

# COMPOUND INFORMATION

## ML-209

**CAS Registry No.:** 1334526-14-5

**Formal Name:** 3-(benzo[d][1,3]dioxol-5-yl)-1-((3S,5R)-3,5-dimethylpiperidin-1-yl)-3-(2-hydroxy-4,6-dimethoxyphenyl)propan-1-one

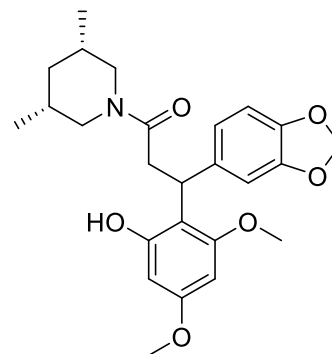
**EUBOPEN ID:** EUB0001161a

**Molecular Formula:** C<sub>25</sub>H<sub>31</sub>NO<sub>6</sub>

**Molecular Weight:** 441.52 g/mol

**Smiles:** COC1=CC(OC)=C(C(C(C(N2C[C@@H](C)C[C@@H](C)C2)=O)C3=C(C=C4OCOC4=C3)C(O)=C1

**Recommended concentration:** 1 & 10 μM



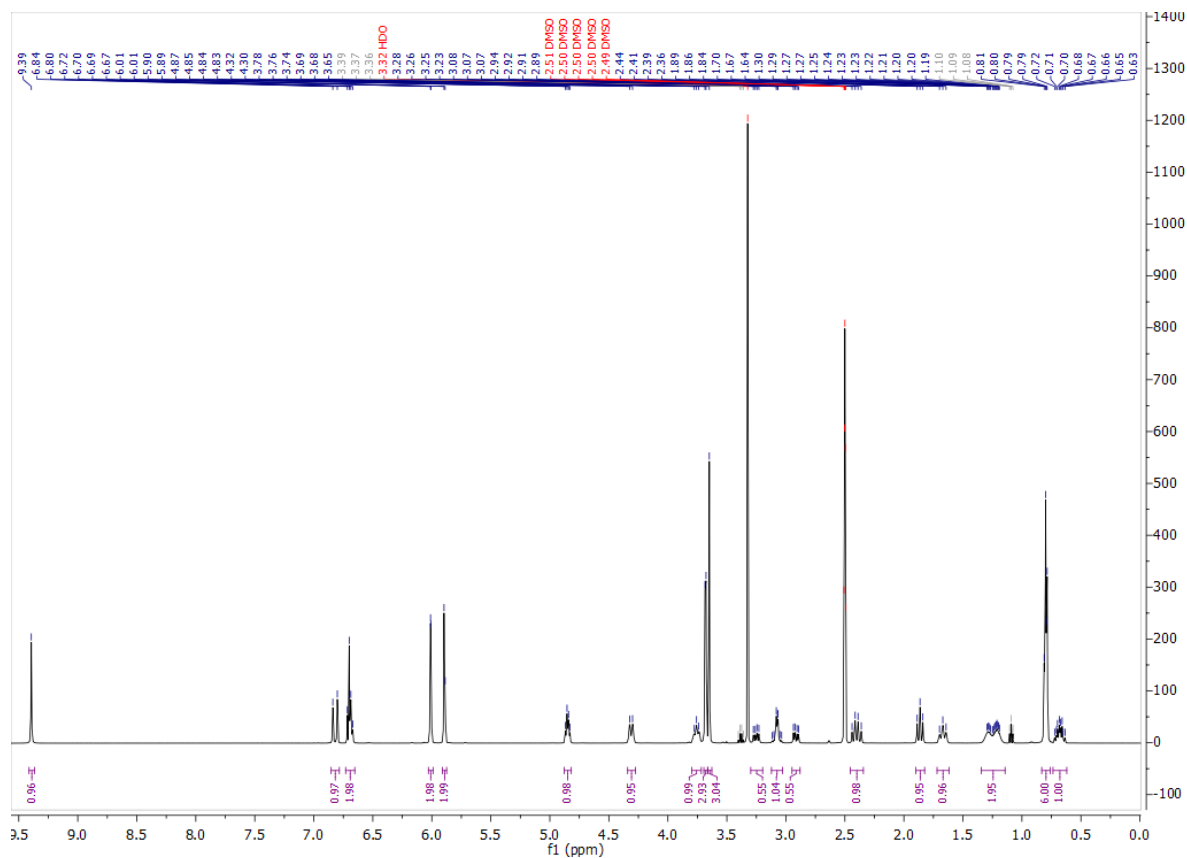
### Biological activity

		Type	IC <sub>50</sub> /EC <sub>50</sub> [μM]	Reference
Main NR target:	NR1F2 (RORβ)	inv. Agonist	1.3	inhouse
	NR1F3 (RORγ)	inv. Agonist	0.20	inhouse
NR off-target @ 10 μM:	NR1F1 (RORα)	inv. Agonist	84% rem. act.	inhouse

