the similar activity as guanine due to the lack of the binding between N7 and protein.<sup>[26- 28]</sup> Further photochemical tests is performed to show their improved photophysical properties than guanine. In order to understand their biochemical reactivity with GDA, both analogs and guanine were reacted with GDA under same conditions for comparison.<sup>[29]</sup>



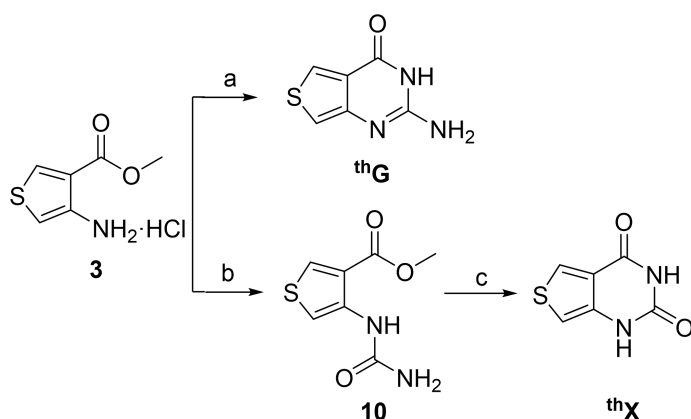
**Figure 3.** Enzymatic conversion of G, <sup>th</sup>G and <sup>tz</sup>G



## 2. Synthesis of the Nucleobases analogues.

### 2.1. Synthesis of <sup>th</sup>G and <sup>th</sup>X

The aminothiophene carboxylate **3** is the key precursor for synthesizing both thieno-Guanine (<sup>th</sup>G) and thieno-Xanthine (<sup>th</sup>X) through 1 and 2 steps, respectively. The synthesis of <sup>th</sup>G was accomplished through the cyclization with carbamimidic chloride. Synthesis of <sup>th</sup>X was started with the treatment of KOCN and furnished **10**, which was subjected to a nucleophilic acyl substitution to give final product <sup>th</sup>X (Figure 2).



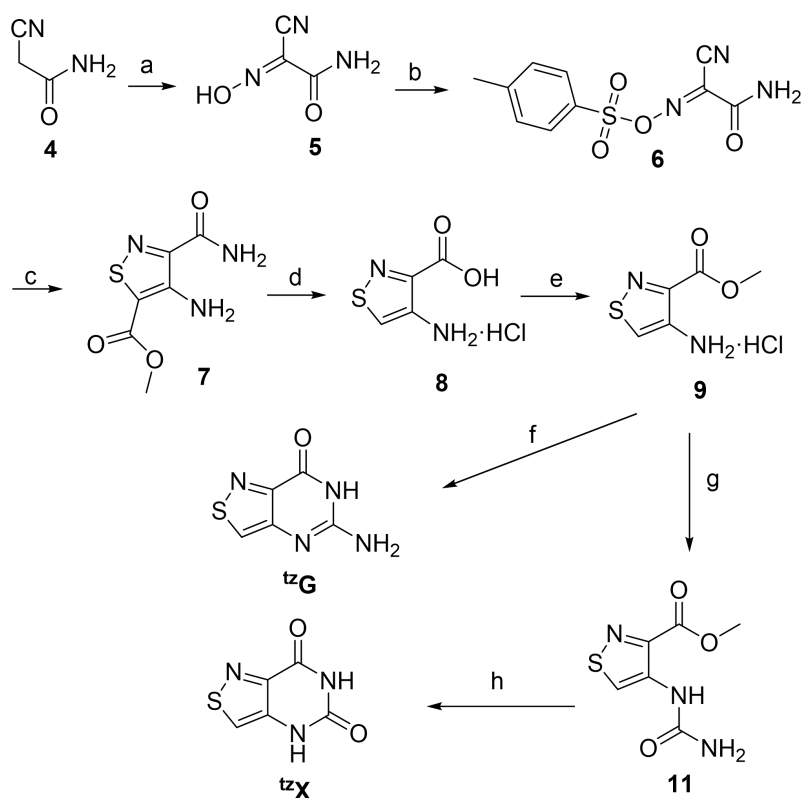
**Figure 4.** Synthesis of <sup>th</sup>G and <sup>th</sup>X. (a) DMSO<sub>2</sub>, H<sub>2</sub>O, 125°C, 1.5h, 76% (b) KOCN, H<sub>2</sub>O, Acetic Acid, rt, overnight, 82%. (c) MeOH, NaOMe, rt, 16h, 74%.

### 2.2. Synthesis of <sup>tz</sup>G and <sup>tz</sup>X

The synthesis of two analogs of Guanine (<sup>tz</sup>G) and Xanthine (<sup>tz</sup>X) was started from 2-cyanoacetamide **4** through 6 and 7 steps, respectively. As a key precursor for the synthesis of both <sup>tz</sup>G and <sup>tz</sup>X, the intermediate isothiazole-core derivative **9** which is the key precursor for the synthesis of <sup>tz</sup>G and <sup>tz</sup>X, a carbamoylisothiazole carboxylate **7** was synthesized,

starting from 2-Cyanoacetamide, which was treated with sodium nitrite to give **5**. Further reaction with tosyl chloride furnished a tosyl cyanide, which was reacted with Methyl-thioglycolate to give the isothiazole-core derivative **7**. After treatment with hydrochloride acid and a esterification reaction, **9** is given in good yield.

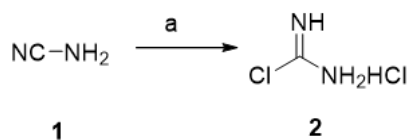
The isothiazole-core derivative **9** is a similar compound as **3**. With this key substrate in hand, the <sup>tz</sup>G and <sup>tz</sup>X can be yielded through the same process of forming the <sup>th</sup>G and <sup>th</sup>X from precursor **3** (Figure 3).



**Figure 5.** Synthesis of <sup>tz</sup>G and <sup>tz</sup>X. (a) NaNO<sub>2</sub>, H<sub>2</sub>O, Acetic Acid, 0°C, overnight, 93%. (b) TosCl, Pyridine, H<sub>2</sub>O, 0°C, 2.5h, 77%. (c) Methyl-thioglycolate, EtOH, Morpholine, 0°C, 40min, 64%. (d) HCl, 110°C, 2h, 72%. (e) MeOH, H<sub>2</sub>SO<sub>4</sub>, NaHCO<sub>3</sub>, Et<sub>2</sub>O, HCl, 80°C, overnight, 69%. (f) DMSO<sub>2</sub>, H<sub>2</sub>O, 125°C, 1.5h, 71%. (g) KOCN, H<sub>2</sub>O, Acetic Acid, rt, over night, 70%. (h) MeOH, NaOMe, rt, overnight, 94%.

### 2.3. Experimental methods

### 2.3.1. Synthesis of the nitrile precursor for the Guanine analogs preparation



**Figure 6.** Synthesis of nitrile precursor 2. (a)  $\text{Et}_2\text{O}$ ,  $\text{H}_2\text{SO}_4$ ,  $\text{NaCl}$ , rt, 2h, 69%.

#### Synthesis of carbamimidic chloride hydrochloride (2)

3g **1** is dissolved in 200 ml  $\text{Et}_2\text{O}$  in a round-bottom flask. To a second round-bottom flask,  $\text{H}_2\text{SO}_4$  is added dropwise on  $\text{NaCl}$  solid and then the generated  $\text{HCl}$  gas is directed into an empty cold trap, a water trap with  $\text{H}_2\text{SO}_4$  for drying and an empty cold trap in order. The dried  $\text{HCl}$  flow is bubbled in the  $\text{Et}_2\text{O}$  solution while stirring for 2 hours. Then the formed solid is filtered on Buchner funnel and washed with  $\text{Et}_2\text{O}$ . After drying process under vacuum overnight, white solid is given (5.6 g, 69%).

### 2.3.2. Synthesis of <sup>th</sup>G

#### Synthesis of 2-aminothieno[3,4-d]pyrimidin-4(3H)-one (<sup>th</sup>G)

A well-mixed fine solid of **3** (1.0 g, 5.2 mmol) and **2** (0.83 g, 7.3 mmol) is dissolved to  $\text{DMSO}_2$  (16.67 g) at 125 °C in a round-bottom flask over 20 minutes and is stirred for 1.5 hours for the reaction. Then the suspension is cooled to room temperature and the mixture is dissolved into 65 ml  $\text{H}_2\text{O}$ . After basified with  $\text{NH}_2\text{OH}$  to pH 10, the mixture is stirred for 1 hour. The formed creamy solid is filtered in a Buchner Funnel and washed by  $\text{H}_2\text{O}$  and  $\text{Et}_2\text{O}$  respectively. After drying under vacuum for 2 days, the yellow solid is given (0.66 g, 76%). ESI-HRMS calculated for  $\text{C}_6\text{H}_5\text{N}_3\text{OS}$   $[\text{M}+\text{H}]^+$  168.0226 found 168.0226.  $^1\text{H}$  NMR (300 MHz,  $\text{DMSO-D}_6$ )  $\delta$  10.57 (s, 1H), 8.23 (d,  $J = 3.3$  Hz, 1H), 6.96 (d,  $J = 3.3$  Hz, 1H), 6.11 (s, 2H).  $^{13}\text{C}$  NMR (126 MHz,  $\text{DMSO}$ )  $\delta$  159.78 (s), 151.61 (s), 149.78 (s), 127.67 (s), 124.26 (s),

108.84 (s).

### **2.3.3. Synthesis of <sup>13</sup>C**

#### **Synthesis of 2-amino-N-hydroxy-2-oxoacetimidoyl cyanide (5)**

To a 250 ml round bottom flask, **4** (10 g, 0.12 mol) and sodium nitrite (10 g, 0.15 mol) is dissolved in 100 ml H<sub>2</sub>O at 0°C. Acetic Acid (14 ml) is added drop by drop in the mixture for 30 minutes. After stirring overnight, the product can be extracted with Ethyl Acetate and evaporate to get the white solid (13 g, 93%).

#### **Synthesis of 2-amino-2-oxo-N-(tosyloxy)acetimidoyl cyanide (6)**

To a 100 ml round bottom flask, **5** (2.5 g, 0.022 mol) is dissolved in 18 ml Pyridine (anhydrous, 99.8%) at 0 °C, and 4-methyl benzene sulfonyl chloride is added in 4 portions every 15 minutes (1.1 g x 4, 0.023 mol). 1 hour after starting adding 4-methyl benzene sulfonyl chloride, 10 ml pyridine is added into the mixture and the solution is stirred for 1 hour at 0 °C. After adding 40 ml H<sub>2</sub>O / ice mixture, the solution is stirred for another 30 minutes and white solid is filtered in Buchner funnel and dried in vacuum overnight (4.6 g, 77%).

#### **Synthesis of methyl 4-amino-3-carbamoylisothiazole-5-carboxylate (7)**

To a 100 ml round bottom flask, **6** (2.49 g, 9.3 mmol) is dissolved in 10 ml EtOH at 0 °C and 1 ml Methyl-thioglycolate is added after preparing a well-mixed solution with 1.4 ml morpholine and 1 ml EtOH, then the mixture is added dropwise over 10 minutes into the

flask. The solution turns from white to yellow during the adding of the morpholine / EtOH mixture and the reaction is stirring for another 30 minutes at 0 °C. After consumption of **6** as monitored by TLC, 40 ml of iced water is added and the suspension is filtered to get a white solid. After drying under vacuum overnight, the product is given (1.2 g, 64%).

### **Synthesis of 4-aminoisothiazole-3-carboxylic acid hydrochloride (8)**

To a 100 ml round bottom flask, **7** (2.5 g, 12 mmol) is dissolved in HCl (32.8 ml) and the solution is boiled at 110 °C for 2 h. After cooling down in ice bath, pink solid can be filtered (1.6 g, 72%).

### **Synthesis of methyl 4-aminoisothiazole-3-carboxylate hydrochloride (9)**

To a 100 ml round bottom flask, **8** (1.6 g, 8.9 mmol) is dissolved in MeOH (31.7 ml) and H<sub>2</sub>SO<sub>4</sub> (1.5 ml) and the solution is refluxed at 80 °C overnight. After cooling down to room temperature, NaHCO<sub>3</sub> is used to neutralize the solution to pH 4. Following with the extraction with Et<sub>2</sub>O, and NaSO<sub>4</sub> is added to remove extra H<sub>2</sub>O. Solvent removed by evaporation, then HCl in dioxane (4 M, 10 ml) is added to the solution and the suspension was evaporated to dryness to give **9** as yellow-brown hydrochloric salt solid (1.2 g, 69%).

### **Synthesis of 5-aminoisothiazolo[4,3-d]pyrimidin-7(6H)-one (<sup>tz</sup>G)**

A well-mixed fine solid of **9** (1.5 g, 7.7 mmol) and **2** (1.32 g, 11.6 mmol) is added slowly and dissolved in melted DMSO<sub>2</sub> (25.5 g) at 125 °C in a round-bottom flask over 20 minutes and reacting at 125 °C for 1.5 hours. Then the suspension is cooled to room

temperature and the mixture is dissolved into 120 ml H<sub>2</sub>O. After basified with NH<sub>2</sub>OH to pH 10, the mixture is followly stirred for 1 hour. The formed creamy solid is filtered in a Buchner funnel and washed by H<sub>2</sub>O and Et<sub>2</sub>O respectively. After drying under vacuum for 2 days, the yellow solid is given (0.92 g, 71%). ESI-HRMS calculated for C<sub>5</sub>H<sub>4</sub>N<sub>4</sub>OS [M+H]<sup>+</sup> 169.0179 found 169.0177. <sup>1</sup>H NMR (300 MHz, DMSO-D<sub>6</sub>) δ 11.06 (s, 1H), 8.62 (s, 1H), 6.32 (s, 2H). <sup>13</sup>C NMR (126 MHz, dmsO) δ 156.70 (s), 151.71 (s), 150.79 (s), 149.01 (s), 136.72 (s).

#### 2.3.4. Synthesis of <sup>th</sup>X

<sup>th</sup>X is synthesized by our lab member, Andrea Fin.

##### Synthesis of methyl 4-ureidothiophene-3-carboxylate (**10**)

**3** (2 g, 10.4 mmol) is dissolved in H<sub>2</sub>O and Acetic Acid (25 ml, 1:1 ratio) in a 100 ml round bottom flask, then KOCN (1.67 g, 20.6 mmol) is dissolved in H<sub>2</sub>O and added dropwise to the solution at room temperature. The reaction is stirred overnight, then the mixture is filtered using the Buchner funnel and washed with H<sub>2</sub>O. The White powder is given (1.708 g, 82%).

##### Synthesis of thieno[3,4-d]pyrimidine-2,4(1H,3H)-dione (<sup>th</sup>X)

To a round bottom flask, **10** (1 g, 5 mmol) is suspended in MeOH (10 ml), and then NaOMe (22 ml) is added slowly to the mixture. After 16 hours stirring, the milky solution is filtered and white solid is washed with MeOH (70 ml) and dried in vacuum. For more purification, the white solid is dissolved in MeOH and H<sub>2</sub>O (1:1 ratio), and acidified with

formic acid. After stirring at room temperature for 1 hour, the white precipitate is given (0.621 g, 74%). ESI-HRMS calculated for C<sub>6</sub>H<sub>4</sub>N<sub>2</sub>O<sub>2</sub>S [M+H]<sup>+</sup> 166.9921 found 166.9922. <sup>1</sup>H NMR (500 MHz, DMSO-D<sub>6</sub>) δ 10.91 (s, 2H), 8.31 (d, J = 3.2 Hz, 1H), 6.78 (d, J = 3.2 Hz, 1H). <sup>13</sup>C NMR (126 MHz, dmsO) δ 159.07 (s), 151.25 (s), 138.20 (s), 130.63 (s), 122.35 (s), 103.17 (s).

### 2.3.5. Synthesis of <sup>tz</sup>X

<sup>tz</sup>X is synthesized by our lab member, Andrea Fin.

#### Synthesis of methyl 4-ureidoisothiazole-3-carboxylate (**11**)

**9** (1.0 g, 5.2 mmol) is dissolved in H<sub>2</sub>O and Acetic Acid (20 ml, 1:1 ratio) in a 100 ml round bottom flask, then KOCN (2.0 g, 24.7 mmol) is dissolved in H<sub>2</sub>O and added dropwise to the solution at room temperature. The reaction is stirred overnight, then the mixture is filtered using the Buchner funnel and washed with H<sub>2</sub>O (70 ml). The White powder is given (0.73 g, 70%).

#### Synthesis of isothiazolo[4,3-d]pyrimidine-5,7(4H,6H)-dione (<sup>tz</sup>X)

To a round bottom flask, **11** (0.710 g, 3.53 mmol) is suspended in MeOH (10 ml), then NaOMe (15 ml) is added dropwise to the mixture and a dense and yellowish solution is observed. After stirring over night at room temperature, the milky solution is filtered and the yellow solid is washed with MeOH (60 ml) and dried in vacuum. For more purification, the yellow solid is dissolved in MeOH / H<sub>2</sub>O (1:1 ratio), and acidified with formic acid to get white precipitate. Solution is stirred for 1 hour at room temperature, then filtered to get white

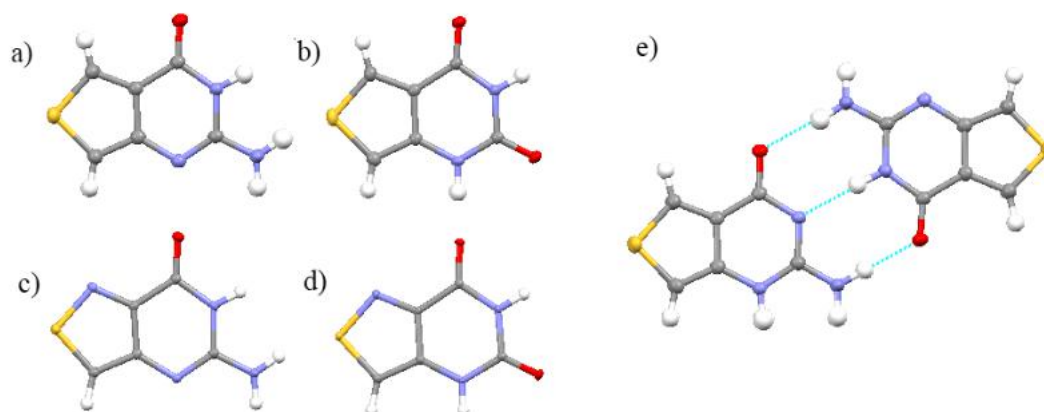
solid (0.56 g, 94%). ESI-HRMS calculated for  $C_5H_3N_3O_2S$   $[M+H]^+$  167.9873 found 167.9874.

