

# COMPOUND INFORMATION

## SR2211

**CAS Registry No.:** 1359164-11-6

**Formal Name:** 1,1,1,3,3,3-hexafluoro-2-(2-fluoro-4'-((4-(pyridin-4-ylmethyl)piperazin-1-yl)methyl)-[1,1'-biphenyl]-4-yl)propan-2-ol

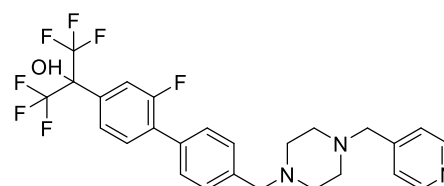
**EUBOPEN ID:** EUB0001160a

**Molecular Formula:** C<sub>26</sub>H<sub>24</sub>F<sub>7</sub>N<sub>3</sub>O

**Molecular Weight:** 527.49 g/mol

**Smiles:** FC1=C(C2=CC=C(CN3CCN(CC4=CC=NC=C4)CC3)C=C2)C=CC(C(F)(F)F)(O)C(F)(F)F=C1

**Recommended concentration:** 1 μM

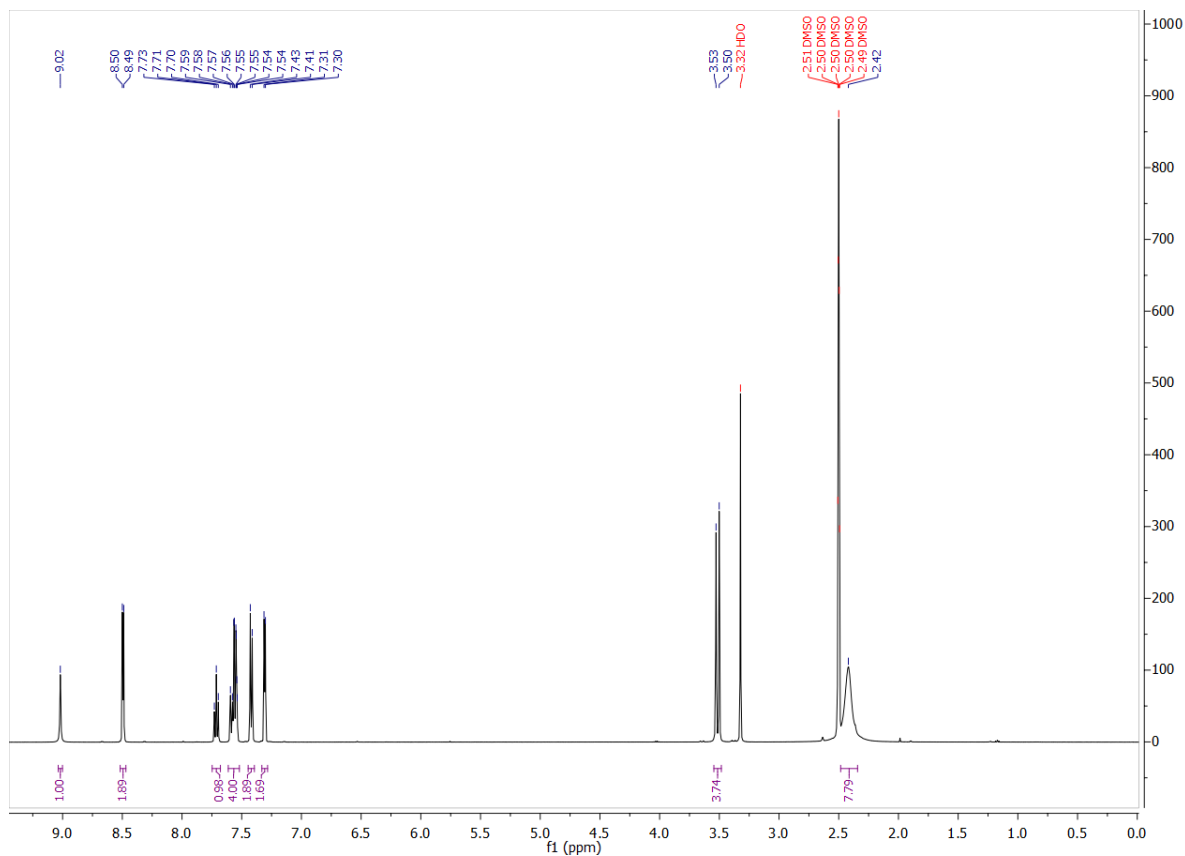


### Biological activity

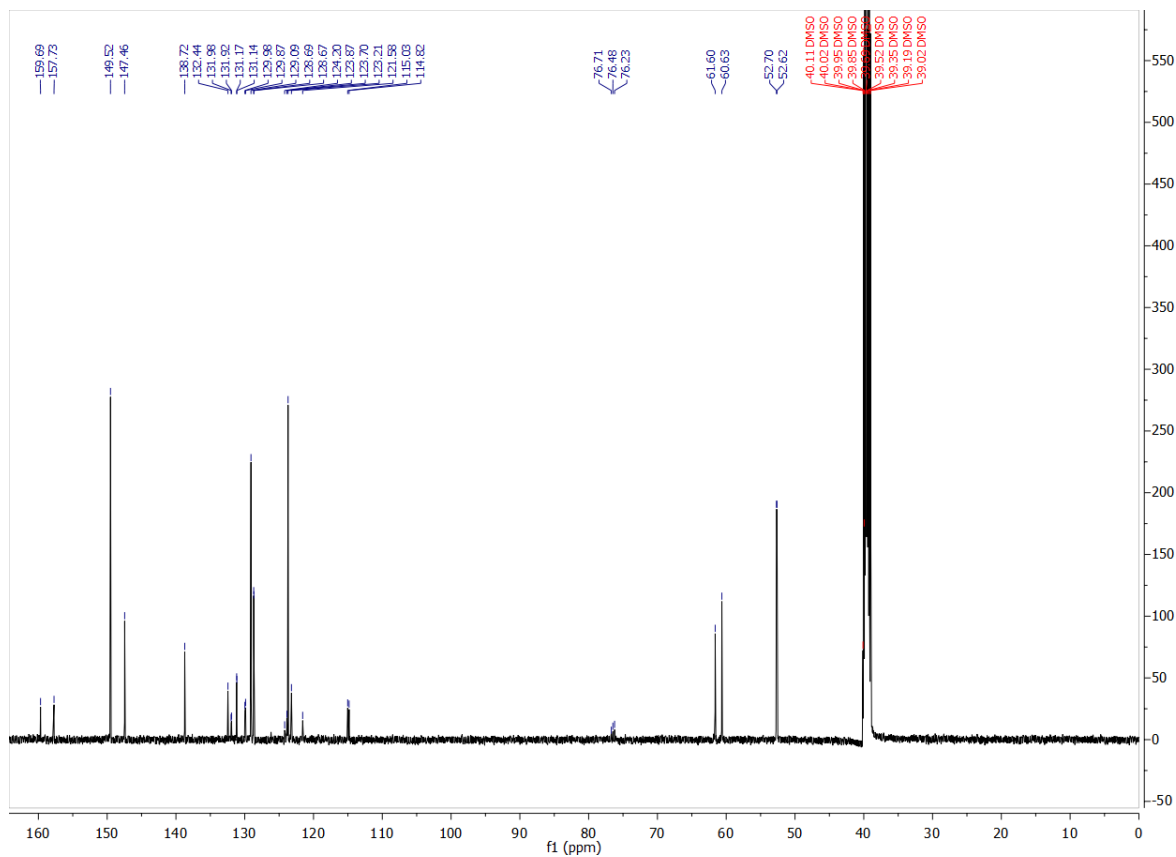
		Type	IC <sub>50</sub> /EC <sub>50</sub> [μM]	Reference
Main NR target:	NR1F3 (RORγ)	inv. Agonist	0.3	<a href="https://doi.org/10.1021/cb200496y">https://doi.org/10.1021/cb200496y</a>
NR off-target:				

