

## GW501516

**CAS Registry No.:** 317318-70-0

**Formal Name:** 2-(2-methyl-4-(((4-methyl-2-(4-(trifluoromethyl)phenyl)thiazol-5-yl)methyl)thio)phenoxy)acetic acid

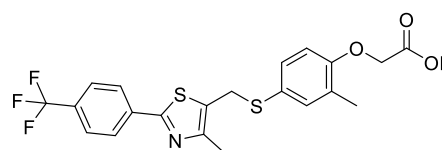
**EUbOPEN ID:** EUB0001148a

**Molecular Formula:** C<sub>21</sub>H<sub>18</sub>F<sub>3</sub>NO<sub>3</sub>S<sub>2</sub>

**Molecular Weight:** 453.49 g/mol

**Smiles:** CC1=C(C=CC(=C1)SCC2=C(N=C(S2)C3=CC=C(C=C3)C(F)(F)F)C)OCC(=O)O

**Recommended concentration:** 1 μM



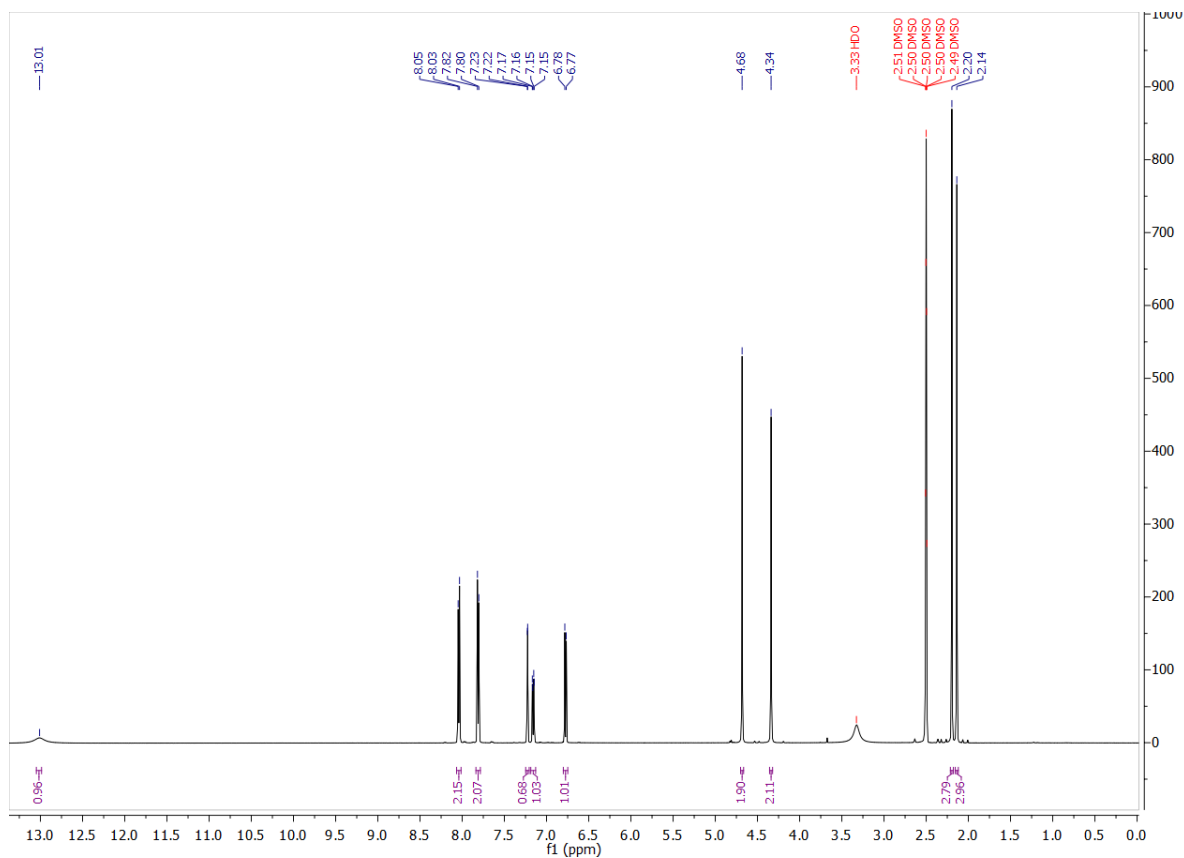
### Biological activity

		Type	IC <sub>50</sub> /EC <sub>50</sub> [μM]	Reference
Main NR target:	NR1C2 (PPARδ)	Agonist	0.001	<a href="https://doi.org/10.1073/pnas.091021198">https://doi.org/10.1073/pnas.091021198</a>
NR off-target:				