$^1H$  NMR (500 MHz, DMSO- $D_6$ )  $\delta$  11.60 (s, 1H), 11.37 (s, 1H), 8.62 (s, 1H).  $^{13}C$  NMR (126 MHz, dmsO)  $\delta$  157.02 (s), 150.81 (s), 146.94 (s), 138.50 (s), 130.74 (s).



## 2. Crystal Structure

Crystal structure determination confirmed the proposed structure (Figure 2) and the structural similarity between two analogues and native guanine. The existence of two structures of <sup>th</sup>G proves the tautomerism in the solid state of <sup>th</sup>G. The H-bonding between the two structures connects them through Watson-Crick face, which is the same face as the connection between guanine and cytosine, further confirmed the structural analogy between the analogue and guanine. [30, 31]



**Figure 7.** X-ray crystal structures of (a) <sup>th</sup>G, (b) <sup>th</sup>X, (c) <sup>tz</sup>G, (d) <sup>tz</sup>X. (e) Tautomerization of <sup>th</sup>G.

### 3. Photophysics of nucleobase analogues.

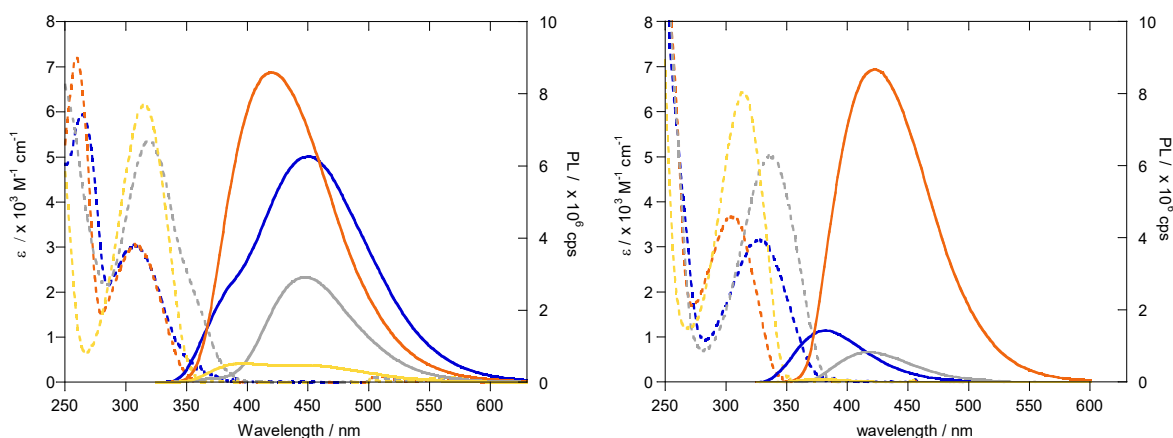
#### 3.1. Photophysics analysis

The fundamental spectroscopic properties, including the molar absorption coefficient, quantum yield, sensitivity toward polarity, and the spectroscopically derived  $pK_a$  values of the nucleobases analogues are listed in Table 1. The red-shifted maxima of the ground-state absorption spectra is showed for the four analogues comparing to the native nucleobases, G (275 nm) and X (273 nm) (Graph 1a). The molar absorption coefficient of <sup>th</sup>G, <sup>th</sup>X, <sup>tz</sup>G, and <sup>tz</sup>X are 3.31, 3.03, 5.09, 6.13, respectively (Table 1). All Emission spectrum were excited at their maximum absorption and the maxima is observed to be ranged from 394 nm (<sup>tz</sup>X) to 446 nm (<sup>tz</sup>G). The emission quantum yield of the two thieno-ring analogues are 0.36 and 0.46 for <sup>th</sup>G and <sup>th</sup>X, respectively, which were higher than the two isothiazole-ring analogues, with 0.07 for <sup>tz</sup>G and 0.02 for <sup>tz</sup>X in water.

**Table 1.** Photophysical Properties of Guanine and Xanthine Analogues.

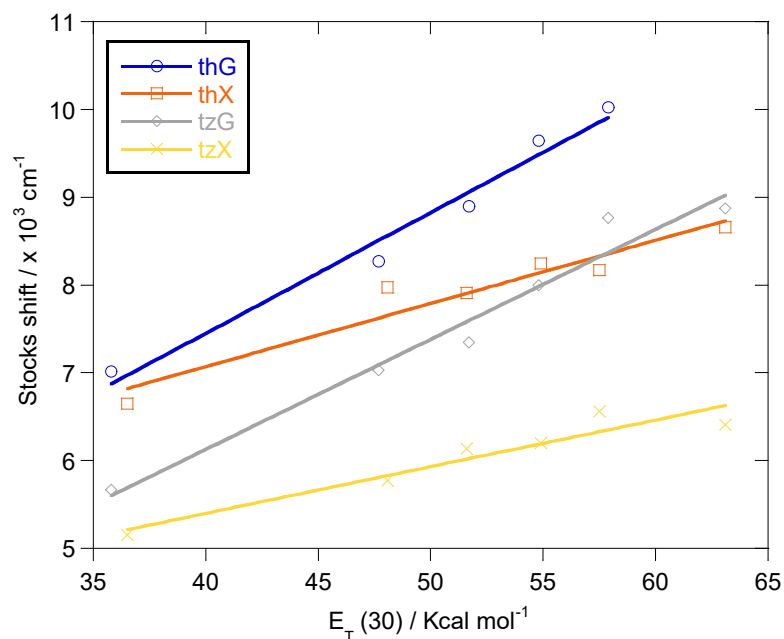
	solvent	$\lambda_{\text{abs}}(\epsilon)^a$	$\lambda_{\text{emi}}(\Phi)^a$	$\Phi\epsilon$	Stokes shift <sup>a</sup>	Polarity sensitivity <sup>b</sup>	pK <sub>a</sub> <sup>c</sup> (abs)
<sup>th</sup> G	water	315 (3.04 ± 0.03)	439 (0.40 ± 0.04)	1179	8.97	137.4	4.41, 10.19
	Dioxane	325 (2.96 ± 0.02)	423 (0.40 ± 0.05)	1182	7.18		
<sup>th</sup> X	water	308 (3.16 ± 0.08)	420 (0.46 ± 0.05)	1406	8.66	71.9	3.28, 8.96
	Dioxane	303 (3.63 ± 0.03)	383 (0.065 ± 0.002)	242	6.95		
<sup>tz</sup> G	water	320 (5.43 ± 0.03)	446 (0.068 ± 0.004)	351	8.88	125.3	8.80
	Dioxane	336 (5.07 ± 0.03)	419 (0.026 ± 0.001)	129	5.90		
<sup>tz</sup> X	water	315 (6.13 ± 0.05)	394 (0.018 ± 0.001)	116	6.42	53.1	9.92
	Dioxane	315 (6.427 ± 0.007)	375 (0.001 ± 0.001)	11	5.08		

a.  $\lambda_{\text{abs}}$ ,  $\epsilon$ ,  $\lambda_{\text{emi}}$ , and Stokes shift are reported in nm,  $10^3\text{M}^{-1}\text{cm}^{-1}$ , nm, and  $10^3\text{cm}^{-1}$  respectively. b. Sensitivity to solvent polarity reported in  $\text{cm}^{-1}/(\text{kcal mol}^{-1})$  is equal to the slope of the linear fit in Graph 2. c. pK<sub>a</sub> values are equal to the inflection point determined by the fitting curves in Graph 3.

**Graph 1.** Emission and Absorbance spectrum for <sup>th</sup>G (blue), <sup>th</sup>X (orange), <sup>tz</sup>G (gray), <sup>tz</sup>X (yellow) in water (left) and dioxane (right).

The absorption spectra measured in dioxane showed bathochromic shifts for <sup>th</sup>G and <sup>tz</sup>G, slight hypochromic shift for <sup>th</sup>X and no significant change for <sup>tz</sup>X compared to the aqueous solutions (Graph 1b). The emission intensity of <sup>th</sup>X, <sup>tz</sup>G, <sup>tz</sup>X is obviously lower in

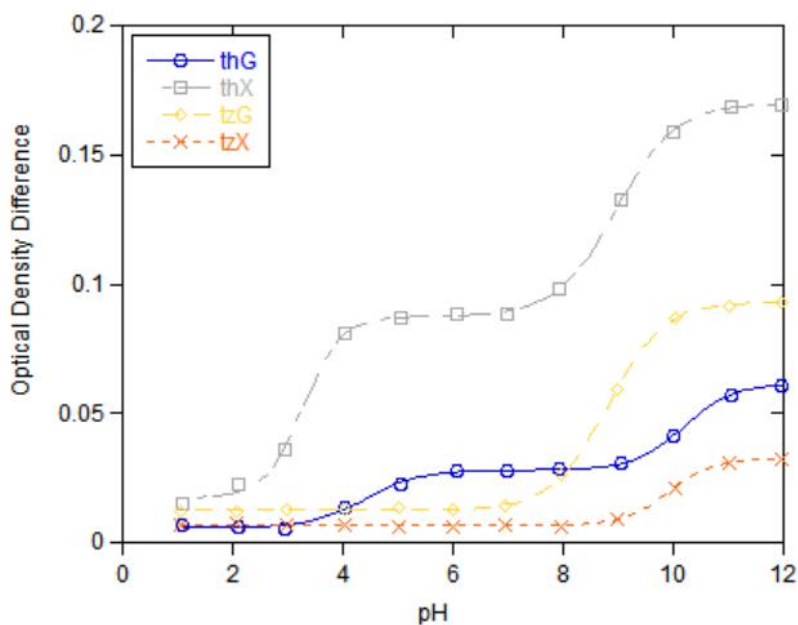
dioxane than in water, except <sup>th</sup>G, for which is significantly higher than in water. The fluorescence maxima displayed a remarkable hypochromic shift for all four nucleobases analogues. Quantum yield in Dioxane for <sup>th</sup>G (0.35) is higher than <sup>tz</sup>G (0.03), and are still closed to their corresponding values in water, while the quantum yield for <sup>th</sup>X is significantly lower than in water and no remarkable change for <sup>tz</sup>X. The sensitivity of the four nucleobase analogues to solvent polarity is reflected by the slope of the linear fitting between the measured Stokes shifts and microscopic solvent polarity parameters,  $E_T(30)$  (Graph 2, Table 1).



**Graph 2.** Stokes shift correlation versus solvent polarity ( $E_T(30)$ ) of water/dioxane mixtures for <sup>th</sup>G (blue), <sup>th</sup>X (orange), <sup>tz</sup>G (gray) and <sup>tz</sup>X (yellow).

All nucleobase analogues display sensitivity toward pH change, as a result, the  $pK_a$  values can be extracted from the dependence of the absorbance on pH (Graph 4). To exclude the influence of concentration, the graph of absorbance differences between two specific wavelengths versus pH is generated for the G analogues and X analogues (Graph 3, Table 1).

The significant change of absorbance characterized the pKa of these four nucleobase analogues, yielding two pKa values for <sup>th</sup>G (4.41 and 10.19) and <sup>tz</sup>X (3.28, 8.96) and one for <sup>tz</sup>G (8.80) and <sup>tz</sup>X (9.92).



**Graph 3.** Optical density difference versus pH for <sup>th</sup>G, <sup>th</sup>X, <sup>tz</sup>G, and <sup>tz</sup>X.

### 3.2. Experimental methods

Spectroscopic grade DMSO and dioxane were obtained from Alfa Aesar and Acros and aqueous solutions were prepared with de-ionized water. All the measurements were carried out in a 1 cm four-sided quartz cuvette from Helma.

Absorption spectra were measured on a Shimadzu UV-2450 spectrophotometer setting the slit at 1 nm and using a resolution of 0.5 nm. All the spectra were corrected for the blank. Steady state emission spectra were measured on a Horiba Fluoromax-4 equipped with a cuvette holder with a stirring system setting both the excitation and the emission slits at 3 nm, the resolution at 1 nm. The steady state fluorescence spectra were performed upon excitation

at the wavelength of the maximum absorption.

All the spectra were corrected for the blank. Both instruments were equipped with a thermostat controlled ethylene glycol-water bath fitted to specially designed cuvette holder and the temperature was kept at  $37.00 \pm 0.10$  °C.

Nucleobases were dissolved in DMSO to prepare highly concentrated stock solutions: X (13.15 mM), <sup>th</sup>G (11.96 mM), <sup>th</sup>X (11.89 mM), <sup>tz</sup>G (11.89 mM) and <sup>tz</sup>X (11.82 mM). G has relatively low solubility in DMSO, then G is dissolved in basic NaOH aqueous solution (pH 12) and the concentration is 11.23 mM. In a typical experiment, aliquots (10 µl) of the concentrated DMSO solution were diluted with air-saturated solvents (3 ml). The solution were mixed with a pipette for 15 seconds and placed in the cuvette holder at  $37.00 \pm 0.10$  °C for 3 minutes before spectra were recorded. All sample except Guanine contain 0.3 v/v % of DMSO.

### 3.2.1. Fluorescence quantum yield

The samples concentration were adjusted to have an optical density lower than 0.1 at the excitation wavelength ( $\lambda_{ex}$ ). The fluorescence quantum yield ( $\Phi$ ) were evaluated based on an external standard, 2-aminopurine (0.68 in water,  $\lambda_{ex}$  320 nm) by using the following equation.

$$\Phi = \Phi_{STD} \frac{I}{I_{STD}} \frac{OD_{STD}}{OD} \frac{n^2}{n_{STD}^2}$$

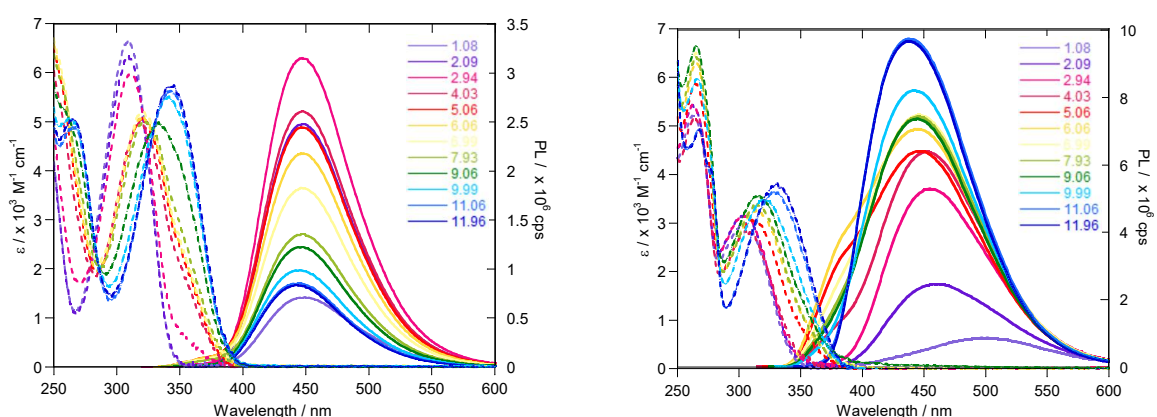
Where  $\Phi_{STD}$  is the fluorescence quantum yield of the standard, I and  $I_{STD}$  are the

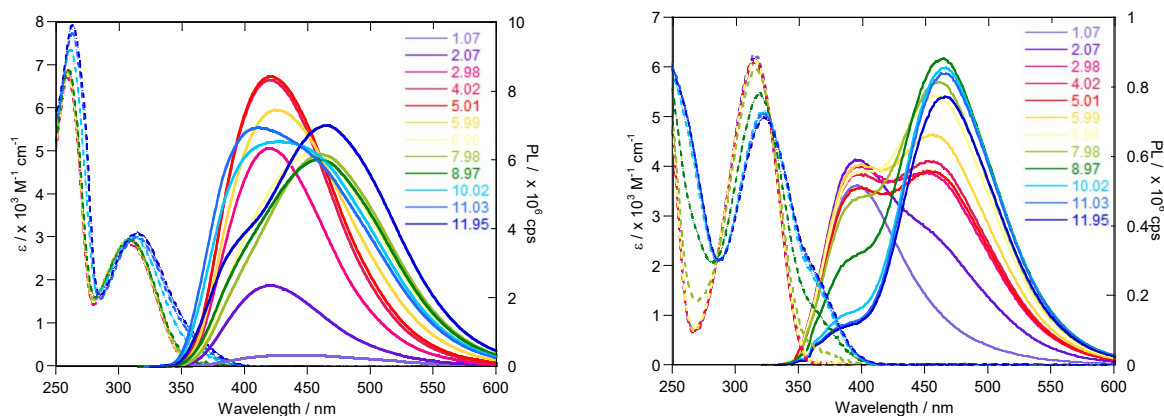
integrated area of the emission band of the sample and the standard respectively, OD and  $OD_{STD}$  are the optical density at the excitation wavelength for the sample and the standard respectively and  $n$  and  $n_{STD}$  are the solvent refractive index of the sample and the standard solutions respectively.

