

Supporting Information

Discovery of an Orally Bioavailable Small-Molecule Inhibitor for the β -Catenin/B-Cell Lymphoma 9 Protein–Protein Interaction

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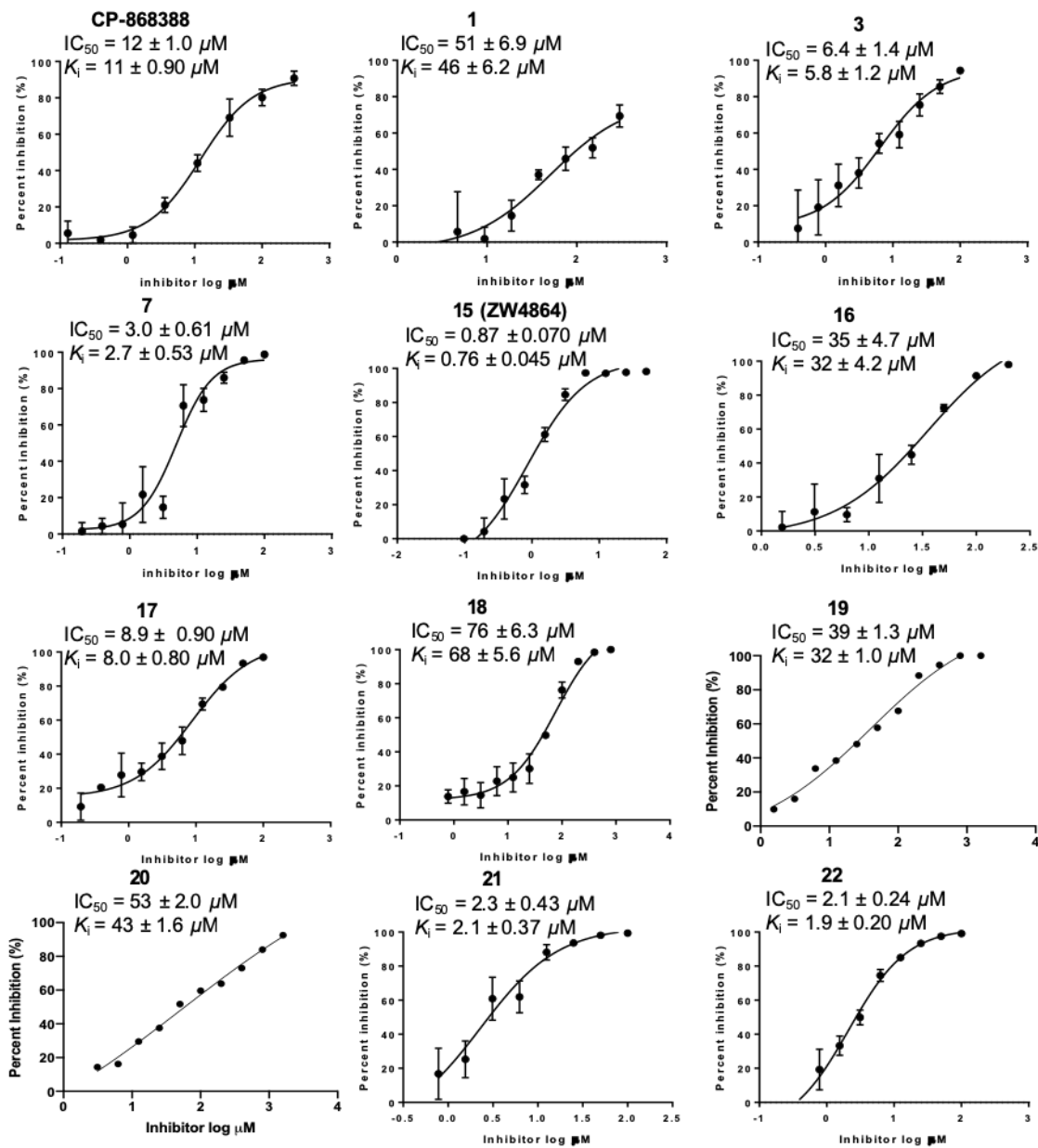
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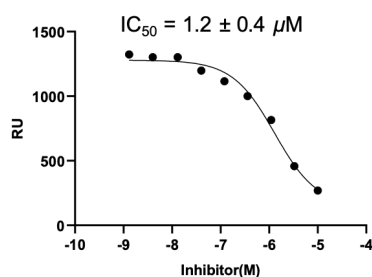
[†]Z.W., M.Z., and V.Q. contributed equally to this work.

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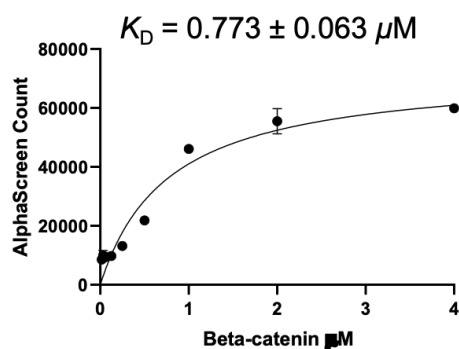


Supplementary Figure S1. The dose-response curves of AlphaScreen assays of CP-868388, 1, 3, 7, 15 (ZW4864), and 16–22 for disruption of the full-length β -catenin/BCL9 PPI. Each set of data is expressed as mean \pm standard deviation ($n = 3$).

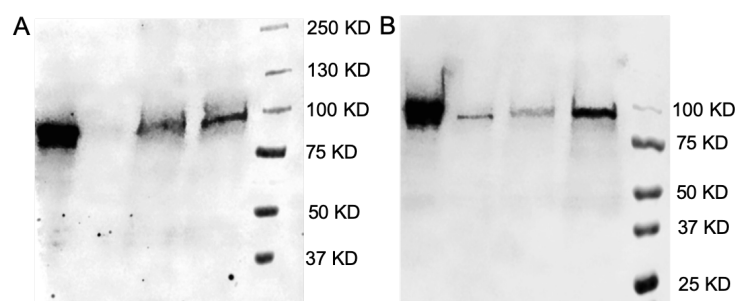


Supplementary Figure S2. The fitting curve that derives the IC_{50} of SPR competitive inhibition assay for disruption of β -catenin/BCL9 PPI. The concentrations of **ZW4864** for curve fitting are 30, 10, 1.1, 0.36, 0.12, 0.04, 0.013, 0.004, and 0.0013 μ M, respectively.

AlphaScreen competitive binding assays
full-length β -catenin/**Biotin-ZW4864**



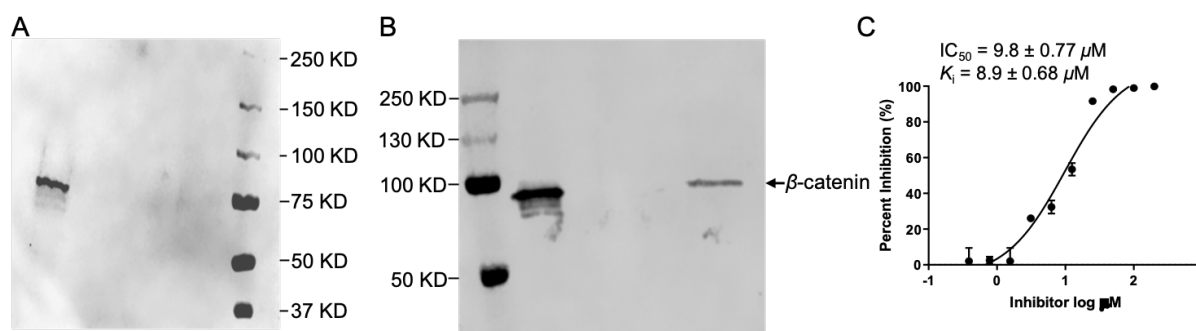
Supplementary Figure S3. The K_D of **Biotin-ZW4864** with full-length β -catenin in AlphaScreen saturation binding assays. Increasing concentrations of His₆-tagged full-length β -catenin were incubated with 10 nM **Biotin-ZW4864** for 1 h. Donor and acceptor beads were then added into each well to a final concentration of 20 μ g/mL in a 25 μ L assay volume. The plate was read after 2 h incubation. Each experiment was performed in triplicate. Data are expressed as mean \pm standard deviation (n = 3).



The normalized densities of the images of the protein pull-down experiment.

ZW4864, μM	full-length β -catenin	SW480 cell lysates
Input	1.00	1.00
0	0.01 ± 0.05	0.07 ± 0.04
1	0.10 ± 0.03	0.05 ± 0.03
10	0.28 ± 0.07	0.29 ± 0.05

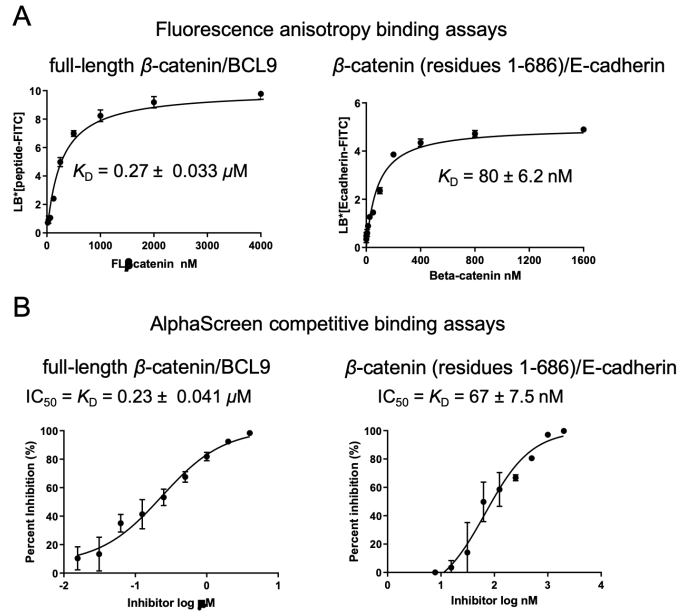
Supplementary Figure S4. Uncropped Western blot gels for the pull-down experiments of **Biotin-ZW4864** using (A) purified full-length human β -catenin. and (B) SW480 cell lysates.



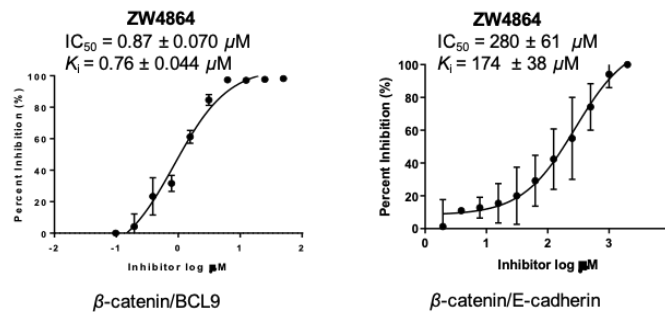
The normalized densities of the images of the protein pull-down experiment.

ZW4864, μM	RIC	ZW4864, μM	RIC
Input	1.00	Input	1.00
0	0.01 ± 0.01	0	0.03 ± 0.01
1	0.03 ± 0.02	5	0.07 ± 0.01
10	0.04 ± 0.02	50	0.27 ± 0.06

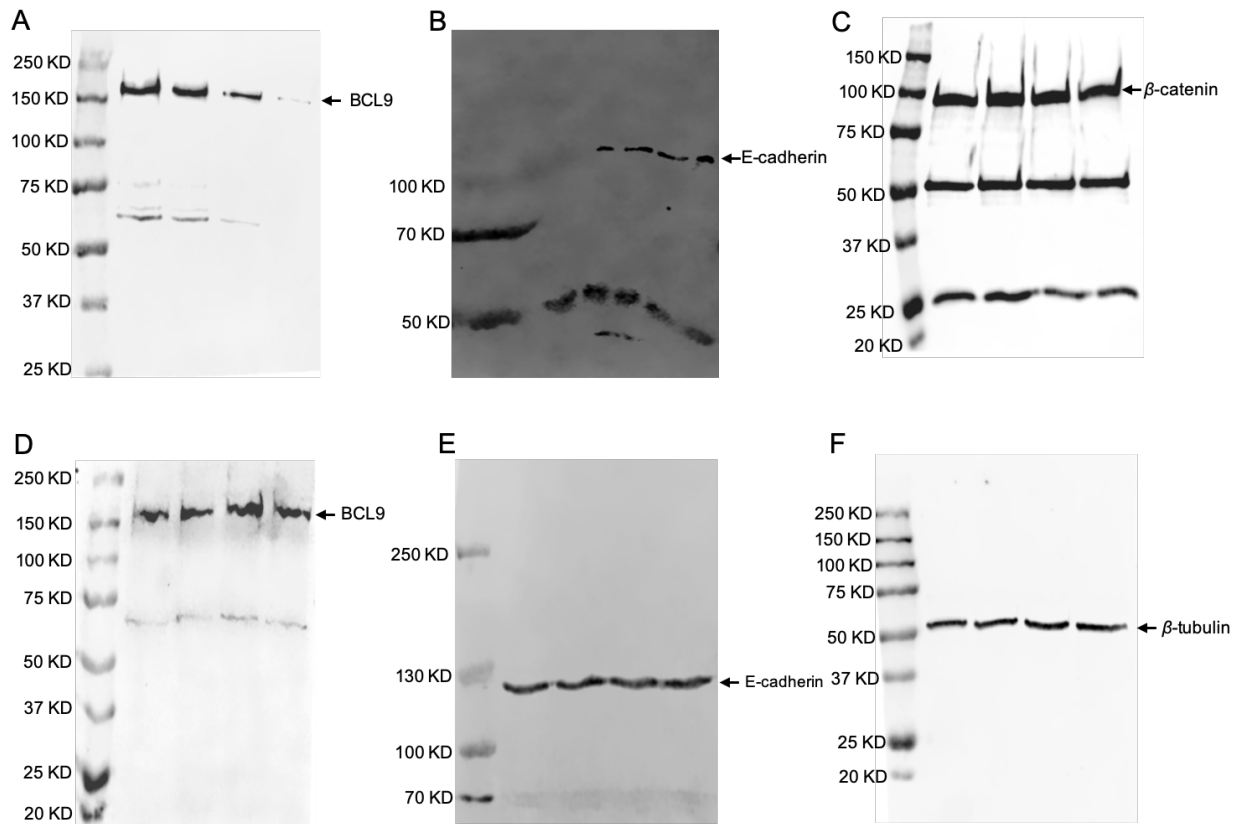
Supplementary Figure S5. Uncropped Western blot gels for the pull-down experiments of **Biotin-ZW4864** at (A) 1 and 10 μM , (B) 5 and 50 μM using β -catenin RIC (residue 138–781). (C) AlphaScreen assay results of **ZW4864** for disruption of β -catenin RIC/BCL9 PPI. Each set of data is expressed as mean \pm standard deviation (n = 3).



Supplementary Figure S6. The K_D s of full-length β -catenin with BCL9 HD2 peptide (residues 350–375) and the K_D s of β -catenin (residues 1-686) with E-cadherin peptide (residues 824–877) using (A) fluorescence anisotropy binding assays and (B) AlphaScreen competitive binding assays. Each set of data is expressed as mean \pm standard deviation ($n = 3$).



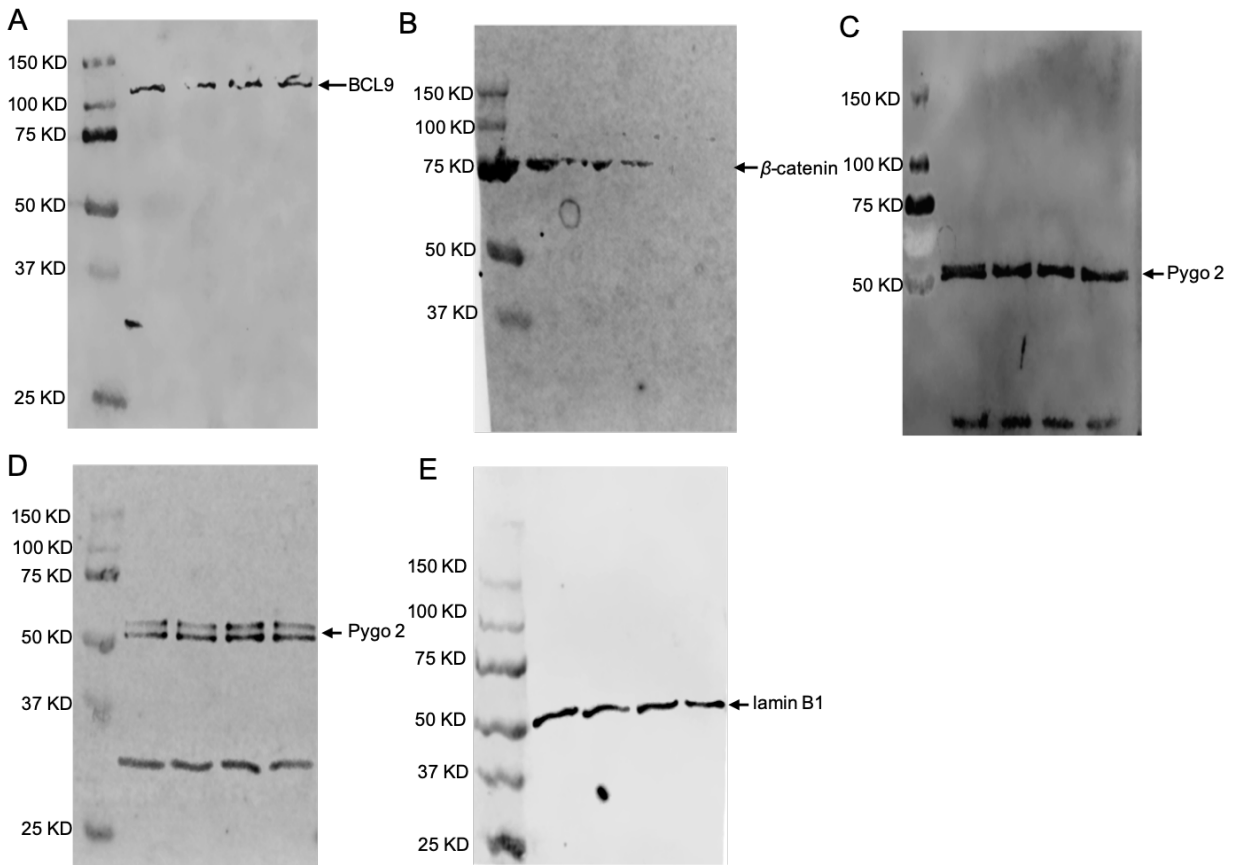
Supplementary Figure S7. AlphaScreen selectivity assay results of **ZW4864** for disruption of full-length β -catenin/BCL9 and full-length β -catenin/E-cadherin PPIs. Each set of data is expressed as mean \pm standard deviation ($n = 3$).



The normalized densities of the images of the co-IP assays.

ZW4864, μM	IP: β -catenin IB: BCL9	IP: β -catenin IB: E-cadherin	IP: β -catenin IB: β -catenin	Input: E-cadherin	Input: BCL9	Input: β -tubulin
0	1.0	1.0	1.0	1.0	1.0	1.0
10	0.73 ± 0.06	1.10 ± 0.05	1.05 ± 0.08	1.10 ± 0.04	1.03 ± 0.05	1.05 ± 0.08
20	0.42 ± 0.05	1.05 ± 0.08	0.99 ± 0.08	1.05 ± 0.06	1.11 ± 0.09	1.10 ± 0.06
40	0.20 ± 0.03	1.15 ± 0.10	0.95 ± 0.06	1.10 ± 0.09	1.02 ± 0.07	1.06 ± 0.05

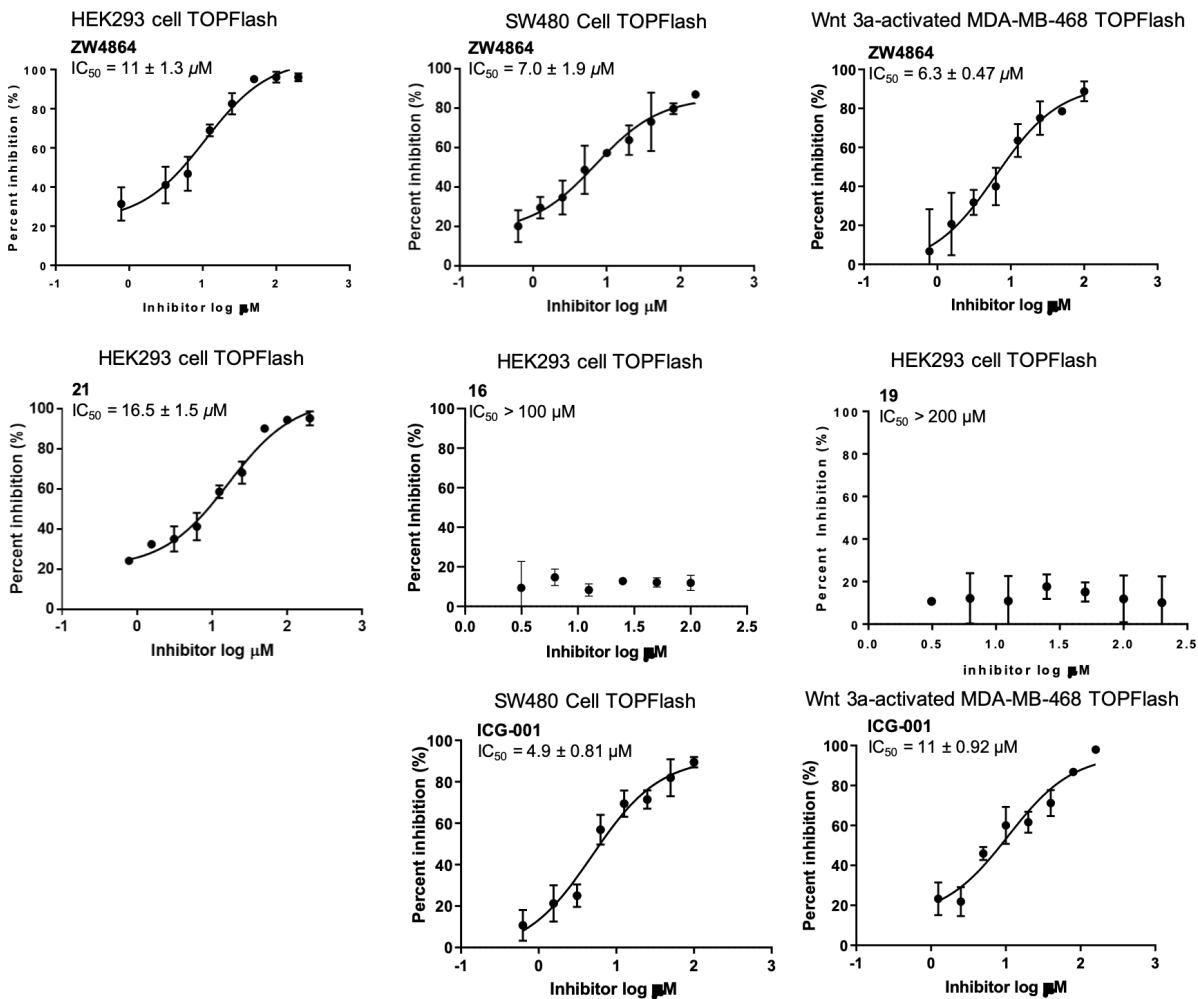
Supplementary Figure S8. Uncropped Western blot gels for the co-IP experiments to evaluate effects of ZW4864 for disruption of the β -catenin/ BCL9 PPI and selectivity for β -catenin/BCL9 over β -catenin/E-cadherin PPIs. (A) β -Catenin immunoprecipitation (IP) and then BCL9 immunoblotting (IB); (B) β -catenin IP and then E-cadherin IB; (C) β -catenin IP and then β -catenin IB; (D) BCL9 IB as the input; (E) E-cadherin IB as the input; and (F) β -tubulin IB as the input.