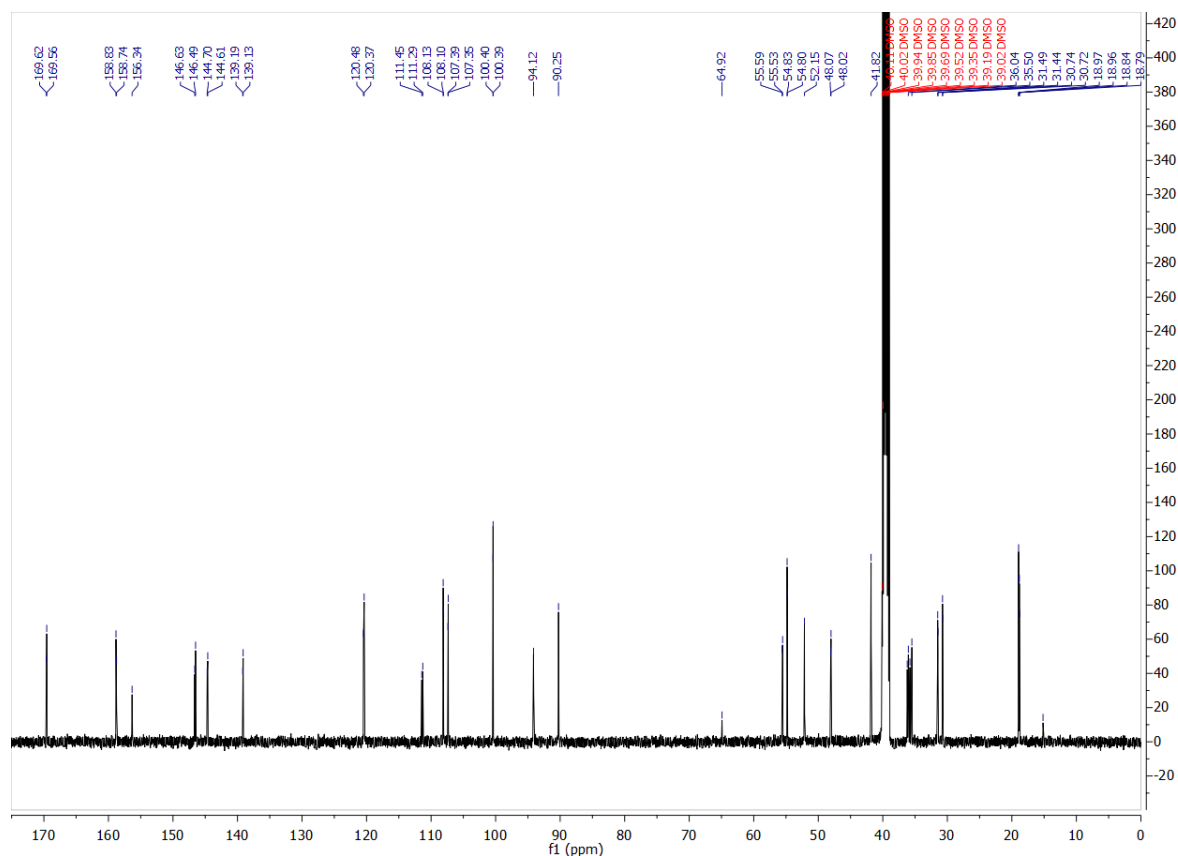
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## Identity