### 3.2.2. Sensitivity to pH

Sodium phosphate buffers (30 ml) were prepared by mixing sodium phosphate dibasic and sodium phosphate monobasic to have a final concentration of 50 mM. The pH of each solution was adjusted to the desire value by adding aliquots of 2 M aqueous HCl or 1 M aqueous NaOH prior to spectral measurements.

The absorption ( $\lambda_{abs}$ ) difference between two selected wavelength were plotted versus the pH and fitted using a Boltzmann sigmoidal curve using Kaleidagraph 3.5 (Graph 3). The  $pK_a$  values were determined by interpolation of the fitting curves. The values and the relative standard deviations are reported in table 1.





**Graph 4.** Absorption (dashed lines) and emission (solid lines) traces in buffer solutions at different pH for <sup>th</sup>G (upper left), <sup>th</sup>X (upper right), <sup>tz</sup>G (bottom left), and <sup>tz</sup>X (bottom right).

### 3.2.3. Sensitivity to polarity

Experiments evaluating the effect of polarity were performed in water, dioxane and their mixtures (20, 40, 60 and 80 v/v % water in dioxane) (Graph 5). The sample  $E_T(30)$  values were determined by dissolving a small amount of Reichardt's dye in the mixture of the same solvent used to dilute the nucleobase's DMSO sample. The observed long wavelength absorption maximum ( $\lambda_{abs}^{max}$ ) was converted to the  $E_T(30)$  values with the following equation (Table 2).

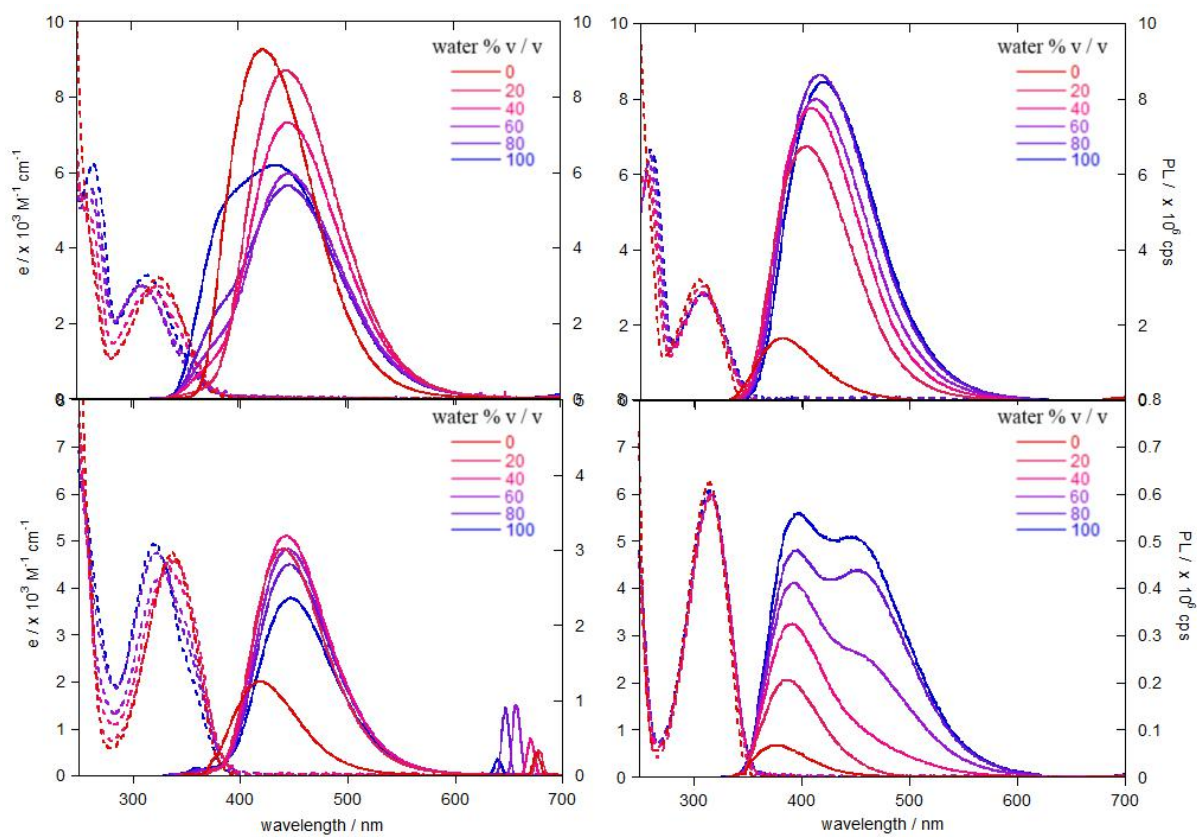
$$E_T(30) = \frac{28591}{\lambda_{abs}^{max}}$$

**Table 2.**  $E_T(30)$  experimental values for water/dioxane mixtures. Due to solubility limitation the water  $E_T(30)$  value was not calculated.

Water:Dioxane	Reported $E_T(30)$ (Kcalmol <sup>-1</sup> )	Experimental $E_T(30)$ (Kcalmol <sup>-1</sup> ) for <sup>th</sup> G and <sup>tz</sup> G	Experimental $E_T(30)$ (Kcalmol <sup>-1</sup> ) for <sup>th</sup> X and <sup>tz</sup> X
1:0	63.1	-	-



4:1	57.5	57.9	57.5
3:2	55.0	54.8	54.9
2:3	51.6	51.7	51.6
1:4	48.3	47.7	48.1
0:1	36.4	35.8	36.5



**Graph 5.** Absorption (dashed lines) and emission (solid lines) traces in water, dioxane and mixture thereof for  ${}^{\text{th}}\text{G}$  (upper left),  ${}^{\text{th}}\text{X}$  (upper right),  ${}^{\text{tz}}\text{G}$  (bottom left), and  ${}^{\text{tz}}\text{X}$  (bottom right).

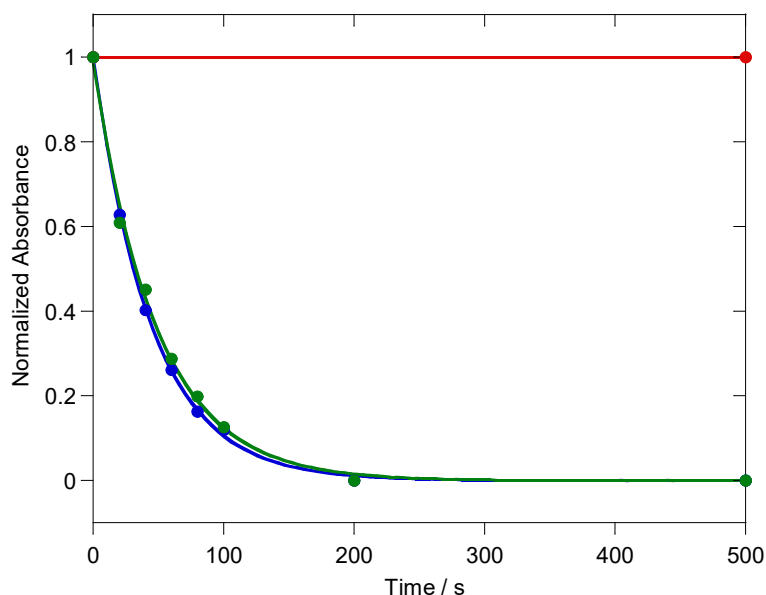
## 4. Enzymatic conversion from G analogue to X analogues.

### 4.1. Enzymatic analysis

To explore the different biochemical activity of the nucleobase analogues, guanine deaminase (GDA) was chosen to react with these potential substrates. The result can help demonstrate the utility of these fluorescent analogues and also the impact of restoring the basic N7 in the enzymatic reaction. It has been previously reported that the enzymatic conversion of a thieno-core Adenosine analogue (<sup>th</sup>A) to the corresponding inosine derivative (<sup>th</sup>I) is significantly lower than the naturally occurring adenosine.<sup>[27, 28]</sup> However, the deamination of isothiazole-core adenosine (<sup>tz</sup>A) to inosine (<sup>tz</sup>I) is at the similar rate as of A to I.<sup>[29]</sup> The H-bonding between N7 and the residue of the protein in the crystal structure of GDA further demonstrate the significance of the N7 in the reaction.<sup>[15- 17]</sup> Therefore, we hypothesized that the Guanine analogues have the similar property, which is that the <sup>tz</sup>G may not have higher activity than the <sup>th</sup>G because of the the restoring of N7.

Firstly, the reaction between GDA and <sup>th</sup>G was tested. Both the steady state absorption and emission didn't change much during the time after adding GDA to the <sup>th</sup>G, which showed that the GDA could barely react with <sup>th</sup>G, while the spectras taken for <sup>tz</sup>G showed a obvious and fast conversion from <sup>tz</sup>G to <sup>tz</sup>X (Graph 10, 11). Real-time continuous absorbance and emission were also performed relying on the photophysical differences between the reagents and expected products (Graph 12, 13, 14). The result confirmed that the GDA cannot deaminate <sup>th</sup>G and showed that the deamination of <sup>tz</sup>G to <sup>tz</sup>X is almost as efficient as G to X.

Assuming a first order reaction, the reaction half-life for  $^{12}\text{G}$  and G deamination were also calculated for comparison ( $t_{1/2} = 33.6 \pm 3.6$  s for  $^{12}\text{G}$  and  $t_{1/2} = 33.7 \pm 3.6$  s for G), which were very closed to each other. This result substantiated my hypothesis, which was further confirmed by HPLC analyses (Graph 6). In summary, the improved functionality of the  $^{12}\text{G}$  over the  $^{13}\text{G}$  was highlighted from these observations, which supported the hypothesis of the significance of the N7 in the enzymatic reaction.



**Graph 6.** Normalized absorbance intensity from HPLC versus reaction time for G (blue),  $^{13}\text{G}$  (red) and  $^{12}\text{G}$  (green).

Since  $^{13}\text{G}$  cannot react with GDA, a hypothesis of being an inhibitor for the enzymatic conversion catalyzed by GDA is further presented. We selected to probe additional HPLC analyses for exploring the activity of G with the well-mixed GDA and  $^{13}\text{G}$ . The unaffected conversion of G to X illustrated  $^{13}\text{G}$  is neither substrate nor an inhibitor of GDA.

#### 4.2. Experimental Methods.

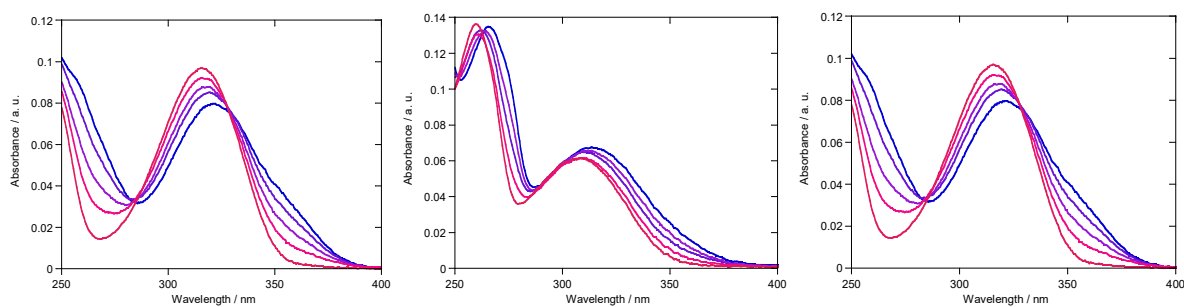
Recombinant human guanine deaminase protein (GDA) was obtained from Novus Biologicals (catalog number NBP2-49692-20ug). Sodium phosphate buffer was prepared to dilute the enzyme solution 10:1 by dissolving an aliquot (2  $\mu$ l) in it (18  $\mu$ l, 50 mM, pH 7.5). The enzyme solution was freshly prepared and kept in ice bath prior to use.

Concentrated stock solutions in DMSO were prepared for X (6.66 mM), <sup>th</sup>G (6.09 mM), <sup>th</sup>X (6.69 mM), <sup>tz</sup>G (4.79 mM), <sup>tz</sup>X (6.54 mM) and in H<sub>2</sub>O (pH 12) for G (9.85 mM). These concentrated stock solution were used for steady state absorbance and emission for clearer graphs and was diluted 20:1 for usage of kinetic measurement and HPLC traces to see the first order reaction process.

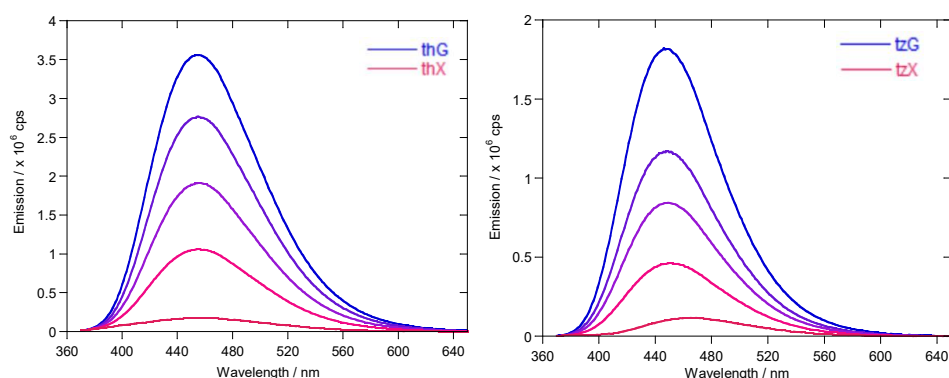
#### **4.2.1 Steady state absorption and emission experiment in presence of GDA**

Steady state absorption spectra over time were measured on a Shimadzu UV-2450 spectrophotometer setting the slit at 1 nm and using a resolution of 0.5 nm. All the spectra were corrected for the blank. Steady state emission spectra were measured on a Horiba Fluoromax-4 equipped with a cuvette holder with a stirring system, setting the excitation and the emission slits at 3 nm, the resolution at 1 nm, and excitation wavelength as it of the maximum absorbance. All the spectra were corrected for the blank.

The steady state absorption and emission of nucleobases (G, <sup>th</sup>G, <sup>tz</sup>G), their expected product with GDA (X, <sup>th</sup>X, <sup>tz</sup>X) and the mixture (25, 50, 75% expected products in reagents) were measured as the standard (Graph 7). The emission of G is absent since G doesn't have fluorescence property. The emission spectrums were excited at 360 nm (Graph 8).

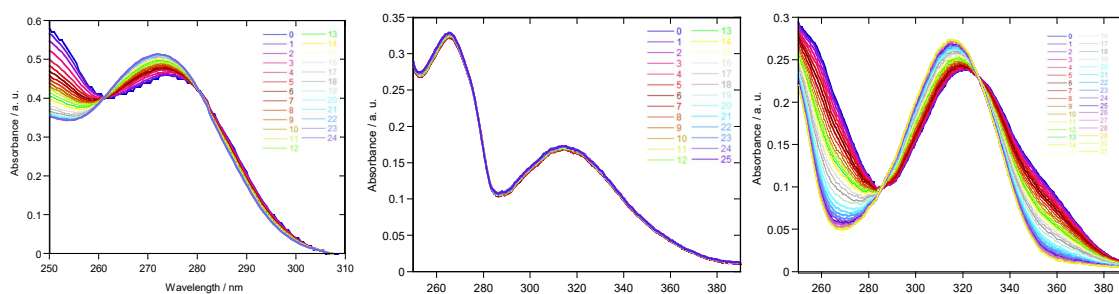


**Graph 7.** Absorbance trace of G, X and the mixture thereof (left);  ${}^{\text{th}}\text{G}$ ,  ${}^{\text{th}}\text{X}$  and the mixture thereof (middle);  ${}^{\text{tz}}\text{G}$ ,  ${}^{\text{tz}}\text{X}$  and the mixture thereof (right) in pH 7.5 Sodium Phosphate buffer at 37 °C. The color for G,  ${}^{\text{th}}\text{G}$ ,  ${}^{\text{tz}}\text{G}$  is blue and for X,  ${}^{\text{th}}\text{X}$ ,  ${}^{\text{tz}}\text{X}$  is red.



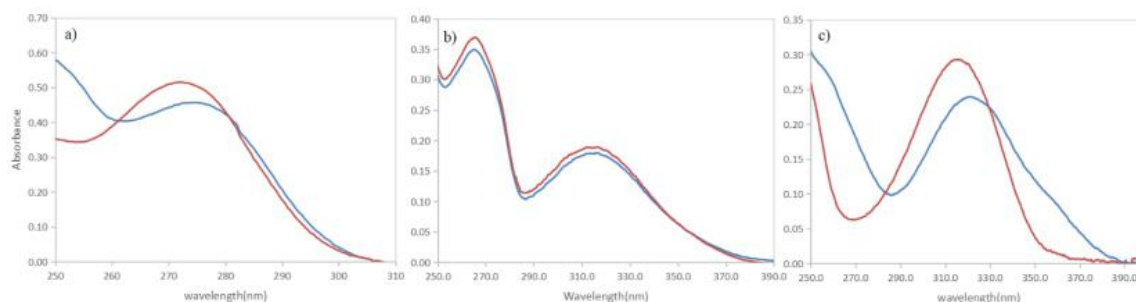
**Graph 8.** Emission trace of a)  ${}^{\text{th}}\text{G}$  (blue),  ${}^{\text{th}}\text{X}$  (red) and the mixture thereof; b)  ${}^{\text{tz}}\text{G}$  (blue),  ${}^{\text{tz}}\text{X}$  (red) and the mixture thereof in pH 7.5 Sodium Phosphate buffer at 37°C.