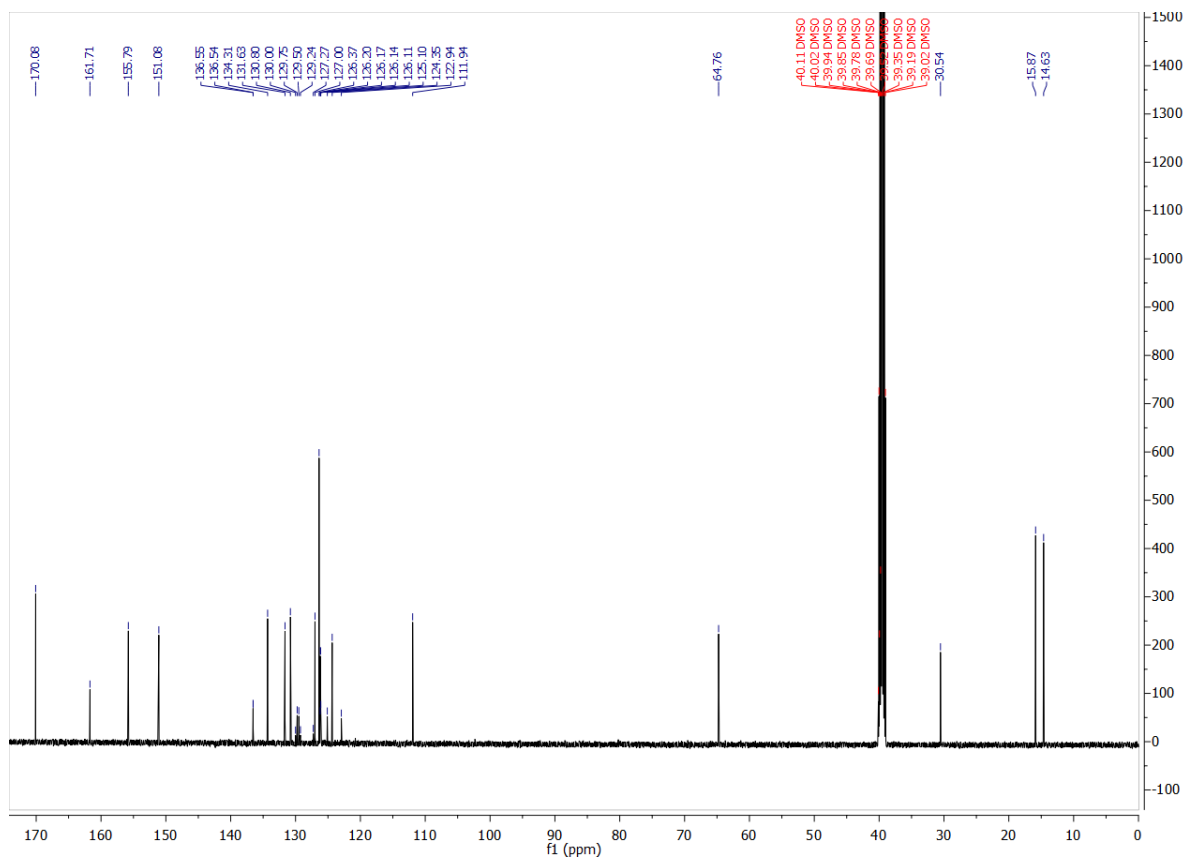
# COMPOUND INFORMATION

## Identity

### <sup>1</sup>H NMR



### <sup>13</sup>C NMR



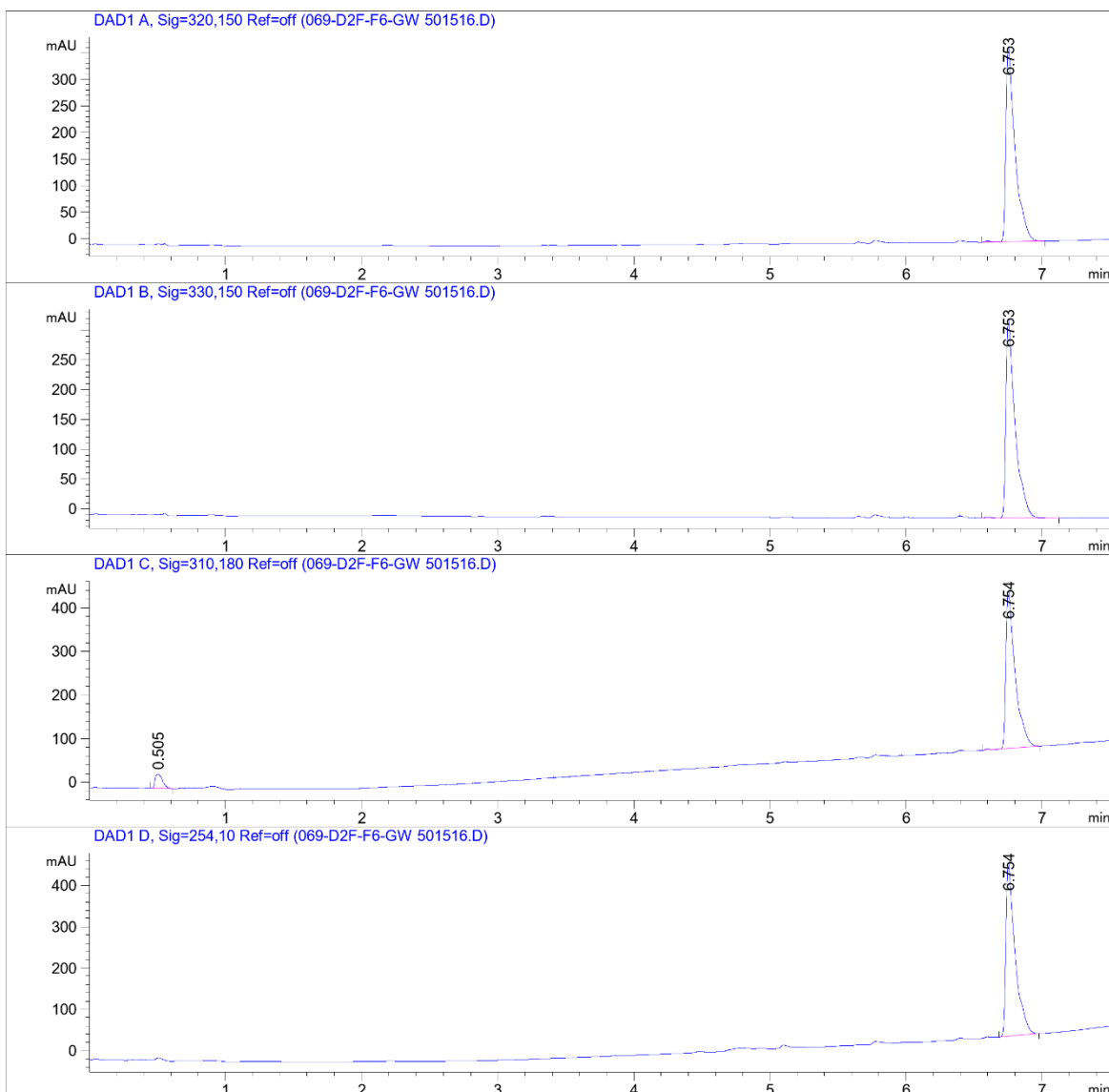
# COMPOUND INFORMATION

## Purity

Data File W:\analyti...\N\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\069-D2F-F6-GW 501516.D