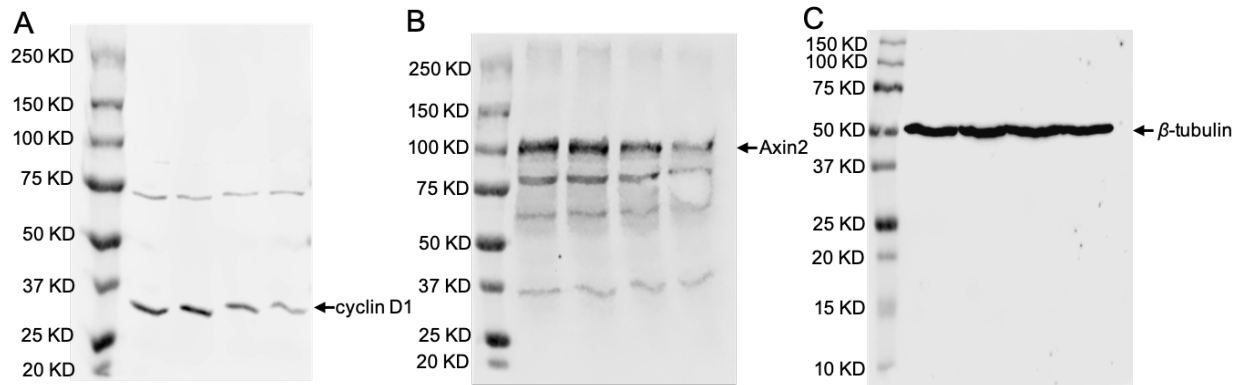
The normalized densities of the images of co-IP experiments to evaluate effects of **ZW4864** for disruption of BCL9/Pygo and β -catenin/Pygo interactions.

ZW4864, μM	IP: Pygo2 IB: BCL9	IP: Pygo2 IB: β -catenin	IP: Pygo2 IB: Pygo2	Input: Pygo2	Input: lamin B1
0	1.0	1.0	1.0	1.0	1.0
10	0.91 \pm 0.06	0.69 \pm 0.05	1.05 \pm 0.05	1.06 \pm 0.06	0.92 \pm 0.05
20	1.01 \pm 0.07	0.36 \pm 0.04	1.05 \pm 0.06	1.19 \pm 0.08	0.96 \pm 0.07
40	1.07 \pm 0.08	0.05 \pm 0.01	0.99 \pm 0.07	1.07 \pm 0.06	0.93 \pm 0.04

Supplementary Figure S9. Uncropped gels for co-IP experiments to evaluate effects of **ZW4864** for disruption of BCL9/Pygo and β -catenin/Pygo interactions. (A) Pygo 2 immunoprecipitation (IP) and then BCL9 immunoblotting (IB); (B) pygo 2 IP and then β -catenin IB; (C) pygo 2 IP and then pygo 2 IB; (D) pygo 2 as the input; and (E) lamin B1 as the input.



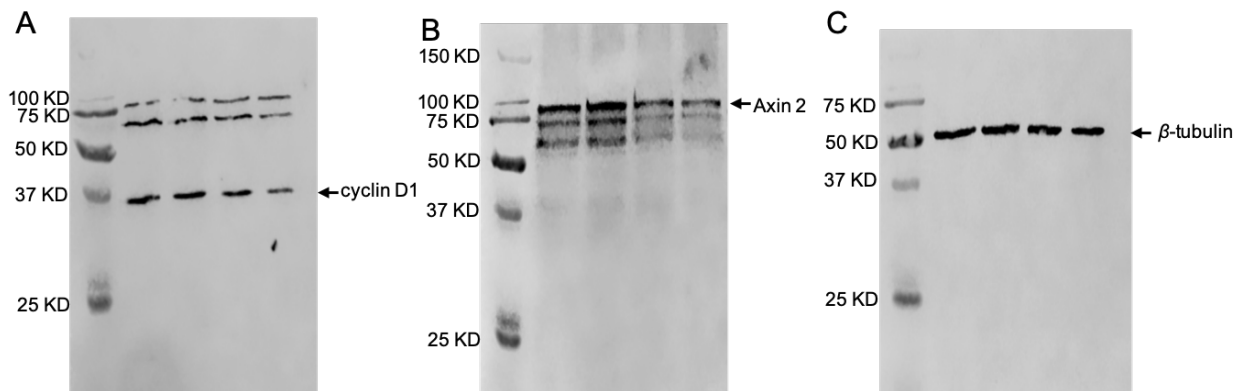
Supplementary Figure S10. The dose-response curves of Wnt-responsive TOPFlash luciferase reporter assay results of compounds **ZW4864 (15)**, **21**, **16**, **19** and **ICG-001**.



The normalized densities of the images of the Western blot analyses

ZW4864, μM	Axin2	cyclin D1	β -tubulin
0	1.0	1.0	1.0
10	1.06 ± 0.08	1.03 ± 0.06	1.04 ± 0.06
20	0.62 ± 0.05	0.56 ± 0.06	1.02 ± 0.08
40	0.35 ± 0.04	0.31 ± 0.05	0.95 ± 0.07

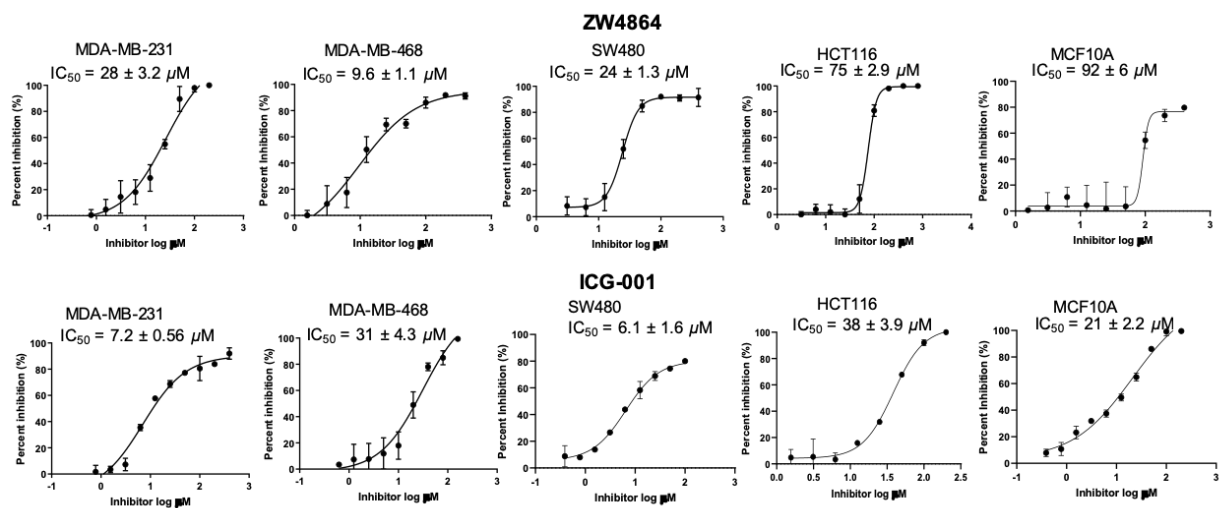
Supplementary Figure S11. Uncropped gels of Western blot experiments of **ZW4864** in SW480 cells, as described in Figure 2G. (A) Cyclin D1; (B) Axin2; and (C) β -tubulin.



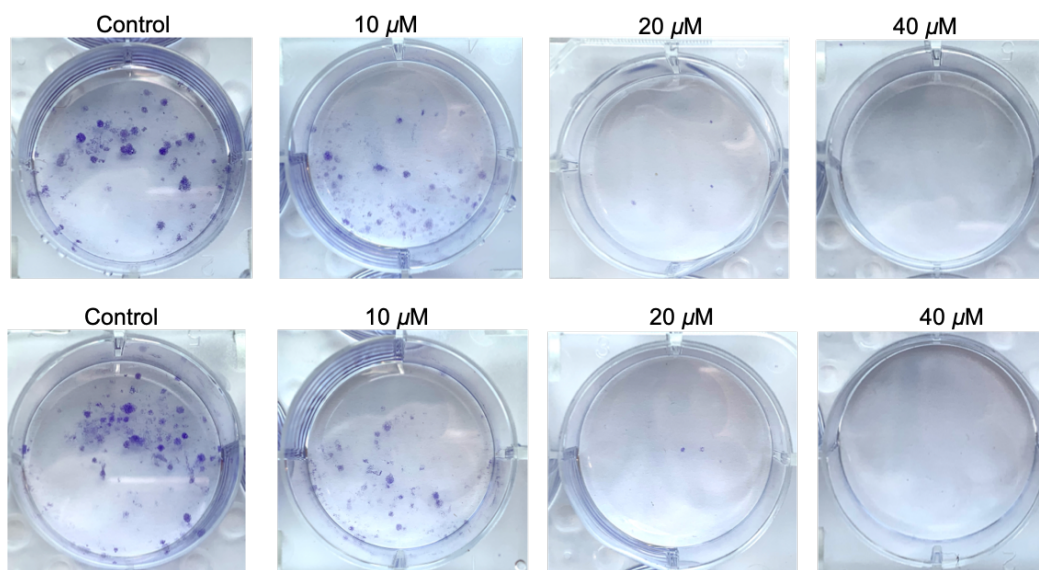
The normalized densities of the images of the Western blot analyses

ZW4864, μM	Axin2	cyclin D1	β -tubulin
0	1.0	1.0	1.0
10	1.08 ± 0.09	0.98 ± 0.08	0.95 ± 0.09
20	0.57 ± 0.06	0.67 ± 0.07	0.92 ± 0.08
40	0.29 ± 0.04	0.32 ± 0.05	0.93 ± 0.08

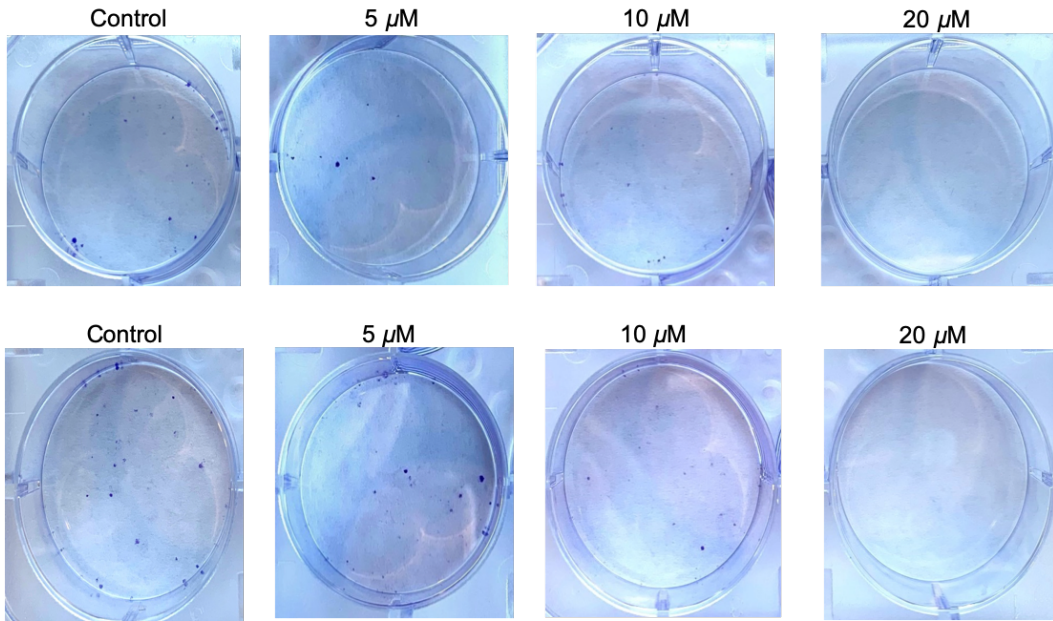
Supplementary Figure S12. Uncropped gels of Western blot experiments of **ZW4864** in Wnt 3a-activated TNBC MDA-MB-231 cells, as described in Figure 2G. (A) Cyclin D1; (B) Axin 2; and (C) β -tubulin.



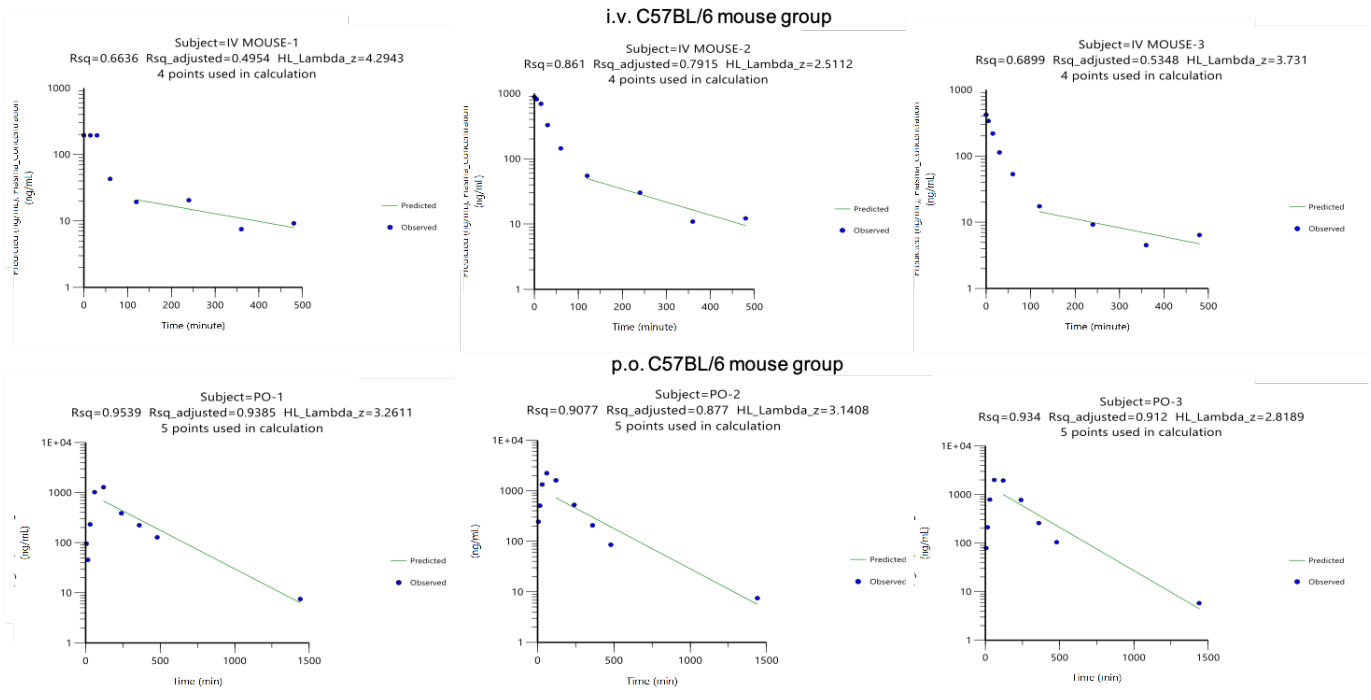
Supplementary Figure S13. MTS growth inhibition assay results of **ZW4864** and **ICG-001** against various TNBC cell lines (MDA-MB-231 and MDA-MB-468), colorectal cancer lines (SW480 and HCT116), and normal mammary epithelial MCF10A cell line after 72-h treatment.



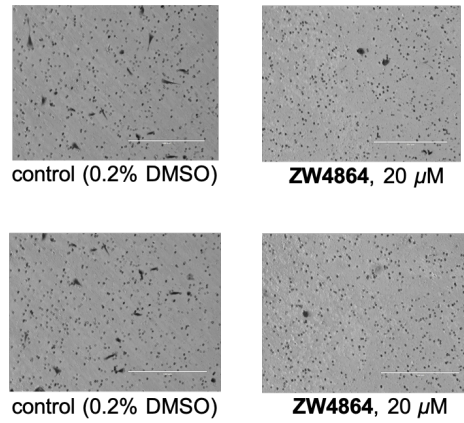
Supplementary Figure S14. The clonogenic assay results of **ZW4864** using MDA-MB-231 cells.



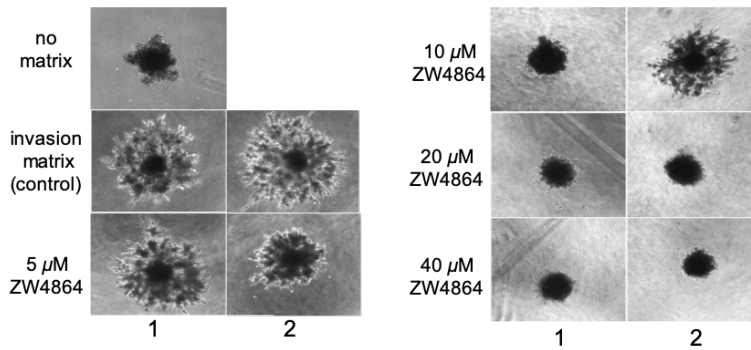
Supplementary Figure S15. Clonogenic assay results of **ZW4864** using MDA-MB-468 cells.



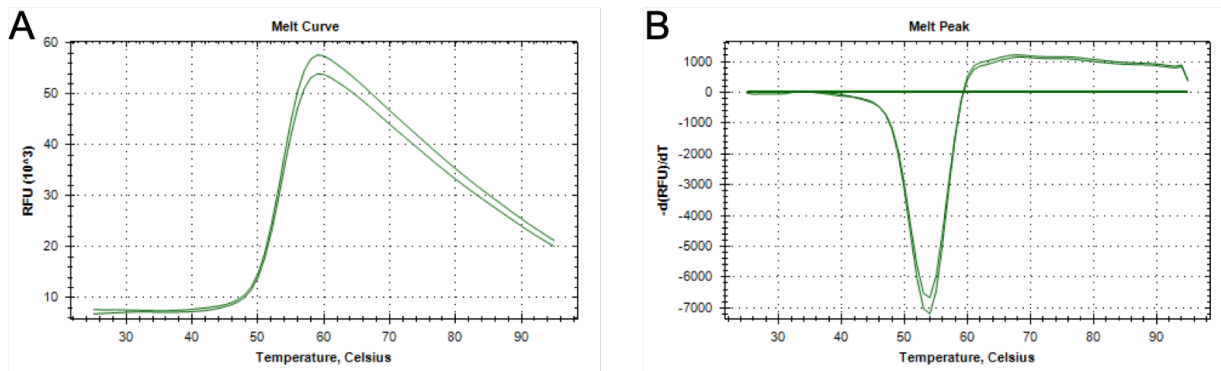
Supplementary Figure S16. Original data plots of C57BL/6 mouse PK studies of **ZW4864**.



Supplementary Figure S17. Original images of the Matrigel invasion assay results of **ZW4864**.



Supplementary Figure S18. Original images of 3D spheroid BME cell invasion assays of **ZW4864**.



Supplementary Figure S19. The quality of full-length β -catenin proteins after purification was examined by thermal shift assays. Here is one example. A temperature increment of $1^\circ/\text{min}$ was applied.

HPLC Condition and Traces

The purity of final compounds was determined by HPLC analysis. The instrument was an Agilent 1260 Infinity II HPLC system with a quaternary pump, a vial sampler, and a DAD detector. A Kromasil 300–5–C18 column (4.6 × 250 mm) was used. The DAD detector was set to 254 nm. The purity of all tested compounds was >95%. Some HPLC traces are shown below.

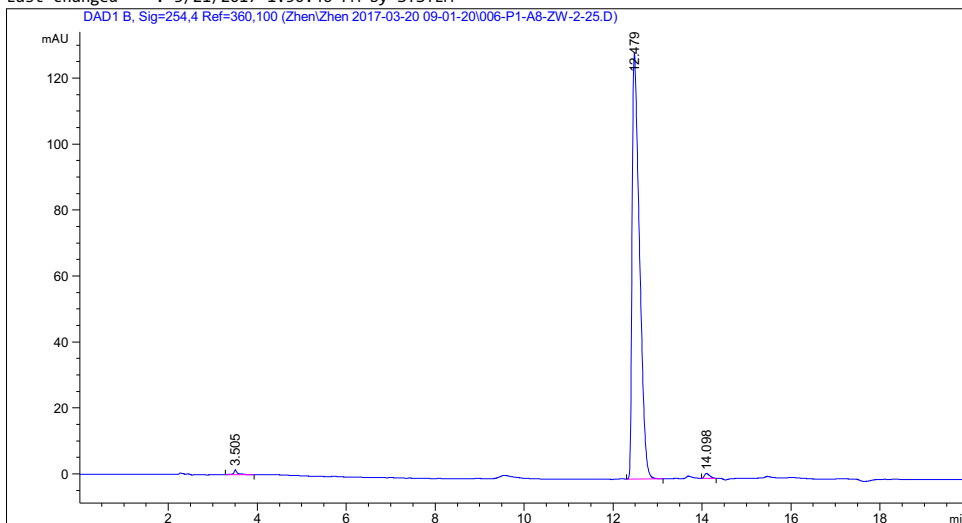
Condition: Elute with gradient starting with 0.1% TFA in water and ending with 0.1% TFA in water and acetonitrile mixture (water with 0.1% TFA : acetonitrile = 1 : 1) for 10 min, and then change to a 10 min-gradient starting with 0.1% TFA in water and acetonitrile 1 : 1 mixture and ending with 100% acetonitrile.