Steady state absorbance during time after the injection of GDA into G,  ${}^{\text{th}}\text{G}$  and  ${}^{\text{tz}}\text{G}$  in the sodium phosphate buffer (pH 7.5) at 37°C, which is in the same condition as the standard, were measured every 1 min and the absorption and emission for reagents and products were also obtained to compare with the standard (Graph 10, 11). All setting is same as the standard measurement.

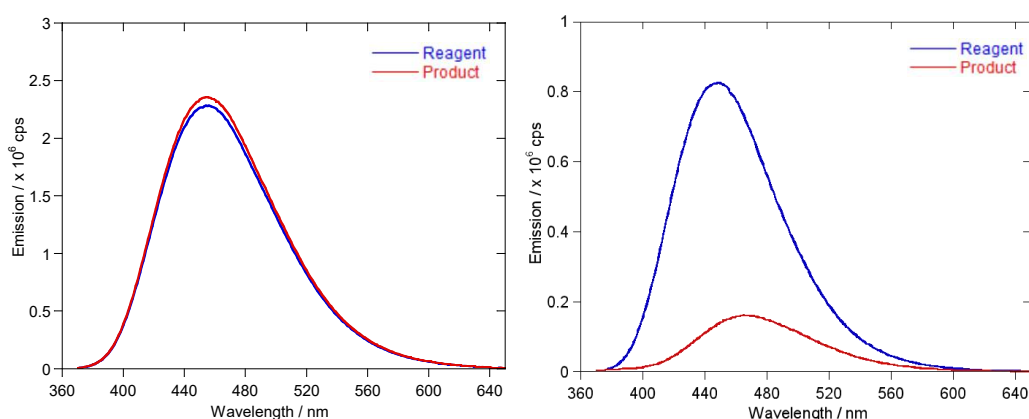


**Graph 9.** Steady state absorbance spectrum during the enzymatic reaction of G (left),  ${}^{\text{th}}\text{G}$  (middle) and  ${}^{\text{tz}}\text{G}$

(right).



**Graph 10.** Absorbance spectrum of reagent (Blue) and product (red) of the enzymatic conversion of GDA with a) G; b) <sup>th</sup>G; c) <sup>tz</sup>G in pH 7.5 Sodium Phosphate buffer at 37°C.



**Graph 11.** Emission spectrum of reagent (Blue) and product (red) of the enzymatic conversion of GDA with a) <sup>th</sup>G; b) <sup>tz</sup>G in pH 7.5 Sodium Phosphate buffer at 37°C.

#### 4.2.2. Absorbance and fluorescence based kinetic assays in presence of ADA

For G the <sup>tz</sup>G traces, the spectroscopy sample were prepared in 1 cm quartz cuvette from Helma to give a nucleobase and enzyme concentration of 2.00  $\mu$ M and 2666 mU/ml in phosphate buffer (50 mM, pH 7.5) and were measured at  $37.00 \pm 0.10^\circ\text{C}$ .

The ADA-mediated enzymatic conversion of guanine and the two guanine analogue were also monitored through the kinetic absorbance and emission base on the difference absorbance and emission between substrates and products at a fixed wavelength.

The enzymatic conversion for G to X was performed on a Shimadzu UV-2450 spectrophotometer setting the slit at 1 nm, taking a point every 2 seconds for total 500

seconds at 270nm upon addition of GDA.

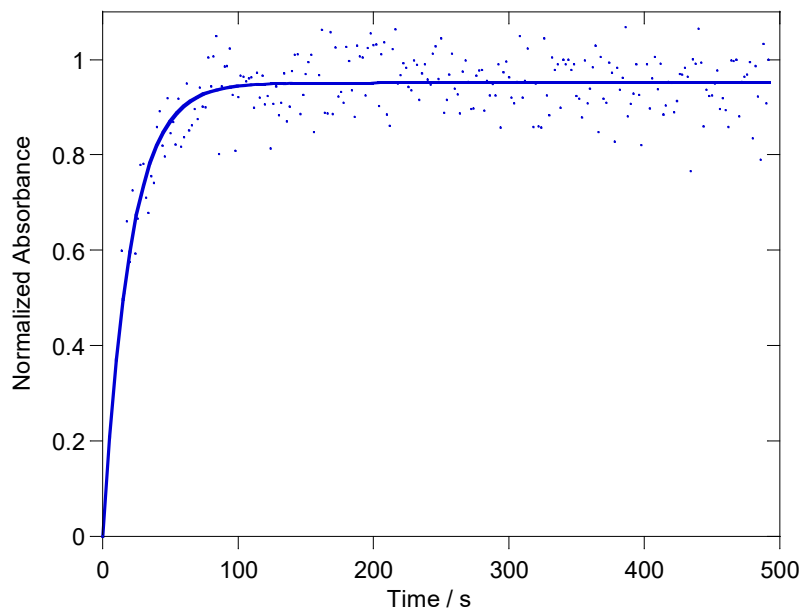
The enzymatic conversion of <sup>th</sup>G to <sup>th</sup>X, <sup>tz</sup>G to <sup>tz</sup>G were performed on a Horiba Fluoromax-4 setting the excitation and the emission slits at 3 nm, excitation wavelength as 360 nm, measured at the wavelength of their maximum emission for every 30 s and 10 s with the total time is 1 hour and 500 seconds for <sup>th</sup>G and <sup>tz</sup>G respectively (Graph 12, 13, 14).

The conversion of G and <sup>tz</sup>G with GDA were performed the first-order reaction. From the curve fit in the KaleidaGraph, the initial rate constant (k) can be obtained and the reaction half-time ( $t_{1/2}$ ) is calculated through following equations.

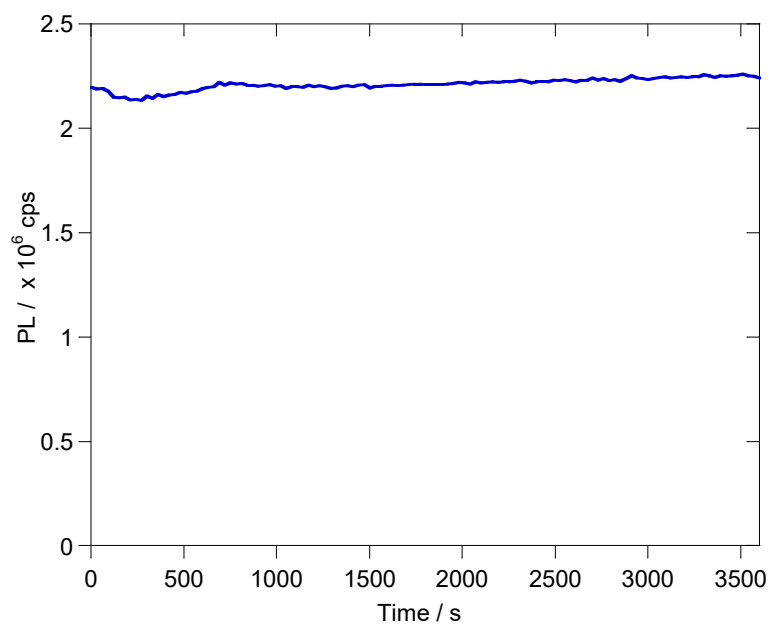
$$[A] = [A]_0 e^{-kt}$$

$$t_{1/2} = \frac{\ln(2)}{k_1}$$

Where [A] is the experimental signal over time,  $[A]_0$  is the initial signal at time zero, t is time, and k is the reaction rate coefficient. The coefficient of determination ( $R^2$ ) for the curve fit were higher than 0.97 for the G and higher than 0.99 for <sup>tz</sup>G traces.

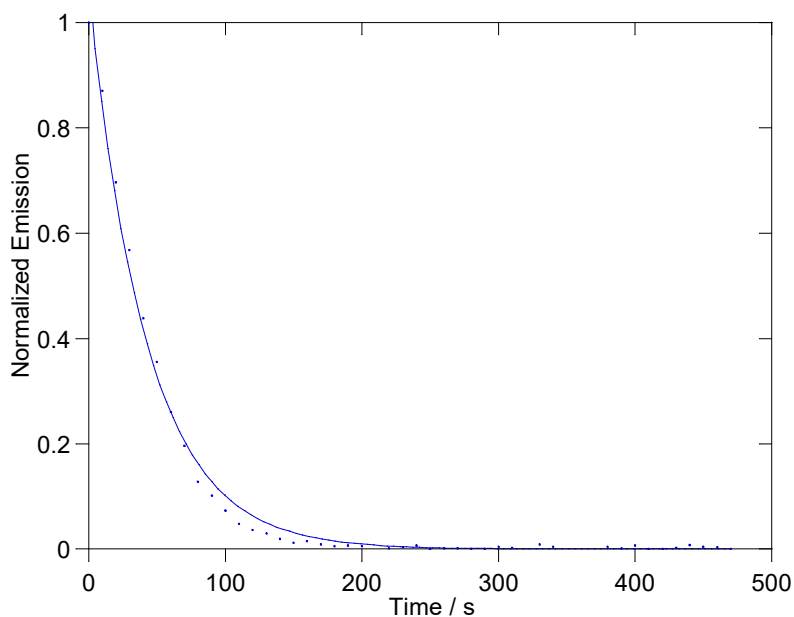


**Graph 12.** Kinetic absorbance spectrum of the enzymatic conversion of GDA with G



**Graph 13.** Kinetic emission spectrum of the enzymatic conversion of GDA with <sup>th</sup>G





**Graph 14.** Kinetic emission spectrum of the enzymatic conversion of GDA with  $^{125}\text{I}\text{G}$

#### 4.2.3. HPLC analysis for the enzymatic conversion

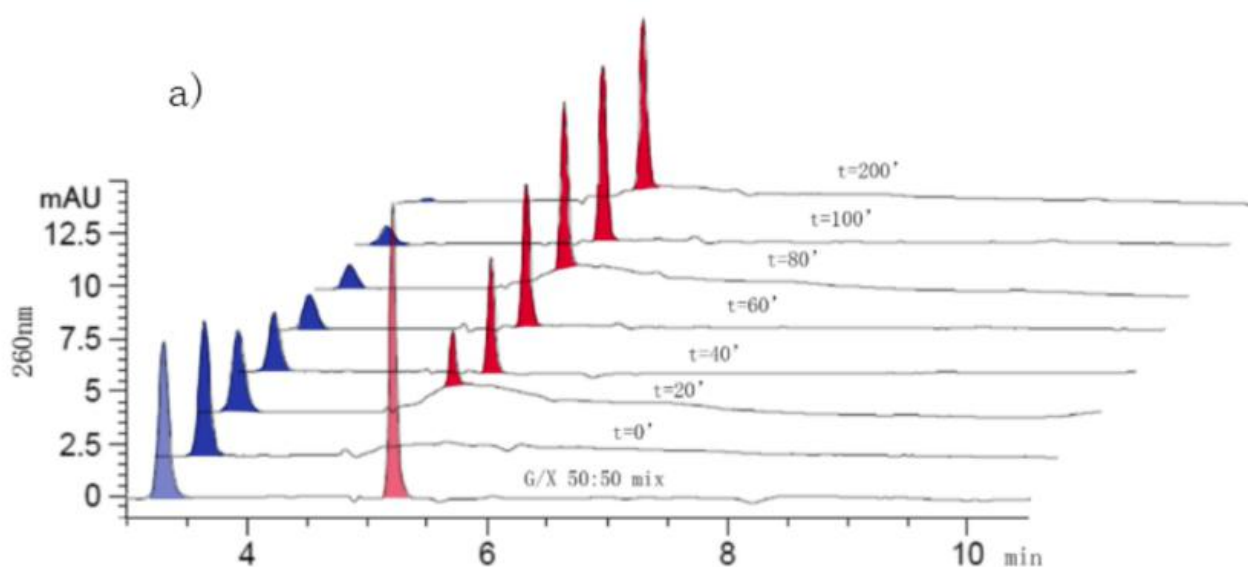
To further illustrate the results we got from last part, HPLC analysis was continued on a Agilent 1260 series system with an Eclipse XDB-C18, 5  $\mu\text{m}$ , 4.6x150 mm column. The condition is same as used in kinetic analysis, while the concentration of nucleobases and enzyme is 2.00  $\mu\text{M}$  and 2666 mU/ml in the solvent of phosphate buffer (50 mM, pH 7.5) and all samples reacted at  $37.00 \pm 0.10^\circ\text{C}$ . Formic acid is used to quench the reaction at 20, 40, 60, 80, 100, 200 and 500 seconds by mixing with the sample solution as the final concentration is 448 mM.<sup>[32-38]</sup> The samples are then centrifuged at 3000g for 5 minutes and filtered to get the supernatant, which is put in ice bath before HPLC analysis. Each injection (40  $\mu\text{l}$ ) of G and  $^{125}\text{I}\text{G}$  reactions was subjected to a gradient (15 minutes for the samples from G and  $^{125}\text{I}\text{G}$  reaction and 17 minutes for  $^3\text{H}\text{G}$ , from 0 to 95% of acetonitrile with 0.1% formic acid in water with 0.1% formic acid) followed by a flush (10 minutes). A flow rate of 1 ml / minute was used

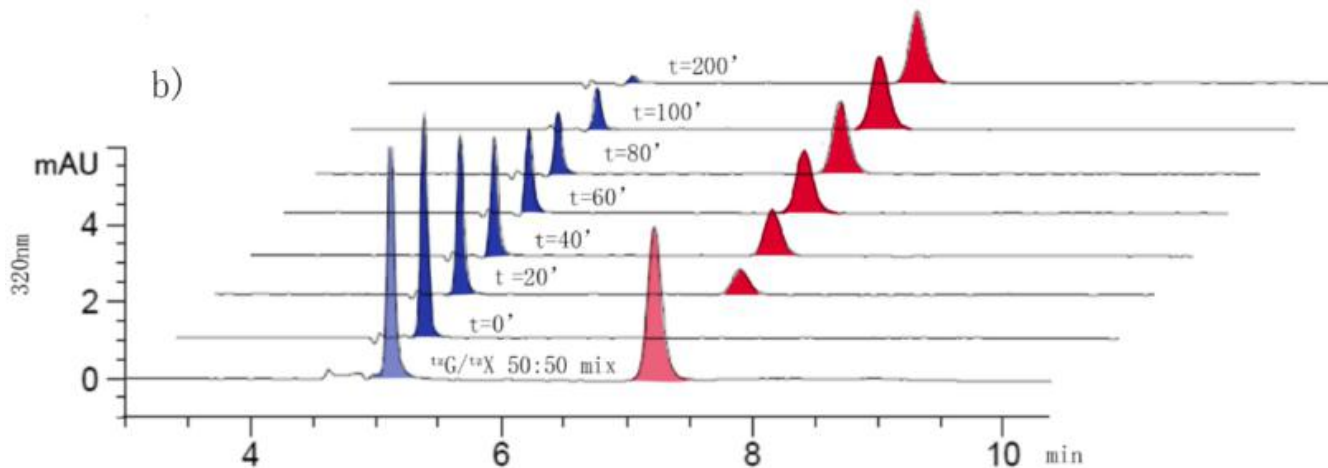
and the run was carried out at  $37.00 \pm 0.10^\circ\text{C}$ . Each run was monitored at 260, 280 and 320 nm with calibrated references at 600 nm and slit set at 1 nm (Graph 15, 17).

The HPLC traces were corrected for the blank and the concentration for each component is calculated from the height (optical density) of the peaks. The concentration changes were plotted as function of time. The trend line represents the loss of substrate (S) and the product (P) formation for a first-order kinetic reaction according to the equation below (Graph 16).

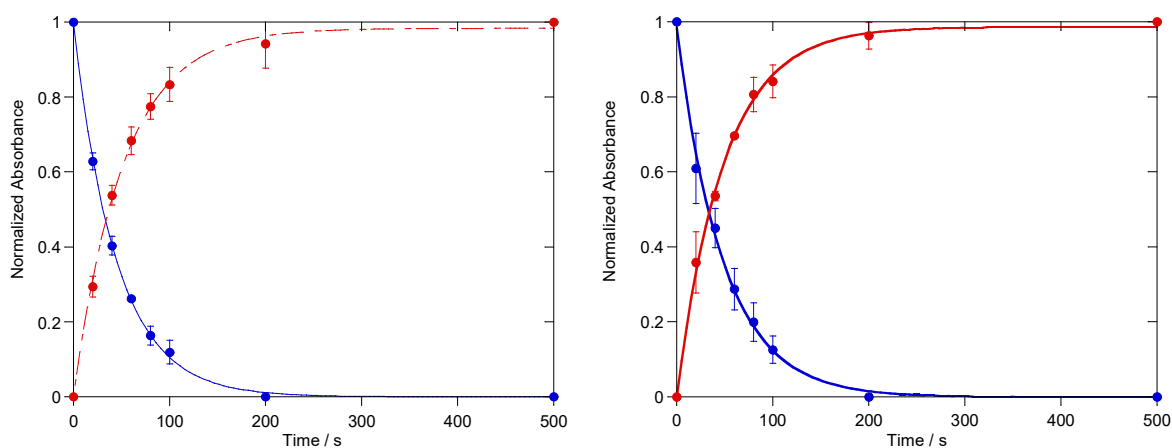
$$[S] = [S]_0 e^{-k_1 t}$$

$$[P] = [P]_\infty (1 - e^{-k_1 t})$$

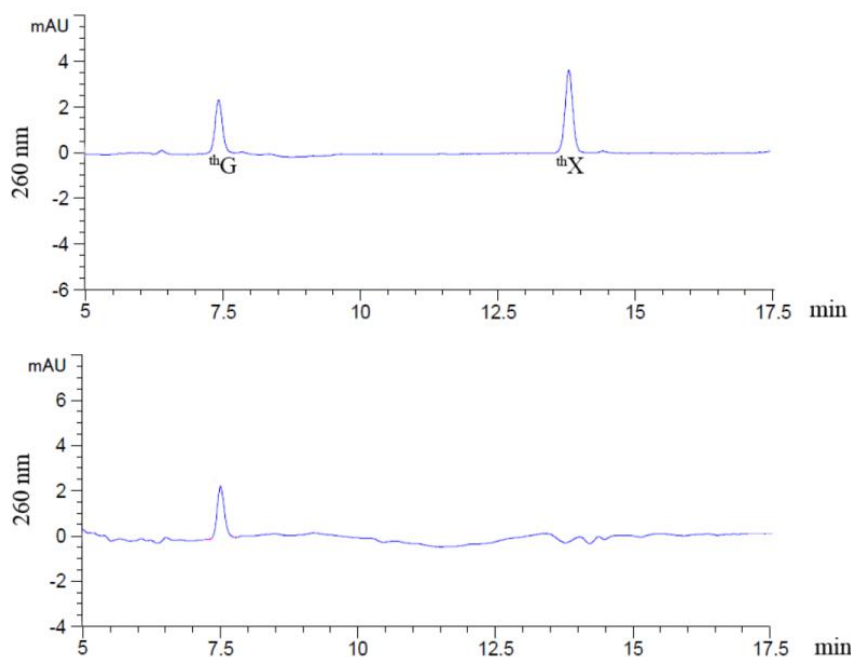




**Graph 15.** GDA-mediated deamination analysis by HPLC. Chromatogram showing the time course of enzymatic conversion of a) G to X b)  $^{12}\text{C}$ G to  $^{12}\text{C}$ X monitored at a) 260 nm b) 320 nm. Each set shows seven time points following the addition of ADA. For reference, an isomolar mixture of a) G and X b)  $^{12}\text{C}$ G and  $^{12}\text{C}$ X is shown at the bottom of each graph (transparent blue/red, respectively).