## Identity

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



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



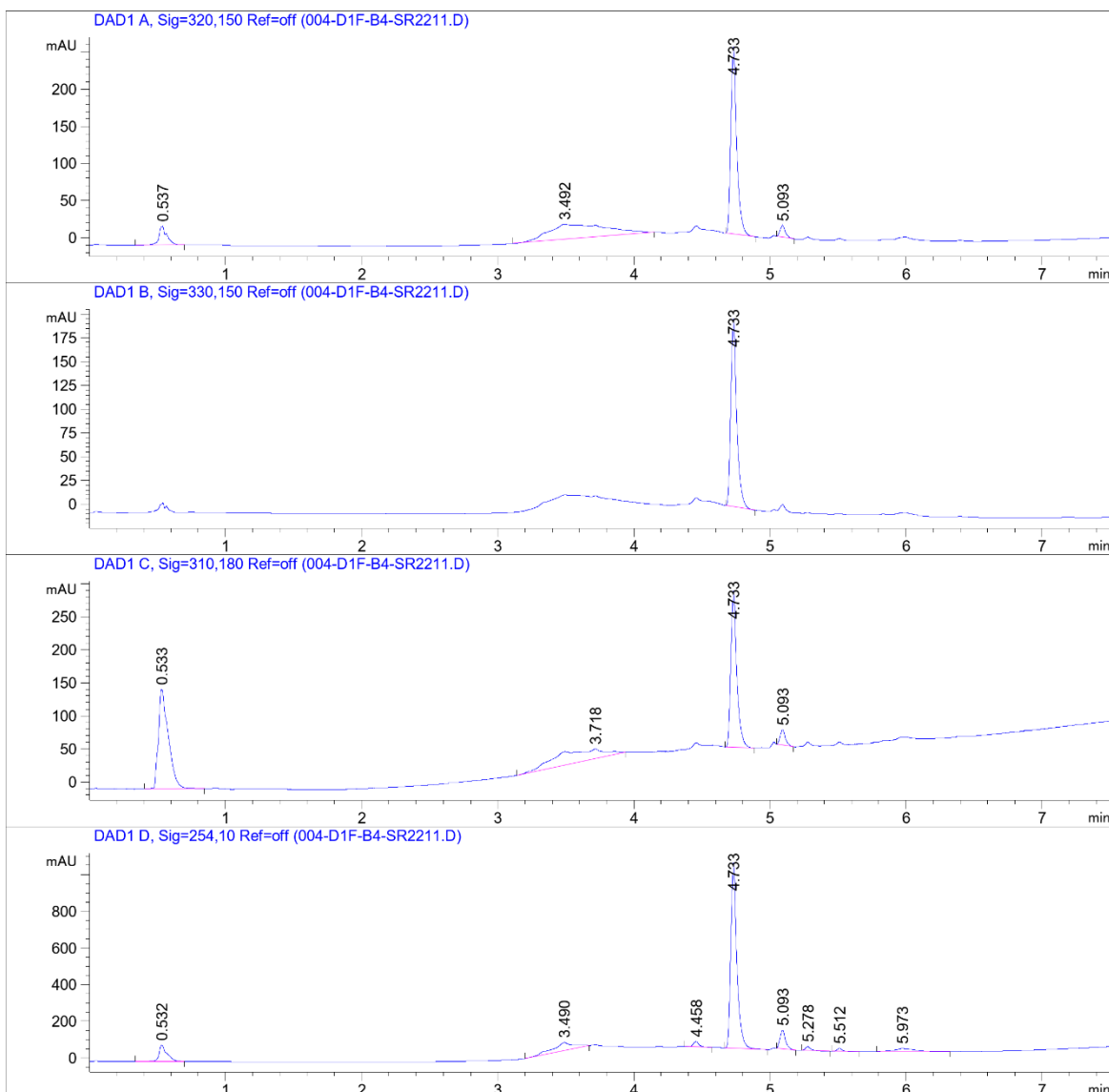
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## Purity

Data File W:\analyti...OPEN\CGC\_wave3\_3rdrun\_vials 2023-04-14 23-47-17\004-D1F-B4-SR2211.D