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



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



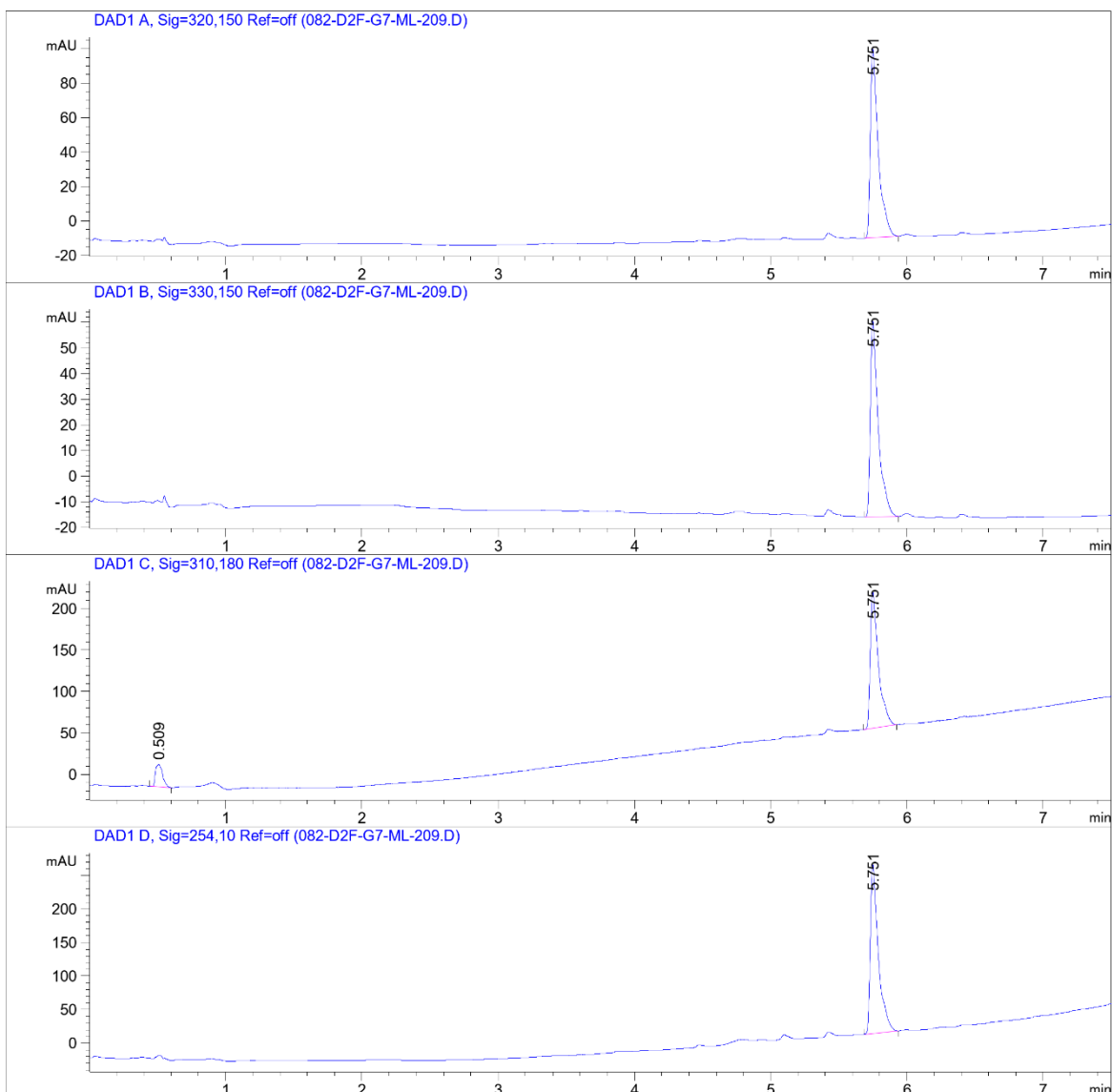
# COMPOUND INFORMATION

## Purity

Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\082-D2F-G7-ML-209.D

Sample Name: ML-209

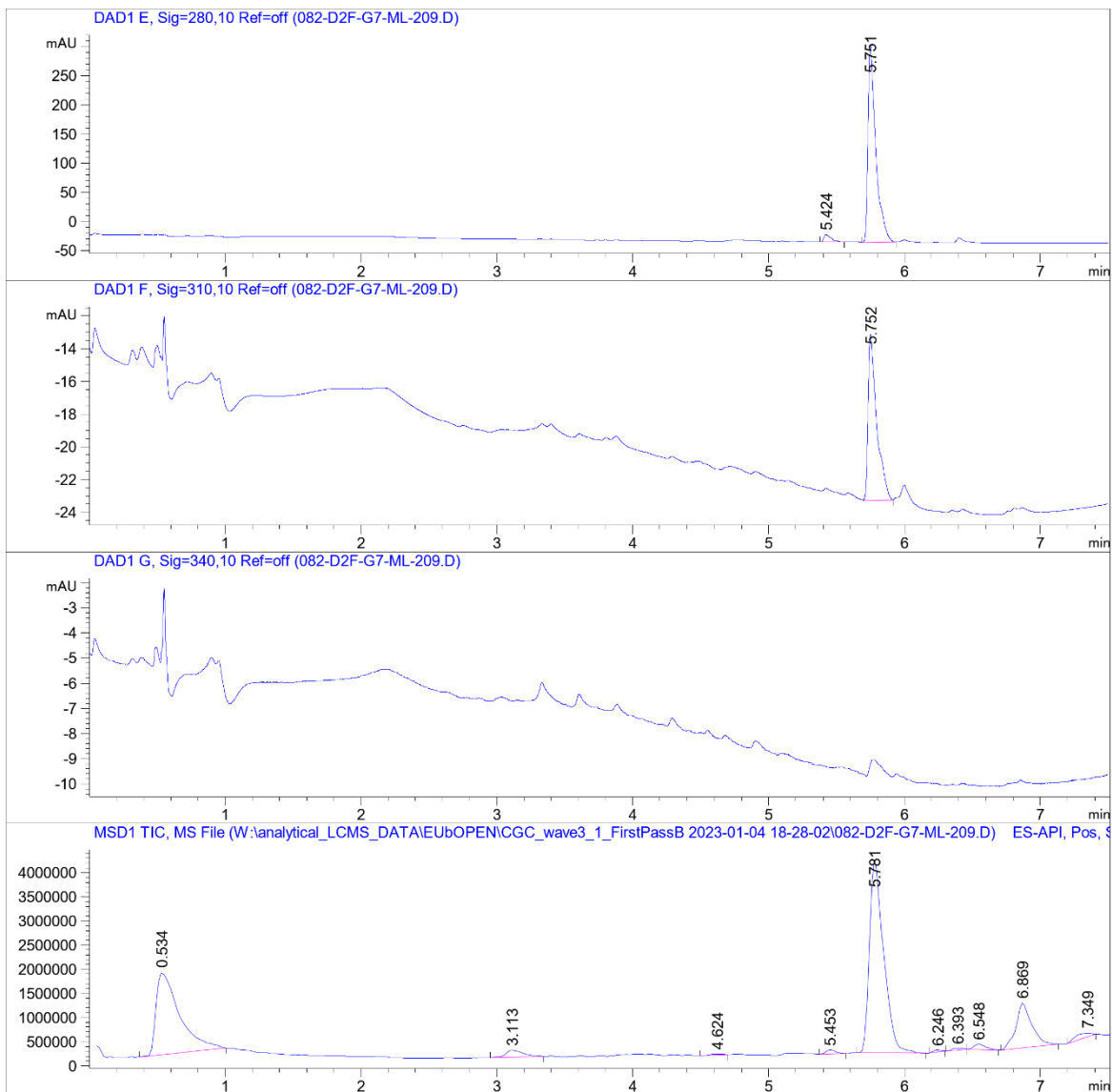
```
=====
Acq. Operator   : SYSTEM                      Seq. Line :   82
Sample Operator : SYSTEM
Acq. Instrument : LCMS test                   Location  : D2F-G7
Injection Date  : 1/5/2023 9:26:32 AM         Inj       :    1
                                           Inj Volume: Inj prog
Sequence File   : 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\082-D2F-G7-ML-209.D

Sample Name: ML-209