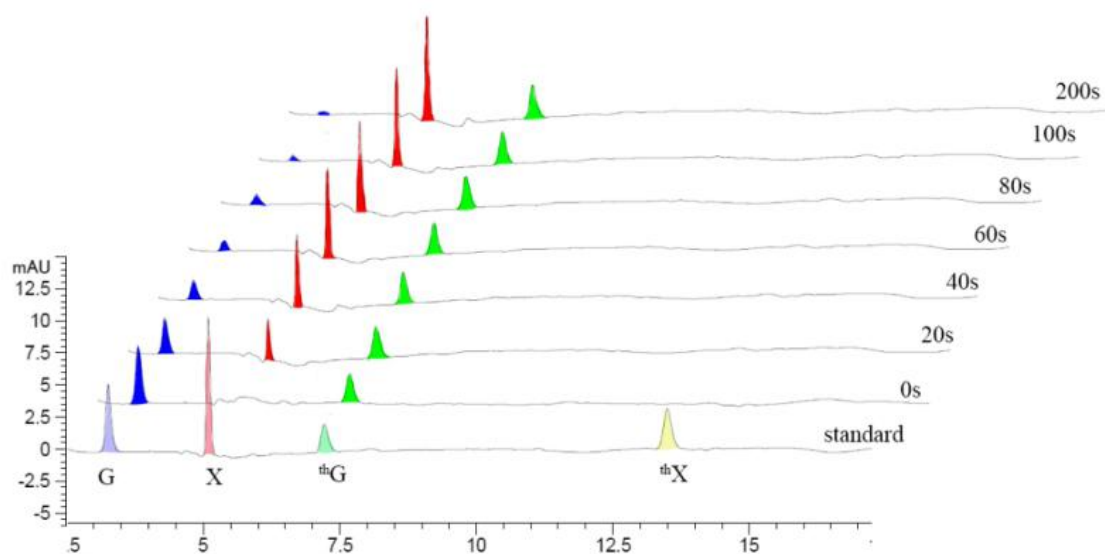
**Graph 16.** Trend line equations for the substrate consumption (solid red) and product formation (solid blue). Normalized absorbance versus time for G and  $^{12}\text{C}$ G (red dots), X and  $^{12}\text{C}$ X (blue dots) based on the HPLC traces.



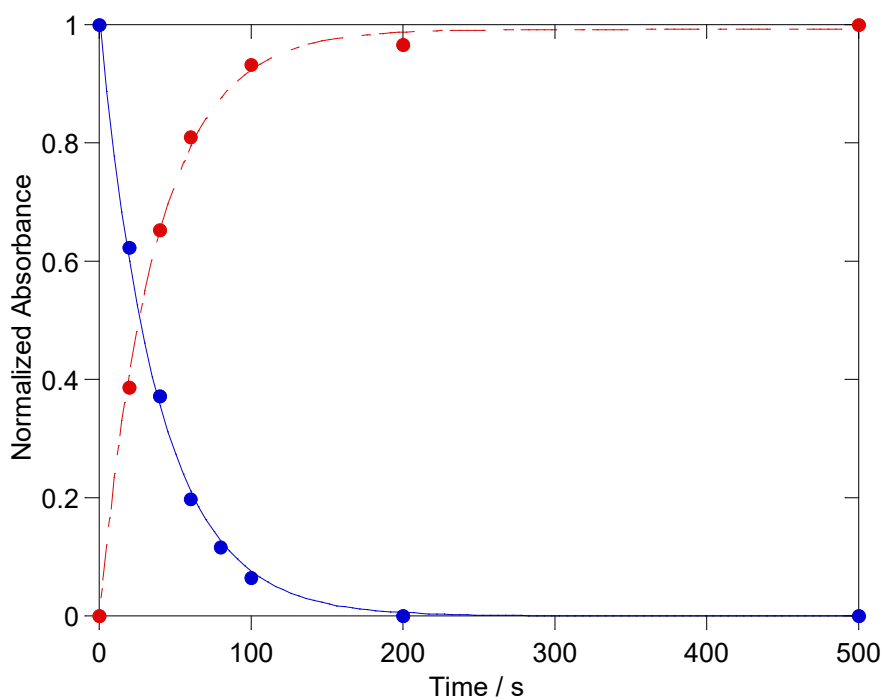
**Graph 17.** GDA-mediated deamination analysis for <sup>th</sup>G to <sup>th</sup>X by HPLC.

#### 4.2.4. Activity of GDA after mixing with <sup>th</sup>G

The GDA-mediated deamination of G to X in the mixture of mixed GDA and <sup>th</sup>G was performed in the same condition of the individual deamination for HPLC analysis. Firstly, the GDA and <sup>th</sup>G is well-mixed and put in the incubator to maintain the temperature of 37°C, then the G is added into the mixture and the reaction is quenched by formic acid at 20s, 40s, 60s, 80s, 100s, 200s and 500s. After the same treatment we discussed before, the samples is analyzed by HPLC (Graph 18, 19).



**Graph 18.** GDA-mediated deamination analysis by HPLC of enzymatic conversion of G to X in the mixture of well mixed GDA and <sup>th</sup>G based on the 260 nm traces. For reference, an isomolar mixture of G, X, <sup>th</sup>G and <sup>th</sup>X is shown at the bottom of each graph (transparent blue/red/green/yellow, respectively).



**Graph 19.** Trend line equations for the substrate consumption (solid red) and product formation (solid blue), Normalized absorbance versus time for G (red dots), X (blue dots) in the mixture of GDA and <sup>th</sup>G based on the HPLC traces at 260 nm.

## 5. Conclusion

In summary, the two emissive Guanine analogues (<sup>th</sup>G, <sup>tz</sup>G) and two xanthine analogues (<sup>th</sup>X, <sup>tz</sup>X), based on a thieno and isothiazole scaffold, respectively, were synthesized and characterized for their improved photophysical properties than native guanine and different biochemical activity due to the structural difference.

First of all, the crystal structure of these nucleobases analogues were measured to make sure the expected compound has been gotten. The result also proved the existence of tautomerization in the solid state of <sup>th</sup>G, which was also revealed by the convolution of emission spectra of <sup>th</sup>G. The similar H-bonding pattern of the analogues confirmed their structural analogy as the native nucleobases, makes them have the ability to be stabilized at the correct position of the active site of the enzyme. All of the nucleobases analogues possess fluorescence properties, illustrate by their emission spectra and sufficient quantum yield, which makes them a valuable surrogates of guanine. Their biochemical function was also tested by monitoring their activity with guanine deaminase (GDA), while the conversion of thienol guanine (<sup>th</sup>G) to thienol xanthine (<sup>th</sup>X) was barely catalyzed, however, the thiazole Guanine (<sup>tz</sup>G) can react with GDA at the almost the same rate as the native guanine. This confirmed the significant role of N7 in the reaction since it's the biggest difference between <sup>th</sup>G and <sup>tz</sup>G. Furthermore, the unaffected activity of native guanine with the mixture of <sup>th</sup>G and GDA showed that <sup>th</sup>G is not an inhibitor in this specific concentration, which further proved that the <sup>th</sup>G cannot bind with GDA because of the absence of N7.

These observations about the novel guanine surrogates display improved structural and functional characteristics and help us better understand the binding and mechanism during the deamination process. What's more, a good fluorescent surrogate of guanine is provided, which facilitate a new fluorescence method to accurately and efficiently monitor the activity of GDA. Furthermore, help identify the inhibitor and activator to manipulate the activity of GDA, which is significant for the research of neurological recovery.