Sample Name: SR2211

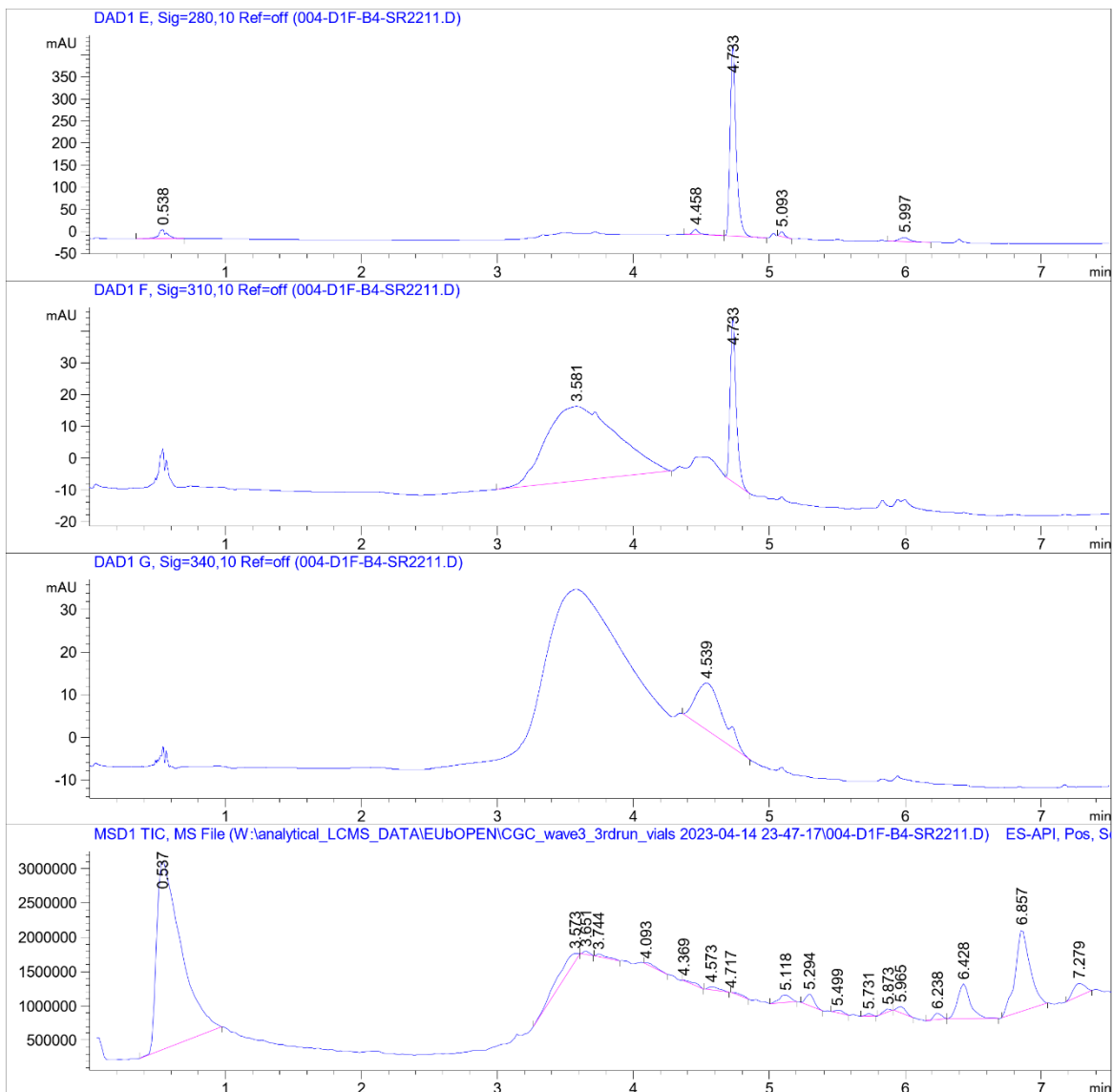
```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    4
Sample Operator : SYSTEM
Acq. Instrument : LCMS test                   Location  : D1F-B4
Injection Date  : 4/15/2023 12:24:54 AM      Inj       :    1
                                           Inj Volume: Inj prog
Sequence File   : W:\analytical_LCMS_DATA\EUBOPEN\CGC_wave3_3rdrun_vials 2023-04-14 23-47-17
                  \CGC_wave3_3rdrun_vials.S
Method          : W:\analytical_LCMS_DATA\EUBOPEN\CGC_wave3_3rdrun_vials 2023-04-14 23-47-17
                  \CGL_FIRSTPASS_GENERALMETHOD_VIAL1+2_VIAL_20220414.M (Sequence Method)
Last changed    : 4/14/2023 5:08:13 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\_3rdrun\_vials 2023-04-14 23-47-17\004-D1F-B4-SR2211.D

Sample Name: SR2211



# COMPOUND INFORMATION



Data File W:\analyti...OPEN\CGC\_wave3\_3rdrun\_vials 2023-04-14 23-47-17\004-D1F-B4-SR2211.D

Sample Name: SR2211

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
0.537	104	0.537	33914356	157.00 I
3.492	593	-	-	
-	-	3.573	2327656	170.80 I
-	-	3.651	162640	170.80 I
-	-	3.744	235333	170.80 I
-	-	4.093	197062	170.80 I
-	-	4.369	213335	170.80 I
-	-	4.573	273748	528.20 I 351.00 I 264.60 I 178.10 I 170.80 I
4.733	763	4.717	127689	528.20 I 351.00 I 264.60 I 178.10 I 170.80 I
5.093	43	5.118	606200	528.20 I 351.00 I 264.60 I 178.10 I
-	-	5.294	698966	528.20 I 398.20 I 376.20 I 351.00 I 264.60 I 178.10 I
-	-	5.499	181000	528.20 I 351.00 I 264.60 I 178.10 I
-	-	5.731	112807	528.10 I 351.00 I 280.20 I 264.60 I 178.10 I
-	-	5.873	152922	318.20 I 296.20 I
-	-	5.965	390011	528.20 I 282.20 I
-	-	6.238	359780	280.20 I 228.20 I

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-	-	6.428	3248994	350.20 I
				282.20 I
-	-	6.857	9042897	282.20 I
-	-	7.279	1142228	400.30 I
				284.30 I
				282.20 I

