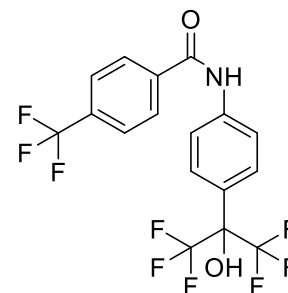


## SR1078

**CAS Registry No.:** 1246525-60-9  
**Formal Name:** N-(4-(1,1,1,3,3,3-hexafluoro-2-hydroxypropan-2-yl)phenyl)-4-(trifluoromethyl)benzamide  
**EUBOPEN ID:** EUB0001156a  
**Molecular Formula:** C<sub>17</sub>H<sub>10</sub>F<sub>9</sub>NO<sub>2</sub>  
**Molecular Weight:** 431.26 g/mol  
**Smiles:** O=C(C1=CC=C(C=C1)C(F)(F)F)NC2=CC=C(C=C2)C(C(F)(F)F)(C(F)(F)F)O  
**Recommended concentration:** 1 μM

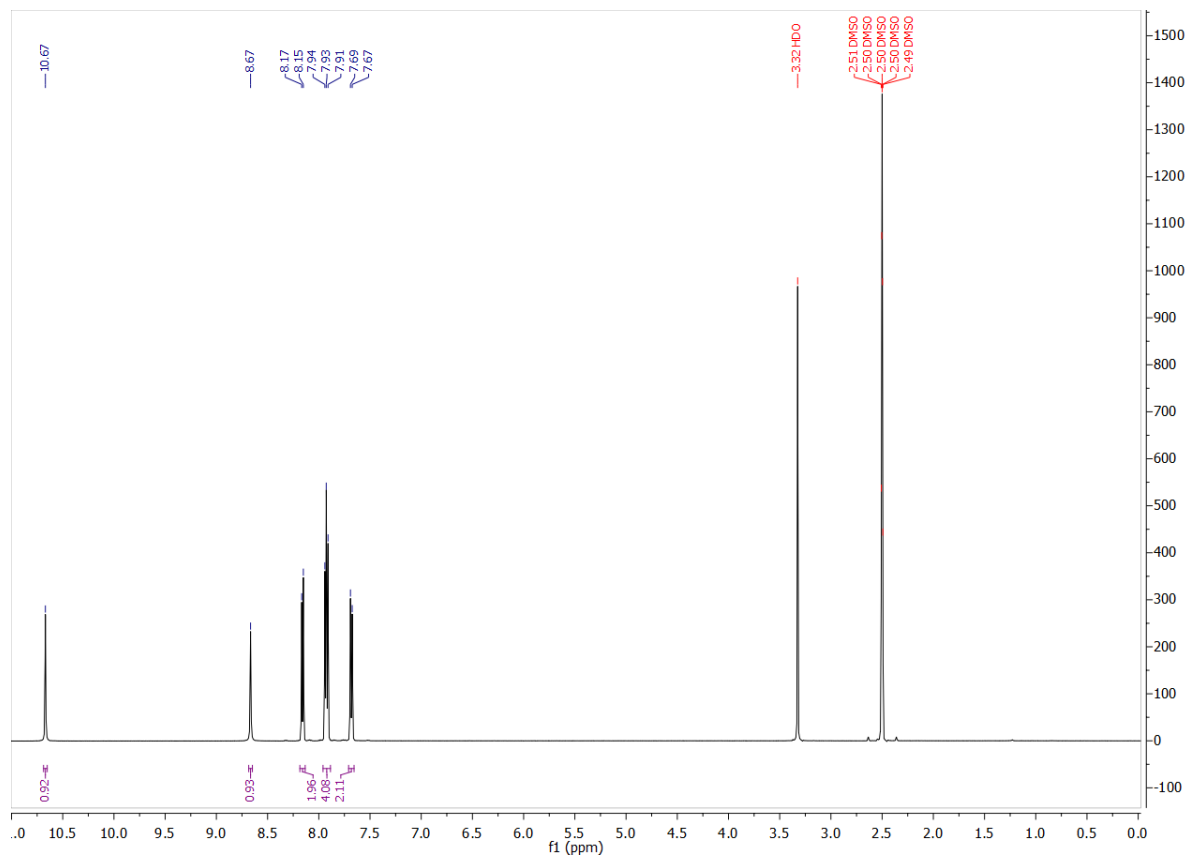


### Biological activity

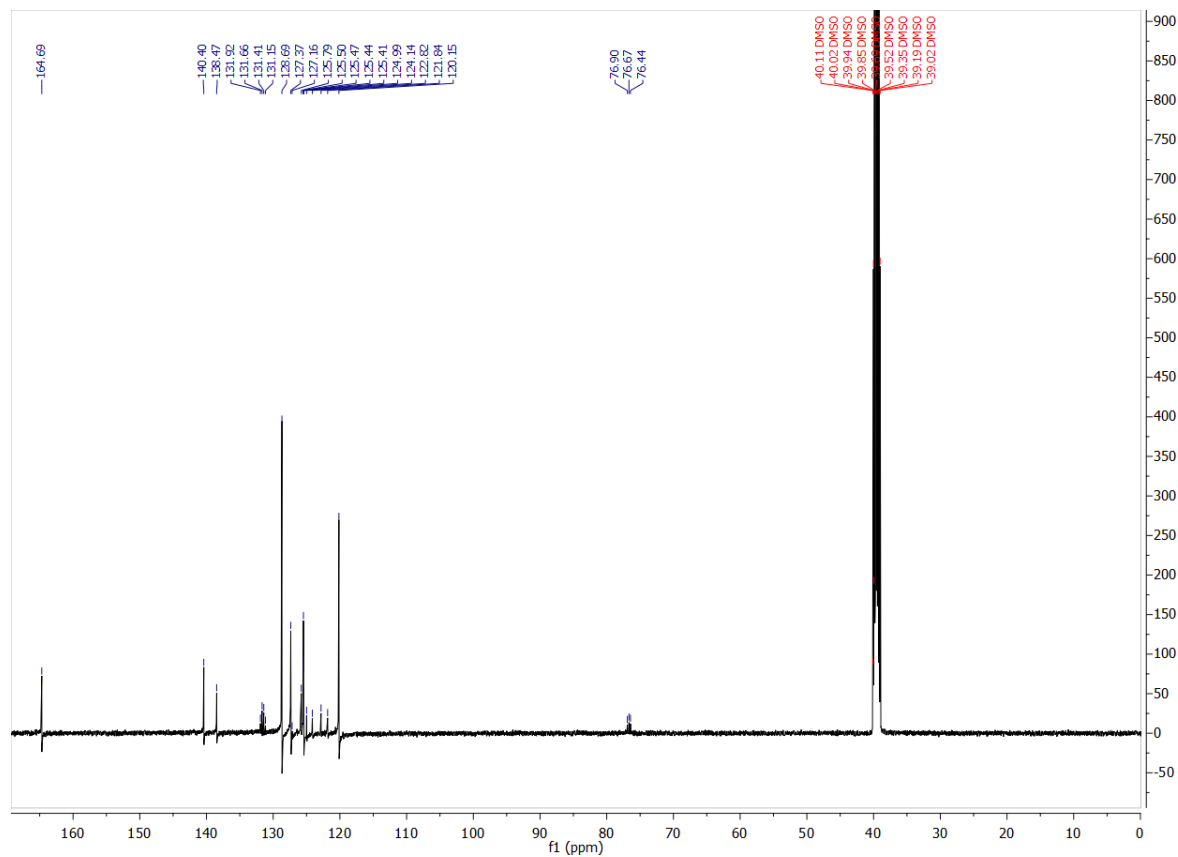
		Type	IC <sub>50</sub> /EC <sub>50</sub> [μM]	Reference
Main NR target:	NR1F3 (RORγ)	inv. Agonist	2	<a href="https://doi.org/10.1021/cb100223d">https://doi.org/10.1021/cb100223d</a>
NR off-target:				