Compound ID	Purity (%)	Retention time (min)
1	98.6	12.48
2	98.9	12.51
3	97.6	10.96
4	100	11.37
5	98.0	11.55
6	100	12.33
7	100	12.15
8	100	12.49
9	99.4	13.42
10	100	12.29
11	99.4	11.90
12	97.1	8.05
13	99.2	8.21
14	98.8	13.07
15	96.8	9.35
16	98.8	9.34
17	99.1	9.41
18	98.0	9.86
19	98.5	9.23
20	99.7	11.52
21	99.9	8.99
22	99.7	9.34
Biotin-ZW4864	96.0	8.83

Compound 1

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Area Percent Report
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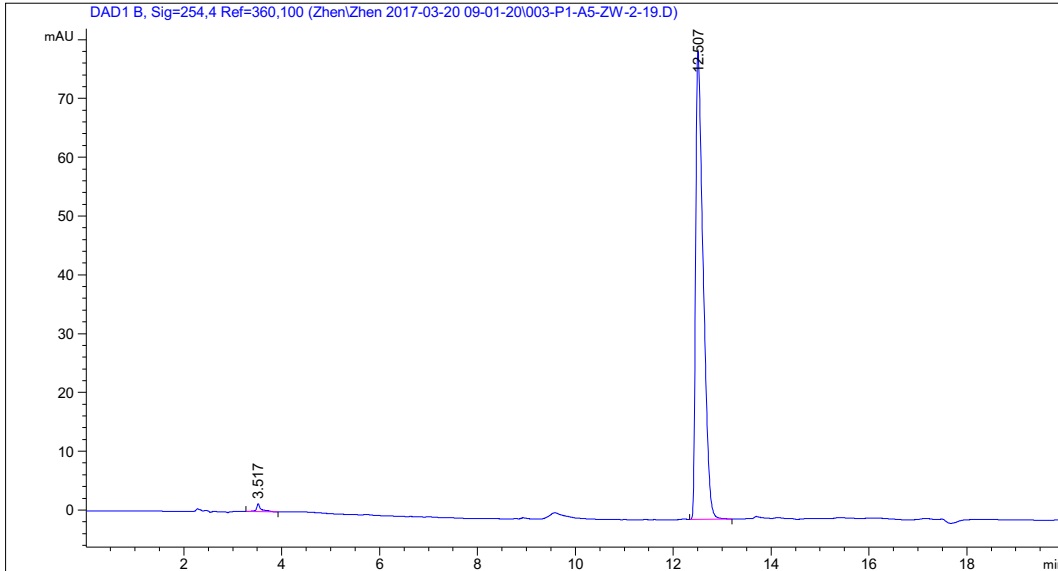
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Area Percent Report
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Use Multiplier & Dilution Factor with ISTDs
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2	12.507	BB	0.1577	855.95752	79.46735	98.9371

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Totals :                      865.15304  80.83259
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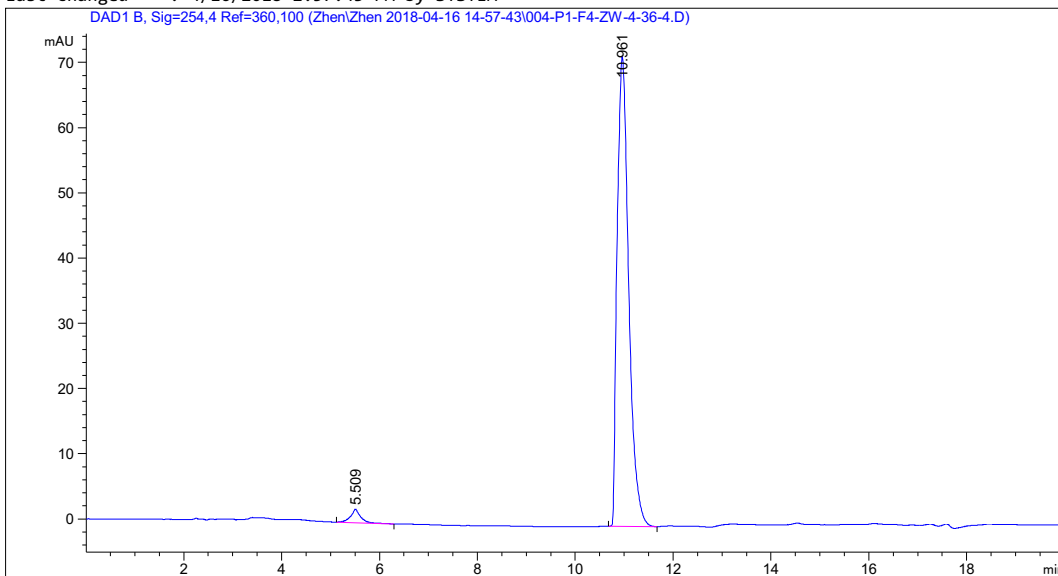
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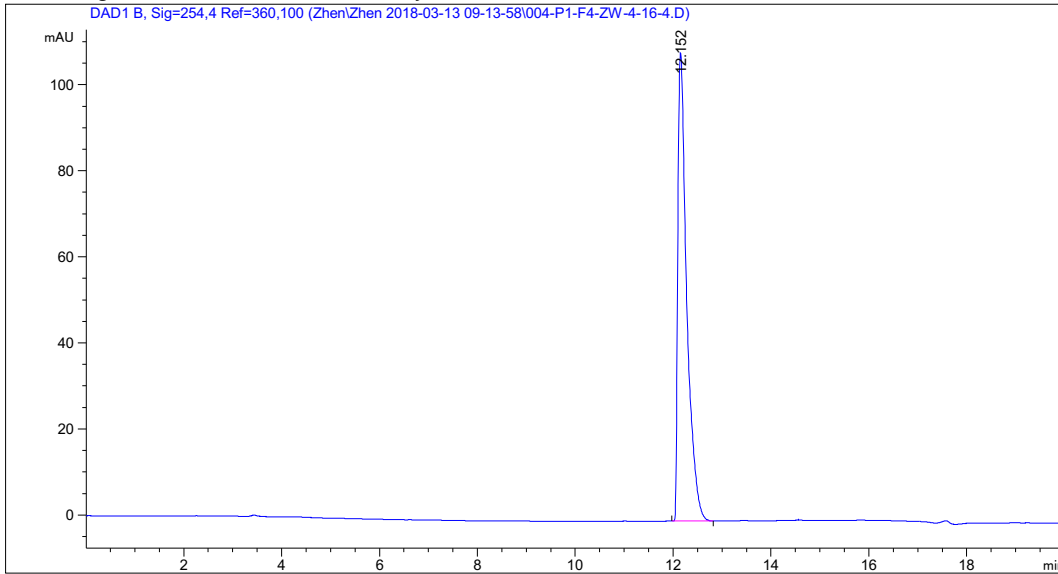
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Compound 7

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Dilution : 1.0000
Use Multiplier & Dilution Factor with ISTDs

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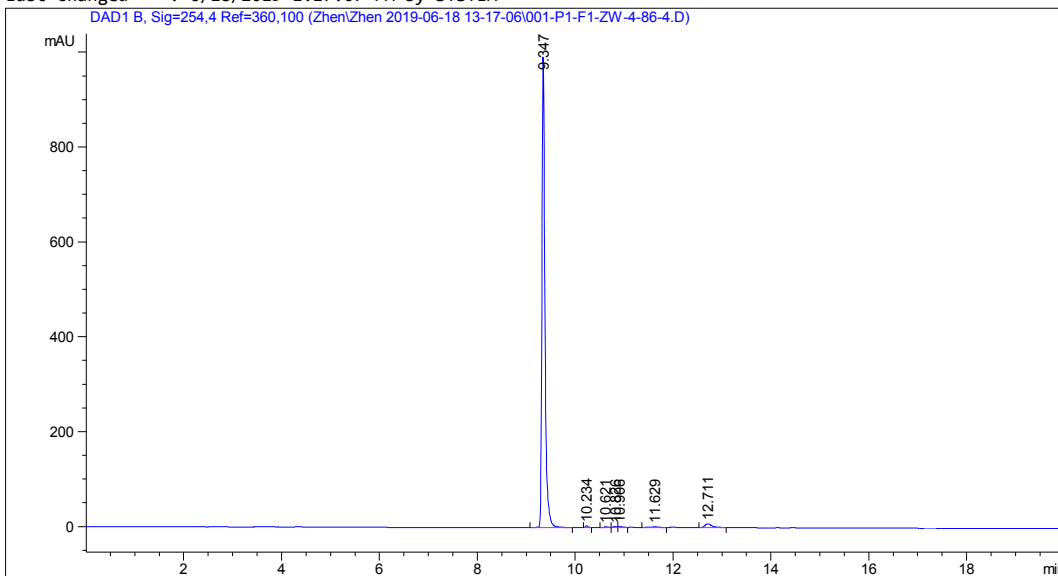
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Compound 15

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Area Percent Report

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Use Multiplier & Dilution Factor with ISTDs
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Signal 1: DAD1 B, Sig=254,4 Ref=360,100

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2	10.234	BB	0.0544	12.00380	3.46852	0.2715
3	10.621	BV	0.0808	10.25903	1.79158	0.2320
4	10.826	VV	0.0862	11.37997	2.00437	0.2574
5	10.906	VB	0.0915	11.55812	1.83584	0.2614
6	11.629	BB	0.1676	20.89855	1.65176	0.4726
7	12.711	BB	0.1467	77.22118	7.85193	1.7464

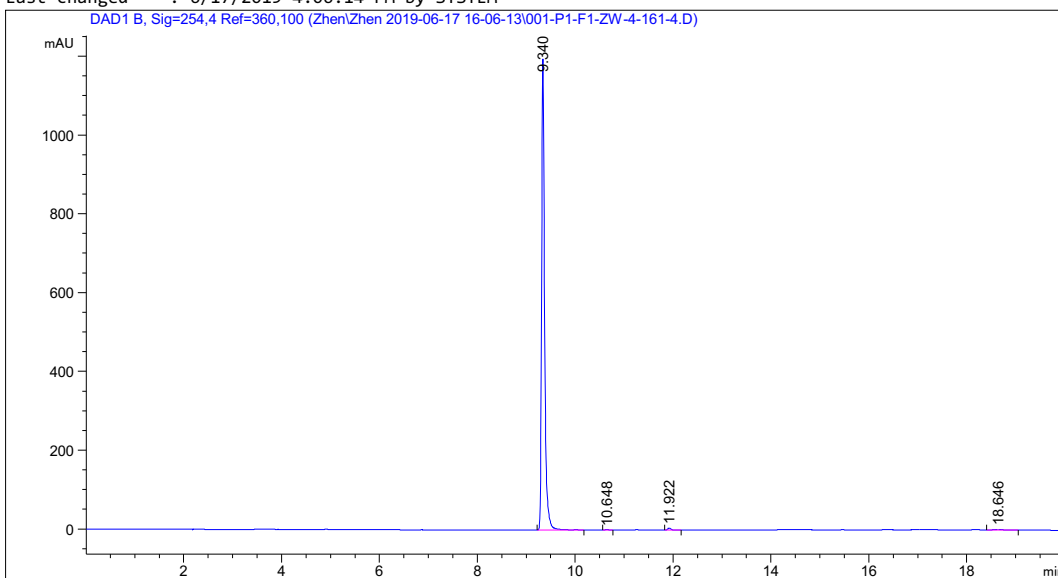
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Compound 16

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Area Percent Report

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3	11.922	BB	0.0694	24.49493	5.35611	0.4725
4	18.646	BB	0.1797	27.60105	2.27311	0.5324

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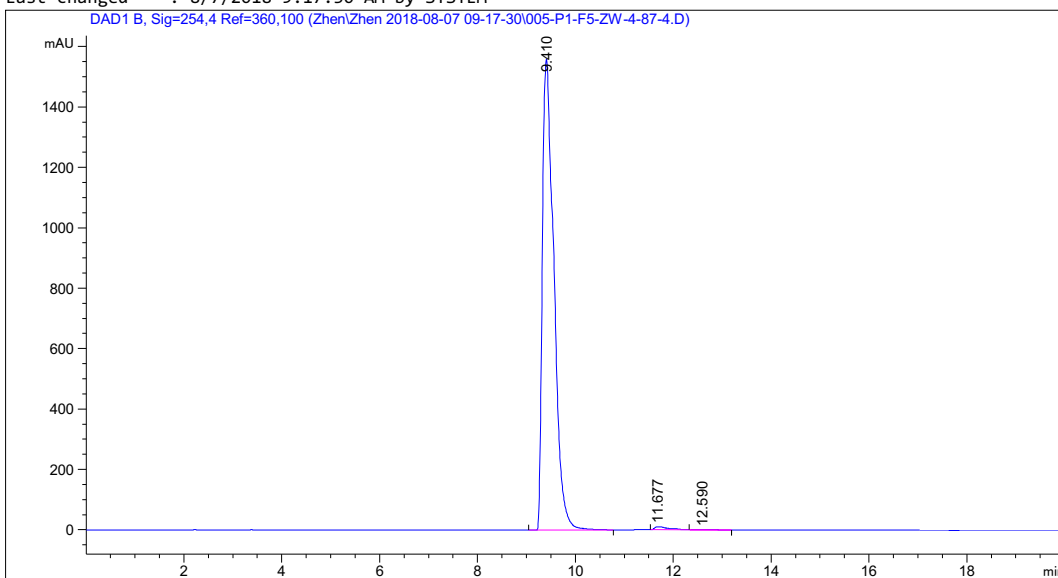
*** End of Report ***

Compound 17

Data File C:\Chem32\1\Data\Zhen\Zhen 2018-08-07 09-17-30\005-P1-F5-ZW-4-87-4.D

Sample Name: ZW-4-87-4

```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    5
Acq. Instrument : HPLC                       Location  : P1-F5
Injection Date  : 8/7/2018 11:07:25 AM      Inj       :    1
                                                Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method         : C:\Chem32\1\Data\Zhen\Zhen 2018-08-07 09-17-30\Zhen.M (Sequence Method)
Last changed   : 8/7/2018 9:17:30 AM by SYSTEM
=====
```



Area Percent Report

```
Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
```

Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	9.410	BB	0.2323	2.54579e4	1560.33081	99.1045
2	11.677	BV R	0.2573	197.27127	9.86483	0.7680
3	12.590	VB E	0.2857	32.75716	1.62133	0.1275

Totals : 2.56879e4 1571.81698

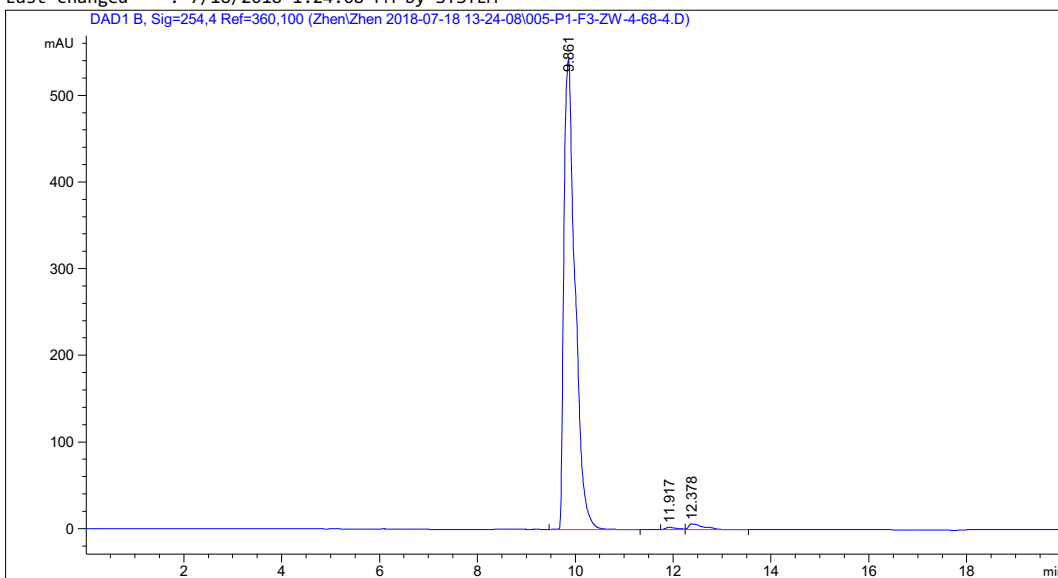
*** End of Report ***

Compound 18

Data File C:\Chem32\1\Data\Zhen\Zhen 2018-07-18 13-24-08\005-P1-F3-ZW-4-68-4.D

Sample Name: ZW-4-68-4

```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    5
Acq. Instrument : HPLC                       Location  : P1-F3
Injection Date  : 7/18/2018 3:20:13 PM       Inj       :    1
                                                Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method          : C:\Chem32\1\Data\Zhen\Zhen 2018-07-18 13-24-08\Zhen.M (Sequence Method)
Last changed    : 7/18/2018 1:24:08 PM by SYSTEM
=====
```



Area Percent Report

```
Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
```

Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	9.861	BB	0.2346	8590.25000	543.04675	97.9676
2	11.917	BV	0.2363	38.83273	2.43210	0.4429
3	12.378	VB	0.2807	139.37563	6.48695	1.5895

Totals : 8768.45835 551.96581

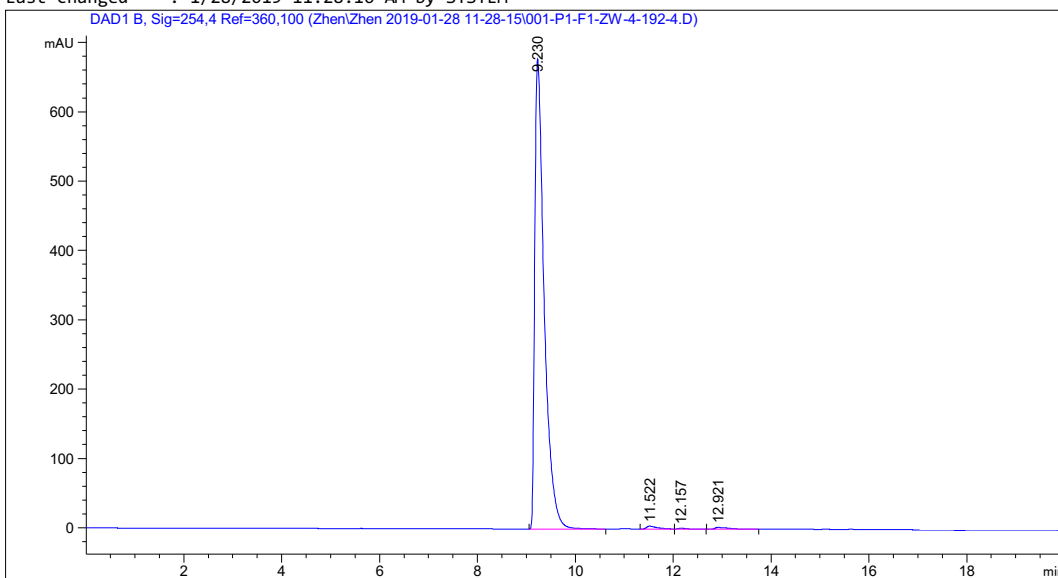
*** End of Report ***

Compound 19

Data File C:\Chem32\1\Data\Zhen\Zhen 2019-01-28 11-28-15\001-P1-F1-ZW-4-192-4.D

Sample Name: ZW-4-192-4

```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    1
Acq. Instrument : HPLC                      Location  : P1-F1
Injection Date  : 1/28/2019 11:37:47 AM      Inj       :    1
                                           Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method         : C:\Chem32\1\Data\Zhen\Zhen 2019-01-28 11-28-15\Zhen.M (Sequence Method)
Last changed   : 1/28/2019 11:28:16 AM by SYSTEM
=====
```



Area Percent Report

```
Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
```

Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	9.230	BB	0.2025	9097.58008	678.28180	98.5312
2	11.522	BV	0.2205	69.13280	4.32439	0.7487
3	12.157	VB	0.1917	17.16881	1.33839	0.1859
4	12.921	BB	0.2458	49.31326	2.64198	0.5341

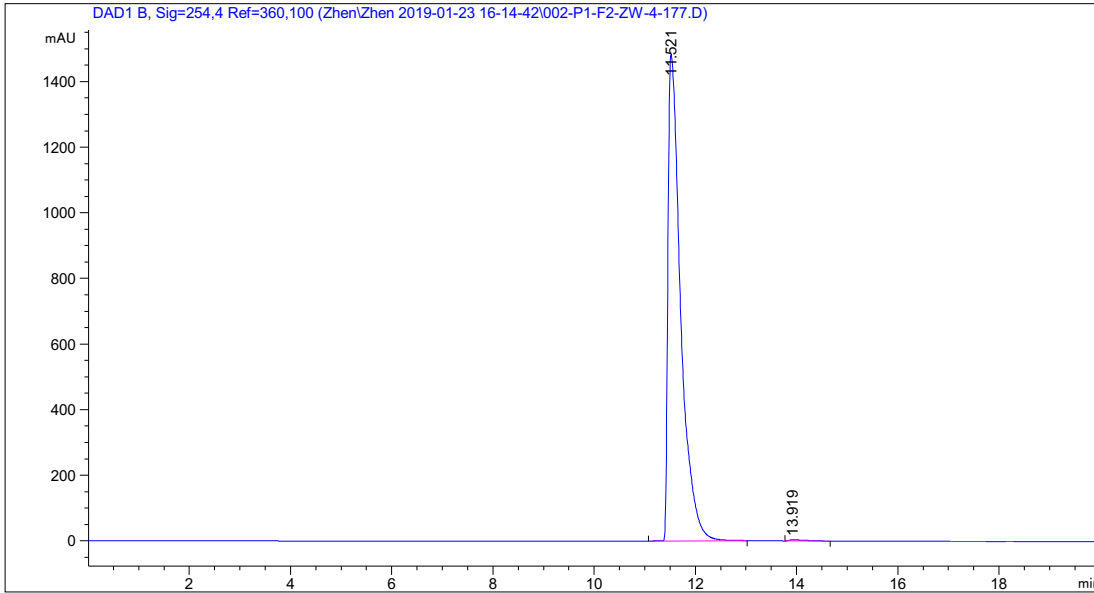
Totals : 9233.19495 686.58656

*** End of Report ***

Compound 20

Data File C:\Chem32\1\Data\Zhen\Zhen 2019-01-23 16-14-42\002-P1-F2-ZW-4-177.D
Sample Name: ZW-4-177

```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    2
Acq. Instrument : HPLC                      Location  : P1-F2
Injection Date  : 1/23/2019 4:47:52 PM      Inj       :    1
                                                Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method          : C:\Chem32\1\Data\Zhen\Zhen 2019-01-23 16-14-42\Zhen.M (Sequence Method)
Last changed    : 1/23/2019 4:14:43 PM by SYSTEM
=====
```



Area Percent Report

```
Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
```

Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	11.521	BB	0.2646	2.57907e4	1484.96643	99.6940
2	13.919	BB	0.2485	79.15220	4.11319	0.3060

Totals : 2.58699e4 1489.07962

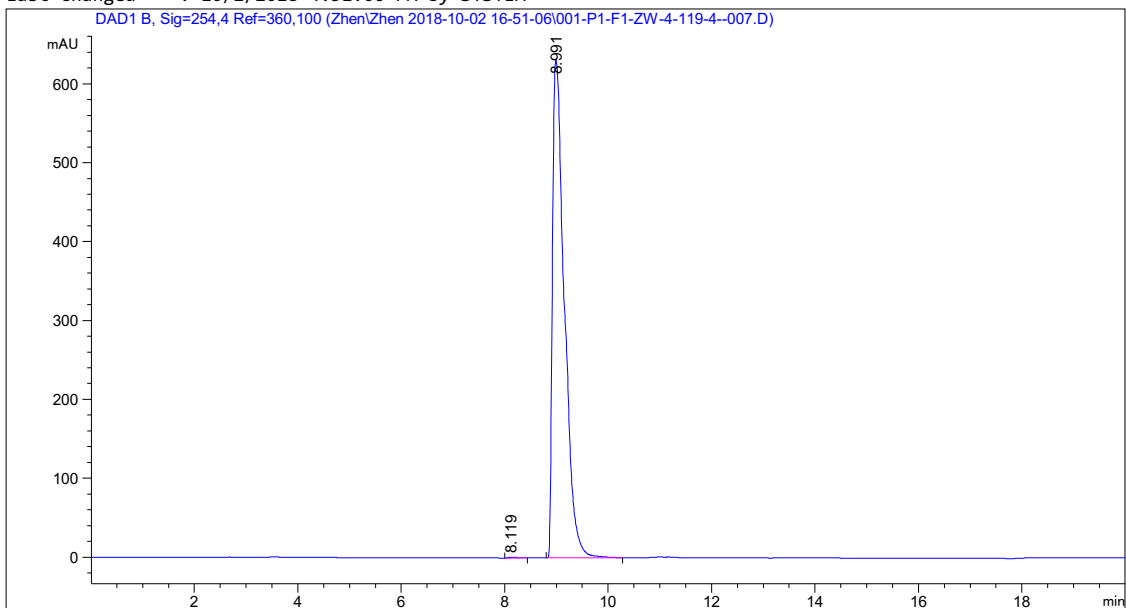
*** End of Report ***

Compound 21

Data File C:\Chem32\1\Data\Zhen\Zhen 2018-10-02 16-51-06\001-P1-F1-ZW-4-119-4--007.D