## 6. Additional information.

### 6.1. NMR Spectrum

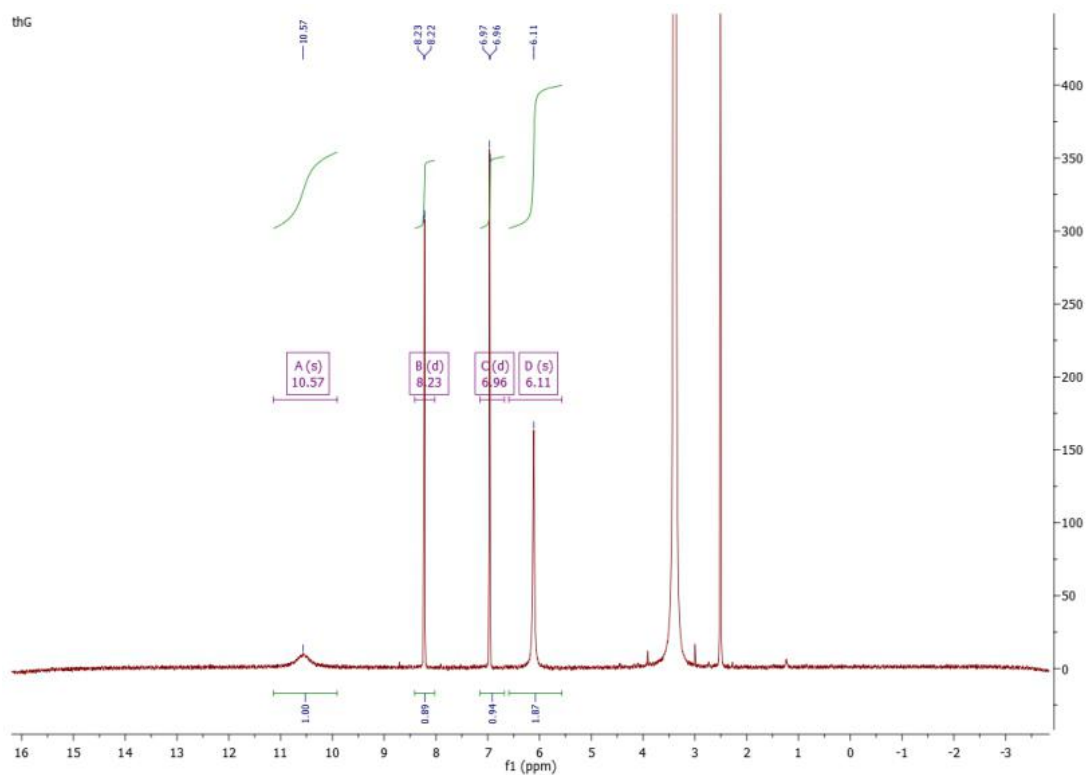


Figure 8. <sup>1</sup>H NMR of thG

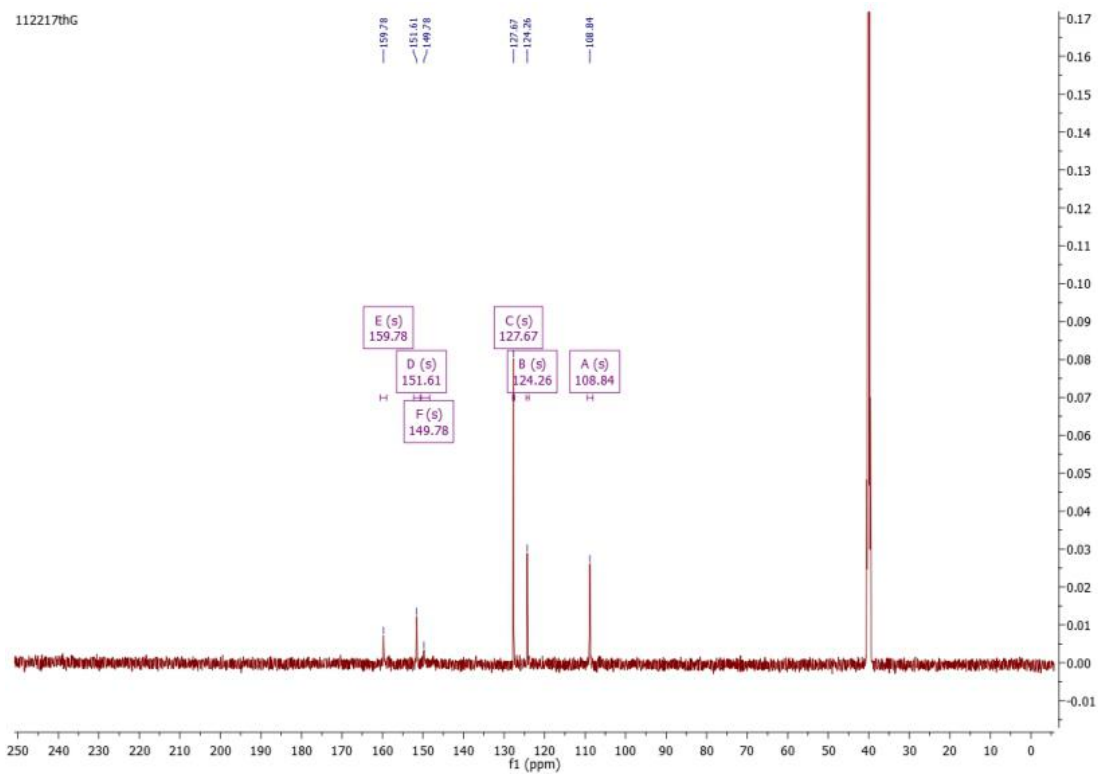


Figure 9. <sup>13</sup>C NMR of thG



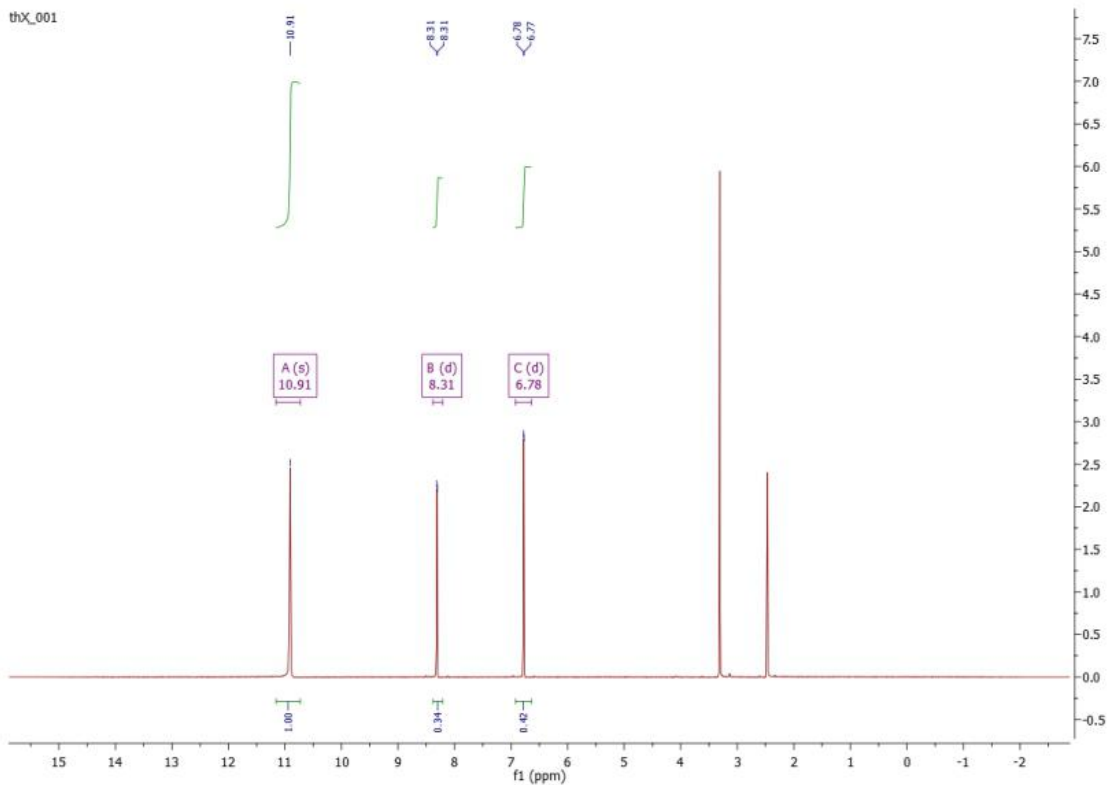


Figure 10.  $^1\text{H}$  NMR of  $^{\text{th}}\text{X}$

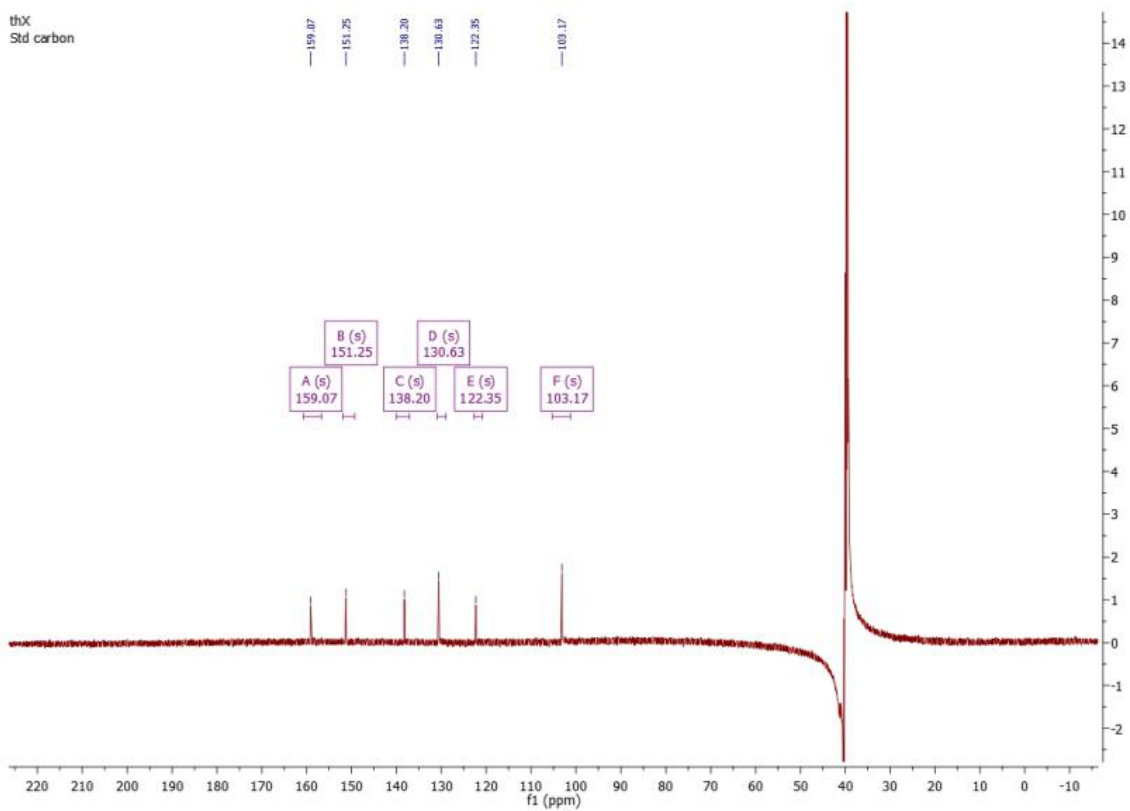


Figure 11.  $^{13}\text{C}$  NMR of  $^{\text{th}}\text{X}$

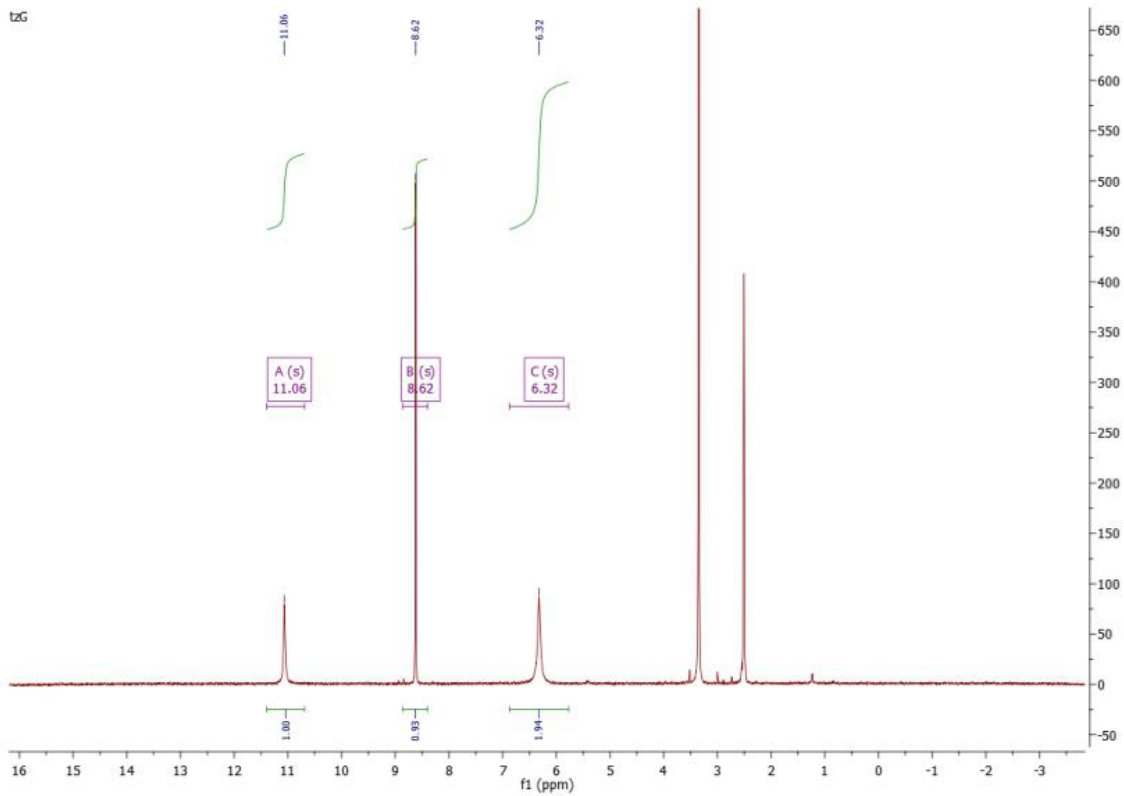


Figure 12.  $^1\text{H}$  NMR of tzG

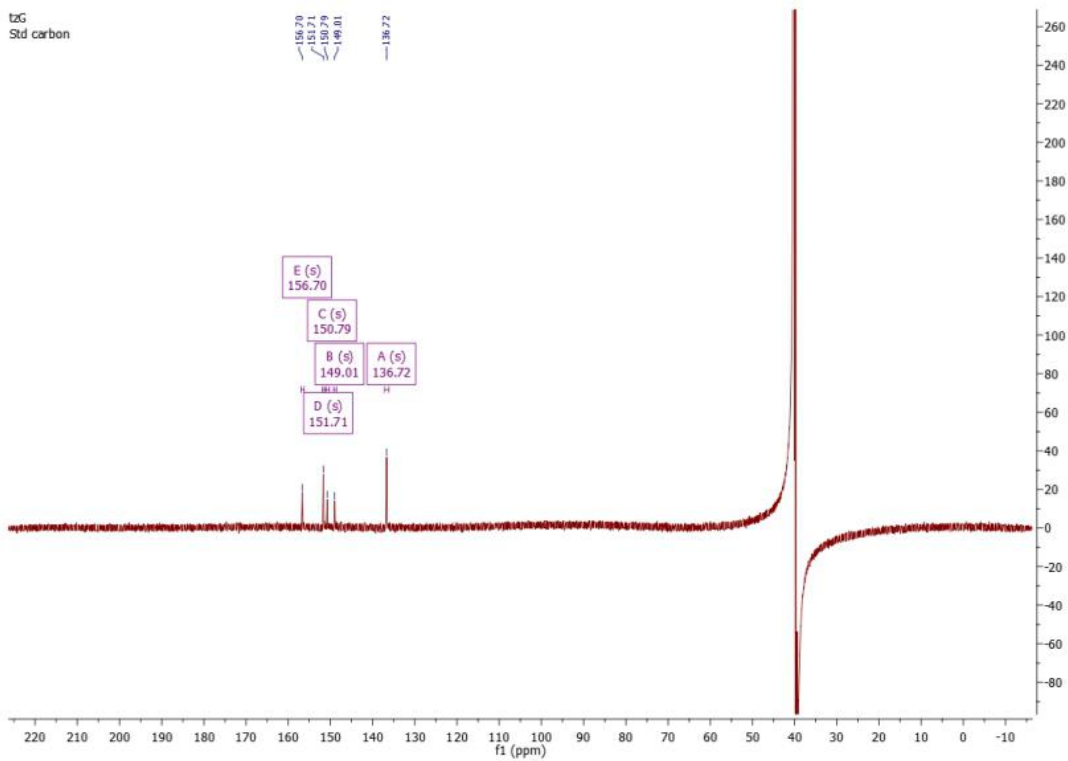


Figure 13.  $^{13}\text{C}$  NMR of tzG

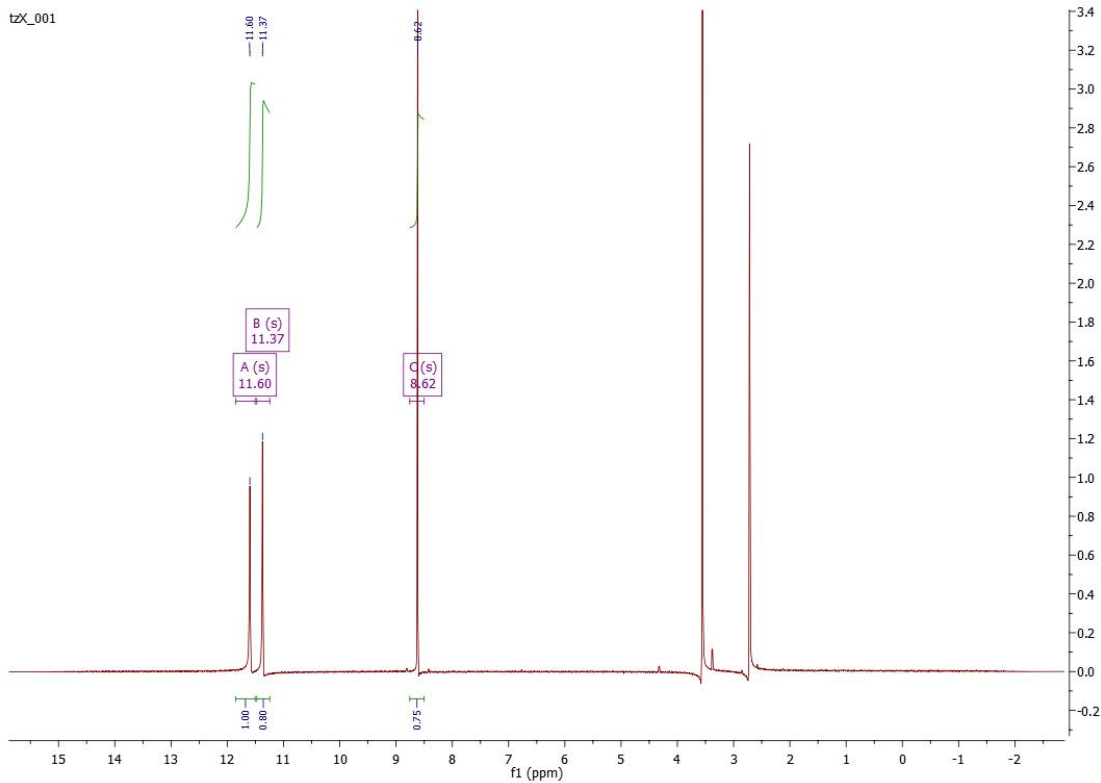


Figure 14.  $^1\text{H}$  NMR of tzX

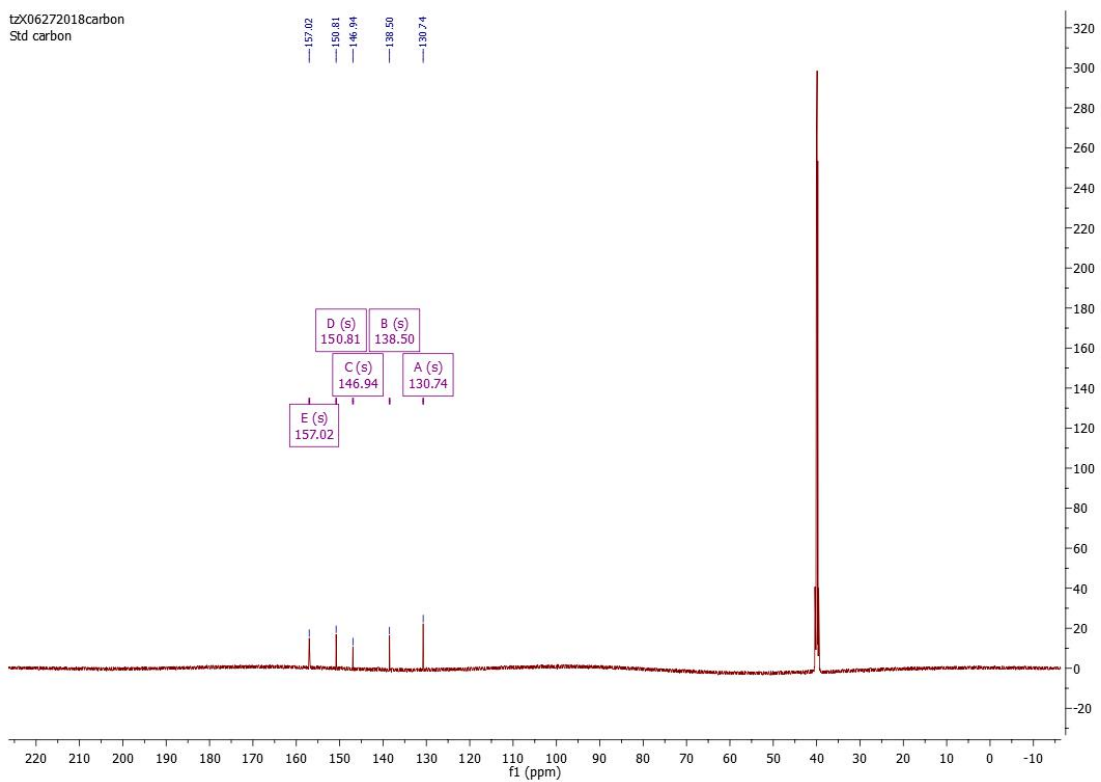
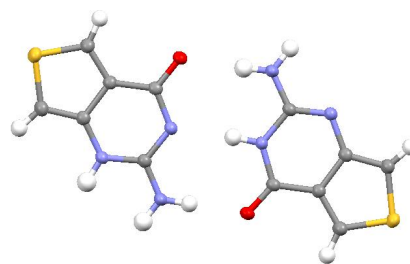


Figure 15.  $^{13}\text{C}$  NMR of tzX

## 6.2. X-ray crystal structure

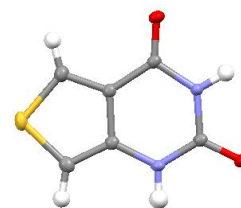
**Table 3.** Crystal data and structure refinement for <sup>th</sup>G.