## Identity

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



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



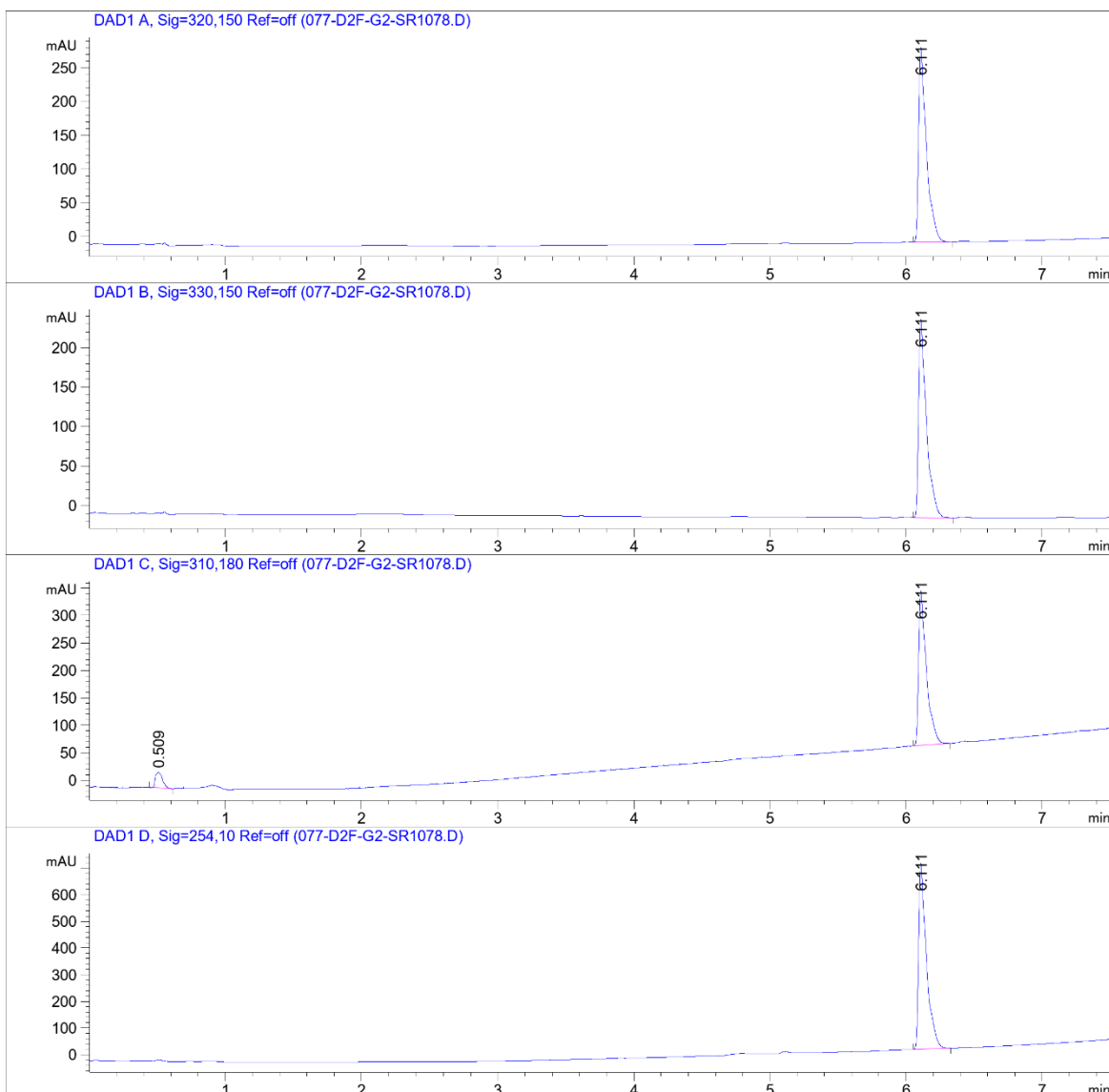
# COMPOUND INFORMATION

## Purity

Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\077-D2F-G2-SR1078.D

Sample Name: SR1078

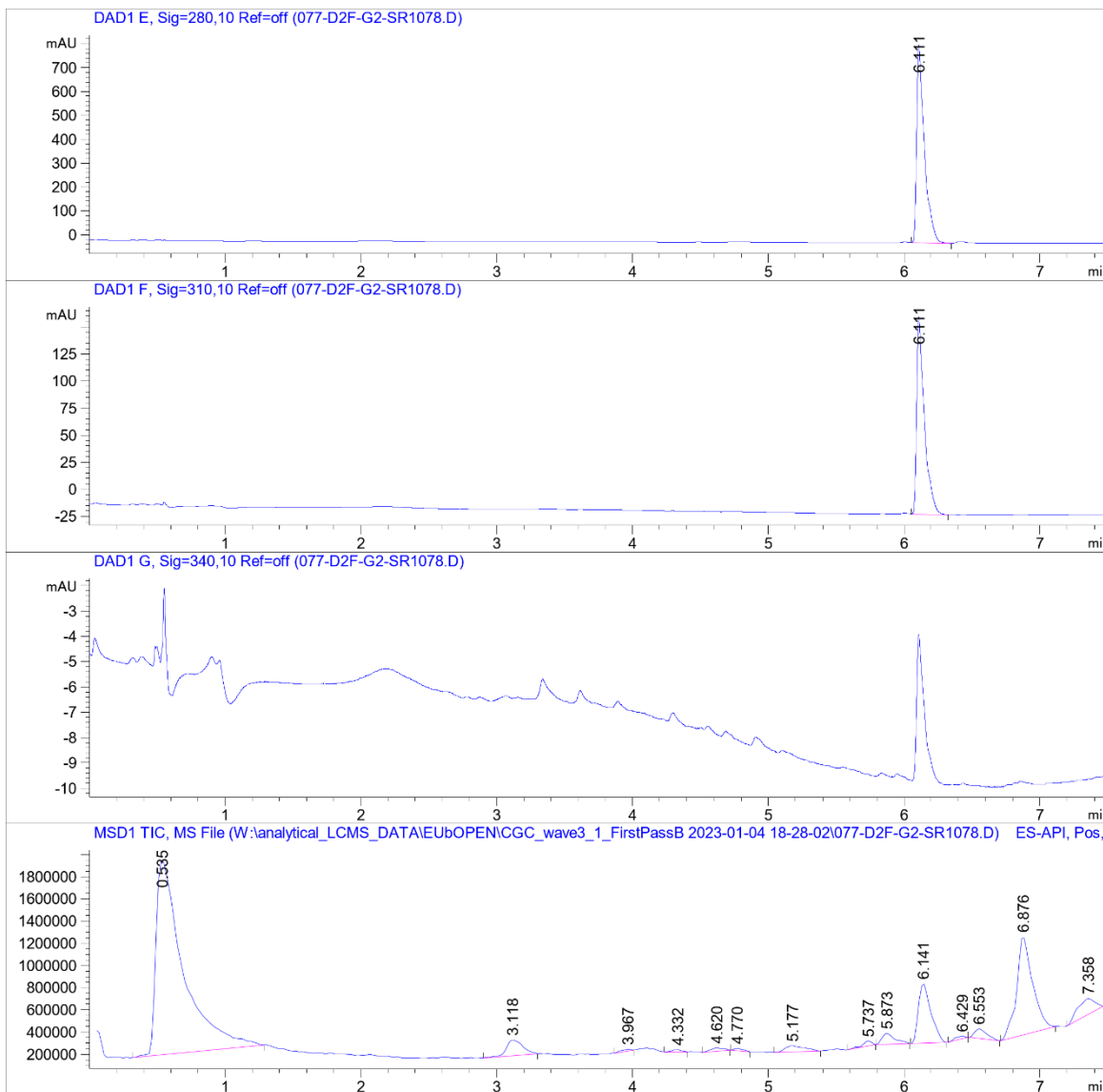
```
=====
Acq. Operator   : SYSTEM                      Seq. Line :   77
Sample Operator : SYSTEM
Acq. Instrument : LCMS test                   Location  : D2F-G2
Injection Date  : 1/5/2023 8:31:14 AM        Inj       :    1
                                           Inj Volume: Inj prog
Sequence File   : 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...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\077-D2F-G2-SR1078.D

Sample Name: SR1078



# COMPOUND INFORMATION

Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\077-D2F-G2-SR1078.D

Sample Name: SR1078

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
-	-	0.535	24433608	157.00 I
-	-	3.118	1196035	239.00 I 217.10 I
-	-	3.967	94859	216.10 I 200.00 I 170.90 I
-	-	4.332	98138	510.40 I 170.90 I
-	-	4.620	184336	510.30 I 170.90 I 158.20 I 137.10 I
-	-	4.770	102449	510.30 I 279.10 I 170.80 I 137.10 I
-	-	5.177	552506	510.30 I 338.20 I 170.80 I 137.10 I
-	-	5.737	179543	280.20 I
-	-	5.873	625801	318.20 I 296.20 I
6.111	1138	6.141	3386224	432.00 I
-	-	6.429	134421	350.20 I 282.20 I 254.20 I 137.20 I
-	-	6.553	539122	507.30 I 280.20 I
-	-	6.876	7225335	282.30 I
-	-	7.358	1309224	400.30 I 282.20 I

