

## SR12813

**CAS Registry No.:** 126411-39-0

**Formal Name:** Tetraethyl (2-(3,5-di-tert-butyl-4-hydroxyphenyl)ethene-1,1-diyl)bis(phosphonate)

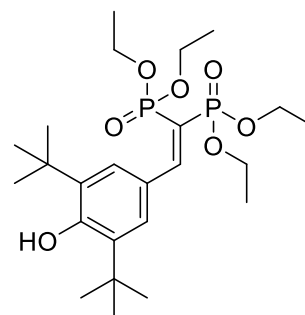
**EUBOPEN ID:** EUB0000576a

**Molecular Formula:** C<sub>24</sub>H<sub>42</sub>O<sub>7</sub>P<sub>2</sub>

**Molecular Weight:** 504.54 g/mol

**Smiles:** CCOP(=O)(OCC)C(=O)C(=C1C(C(C)C)=C(O)C(C(C)C)=C1)OCC=O

**Recommended concentration:** 1 μM



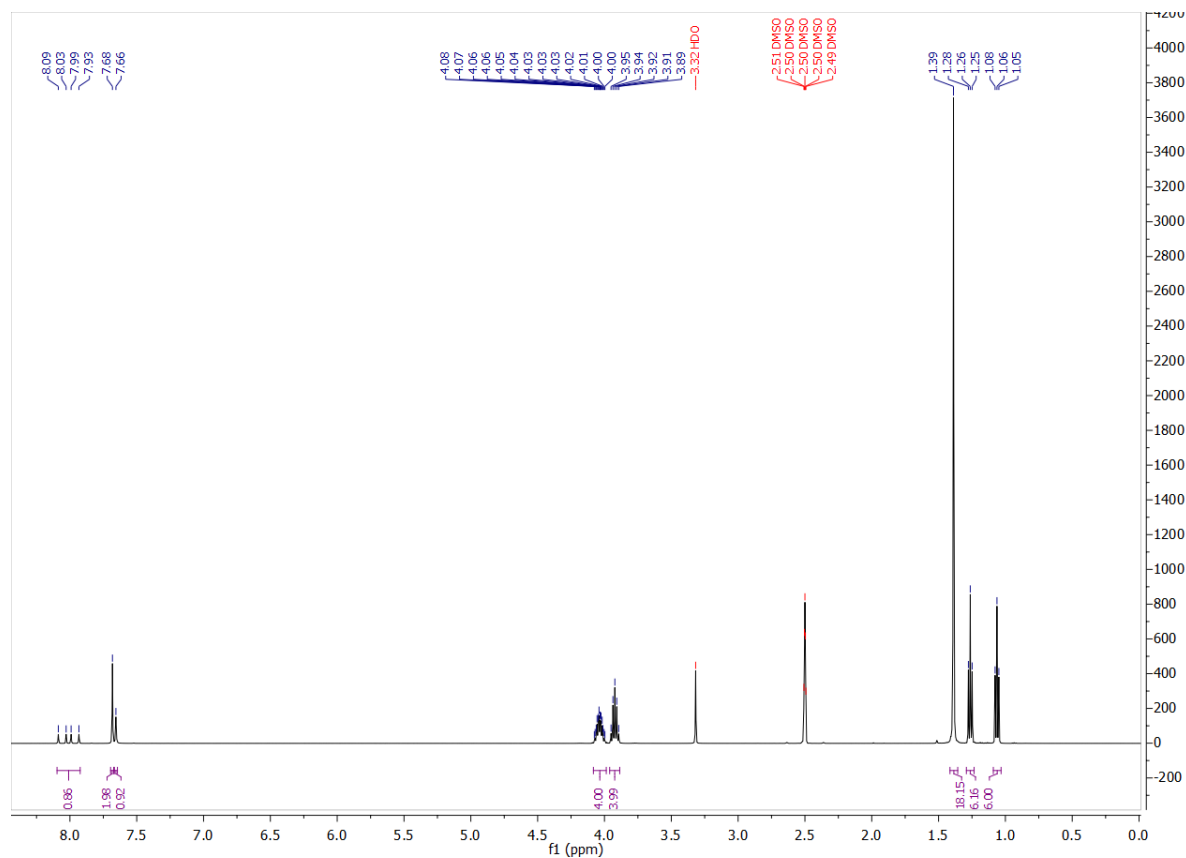
### Biological activity

		Type	IC <sub>50</sub> /EC <sub>50</sub> [μM]	Reference
Main NR target:	NR112 (PXR)	Agonist	0.14	<a href="https://doi.org/10.1124/mol.106.033415">https://doi.org/10.1124/mol.106.033415</a>
NR off-target:				