Report date	2018-03-21	
Identification code	thG	
Empirical formula	C <sub>6</sub> H <sub>5</sub> N <sub>3</sub> O S	
Molecular formula	C <sub>6</sub> H <sub>5</sub> N <sub>3</sub> O S	
Formula weight	167.19	
Temperature	100.0 K	
Wavelength	1.54178 Å	
Crystal system	Monoclinic	
Space group	P 1 21/c 1	
Unit cell dimensions	a = 7.2425(6) Å	= 90°.
	b = 14.8279(12) Å	= 94.751(3)°.
	c = 12.8685(11) Å	= 90°.
Volume	1377.2(2) Å <sup>3</sup>	
Z	8	
Density (calculated)	1.613 Mg/m <sup>3</sup>	
Absorption coefficient	3.680 mm <sup>-1</sup>	
F(000)	688	
Crystal size	0.231 x 0.157 x 0.084 mm <sup>3</sup>	
Crystal color, habit	Colorless Block	
Theta range for data collection	4.558 to 68.321°.	
Index ranges	-8<=h<=8, -17<=k<=17, -15<=l<=14	
Reflections collected	25913	
Independent reflections	2519 [R(int) = 0.0310, R(sigma) = 0.0171]	
Completeness to theta = 68.000°	99.8 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.3201 and 0.2149	
Refinement method	Full-matrix least-squares on F <sup>2</sup>	
Data / restraints / parameters	2519 / 6 / 223	
Goodness-of-fit on F <sup>2</sup>	1.062	
Final R indices [I>2sigma(I)]	R1 = 0.0301, wR2 = 0.0833	
R indices (all data)	R1 = 0.0319, wR2 = 0.0849	
Extinction coefficient	n/a	
Largest diff. peak and hole	0.256 and -0.267 e.Å <sup>-3</sup>	

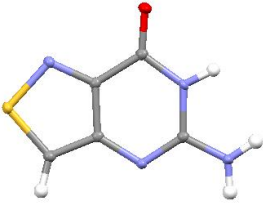


**Table 4.** Crystal data and structure refinement for <sup>th</sup>X.

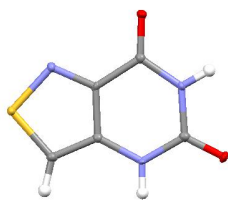
Report date	2018-04-04	
Identification code	THX01	
Empirical formula	C6 H4 N2 O2 S	
Molecular formula	C6 H4 N2 O2 S	
Formula weight	168.17	
Temperature	100.0 K	
Wavelength	0.71073 Å	
Crystal system	Monoclinic	
Space group	P 1 21/c 1	
Unit cell dimensions	a = 10.1968(16) Å	= 90°.
	b = 5.3149(8) Å	= 102.119(4)°.
	c = 12.257(2) Å	= 90°.
Volume	649.48(18) Å <sup>3</sup>	
Z	4	
Density (calculated)	1.720 Mg/m <sup>3</sup>	
Absorption coefficient	0.436 mm <sup>-1</sup>	
F(000)	344	
Crystal size	0.271 x 0.117 x 0.093 mm <sup>3</sup>	
Crystal color, habit	Colorless Block	
Theta range for data collection	2.043 to 28.278°.	
Index ranges	-13<=h<=13, -6<=k<=7, -16<=l<=13	
Reflections collected	8784	
Independent reflections	1596 [R(int) = 0.0582, R(sigma) = 0.0463]	
Completeness to theta = 25.000°	100.0 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.2482 and 0.2096	
Refinement method	Full-matrix least-squares on F <sup>2</sup>	
Data / restraints / parameters	1596 / 2 / 108	
Goodness-of-fit on F <sup>2</sup>	1.075	
Final R indices [I>2sigma(I)]	R1 = 0.0376, wR2 = 0.0963	
R indices (all data)	R1 = 0.0435, wR2 = 0.1004	
Extinction coefficient	n/a	
Largest diff. peak and hole	0.361 and -0.331 e.Å <sup>-3</sup>	



**Table 5.** Crystal data and structure refinement for <sup>12</sup>G.

Report date	2018-07-10		
Identification code	tor119		
Empirical formula	C <sub>5</sub> H <sub>4</sub> N <sub>4</sub> O S		
Molecular formula	C <sub>5</sub> H <sub>4</sub> N <sub>4</sub> O S		
Formula weight	168.18		
Temperature	100.0 K		
Wavelength	0.71073 Å		
Crystal system	Monoclinic		
Space group	P 1 21/c 1		
Unit cell dimensions	a = 3.6599(2) Å		α = 90°.
	b = 11.2567(5) Å		β = 96.825(2)°.
	c = 15.0364(7) Å		γ = 90°.
Volume	615.09(5) Å <sup>3</sup>		
Z	4		
Density (calculated)	1.816 Mg/m <sup>3</sup>		
Absorption coefficient	0.457 mm <sup>-1</sup>		
F(000)	344		
Crystal size	0.03 x 0.005 x 0.005 mm <sup>3</sup>		
Crystal color, habit	light yellow needle		
Theta range for data collection	2.266 to 26.720°.		
Index ranges	-4 ≤ h ≤ 4, -13 ≤ k ≤ 14, -18 ≤ l ≤ 18		
Reflections collected	9691		
Independent reflections	1303 [R(int) = 0.0555]		
Completeness to theta = 25.242°	100.0 %		
Absorption correction	Semi-empirical from equivalents		
Max. and min. transmission	0.7455 and 0.6591		
Refinement method	Full-matrix least-squares on F <sup>2</sup>		
Data / restraints / parameters	1303 / 0 / 100		
Goodness-of-fit on F <sup>2</sup>	1.046		
Final R indices [I > 2σ(I)]	R1 = 0.0337, wR2 = 0.0733		
R indices (all data)	R1 = 0.0471, wR2 = 0.0784		
Largest diff. peak and hole	0.332 and -0.314 e.Å <sup>-3</sup>		

**Table 6.** Crystal data and structure refinement for <sup>12</sup>X.

Report date	2018-07-11	
Identification code	tor120	
Empirical formula	C <sub>5</sub> H <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	
Molecular formula	C <sub>5</sub> H <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	
Formula weight	169.16	
Temperature	100.0 K	
Wavelength	0.71073 Å	
Crystal system	Monoclinic	
Space group	P 1 21/c 1	
Unit cell dimensions	a = 6.9522(10) Å	α = 90°.
	b = 12.2384(17) Å	β = 93.431(3)°.
	c = 6.9021(10) Å	γ = 90°.
Volume	586.20(14) Å <sup>3</sup>	
Z	4	
Density (calculated)	1.917 Mg/m <sup>3</sup>	
Absorption coefficient	0.488 mm <sup>-1</sup>	
F(000)	344	
Crystal size	0.12 x 0.08 x 0.04 mm <sup>3</sup>	
Crystal color, habit	colorless block	
Theta range for data collection	2.935 to 26.736°.	
Index ranges	-8 ≤ h ≤ 8, 0 ≤ k ≤ 15, 0 ≤ l ≤ 8	
Reflections collected	1244	
Independent reflections	1244 [R(int) = 0.0362]	
Completeness to theta = 25.242°	100.0 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.491 and 0.402	
Refinement method	Full-matrix least-squares on F <sup>2</sup>	
Data / restraints / parameters	1244 / 0 / 101	
Goodness-of-fit on F <sup>2</sup>	1.036	
Final R indices [I > 2σ(I)]	R1 = 0.0362, wR2 = 0.0873	
R indices (all data)	R1 = 0.0456, wR2 = 0.0927	
Largest diff. peak and hole	0.409 and -0.459 e.Å <sup>-3</sup>	

### 6.3. Raw HPLC Traces

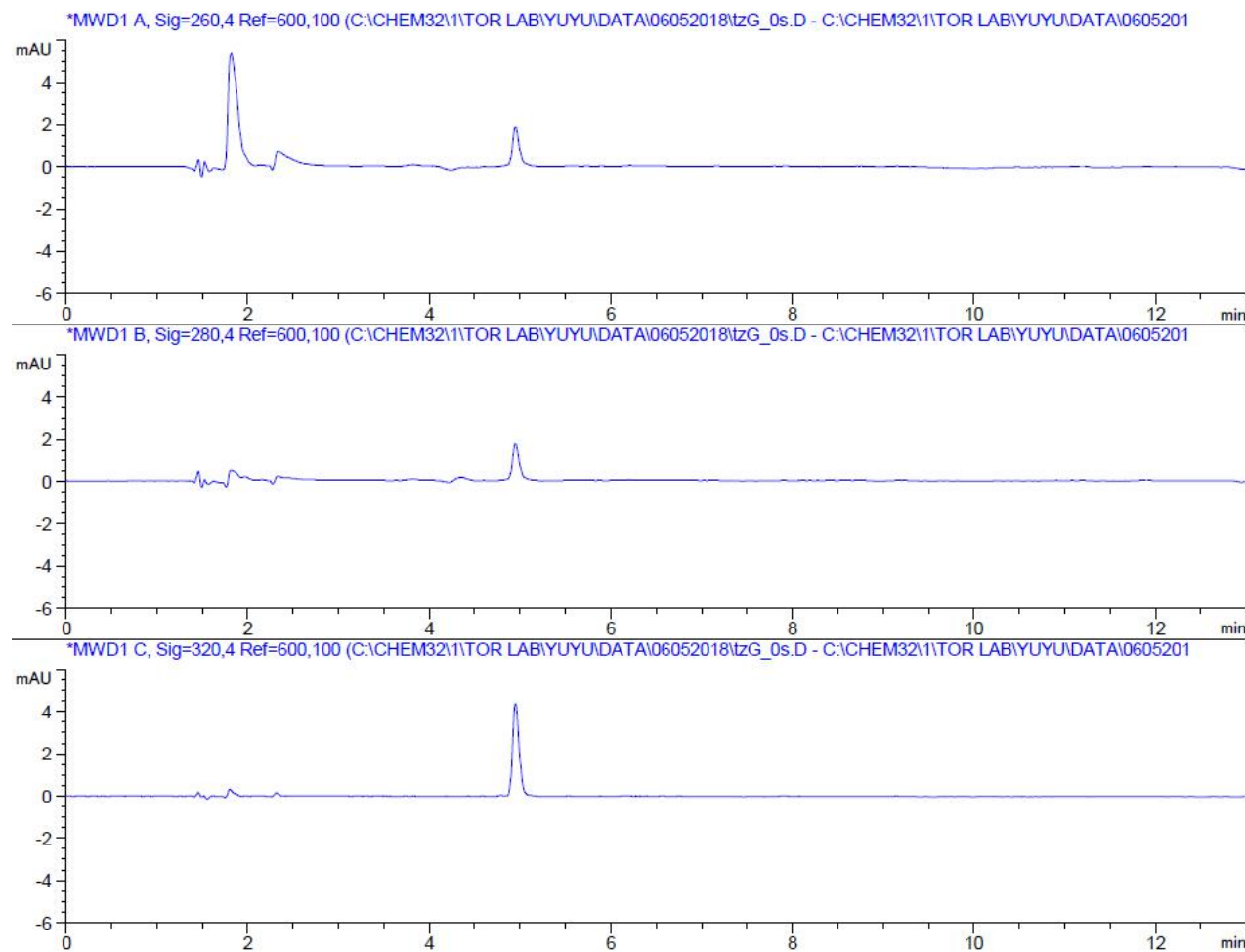


Figure 16. GDA-mediated  $^{125}\text{I}$ G to  $^{125}\text{I}$ X conversion at  $t = 0$  s.



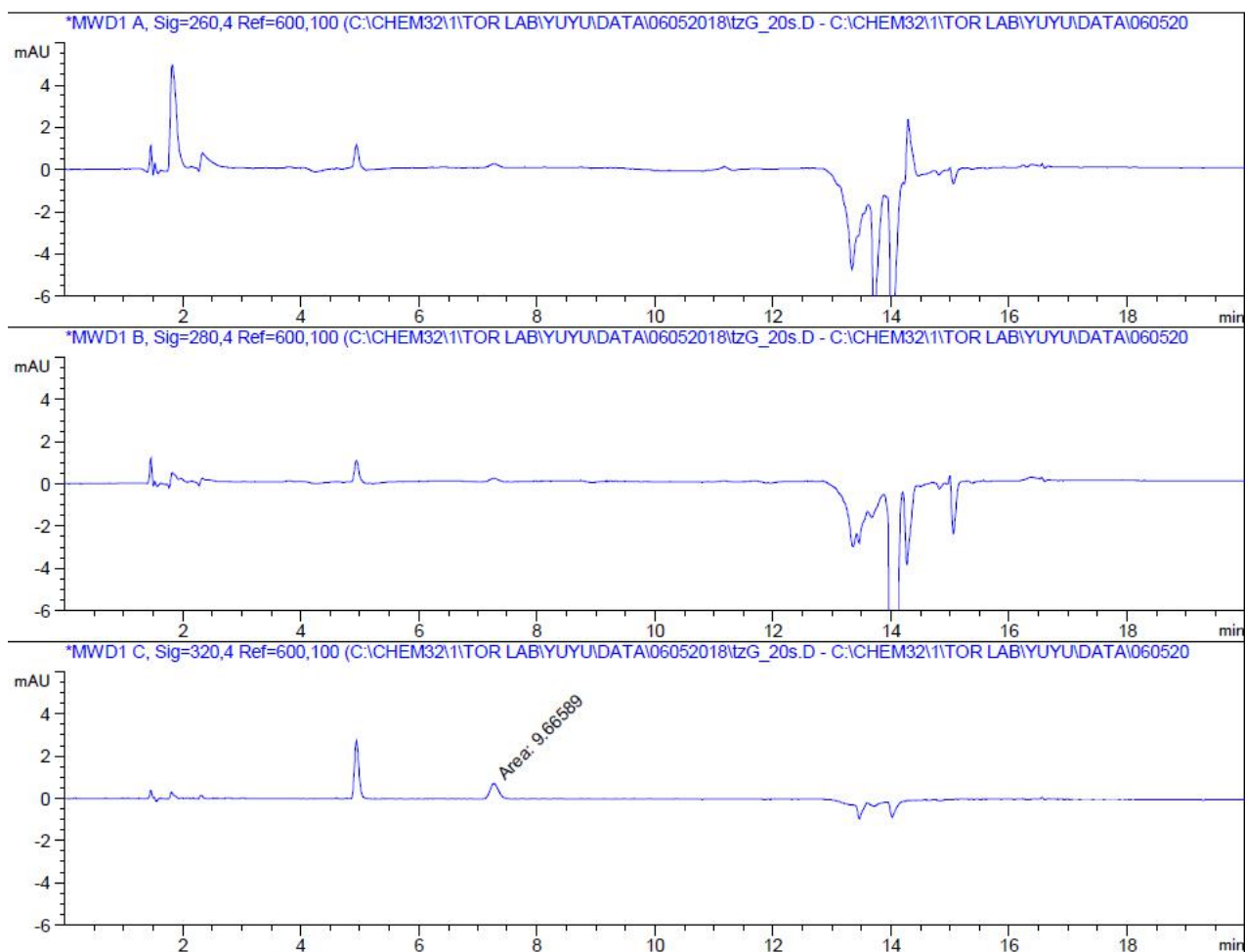


Figure 17. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 20 s.

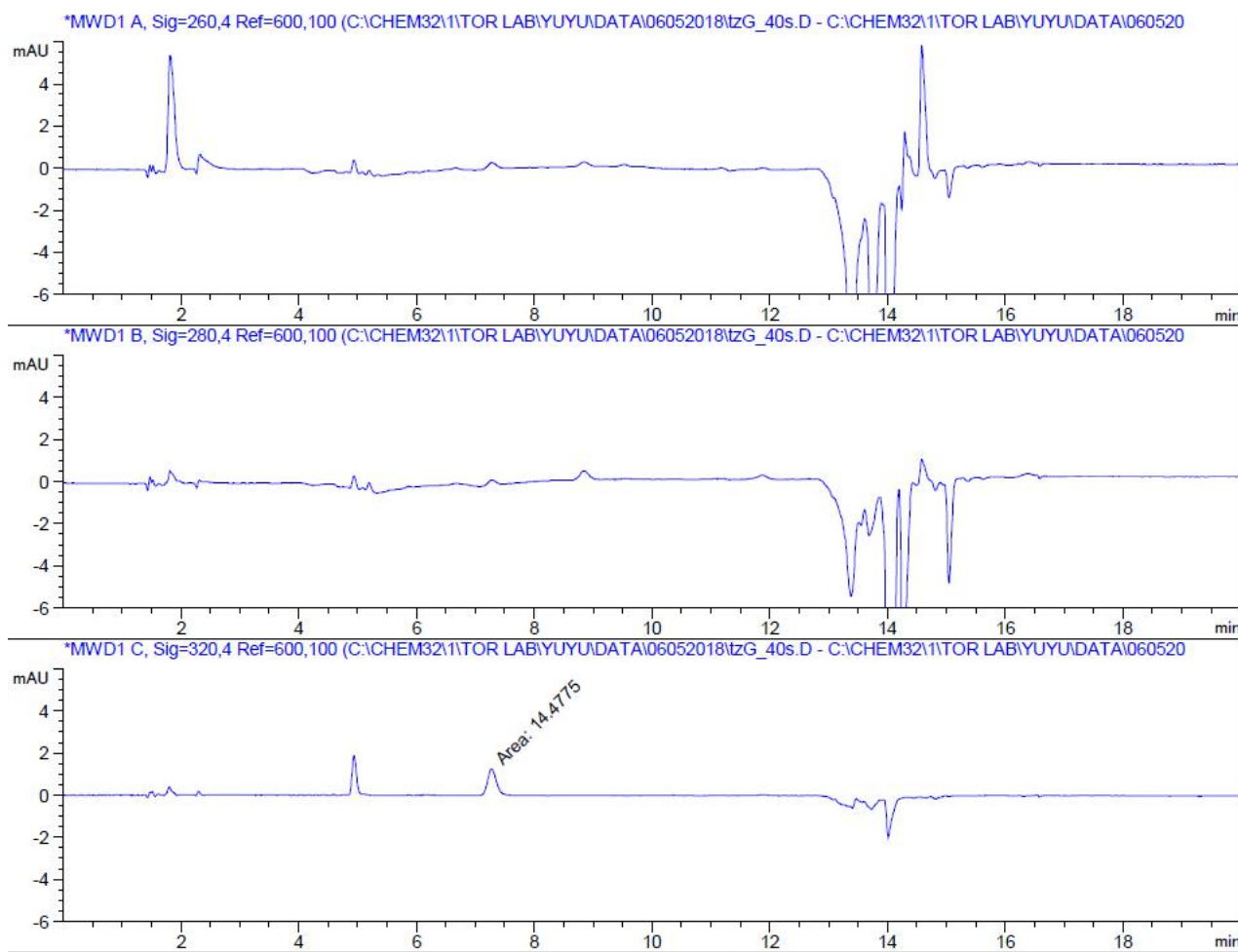


Figure 18. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 40 s.

