

# COMPOUND INFORMATION

## SR1001

**CAS Registry No.:** 1335106-03-0

**Formal Name:** N-(5-(N-(4-(1,1,1,3,3,3-hexafluoro-2-hydroxypropan-2-yl)phenyl)sulfamoyl)-4-methylthiazol-2-yl)acetamide

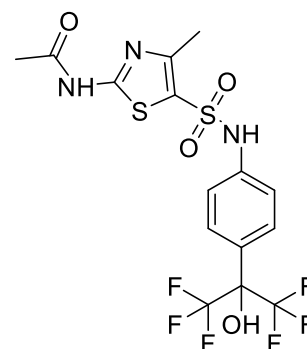
**EUBOPEN ID:** EUB0001158a

**Molecular Formula:** C<sub>15</sub>H<sub>13</sub>F<sub>6</sub>N<sub>3</sub>O<sub>4</sub>S<sub>2</sub>

**Molecular Weight:** 477.40 g/mol

**Smiles:** CC(NC1=NC(C)=C(S1)S(=O)(=O)NC2=CC=C(C(C2)C(F)(F)F)C(F)(F)F)O)=O

**Recommended concentration:** 1  $\mu$ M



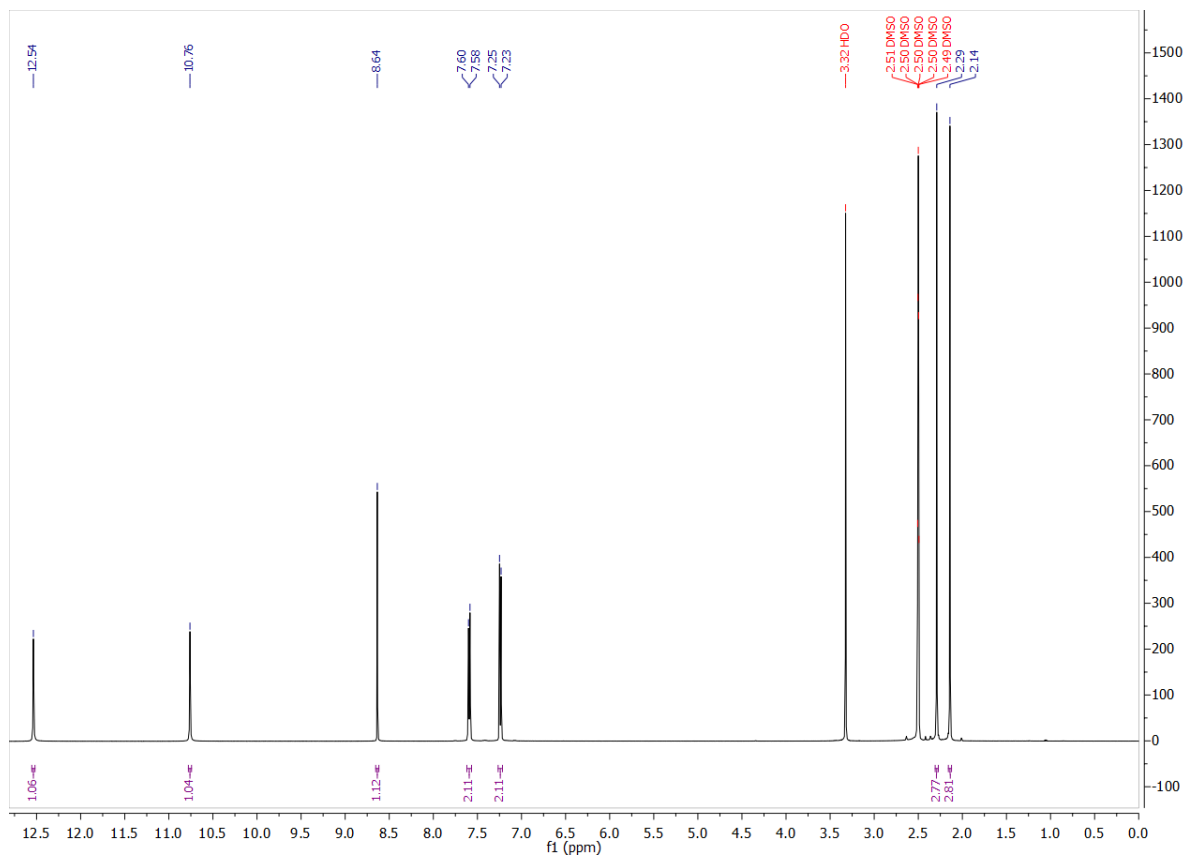
### Biological activity

		Type	IC <sub>50</sub> /EC <sub>50</sub> [ $\mu$ M]	Reference
Main NR target:	NR1F1 (ROR $\alpha$ )	inv. Agonist	0.9	inhouse
	NR1F2 (ROR $\beta$ )	inv. Agonist	1.1	inhouse
	NR1F3 (ROR $\gamma$ )	inv. Agonist	0.11	inhouse
NR off-target:				