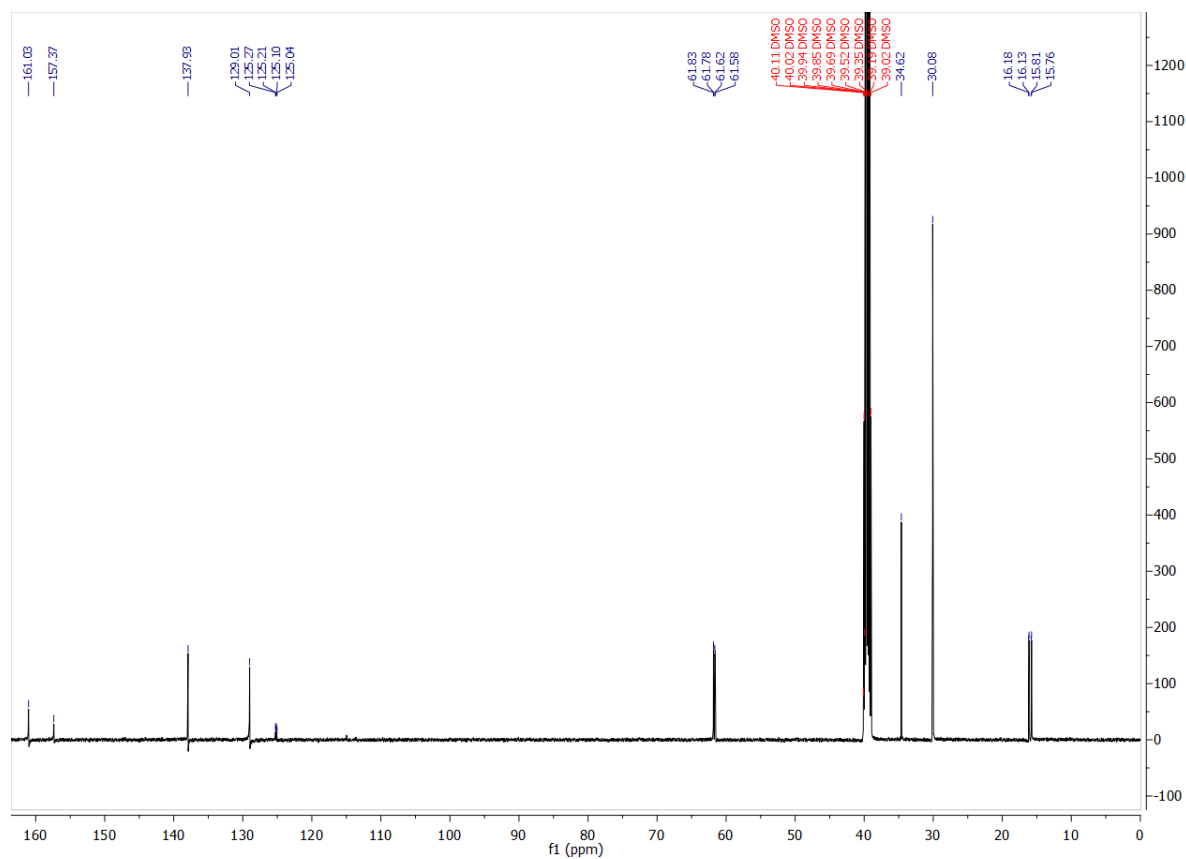
# COMPOUND INFORMATION

## Identity

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



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