Sample Name: ZW-4-114-4

```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    7
Acq. Instrument : HPLC                      Location  : P1-F7
Injection Date  : 10/2/2018 7:40:15 PM      Inj       :    1
                                           Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method          : C:\Chem32\1\Data\Zhen\Zhen 2018-10-02 16-51-06\Zhen.M (Sequence Method)
Last changed    : 10/2/2018 4:51:06 PM by SYSTEM
=====
```



Area Percent Report

```
=====
Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
=====
```

Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.119	BB	0.1395	11.79428	1.15122	0.1169
2	8.991	BB	0.2364	1.00763e4	630.53009	99.8831

Totals : 1.00881e4 631.68131

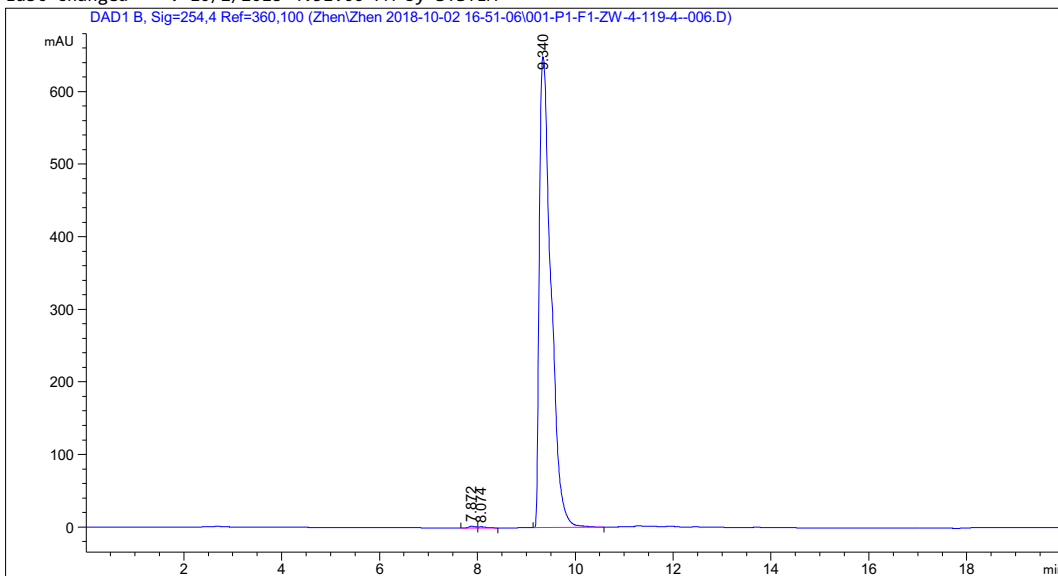
*** End of Report ***

Compound 22

Data File C:\Chem32\1\Data\Zhen\Zhen 2018-10-02 16-51-06\001-P1-F1-ZW-4-119-4--006.D

Sample Name: ZW-4-112-4

```
=====
Acq. Operator   : SYSTEM                      Seq. Line :    6
Acq. Instrument : HPLC                      Location  : P1-F6
Injection Date  : 10/2/2018 7:13:39 PM      Inj       :    1
                                                Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method         : C:\Chem32\1\Data\Zhen\Zhen 2018-10-02 16-51-06\Zhen.M (Sequence Method)
Last changed   : 10/2/2018 4:51:06 PM by SYSTEM
=====
```



Area Percent Report

```
Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
```

Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	7.872	BV	0.1227	20.46256	2.32472	0.1909
2	8.074	VB	0.1309	14.41464	1.48991	0.1345
3	9.340	BB	0.2442	1.06845e4	648.63574	99.6746

Totals : 1.07194e4 652.45038

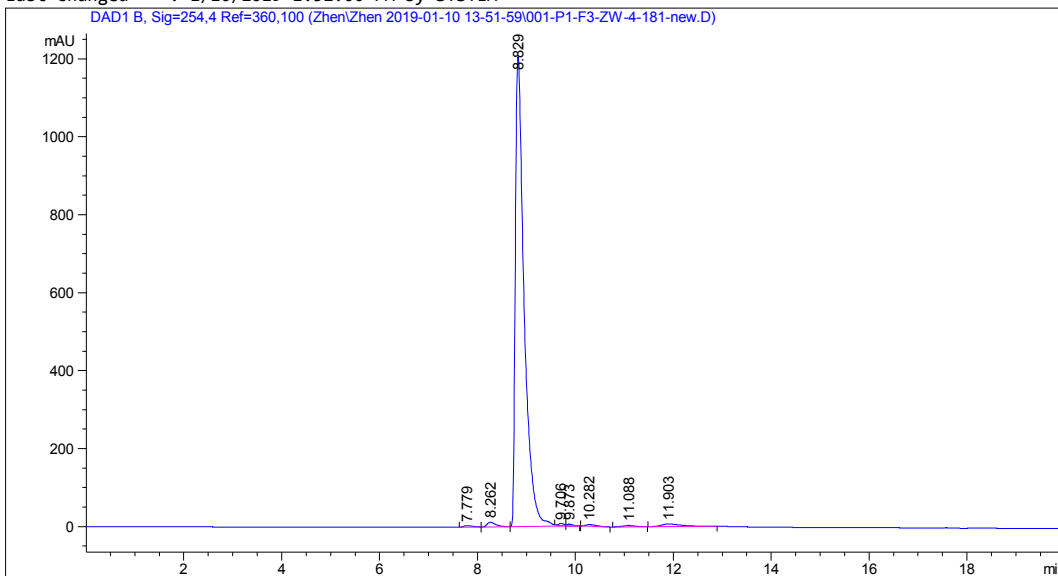
*** End of Report ***

Biotin-ZW4864

Data File C:\Chem32\1\Data\Zhen\Zhen 2019-01-10 13-51-59\001-P1-F3-ZW-4-181-new.D
 Sample Name: ZW-4-181-new

```

=====
Acq. Operator   : SYSTEM                      Seq. Line :    1
Acq. Instrument : HPLC                      Location  : P1-F3
Injection Date  : 1/10/2019 1:58:38 PM      Inj       :    1
                                           Inj Volume: 80.000 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 20.000 µl
Method          : C:\Chem32\1\Data\Zhen\Zhen 2019-01-10 13-51-59\Zhen.M (Sequence Method)
Last changed    : 1/10/2019 1:52:00 PM by SYSTEM
  
```



Area Percent Report

```

Sorted By      : Signal
Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
  
```

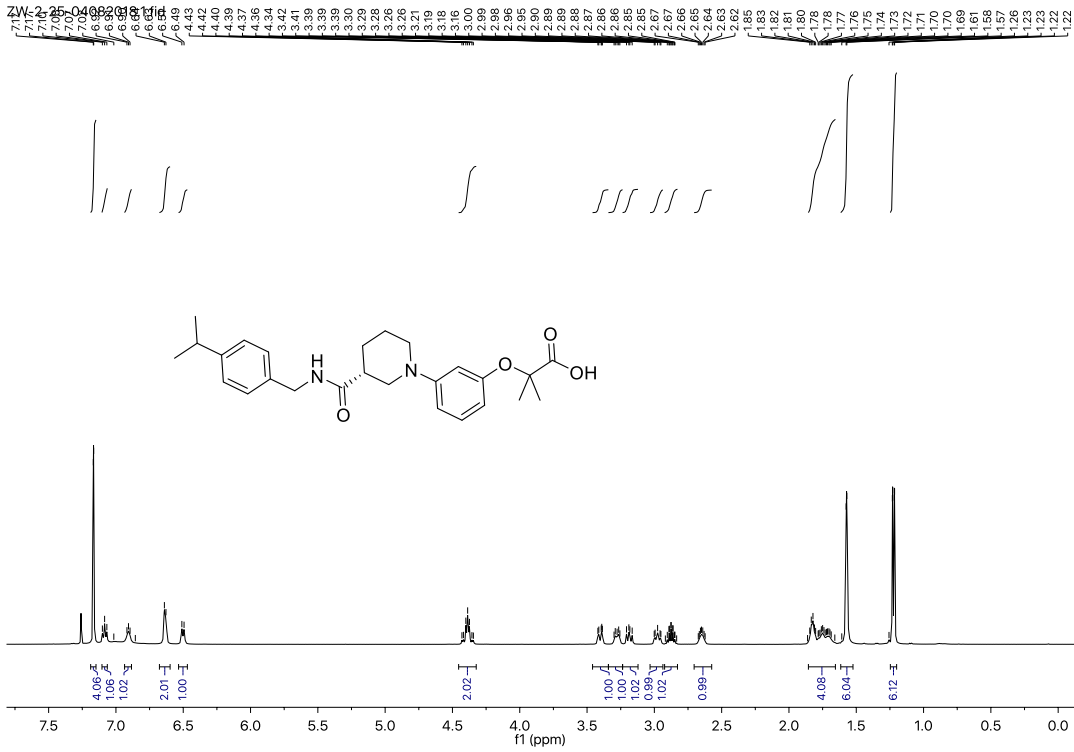
Signal 1: DAD1 B, Sig=254,4 Ref=360,100

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	7.779	BV	0.1983	41.73529	3.42166	0.2643
2	8.262	VB	0.1851	150.59569	12.10558	0.9536
3	8.829	BV R	0.1867	1.51688e4	1205.94653	96.0505
4	9.706	VV E	0.1159	44.89343	5.82460	0.2843
5	9.873	VB E	0.1318	42.79675	4.81042	0.2710
6	10.282	BB	0.1951	64.29467	4.54352	0.4071
7	11.088	BB	0.2791	65.54153	3.52163	0.4150
8	11.903	BB	0.4381	213.86261	6.96707	1.3542