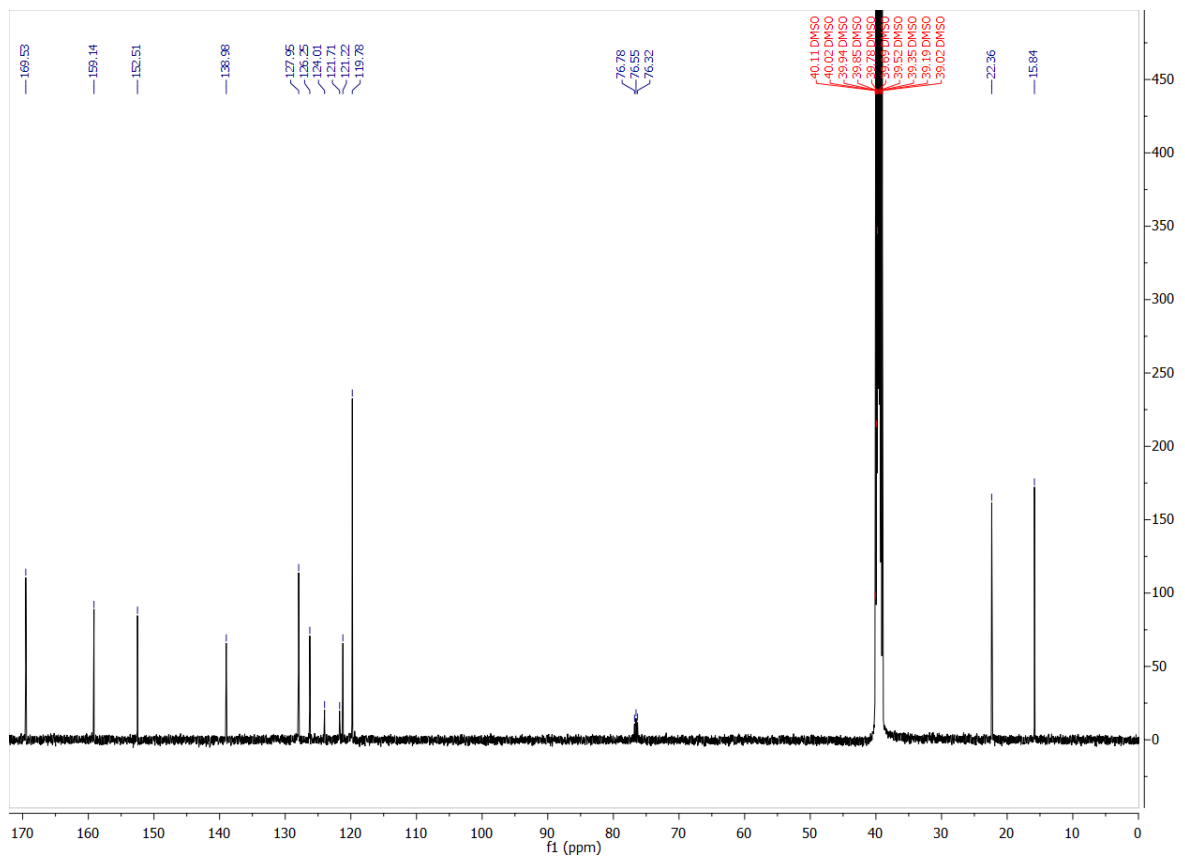
# COMPOUND INFORMATION

## Identity

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



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



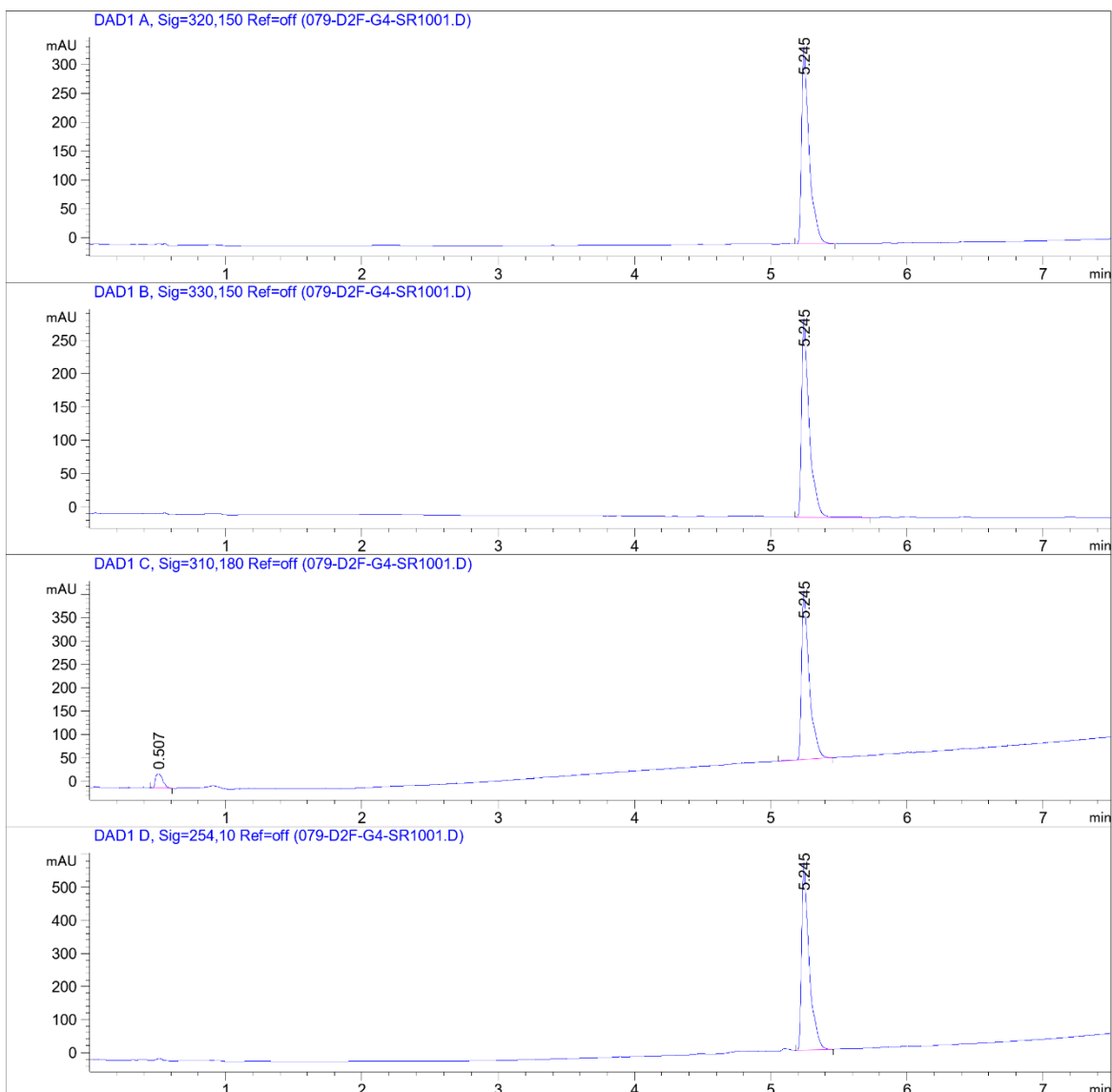
# COMPOUND INFORMATION

## Purity

Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\079-D2F-G4-SR1001.D

Sample Name: SR1001

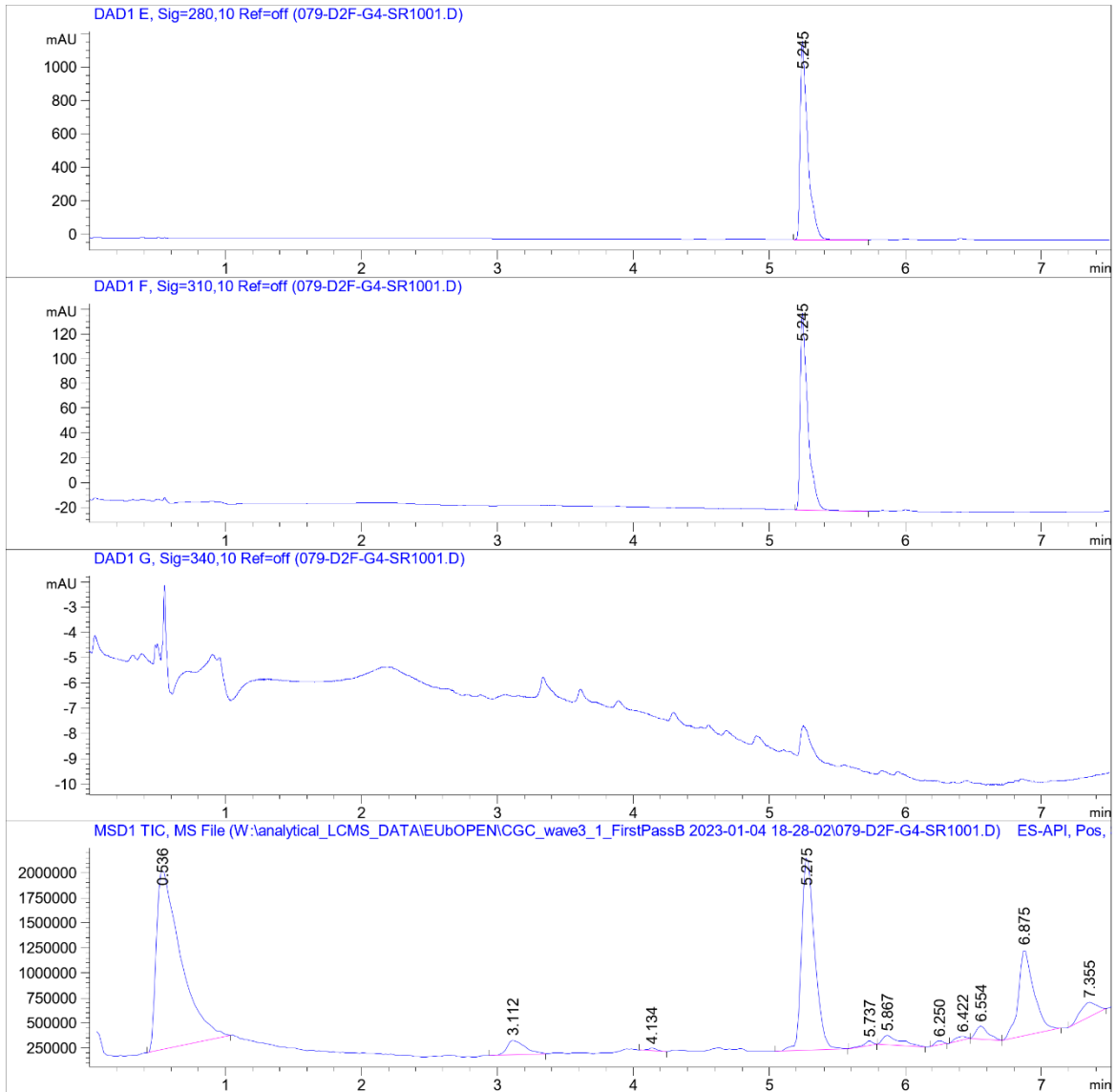
```
=====
Acq. Operator   : SYSTEM                      Seq. Line :   79
Sample Operator : SYSTEM
Acq. Instrument : LCMS test                   Location  : D2F-G4