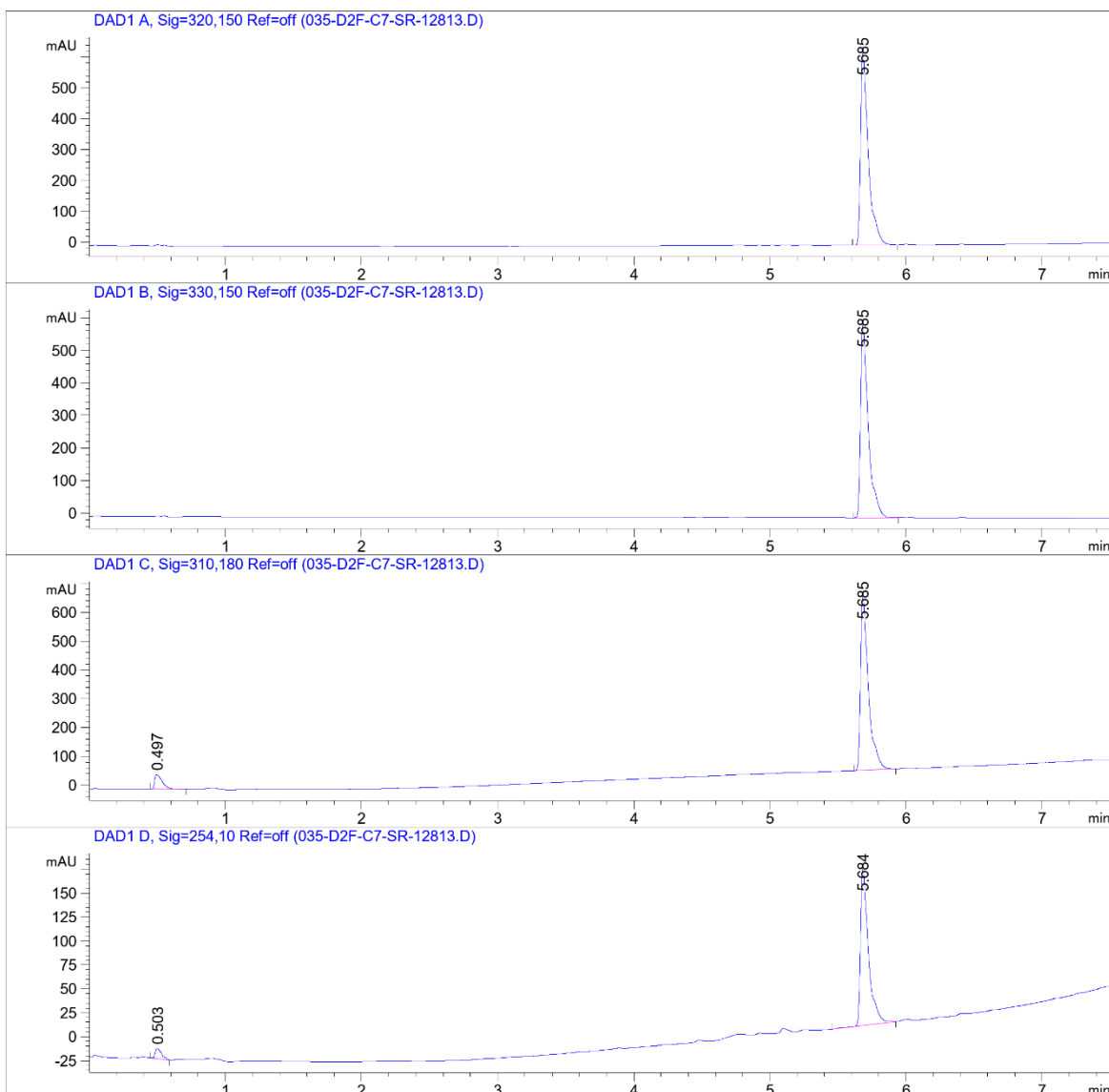
# COMPOUND INFORMATION

## Purity

Data File W:\analyti...PEN\CGC\_wave3\_2\_FirstPass 2023-01-05 22-10-41\035-D2F-C7-SR-12813.D