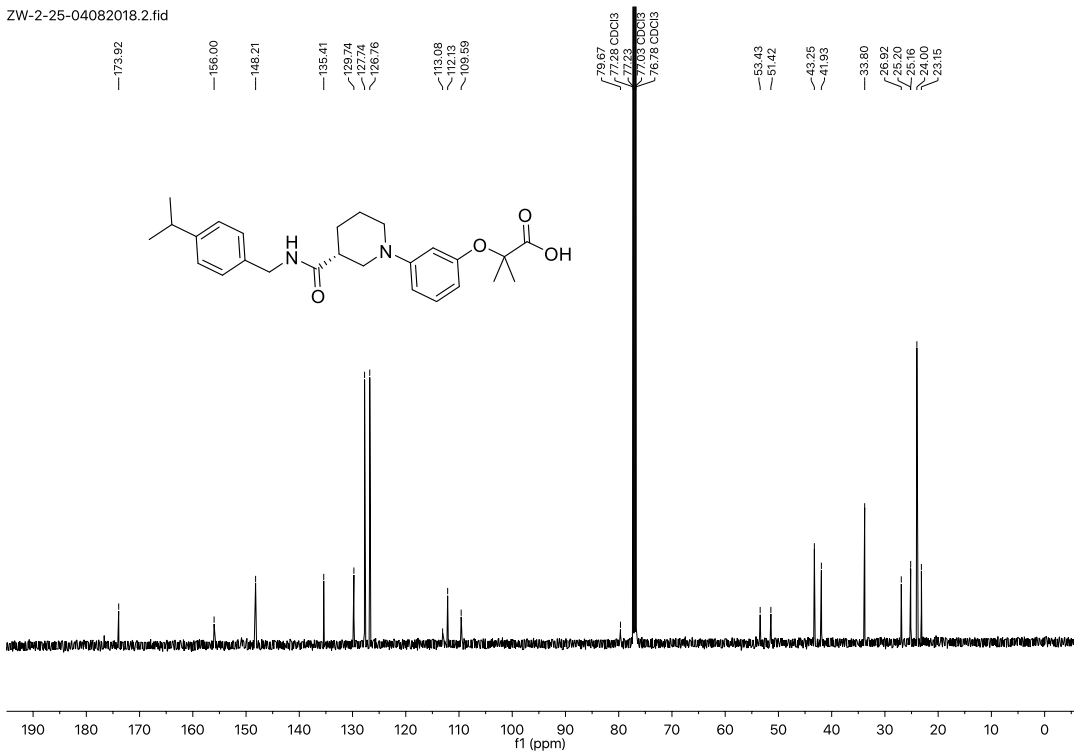
Totals : 1.57925e4 1247.14102

NMR Spectra

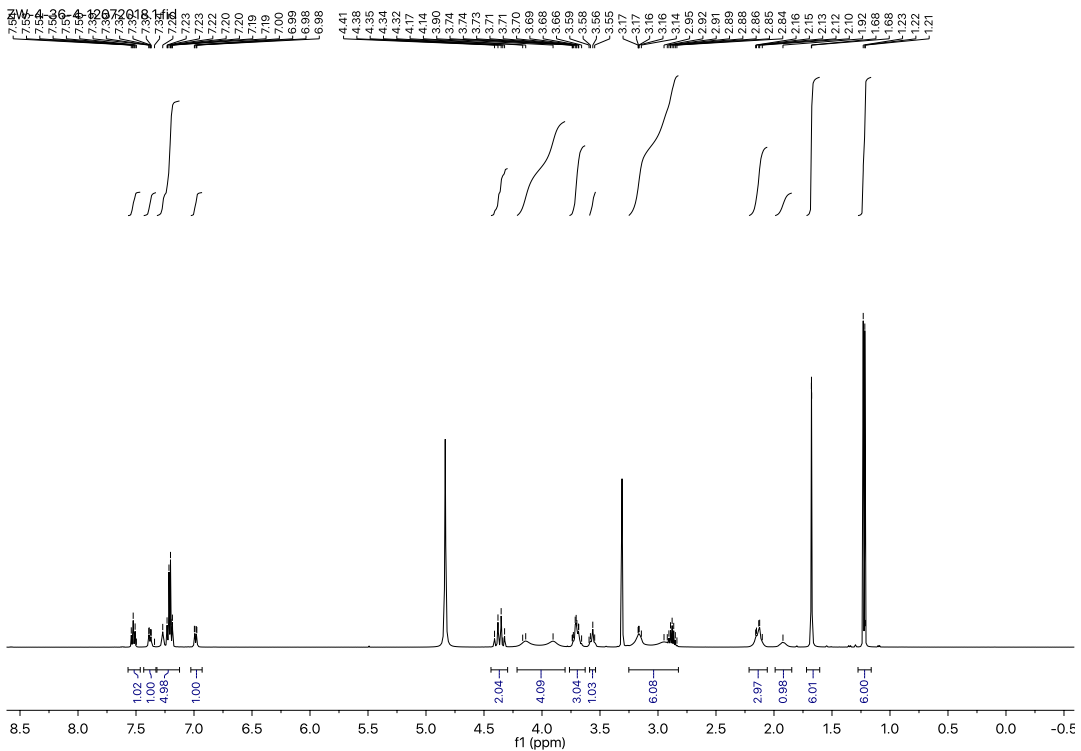
¹H NMR for compound 1



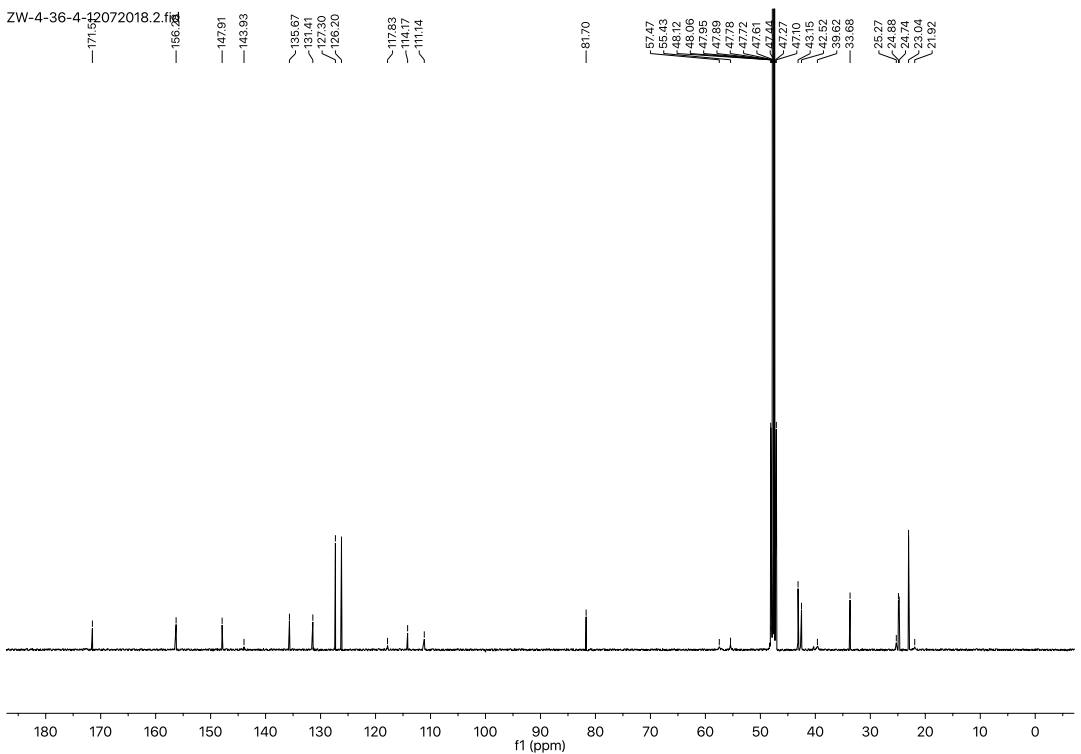
¹³C NMR for compound 1



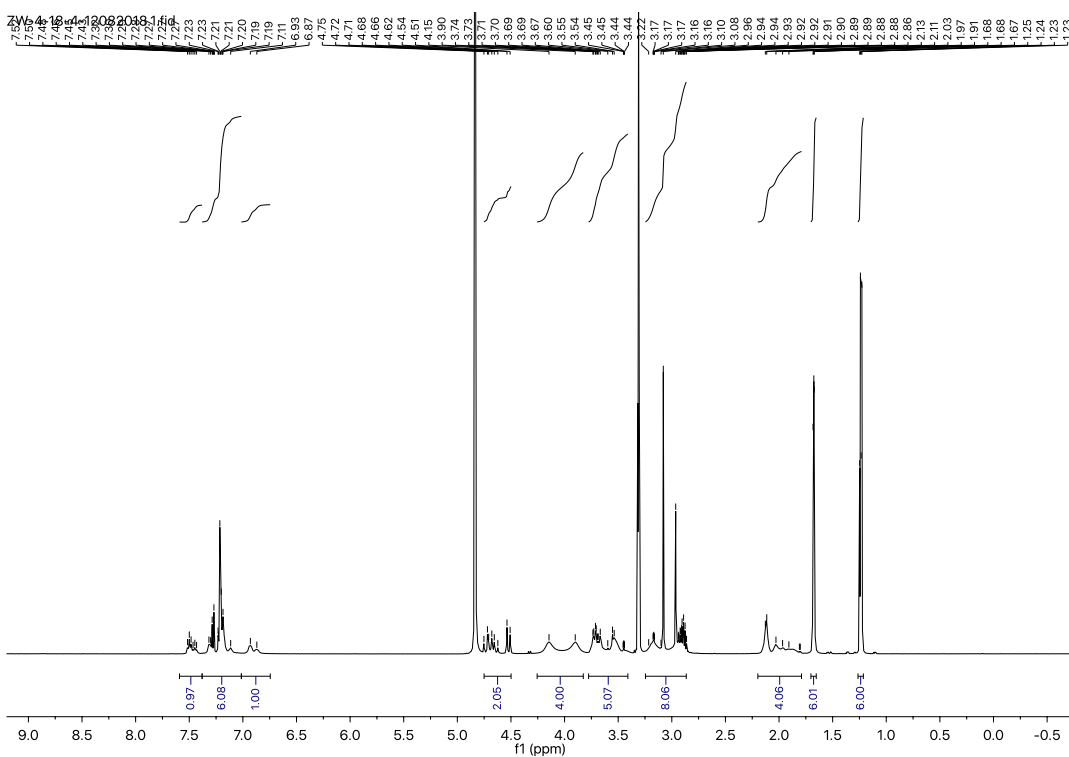
¹H NMR for compound 3



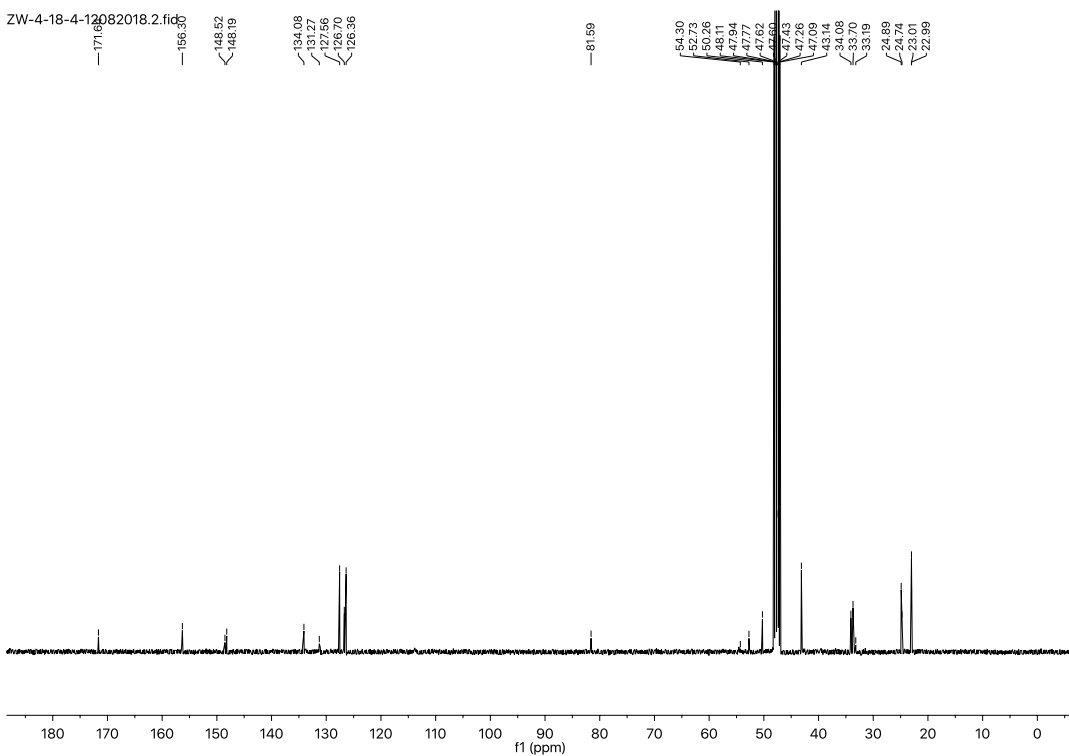
¹³C NMR for compound 3



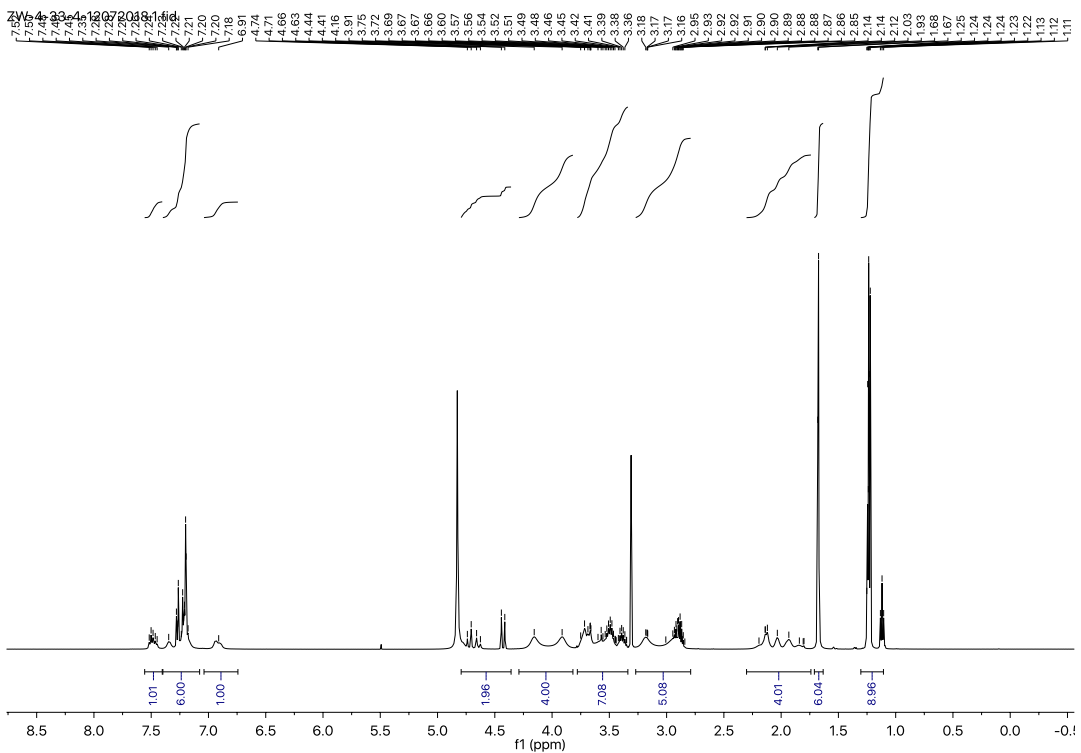
¹H NMR for compound 4



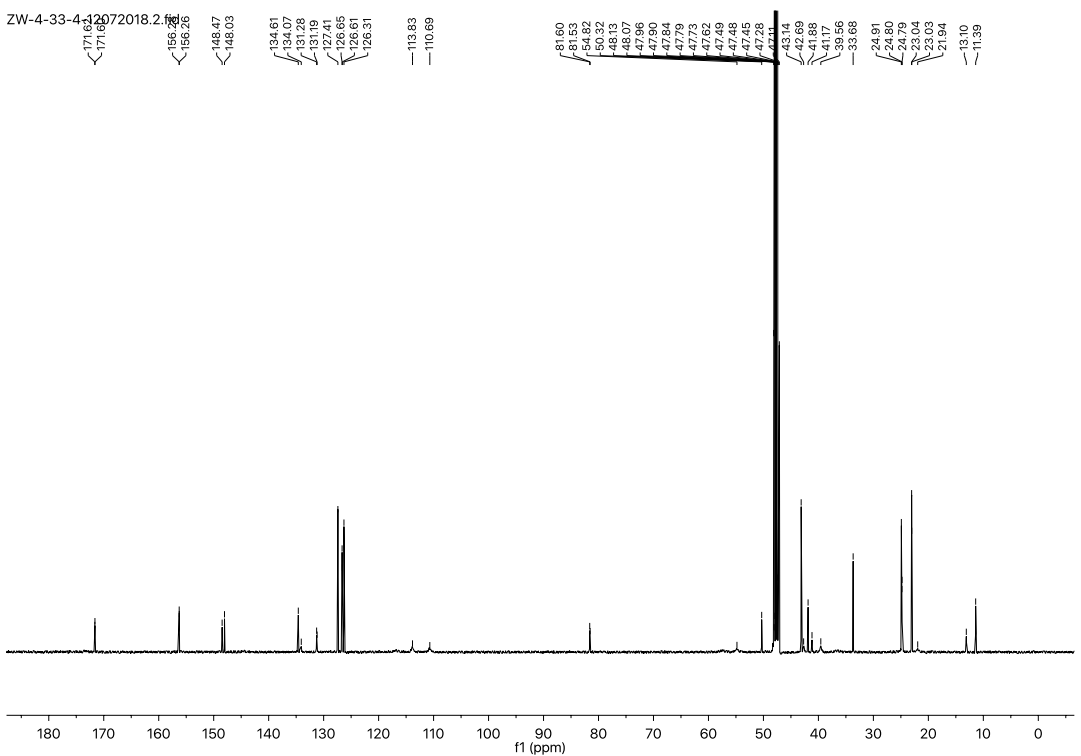
¹³C NMR for compound 4



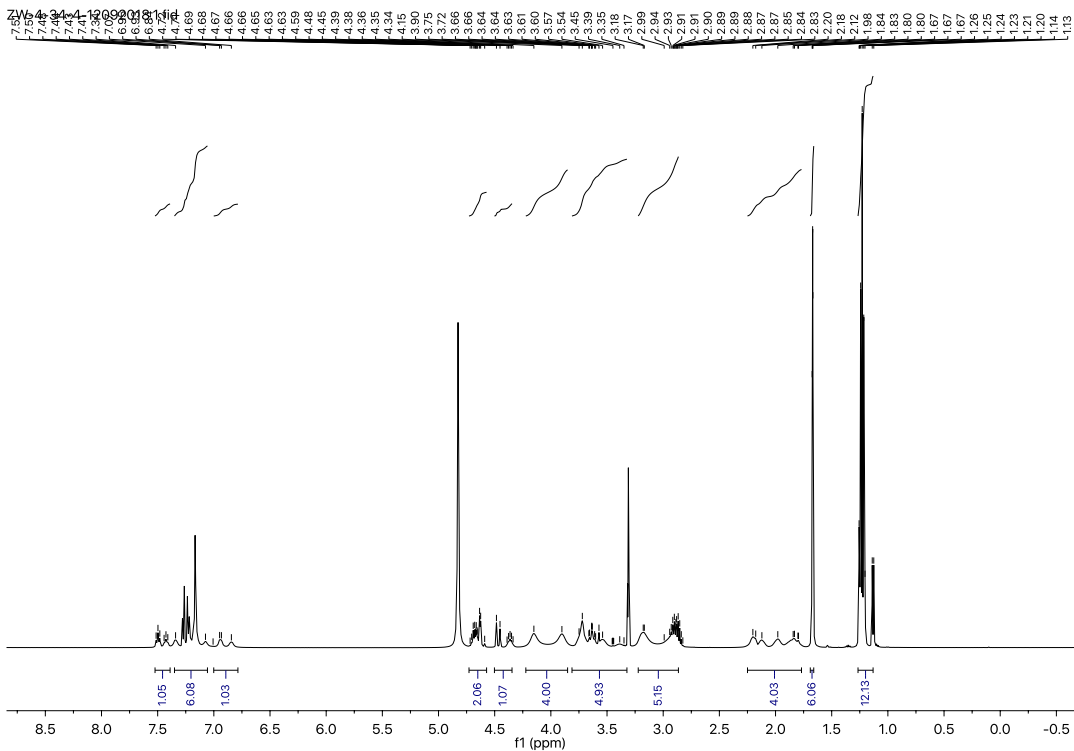
¹H NMR for compound 5



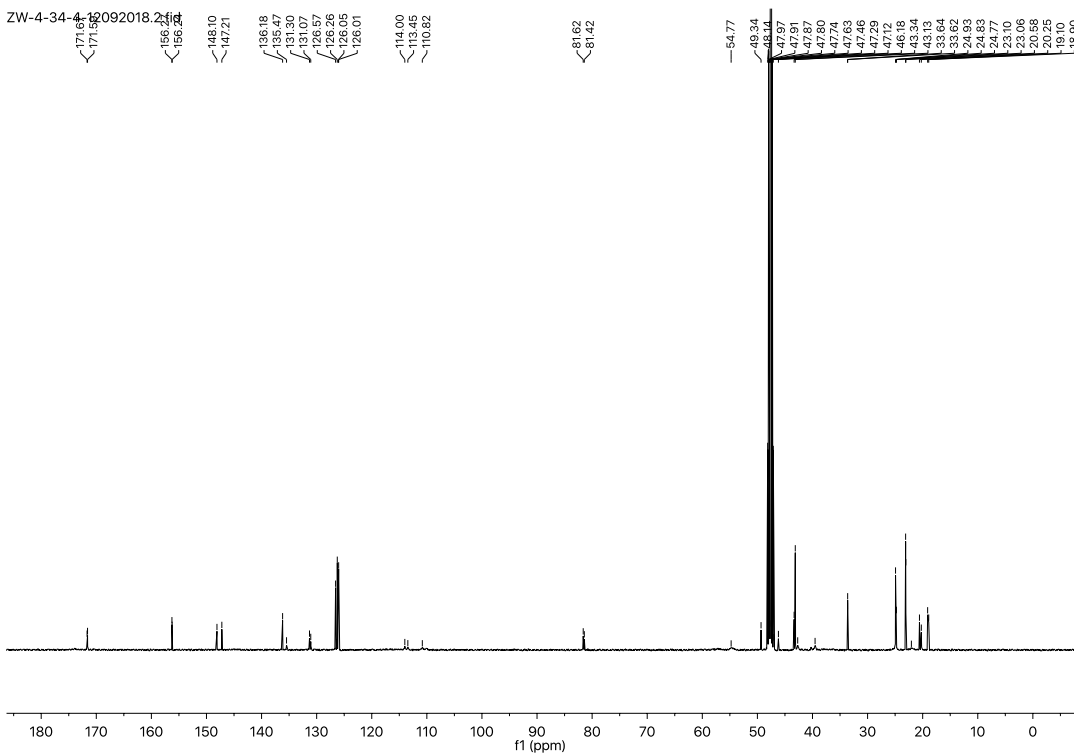
¹³C NMR for compound 5



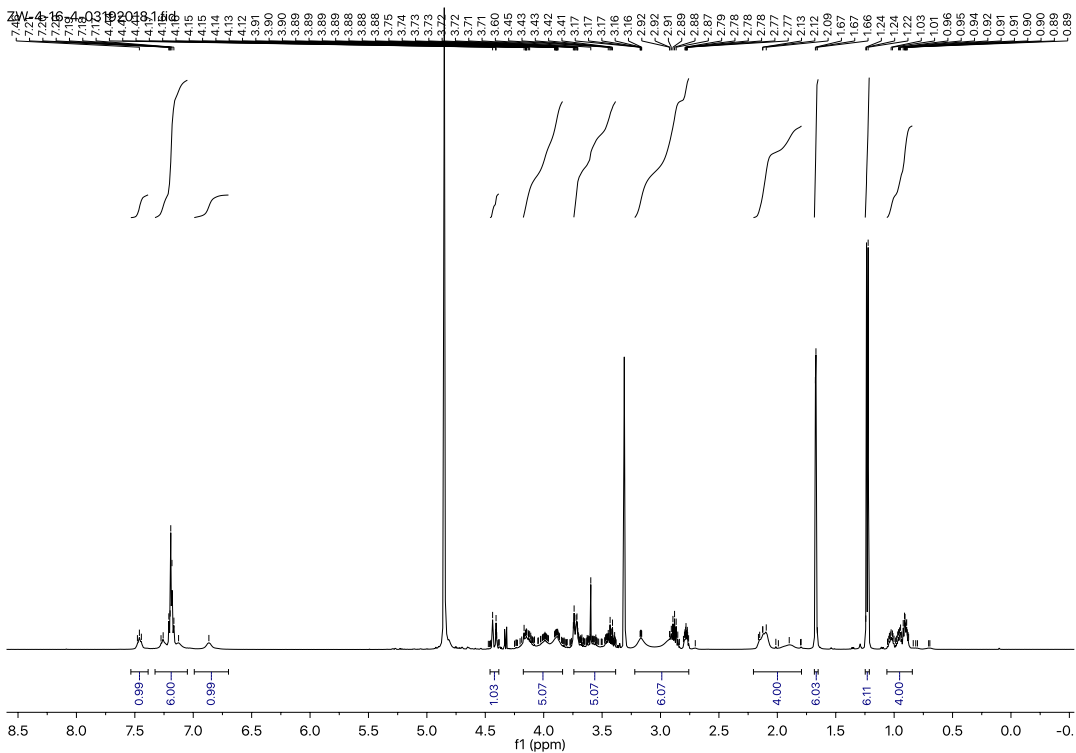
¹H NMR for compound 6



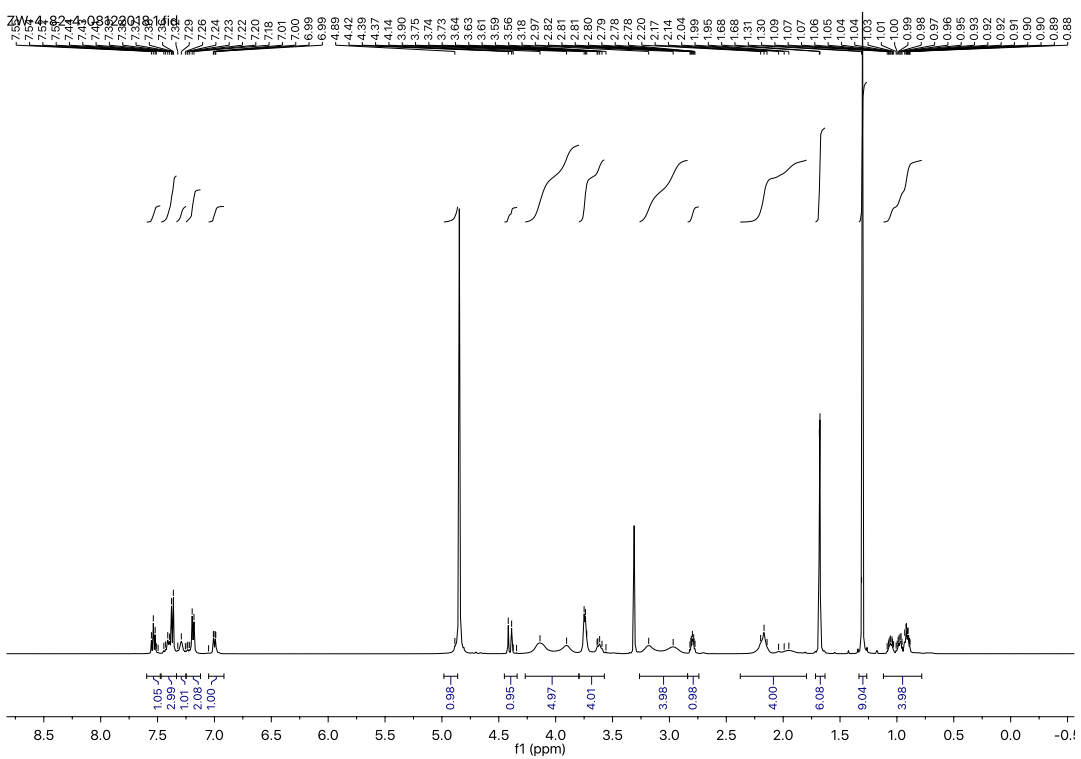
¹³C NMR for compound 6