Injection Date  : 1/5/2023 8:53:21 AM        Inj       :    1
                                           Inj Volume: Inj prog
Sequence File   : 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
=====
```



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Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\079-D2F-G4-SR1001.D

Sample Name: SR1001



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Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\079-D2F-G4-SR1001.D

Sample Name: SR1001

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

Spectra from peak tops.

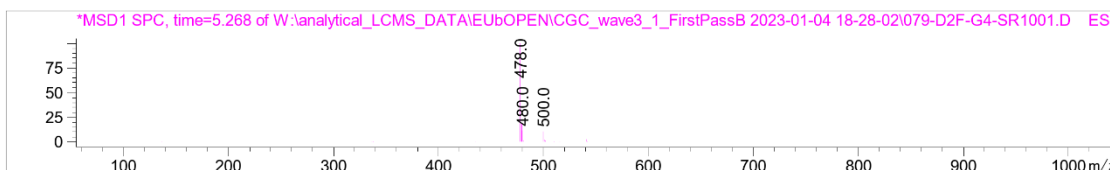
Noise Cutoff: 1000 counts.

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	22368776	157.10 I
-	-	3.112	1362420	239.00 I 217.10 I
-	-	4.134	111615	510.40 I 258.10 I 170.80 I
5.245	1299	5.275	12884487	478.00 I
-	-	5.737	173390	280.20 I
-	-	5.867	851386	318.20 I 296.20 I
-	-	6.250	161489	228.20 I 137.20 I
-	-	6.422	219314	350.20 I 282.20 I 254.20 I 137.10 I
-	-	6.554	787709	507.30 I 485.30 I 280.20 I
-	-	6.875	6840838	282.20 I
-	-	7.355	1244306	400.30 I 282.20 I



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## Biological activity

