# Synthesis and evaluation of 3'-[<sup>18</sup>F]fluorothymidine-5'-squaryl as a bioisostere of 3'-[<sup>18</sup>F]fluorothymidine-5'-monophosphate

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#### 1.0 Chemistry



**Scheme 1.** Synthesis of compound **1**. *Reaction conditions:* i) thymidine, DMTrCl, pyridine, RT, 16 h; ii) MsCl, 0 °C, 3 h; iii) NaOH (10 N), EtOH, 80 °C, 1.5 h.

Compound 1 was synthesised according to literature procedures.<sup>1,2</sup>





Figure 2. <sup>13</sup>C-NMR spectra of compound 1.



Figure 4. <sup>13</sup>C-NMR spectra of compound 2.





Figure 8. <sup>13</sup>C-NMR spectra of compound 4.



Figure 10. <sup>13</sup>C-NMR spectra of compound 5.



Figure 12. <sup>13</sup>C-NMR spectra of compound 6.



Figure 14. <sup>1</sup>H-NMR spectra of compound 8.



**Figure 16.** DEPT-NMR spectra of compound **8**, showing carbon peaks that would otherwise be hidden by the DMSO solvent peak.





Figure 18. <sup>1</sup>H-NMR spectra of compound 9.

---175.21











<sup>10</sup> 0 -10 -20 -30 -40 -50 -60 -70 -80 -90 -100 -110 -120 -130 -140 -150 -160 -170 -180 -190 -200 -210 **Figure 26.** <sup>19</sup>F-NMR spectra of compound **11**.

# 3.0 Radiochemistry



- 12 Precursor: Compound 6 (12 mg) dissolved in 2-methyl-2-butanol/MeCN (1:1 v/v, 1.2 mL)
- 15 Water Bag
- **17 Dilution Vial:** Water (40 mL)
- 18 19 HLB Plus SPE Cartridge to trap compound [<sup>18</sup>F]7

Figure 27. A schematic representation of the GE FASTLab<sup>™</sup> cassette used to synthesise [<sup>18</sup>F]7

# 4.0 HPLC chromatograms

Analytical HPLC chromatograms were produced using an Agilent 1200 series or a Shimadzu LC-10Ai Prominence instrument connected to a flow-ram detector (Lablogic, Sheffield, UK).

Variable polarity of the radioactive intermediates in the synthesis of [<sup>18</sup>F]SqFLT warranted the use two different analytical columns; the same solvent gradient was used for both columns.

Compound [<sup>18</sup>F]7 was analysed using a Phenomenex Gemini 5  $\mu$ m C18 110A, 150 x 4.6 mm column with solvents A) H<sub>2</sub>O and B) CH<sub>3</sub>CN. Compounds [<sup>18</sup>F]9, [<sup>18</sup>F]10 and [<sup>18</sup>F]SqFLT were analysed using a Phenomenex Luna 5  $\mu$ m Phenyl-Hexyl, 150 x 4.6 mm with solvents A) 0.1% TFA in H<sub>2</sub>O and B) CH<sub>3</sub>CN.

The gradient is described below:

Flow	A (%)	B (%)
mL/min		
1	95	5
1	95	5
1	5	95
1	95	5
1	95	5
	Flow nL/min 1 1 1 1 1	Flow A (%)   mL/min 95   1 95   1 5   1 95   1 5   1 95   1 95   1 95   1 95   1 95   1 95



**Figure 28.** Representative radio-HPLC chromatogram of  $[^{18}F]7$  (t<sub>R</sub> = 10:41 mm:ss) after purification by HLB-SPE.



**Figure 29.** Representative HPLC chromatogram of [<sup>18</sup>F]10: A) radio-HPLC ( $t_R$  = 7:47 mm:ss); B) UV-HPLC (254 nm) spiked with reference standard 10. C) Representative preparative-HPLC chromatogram showing the purification of [<sup>18</sup>F]10 ( $t_R$  = ca 16 min) Red: radioactive counts; Blue: UV<sub>254</sub>.





## 5.0 LogD<sub>7.5</sub> determination

**Table 1.** Calculated LogP (cLogP) values and measured LogD<sub>7.5</sub> for [<sup>18</sup>F]FLT, [<sup>18</sup>F]FLTMP and [<sup>18</sup>F]SqFLT.

	Partition coefficient				
Method	[ <sup>18</sup> F]FLT	[ <sup>18</sup> F]FLTMP	[ <sup>18</sup> F]SqFLT		
LogD <sub>7.5</sub>	ND	 ND	$-2.90 \pm 0.24^{b}$		
cLogP	-0.74	-1.15	-1.20		

ND = not determined.

<sup>a</sup> Calculated using Chemdraw 16.0 (Cambridgesoft, USA)

<sup>b</sup> Performed n = 3 with triplicate measurements, represented as Mean ± SD.

## 6.0 Metabolite analysis in microsomes



**Figure 31.** Representative HPLC chromatograms showing **A**) parent [<sup>18</sup>F]SqFLT, and incubation of [<sup>18</sup>F]SqFLT with HLM for **B**) 30 mins and **C**) 60 mins. The extraction efficiency was 96.1  $\pm$  0.4 %. The experiment was performed in triplicate (n = 3)

# 7.0 Gene structure and positions of CRISPR and PCR primers.



**Figure 32.** Diagram (not to scale) indicates overall gene structure and positions of the CRISPRS and PCR primers. DNA sequence heterogeneity resulting from Cas9 targeting by CRISPR 25926363 – 25926373.

# 8.0 Determining if [<sup>18</sup>F]SqFLT is a substrate for TK1 and ENT1/2



**Figure 33.** Competition assay using pharmacological doses ( $10^{-3} - 10^1 \mu g$ ) of **A**) thymidine and **B**) [<sup>19</sup>F]SqFLT to determine the effect on [<sup>18</sup>F]FLT uptake. HCT116 cells (at 60 – 70 % confluence) were incubated with either thymidine or [<sup>19</sup>F]SqFLT at multiple concentrations for 20 min prior to co-incubation with [<sup>18</sup>F]FLT (0.74 MBq). Radioactive uptake normalised to total protein was plotted against log thymidine/[<sup>19</sup>F]SqFLT concentrations. (n = 5, Mean ± SEM)



9.0 Determining if [<sup>18</sup>F]SqFLT is a substrate for multidrug resistance proteins

**Figure 34.** Uptake of [<sup>18</sup>F]SqFLT in HCT116 cells following treatment with Verapamil. HCT 116 cells were pre-incubated with Verapamil (3  $\mu$ M) for 60 min followed by co-incubated with [<sup>18</sup>F]SqFLT (0.74 MBq). One-way Anova statistical analysis and Dunnets post hoc was performed for n = 5 samples.





**Figure 35.** Time-activity curves derived from region-of-interest analysis of PET imaging data (including detail of uptake kinetics between 0-5 min of [<sup>18</sup>F]SqFLT injection) in bladder, kidneys and gallbladder. Data represent mean ± SEM (n>5) and is expressed as % of injected dose per gram (%ID/g).

#### 11.0 References

- 1 M. Yun, S. J. Oh, H. J. Ha, J. S. Ryu and D. H. Moon, *Nucl. Med. Biol.*, 2003, **30**, 151– 157.
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