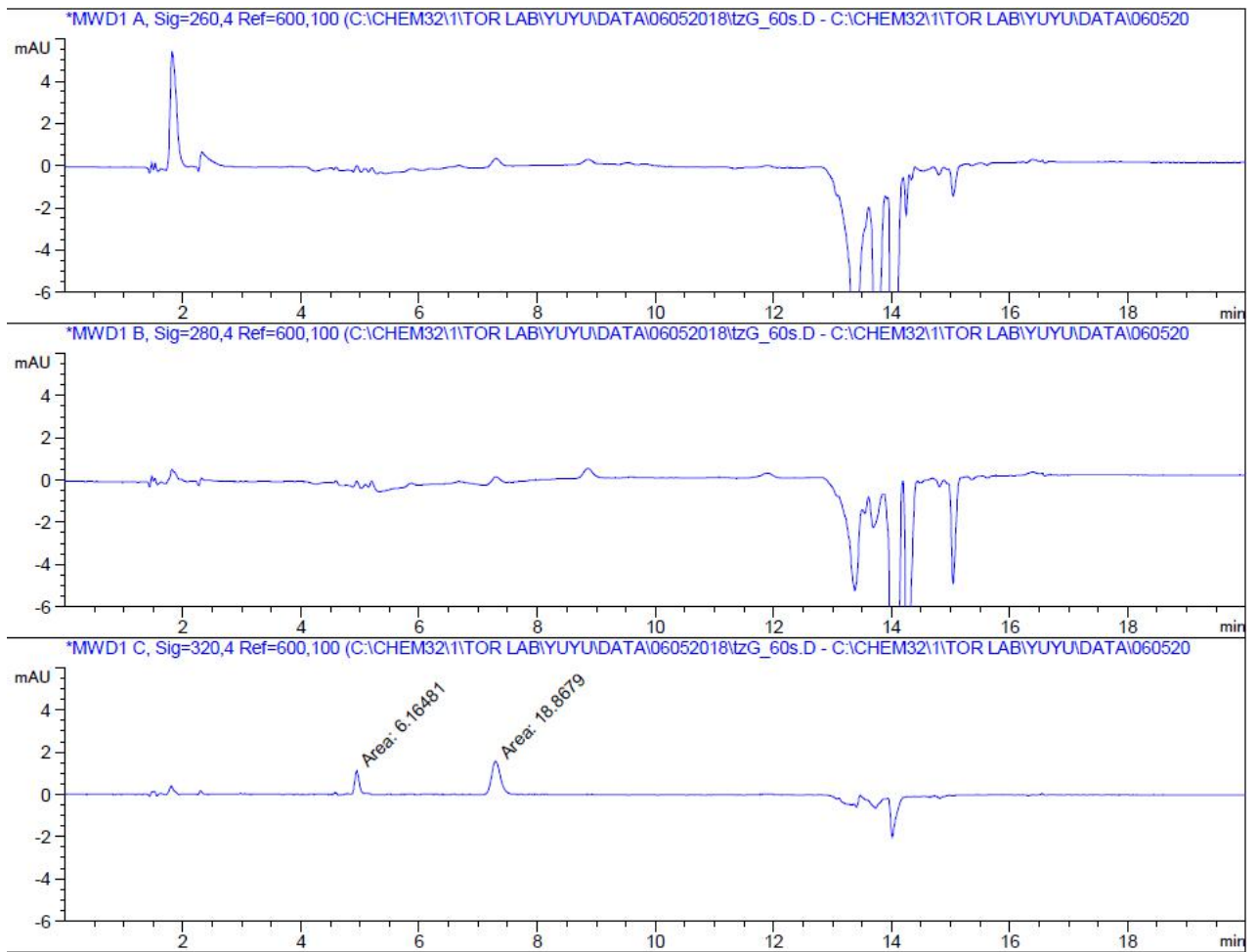


Figure 19. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 60 s.

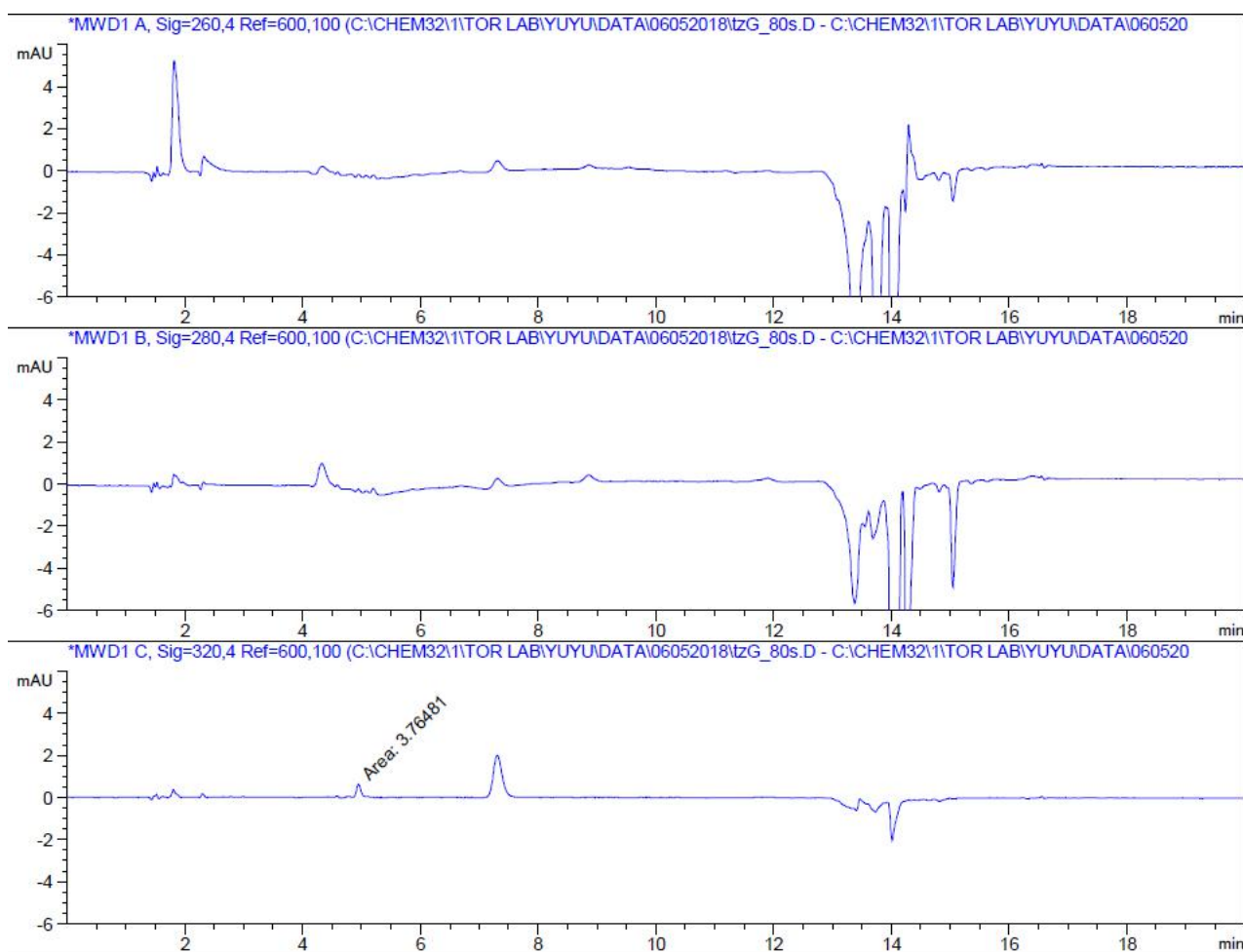


Figure 20. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 80 s.

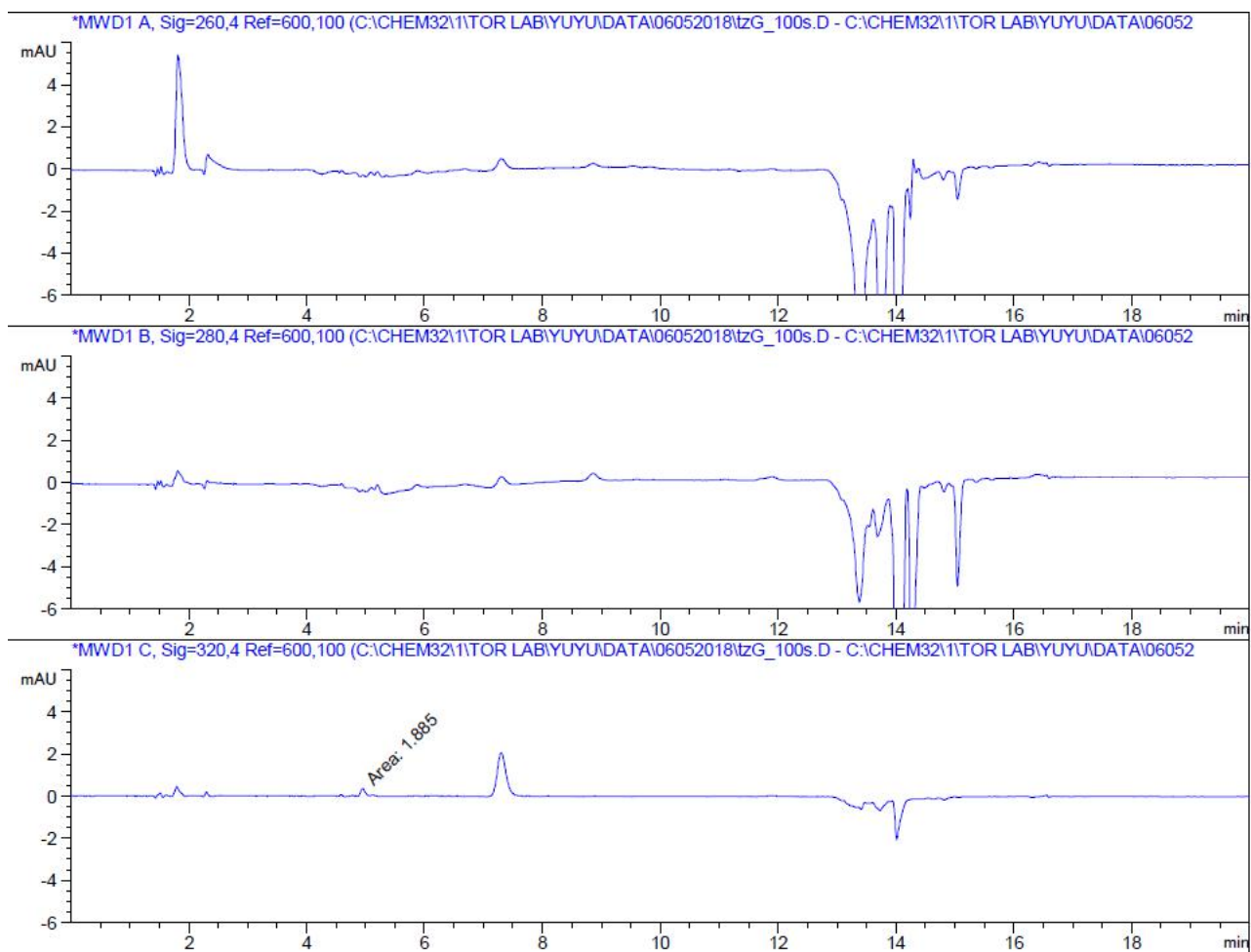


Figure 21. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 100 s.

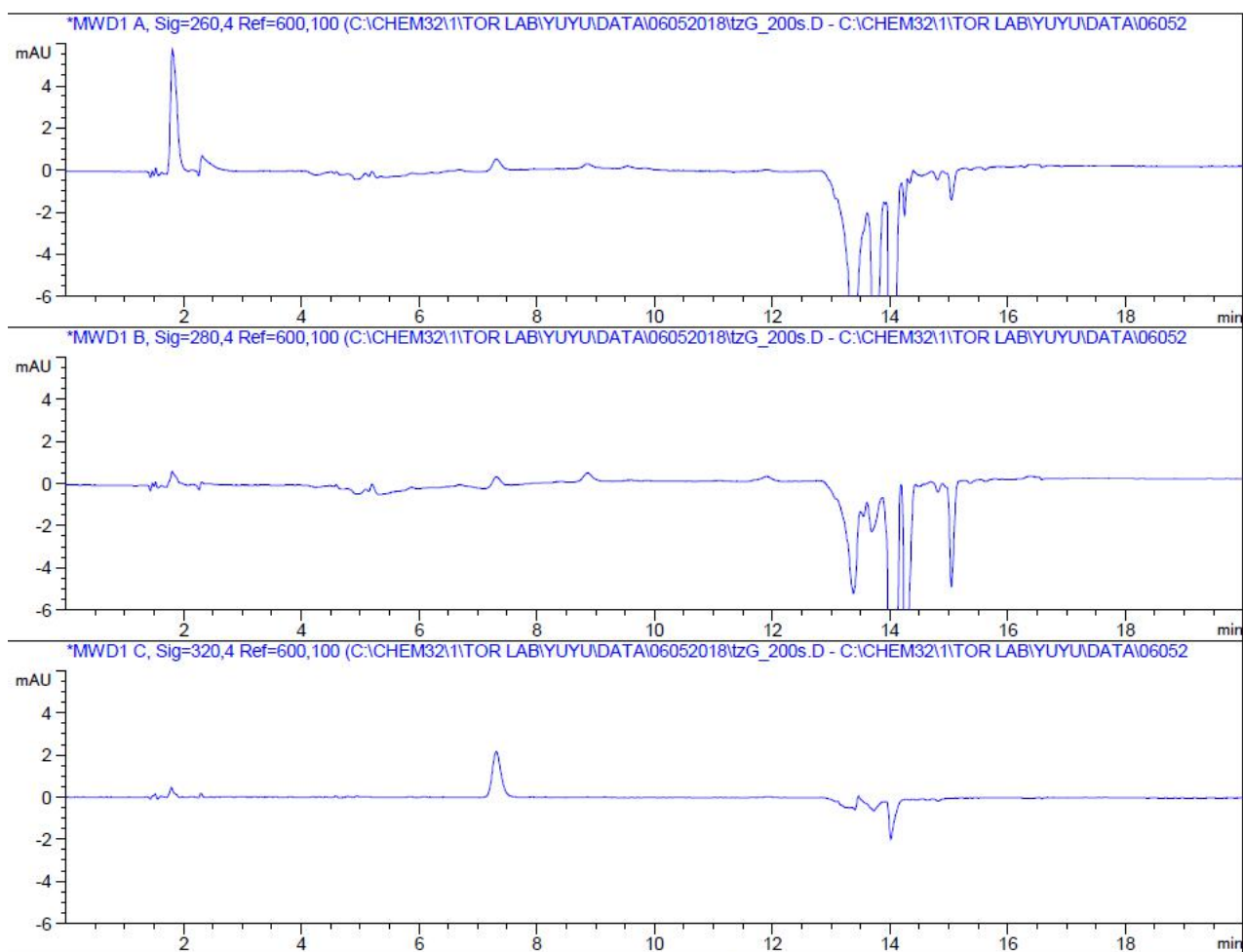


Figure 22. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 200 s.

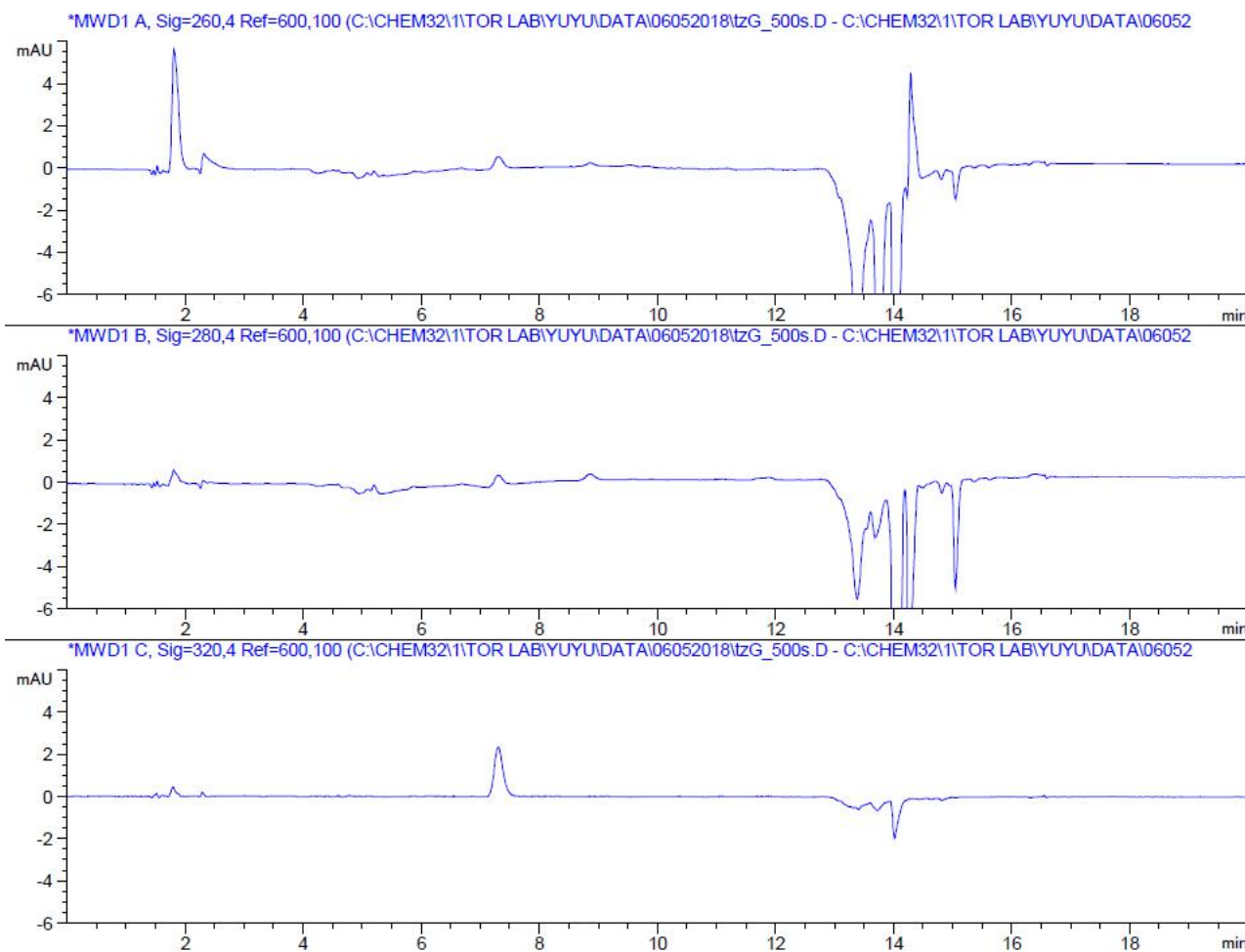


Figure 23. GDA-mediated <sup>12</sup>G to <sup>12</sup>X conversion at t = 500 s.

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