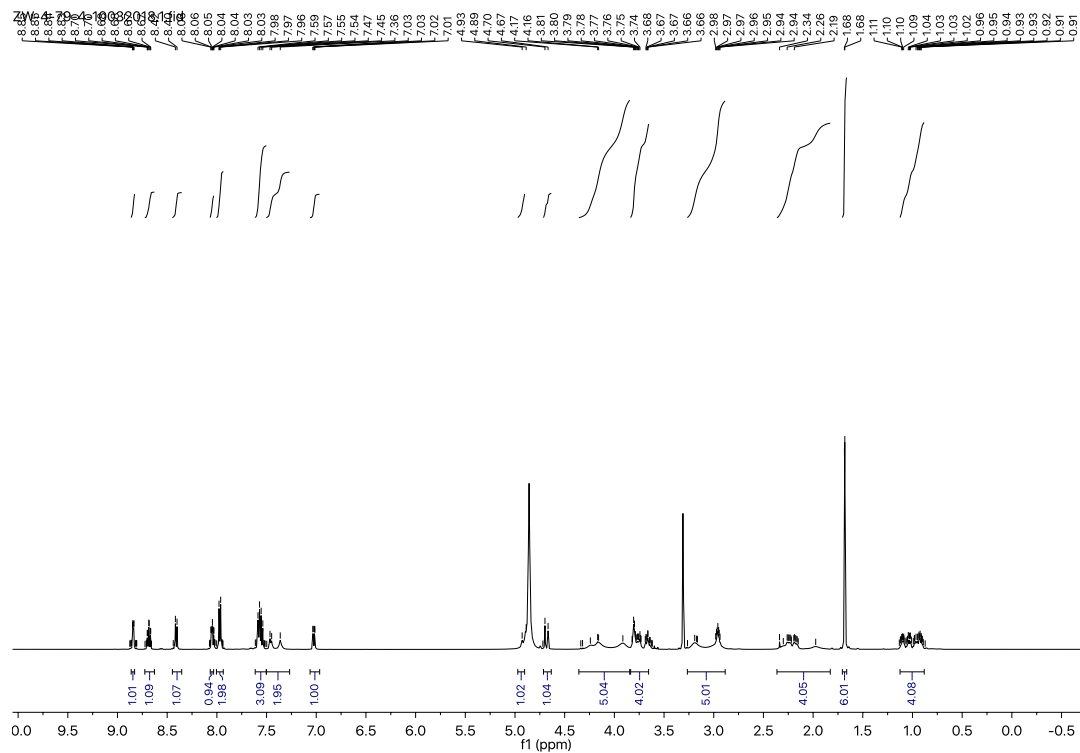
¹H NMR for compound 7



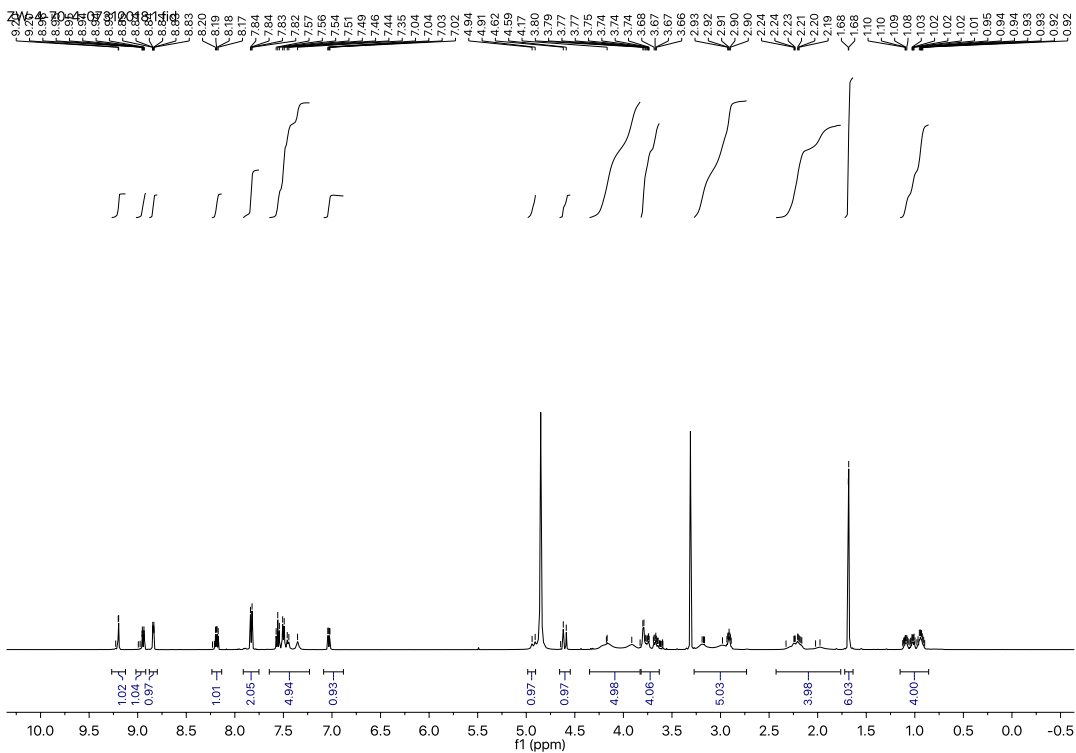
¹H NMR for compound **8**



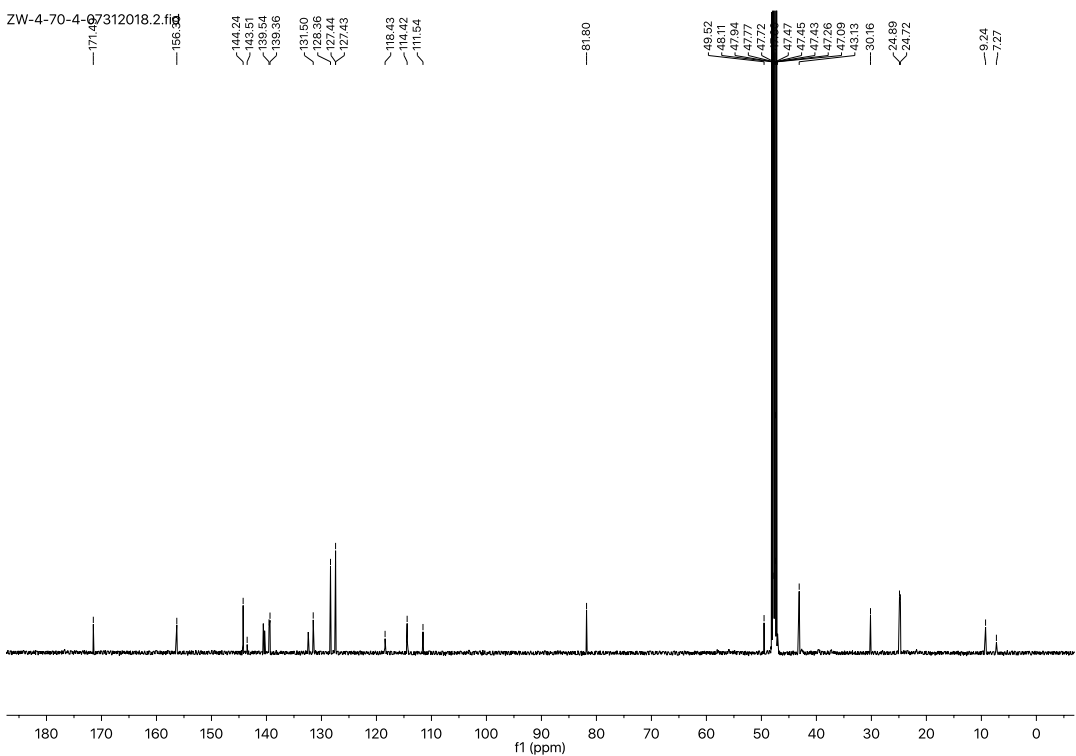
¹H NMR for compound 12



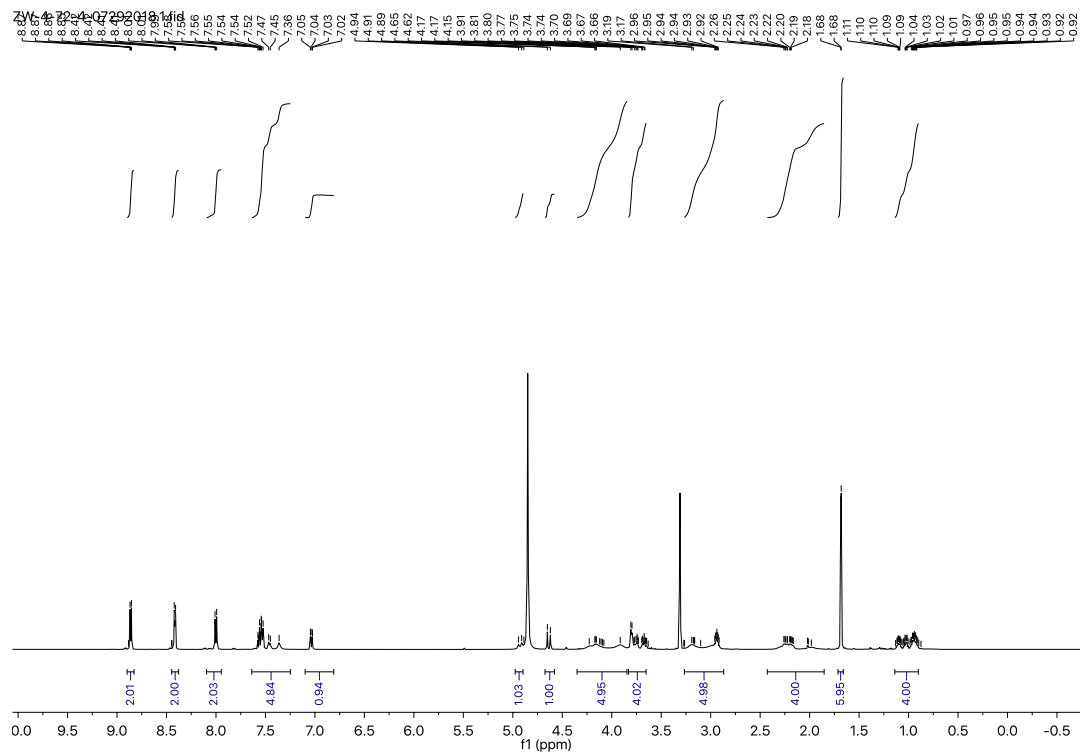
¹H NMR for compound 13



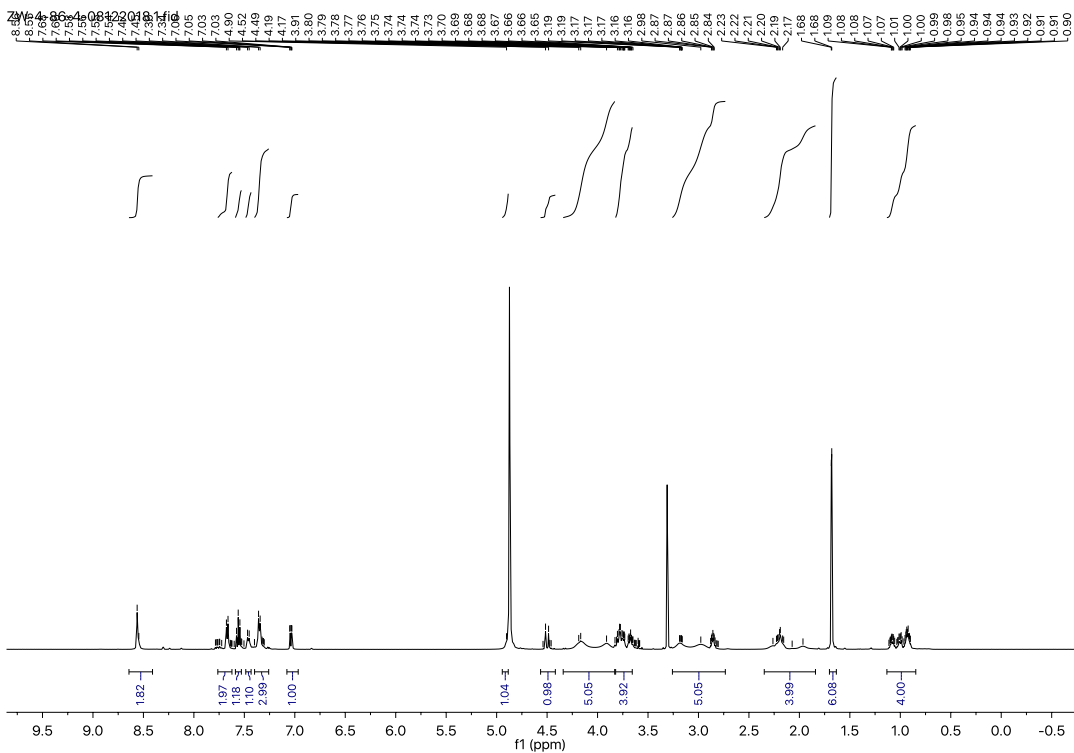
¹³C NMR for compound 13



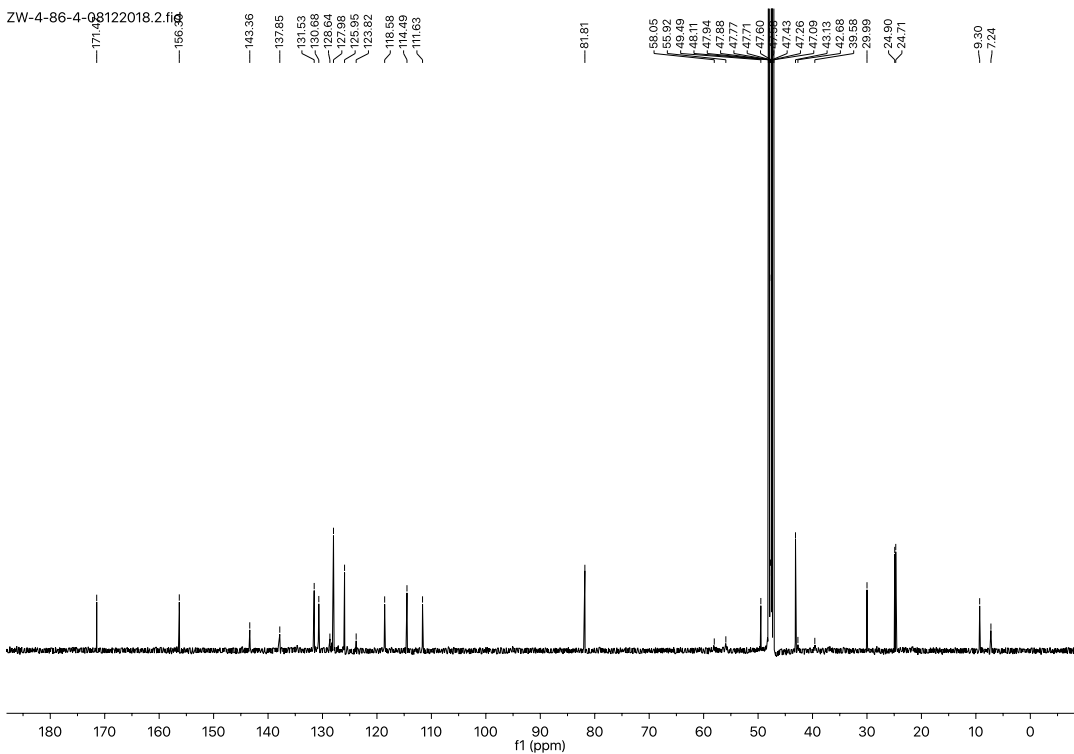
¹H NMR for compound 14



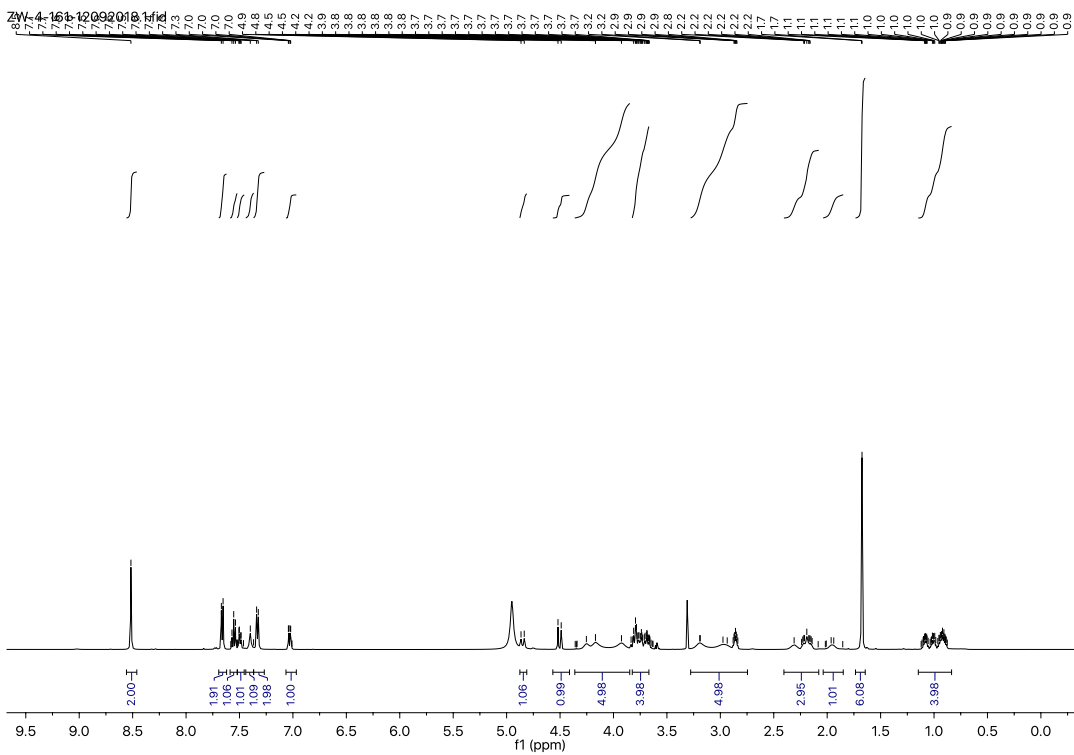
¹H NMR for compound 15



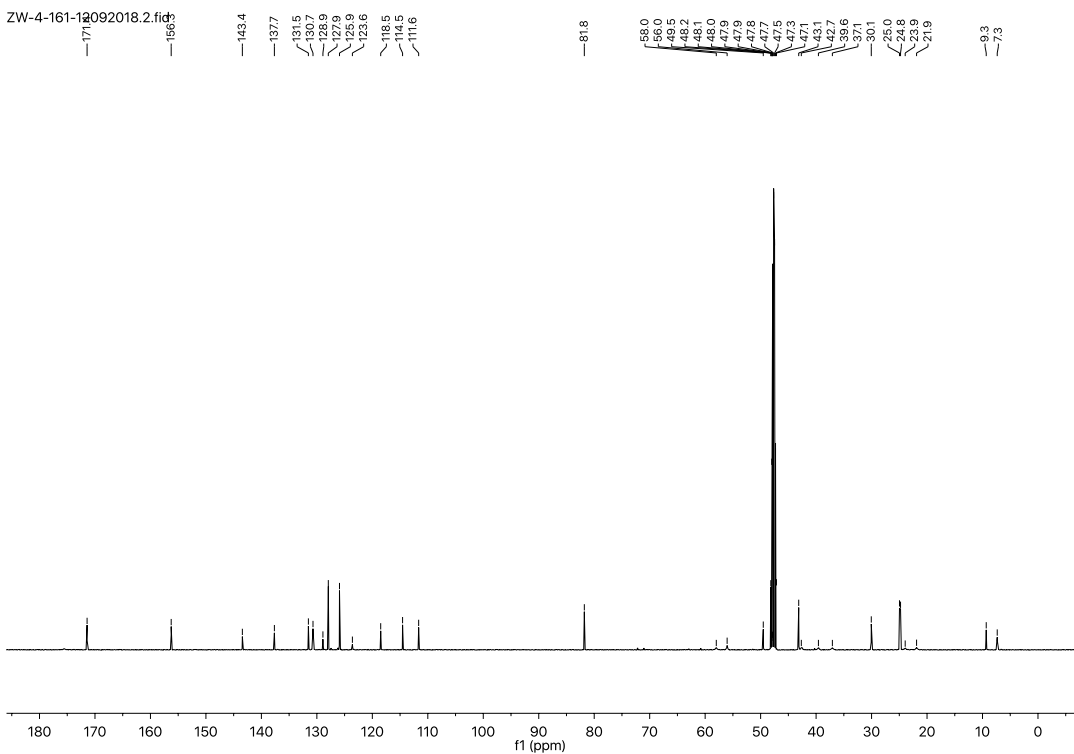
¹³C NMR for compound 15



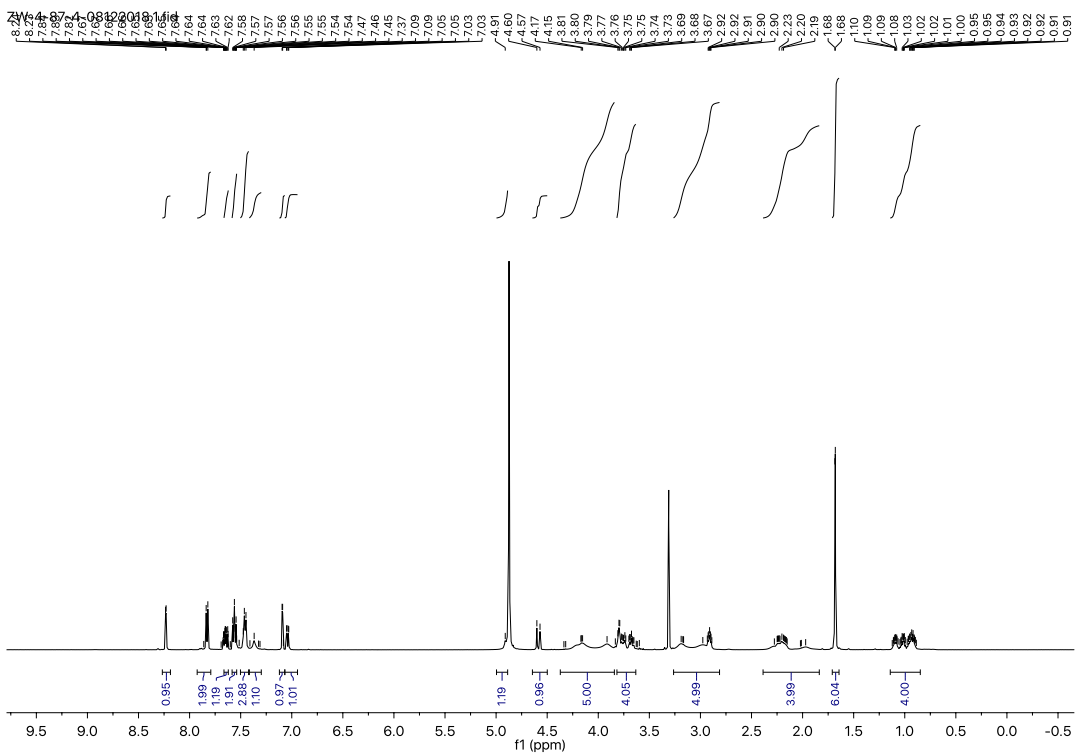
¹H NMR for compound 16



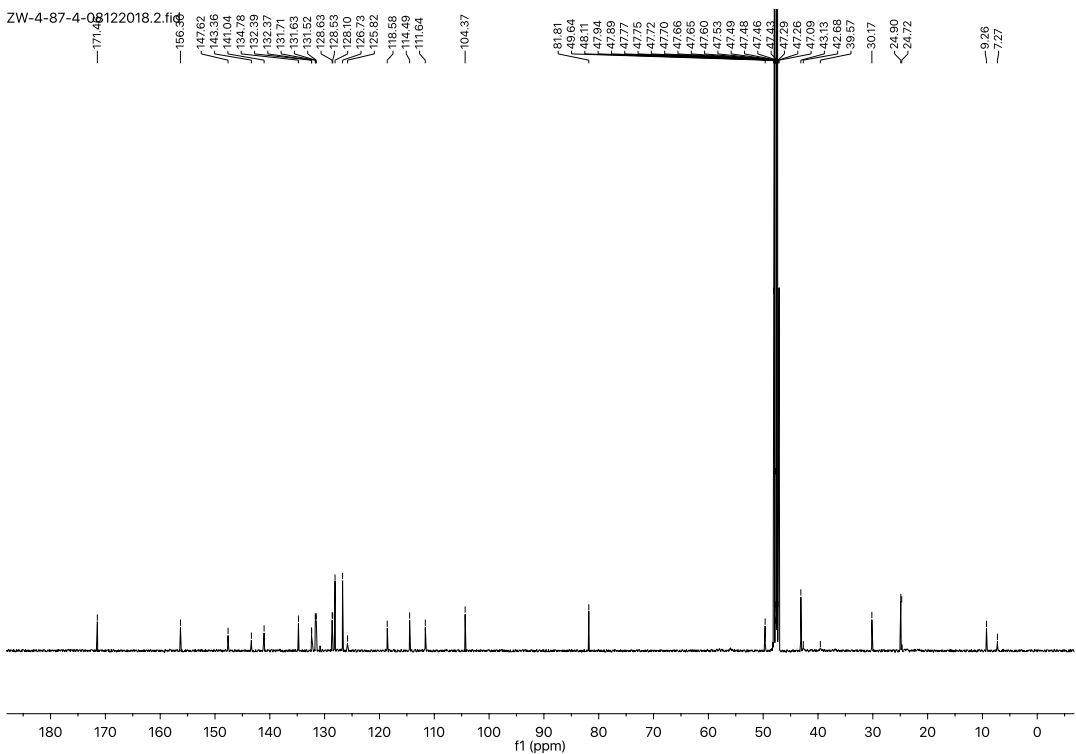
¹³C NMR for compound 16



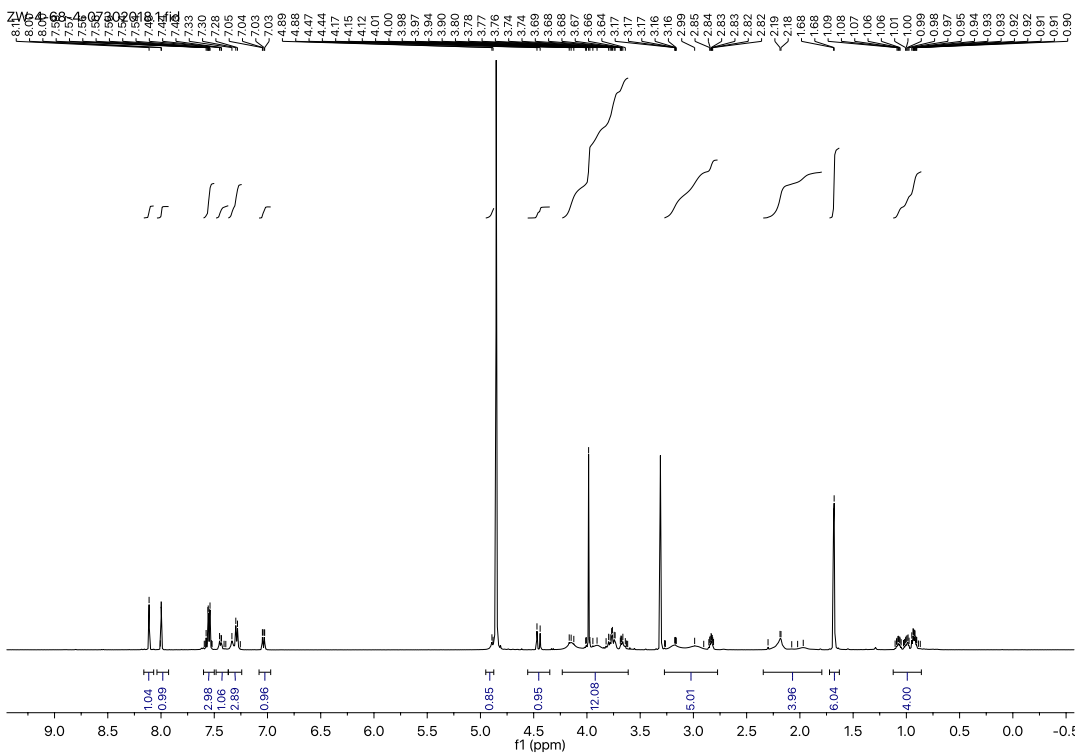
¹H NMR for compound 17



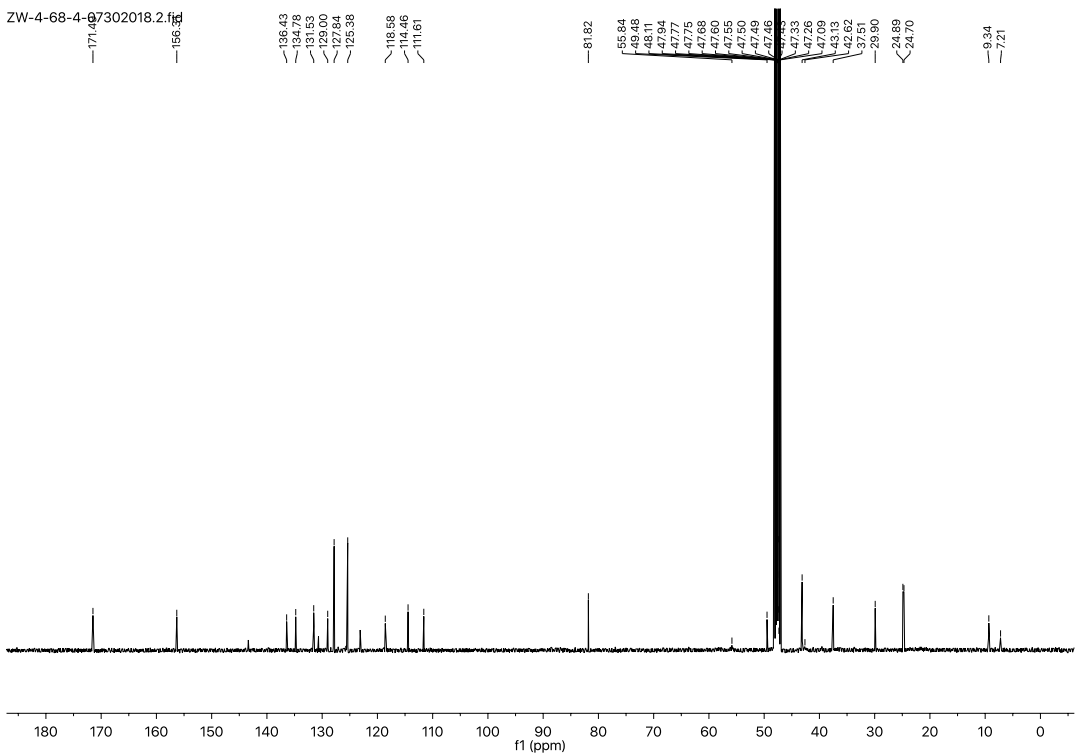