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Data File W:\analyti...OPEN\CGC\_wave3\_1\_FirstPassB 2023-01-04 18-28-02\082-D2F-G7-ML-209.D

Sample Name: ML-209

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

Spectra from peak tops.

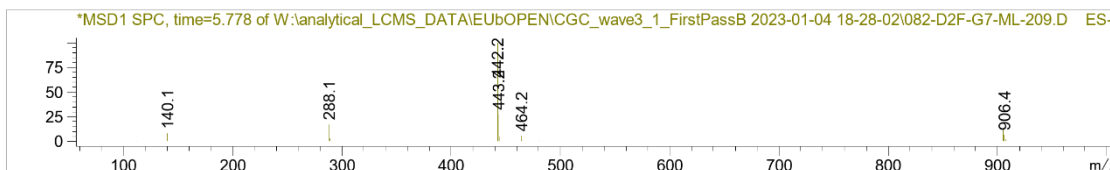
Noise Cutoff: 1000 counts.

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
-	-	0.534	20281808	157.00 I
-	-	3.113	1294489	239.10 I 217.10 I
-	-	4.624	134280	510.30 I 170.90 I 158.10 I 137.10 I
-	-	5.453	424767	510.30 I 351.00 I 329.10 I 207.00 I 137.10 I
5.751	441	5.781	28893416	442.20 I
-	-	6.246	143603	228.20 I 137.10 I
-	-	6.393	195335	254.20 I
-	-	6.548	657064	507.30 I 280.20 I
-	-	6.869	7581119	282.30 I
-	-	7.349	862882	400.30 I 282.20 I



## Biological activity