Sample Name: GW 501516

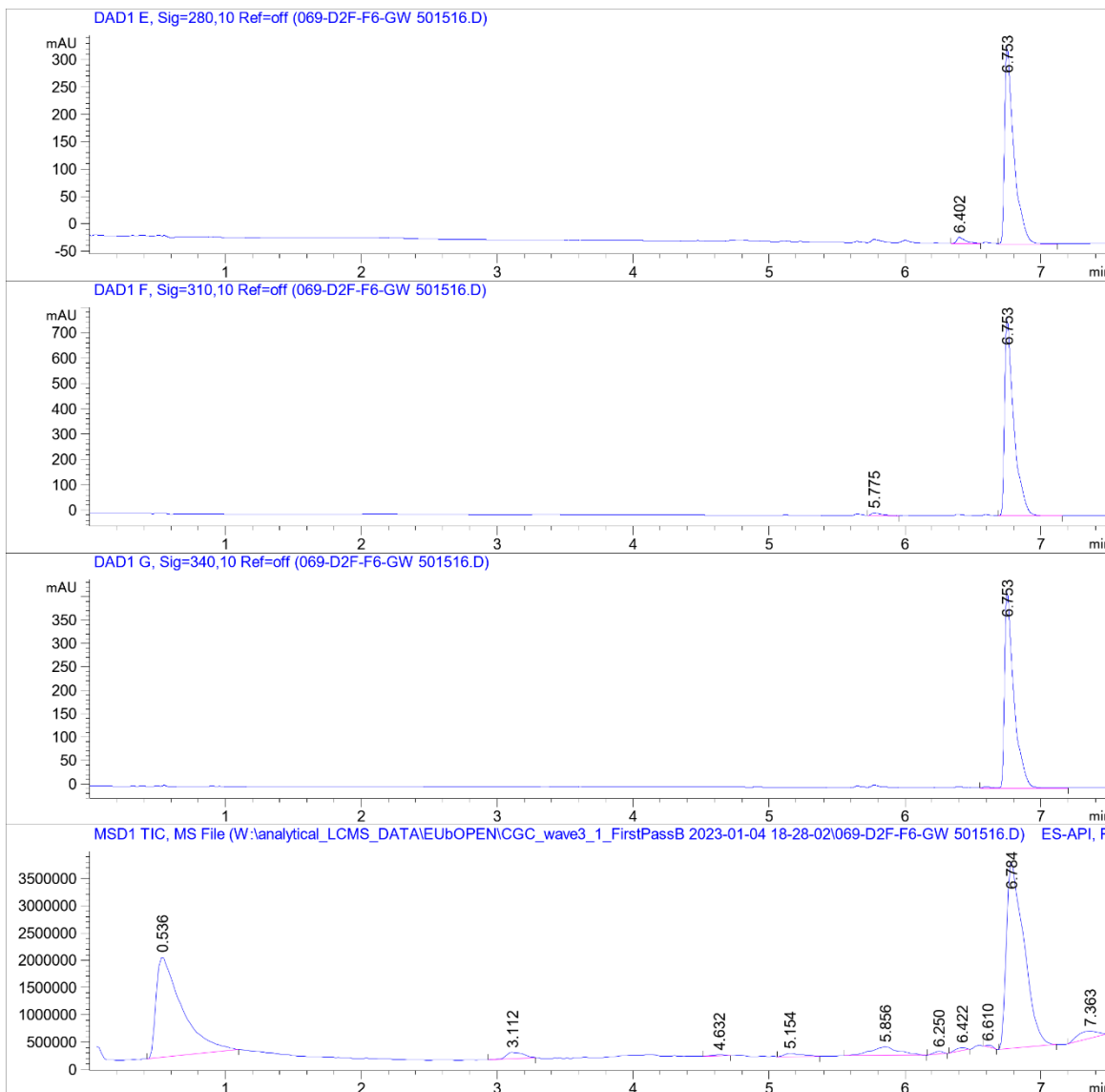
```
=====
Acq. Operator   : SYSTEM                      Seq. Line :   69
Sample Operator : SYSTEM
Acq. Instrument : LCMS test                   Location  : D2F-F6
Injection Date  : 1/5/2023 7:02:46 AM        Inj       :    1
                                           Inj Volume: Inj prog
Sequence File   : W:\analytical_LCMS_DATA\EUBOPEN\CGC_wave3_1_FirstPassB 2023-01-04 18-28-02
                  \CGC_wave3_1_FirstPassB.S
Method          : W:\analytical_LCMS_DATA\EUBOPEN\CGC_wave3_1_FirstPassB 2023-01-04 18-28-02
                  \CGL_FIRSTPASS_GENERALMETHOD_VIAL1+2_20210319.M (Sequence Method)
Last changed    : 1/25/2022 4:36:18 PM by SYSTEM
Method Info     : CGL wellplate, 0.5 uL of 10 mM DMSO, general method
=====
```



# COMPOUND INFORMATION

Data File W:\analyti...N\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\069-D2F-F6-GW 501516.D

Sample Name: GW 501516



# COMPOUND INFORMATION

Data File W:\analyti...\N\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\069-D2F-F6-GW 501516.D

Sample Name: GW 501516

MS Signal: MSD1 TIC, MS File, ES-API, Pos, Scan, Frag: 70, "POS Scan"

Spectra from peak tops.

Noise Cutoff: 1000 counts.

Reportable Ion Abundance: > 50%.

LC Signal: DAD1 A, Sig=320,150 Ref=off

Peak matching window: 0.1 min

Retention Time (LC)	LC Area	Retention Time (MS)	MS Area	Mol. Weight or Ion
-	-	0.536	24070416	157.00 I
-	-	3.112	1019263	239.00 I 217.10 I
-	-	4.632	164508	510.40 I 170.80 I 158.10 I
-	-	5.154	578058	510.30 I 274.00 I 170.90 I 137.10 I
-	-	5.856	2261523	318.20 I 296.20 I
-	-	6.250	216604	228.20 I 137.10 I
-	-	6.422	326089	436.10 I 350.20 I 282.30 I 254.20 I 137.20 I
-	-	6.610	133779	507.30 I 468.00 I 280.20 I
6.753	1709	6.784	30612550	454.10 I
-	-	7.363	1397508	400.30 I 282.30 I