¹³C NMR for compound 17



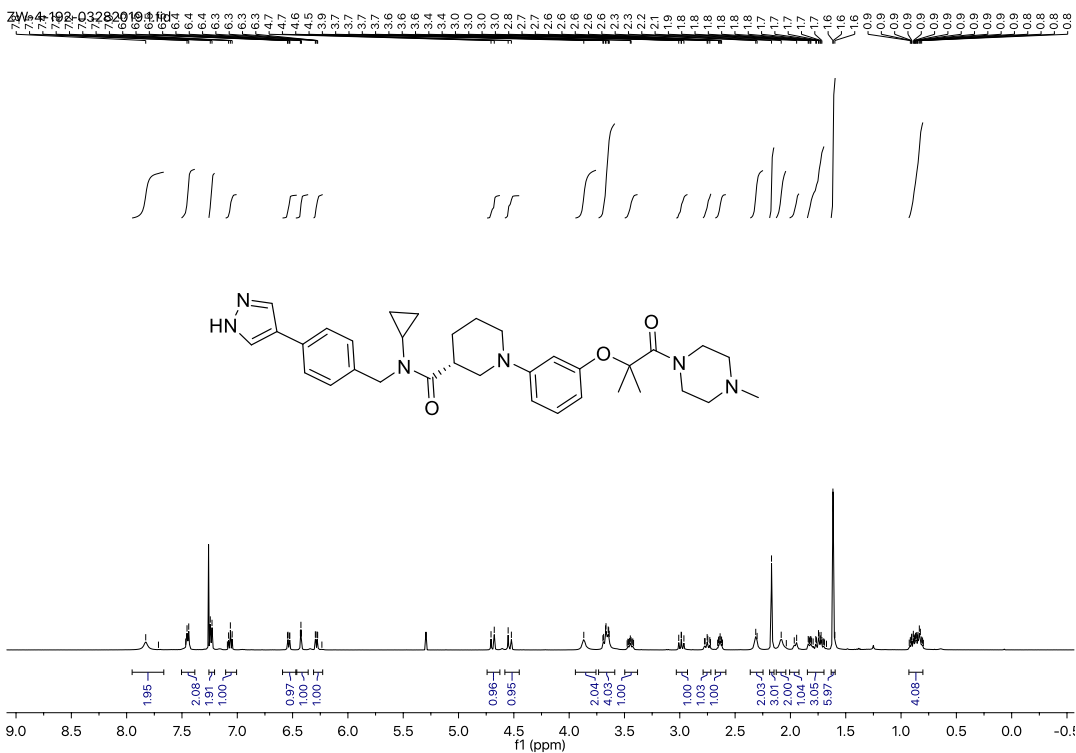
¹H NMR for compound 18



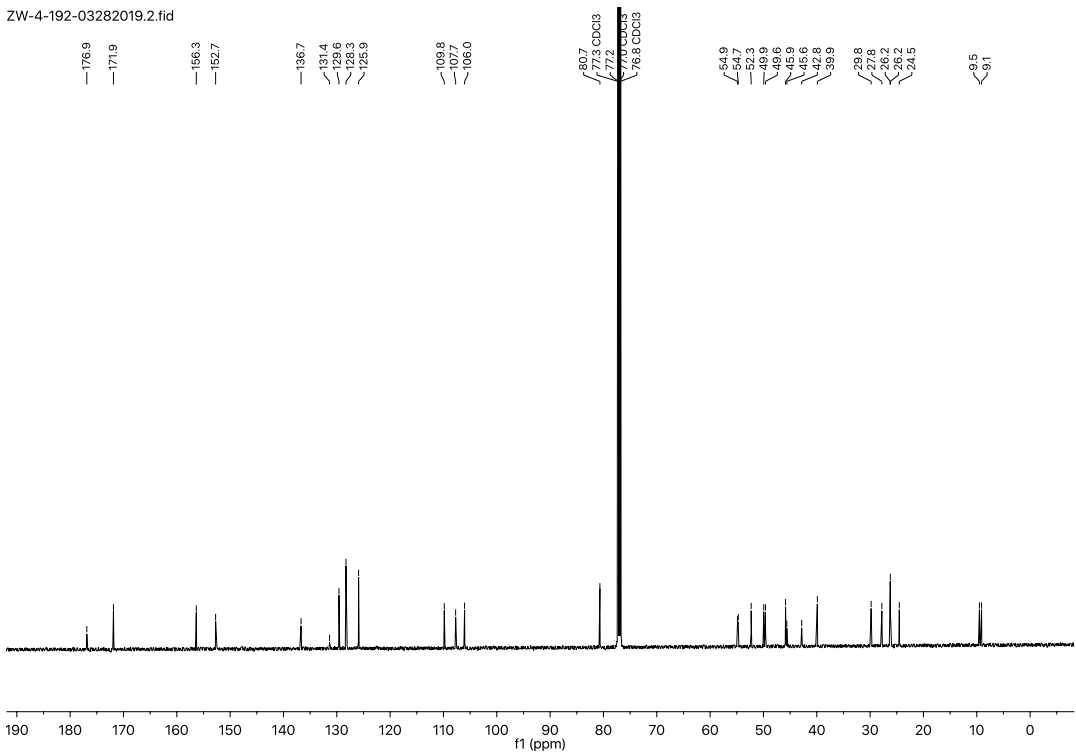
¹³C NMR for compound 18



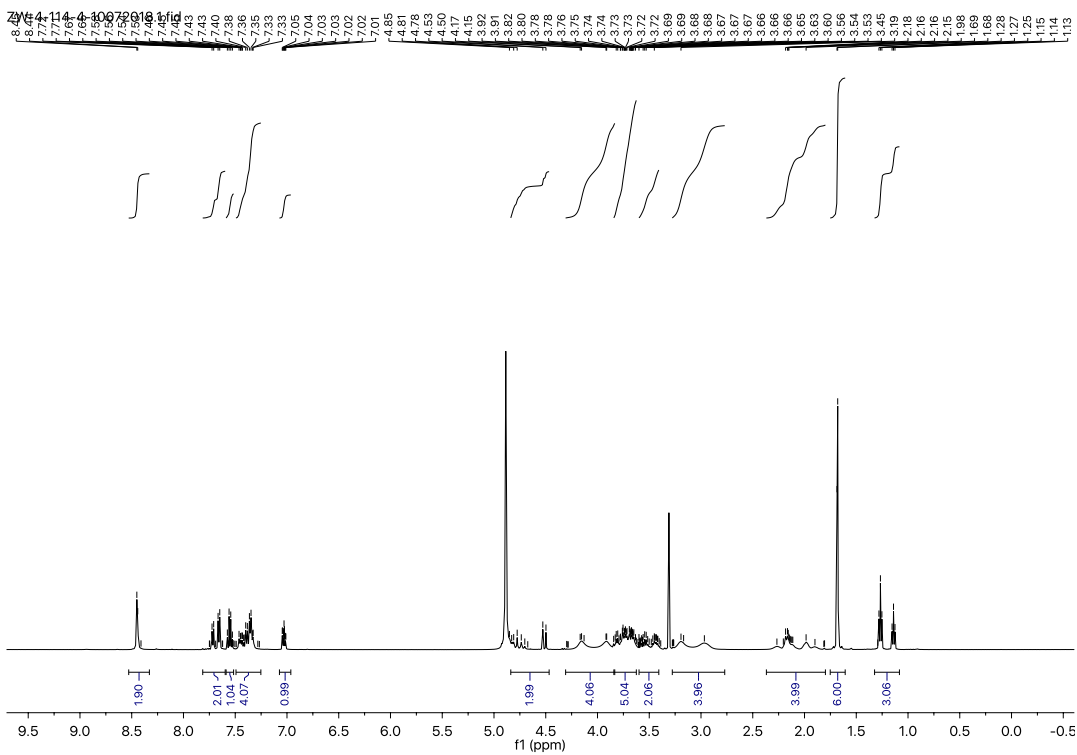
¹H NMR for compound 19



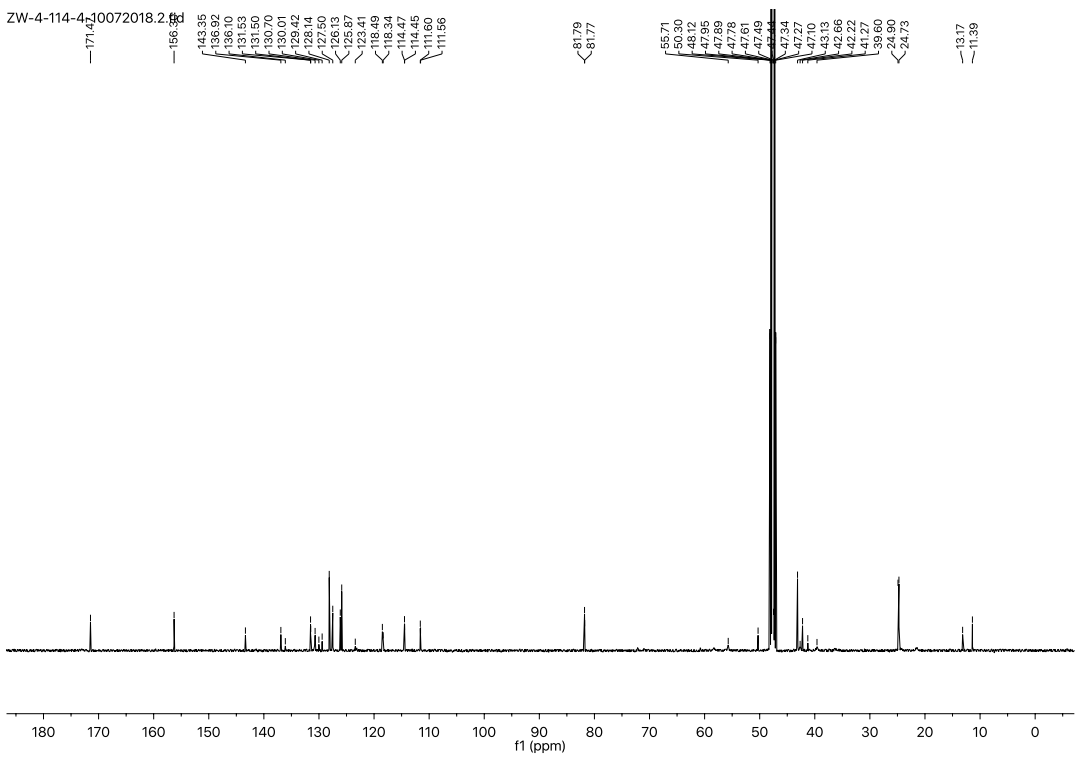
¹³C NMR for compound 19



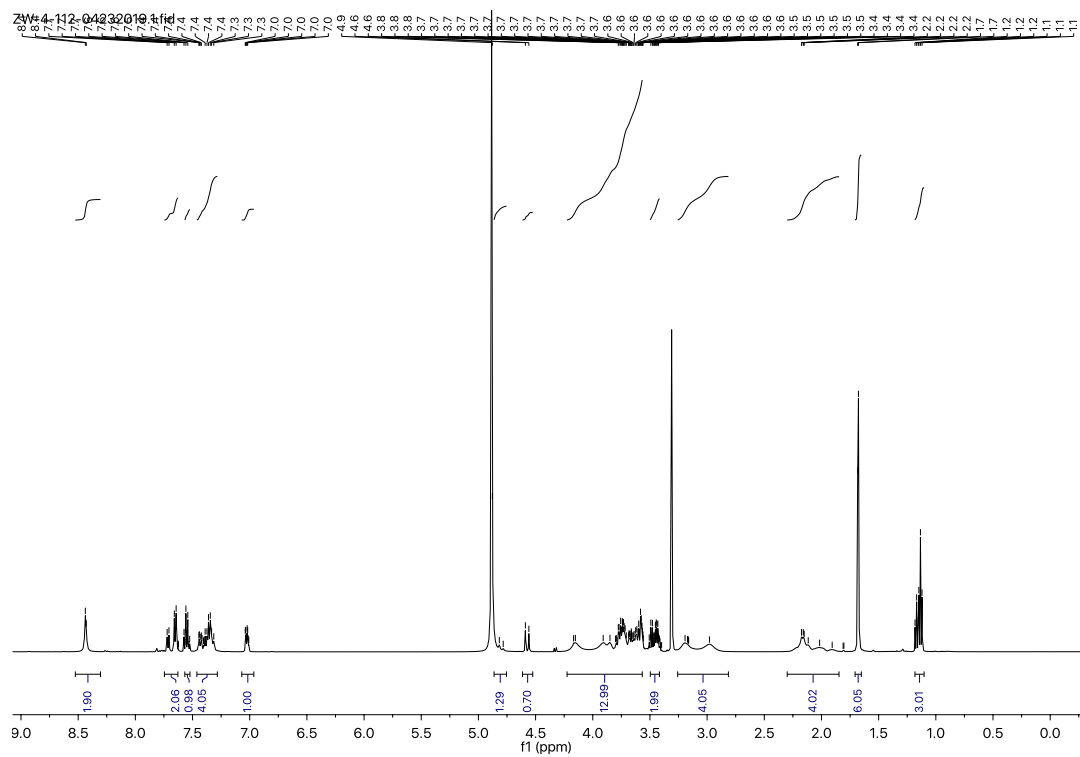
¹H NMR for compound 21



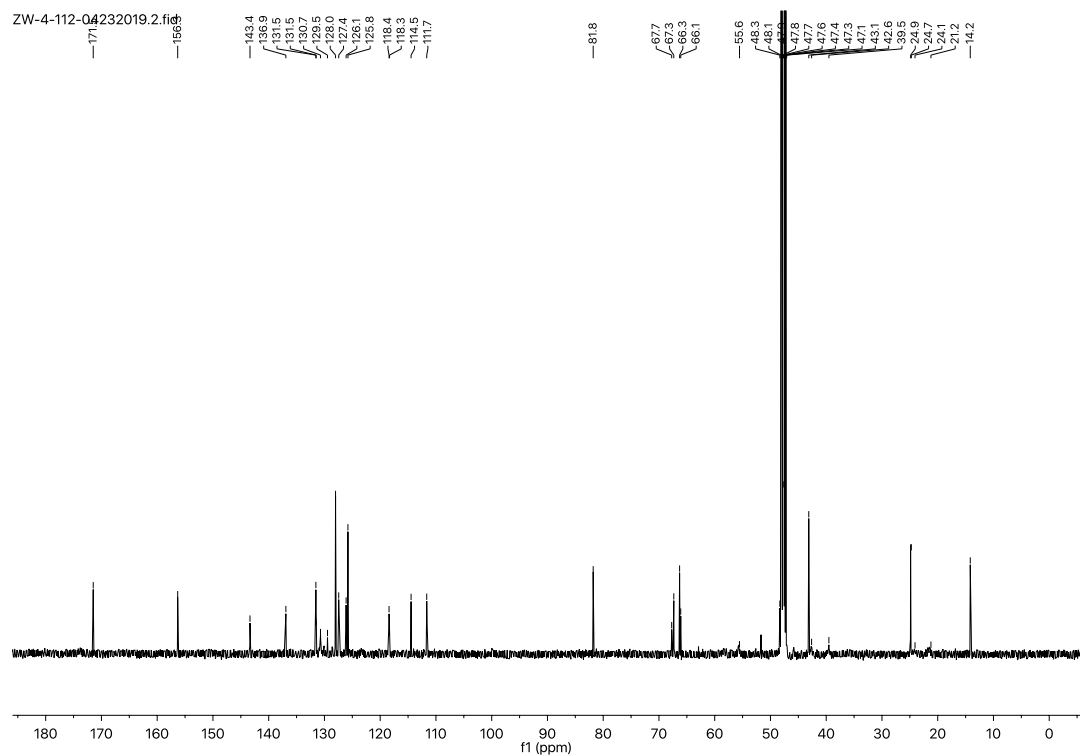
¹³C NMR for compound 21



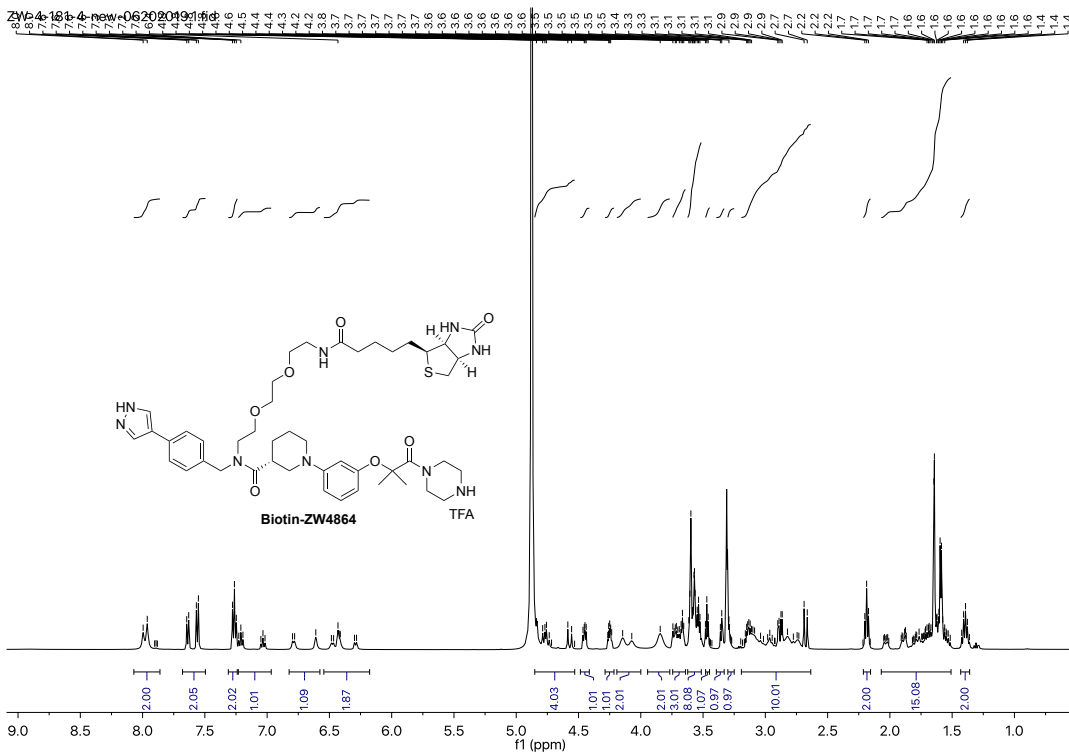
¹H NMR for compound 22



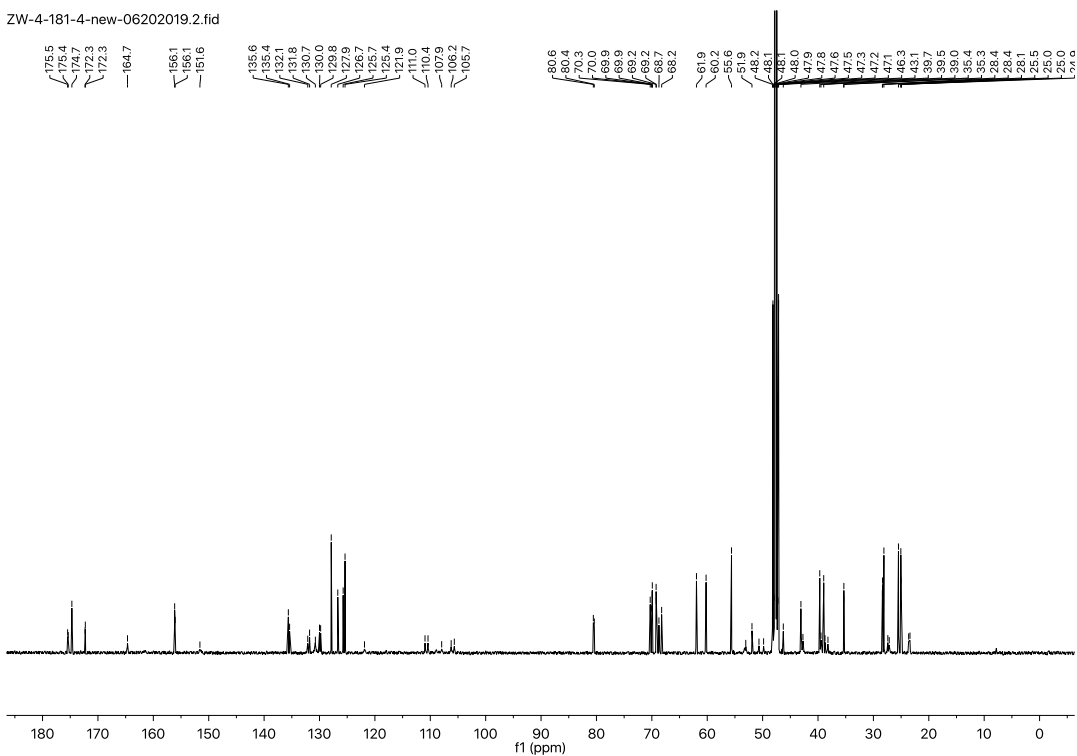
¹³C NMR for compound 22



¹H NMR for compound **Biotin-ZW4864**

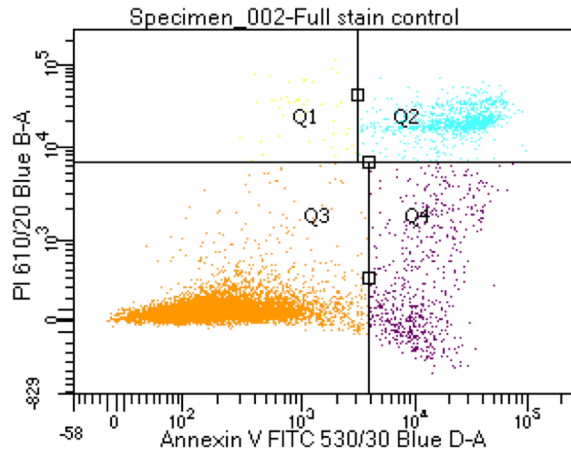
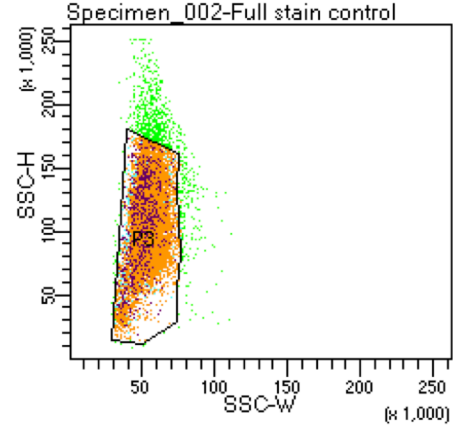
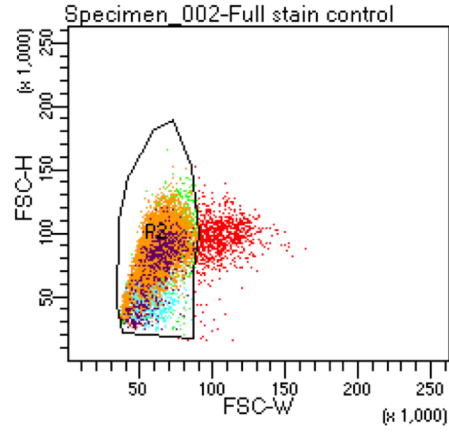
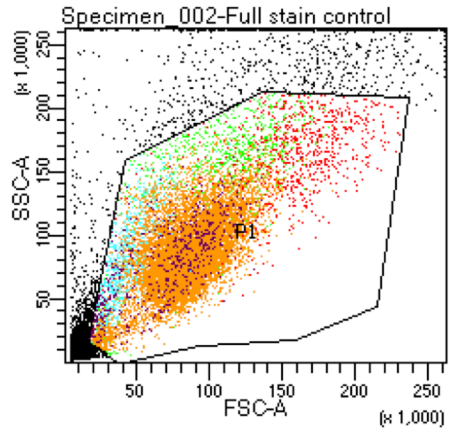