Sample Name: SR-12813

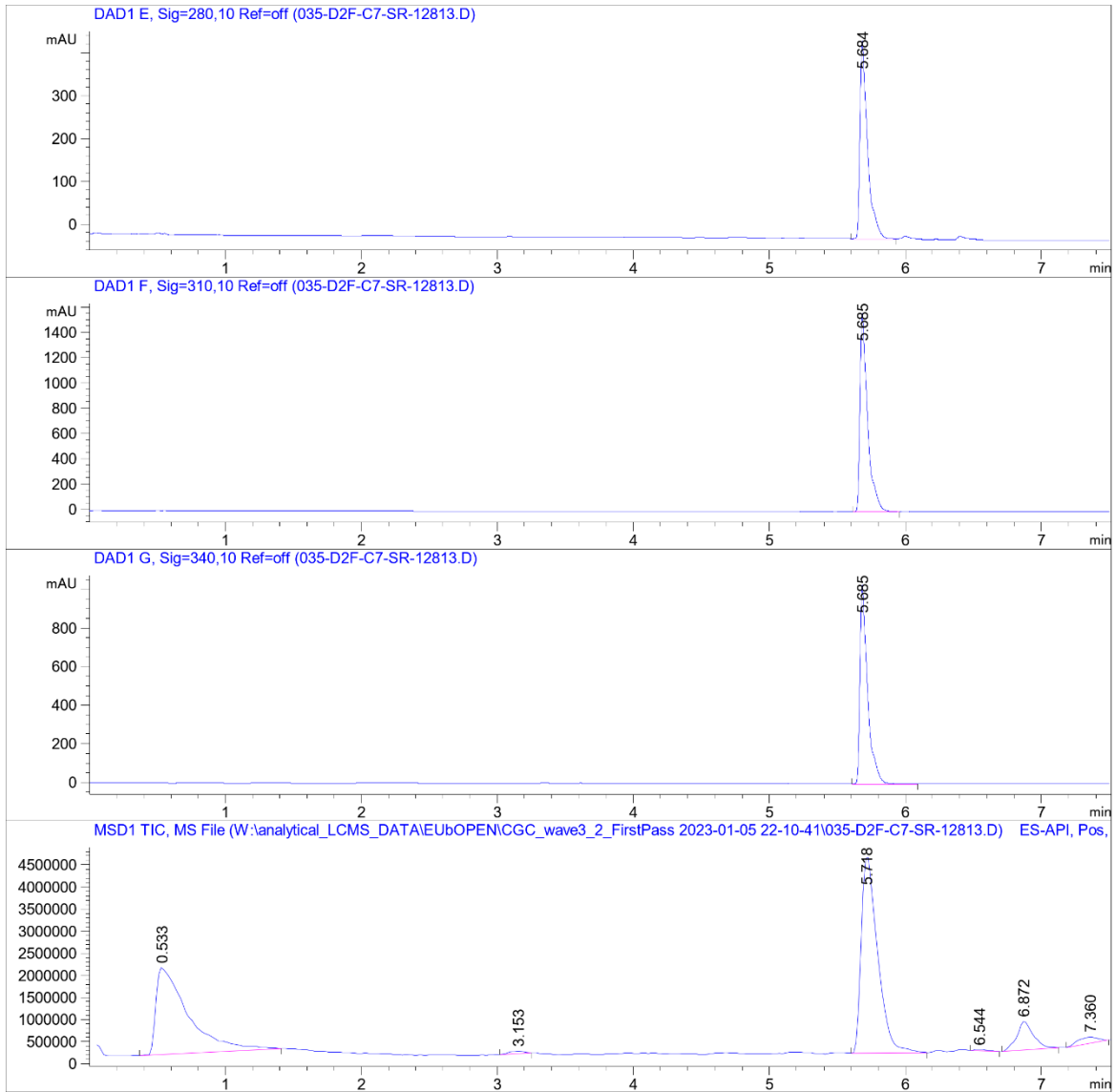
```
=====
Acq. Operator   : SYSTEM                      Seq. Line :   35
Sample Operator : SYSTEM
Acq. Instrument : LCMS test                   Location  : D2F-C7
Injection Date  : 1/6/2023 4:29:43 AM        Inj       :    1
                                           Inj Volume: Inj prog
Sequence File   : W:\analytical_LCMS_DATA\EUBOPEN\CGC_wave3_2_FirstPass 2023-01-05 22-10-41
                  \CGC_wave3_2_FirstPass.S
Method          : W:\analytical_LCMS_DATA\EUBOPEN\CGC_wave3_2_FirstPass 2023-01-05 22-10-41
                  \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...PEN\CGC\_wave3\_2\_FirstPass 2023-01-05 22-10-41\035-D2F-C7-SR-12813.D

Sample Name: SR-12813



# COMPOUND INFORMATION



Data File W:\analyti...PEN\CGC\_wave3\_2\_FirstPass 2023-01-05 22-10-41\035-D2F-C7-SR-12813.D

Sample Name: SR-12813

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
-	-	0.533	32099910	157.00 I
-	-	3.153	457253	188.10 I 170.10 I
5.685	2483	5.718	37039984	505.30 I
-	-	6.544	209209	507.30 I 280.20 I
-	-	6.872	5459114	282.20 I
-	-	7.360	1514067	400.30 I 282.20 I