¹³C NMR for compound **Biotin-ZW4864**



MDA-MB-231 vehicle control

BD FACSDiva 8.0.1

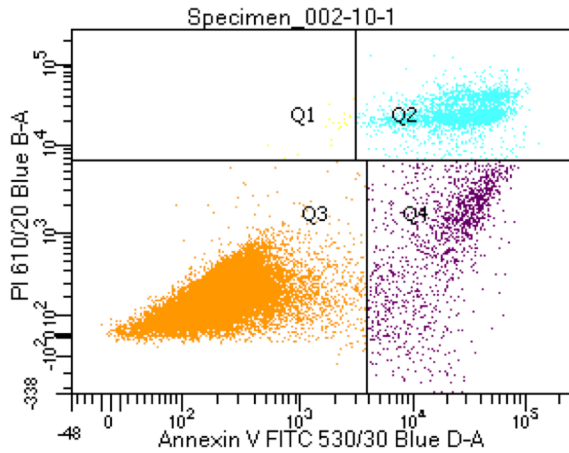
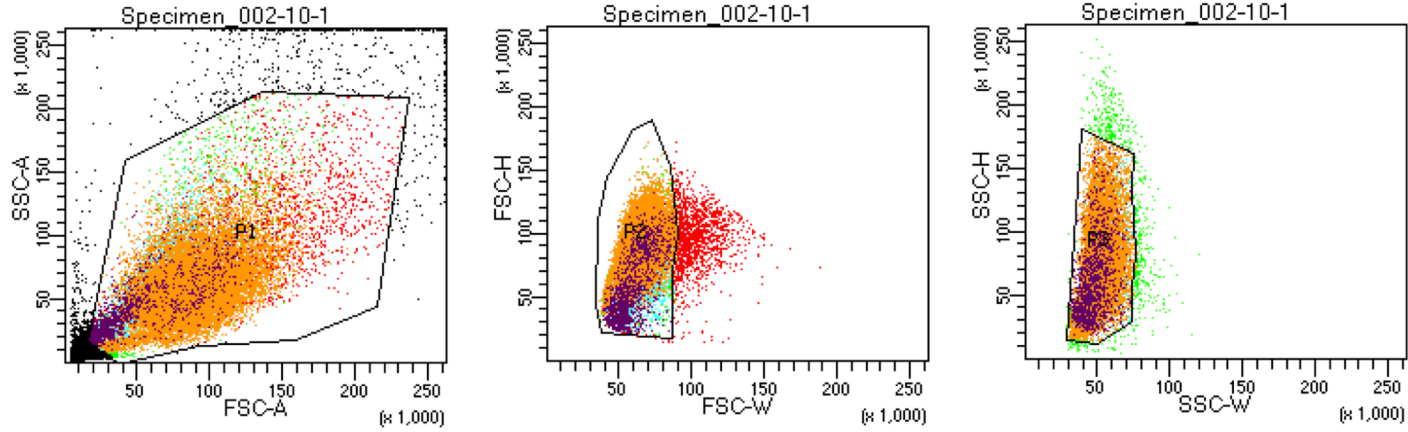


Tube: Full stain control

Population	#Events	%Parent	%Total
All Events	19,051	####	100.0
P1	13,269	69.6	69.6
P2	12,417	93.6	65.2
P3	11,557	93.1	60.7
Q1	45	0.4	0.2
Q2	1,019	8.8	5.3
Q3	9,883	85.5	51.9
Q4	610	5.3	3.2

MDA-MB-231 10 uM ZW4864

BD FACSDiva 8.0.1

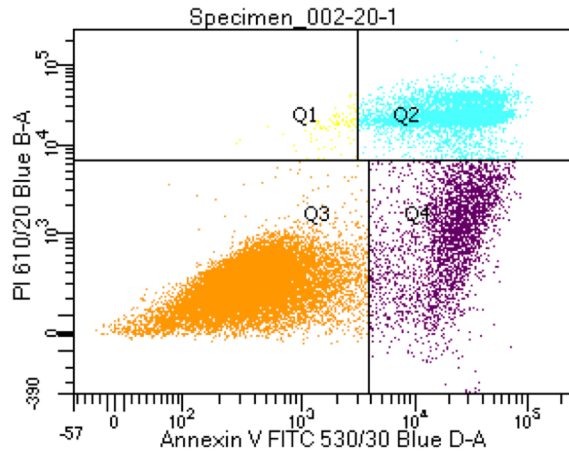
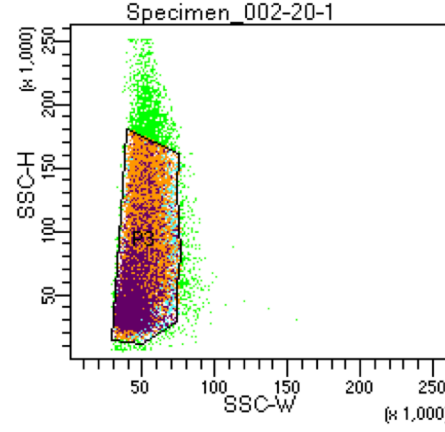
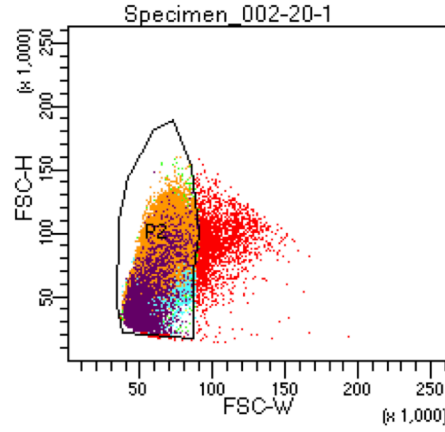
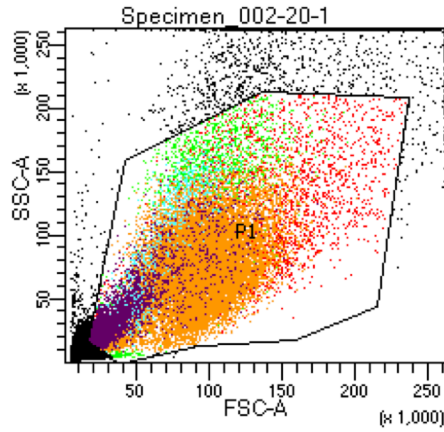


Tube: 10-1

Population	#Events	%Parent	%Total
All Events	35,804	####	100.0
P1	26,380	73.7	73.7
P2	24,902	94.4	69.6
P3	24,254	97.4	67.7
Q1	22	0.1	0.1
Q2	2,961	12.2	8.3
Q3	19,965	82.3	55.8
Q4	1,306	5.4	3.6

MDA-MB-231 20 uM ZW4864

BD FACSDiva 8.0.1

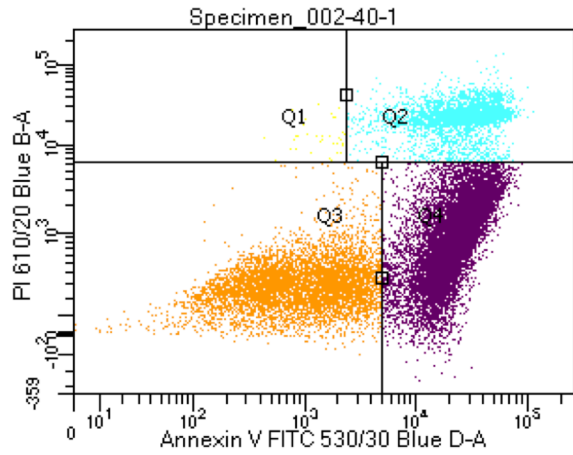
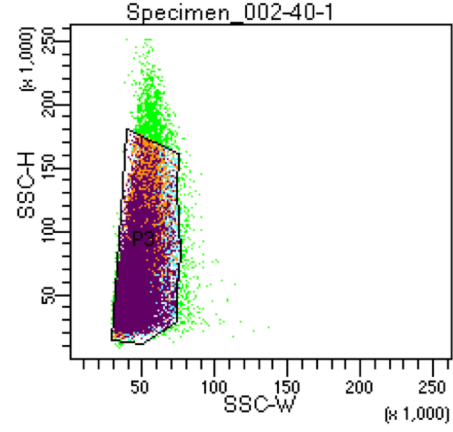
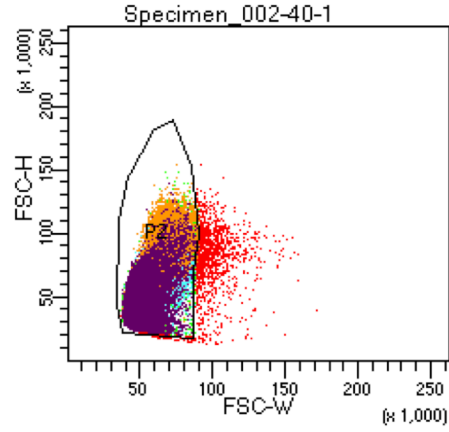
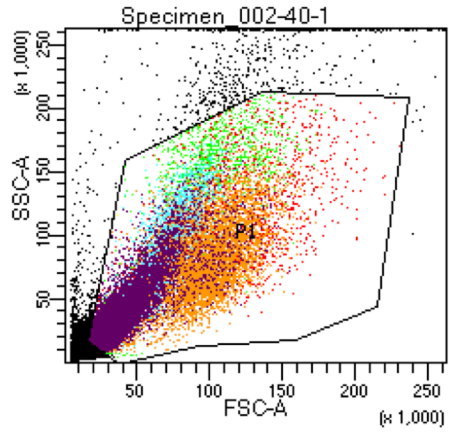


Tube: 20-1

Population	#Events	%Parent	%Total
All Events	43,344	####	100.0
P1	26,462	61.1	61.1
P2	24,116	91.1	55.6
P3	22,351	92.7	51.6
Q1	92	0.4	0.2
Q2	5,147	23.0	11.9
Q3	14,116	63.2	32.6
Q4	2,996	13.4	6.9

MDA-MB-231 40 uM ZW4864

BD FACSDiva 8.0.1

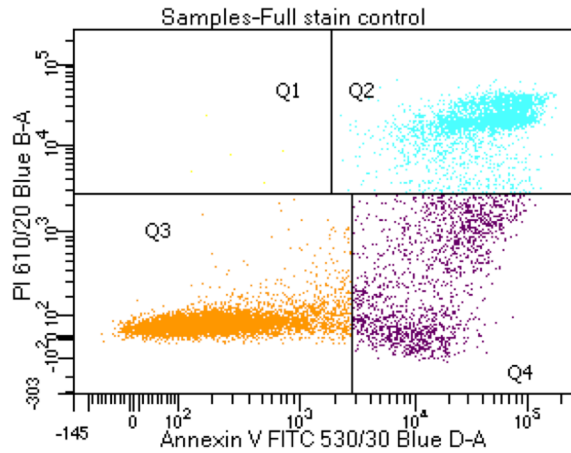
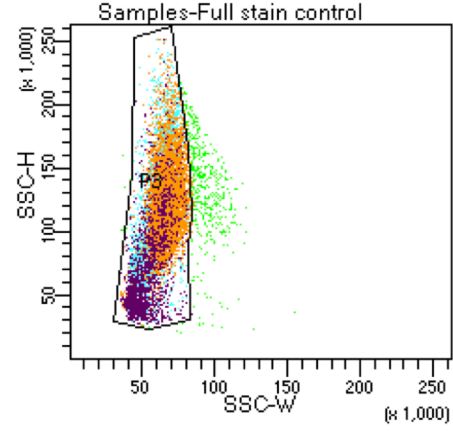
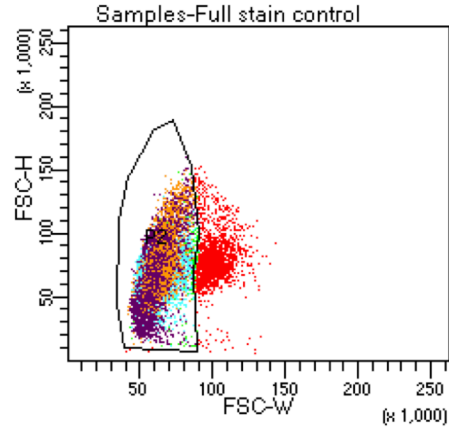
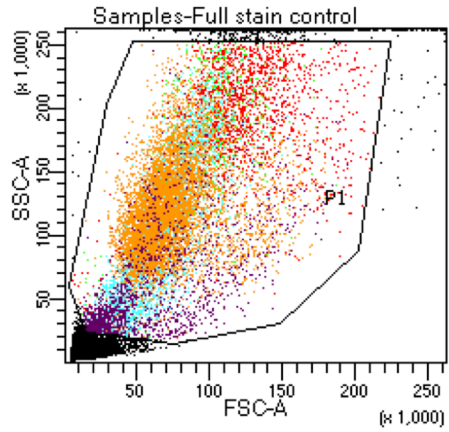


Tube: 40-1

Population	#Events	%Parent	%Total
All Events	76,090	####	100.0
P1	25,557	33.6	33.6
P2	24,118	94.4	31.7
P3	22,555	93.5	29.6
Q1	35	0.2	0.0
Q2	4,252	18.9	5.6
Q3	6,281	27.8	8.3
Q4	11,987	53.1	15.8

MDA-MB-468 vehicle control

BD FACSDiva 8.0.1

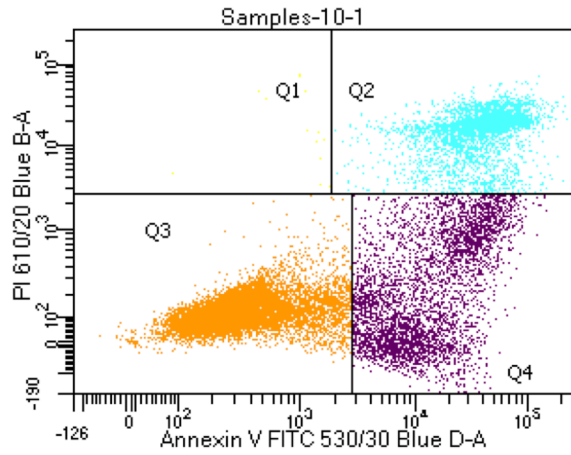
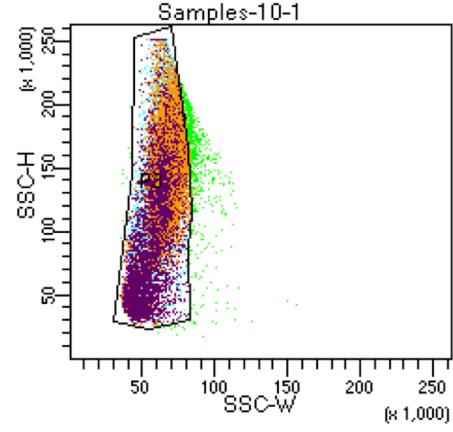
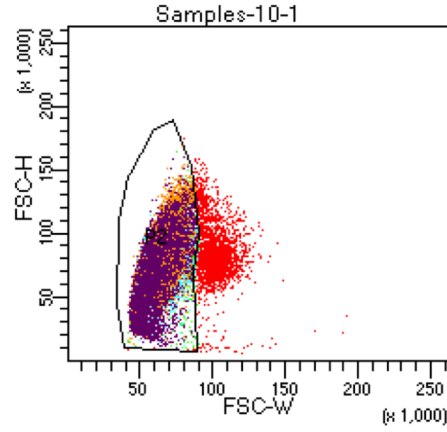
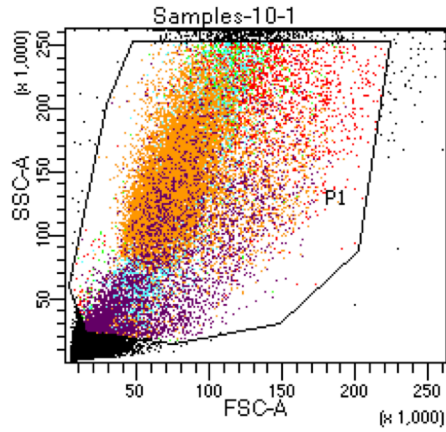


Tube: Full stain control

Population	#Events	%Parent	%Total
All Events	22,537	####	100.0
P1	13,648	60.6	60.6
P2	11,824	86.6	52.5
P3	11,370	96.2	50.5
Q1	5	0.0	0.0
Q2	3,288	28.9	14.6
Q3	6,492	57.1	28.8
Q4	1,585	13.9	7.0

MDA-MB-468 10 uM ZW4864

BD FACSDiva 8.0.1

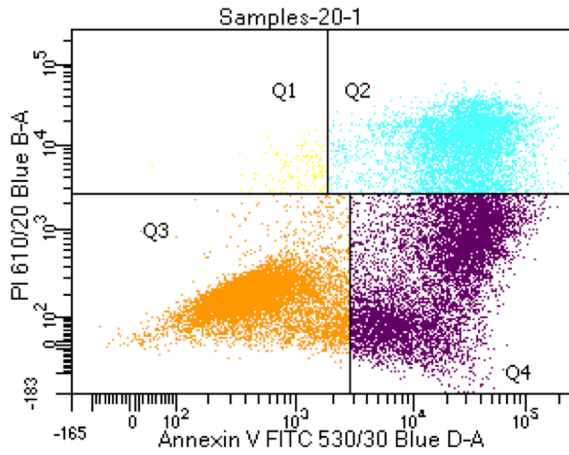
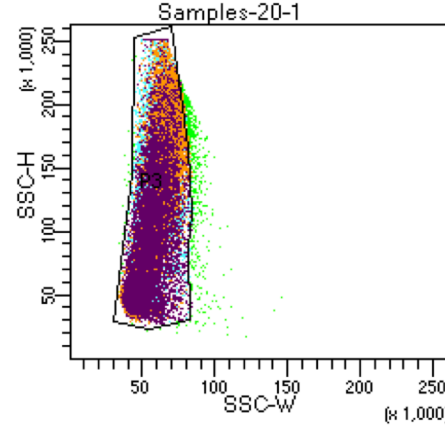
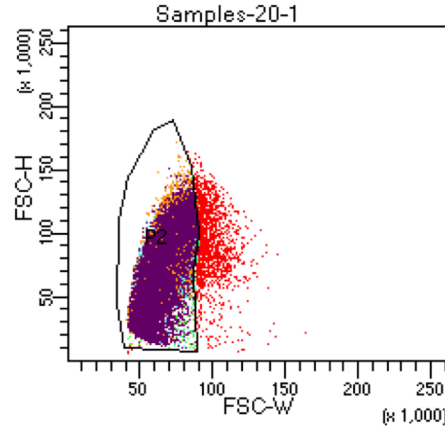
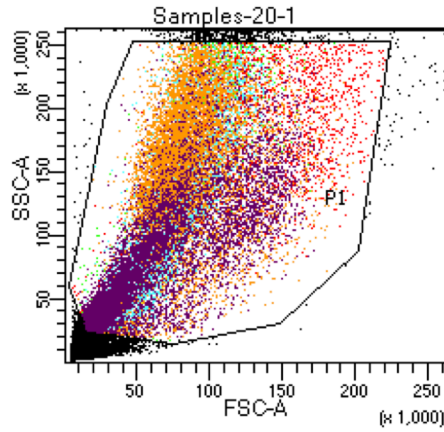


Tube: 10-1

Population	#Events	%Parent	%Total
All Events	44,726	####	100.0
P1	25,283	56.5	56.5
P2	22,969	90.8	51.4
P3	22,306	97.1	49.9
Q1	13	0.1	0.0
Q2	3,925	17.6	8.8
Q3	14,460	64.8	32.3
Q4	3,908	17.5	8.7

MDA-MB-468 20 uM ZW4864

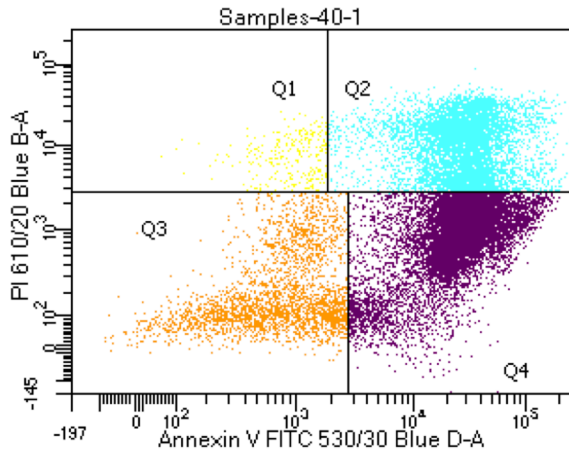
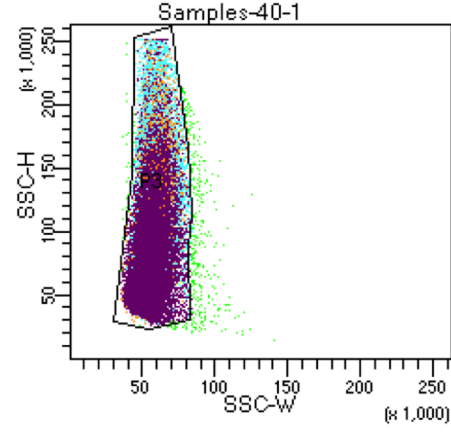
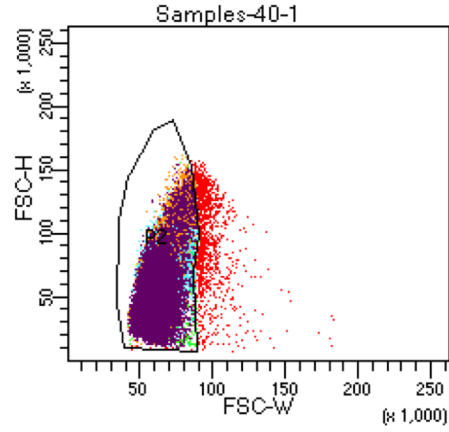
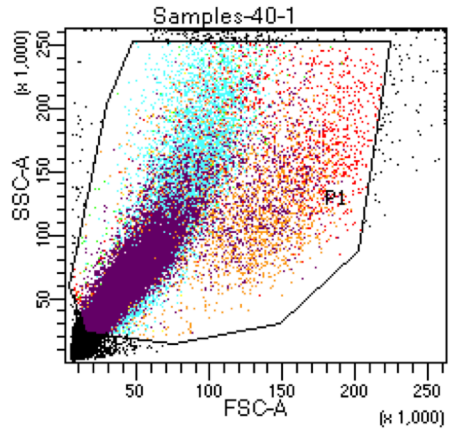
BD FACSDiva 8.0.1



Tube: 20-1			
Population	#Events	%Parent	%Total
All Events	46,880	####	100.0
P1	25,581	54.6	54.6
P2	23,954	93.6	51.1
P3	23,353	97.5	49.8
Q1	126	0.5	0.3
Q2	5,735	24.6	12.2
Q3	9,406	40.3	20.1
Q4	8,086	34.6	17.2

MDA-MB-468 40 uM ZW4864

BD FACSDiva 8.0.1

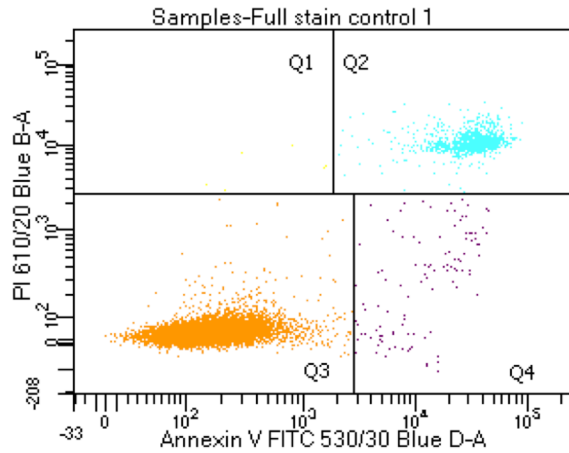
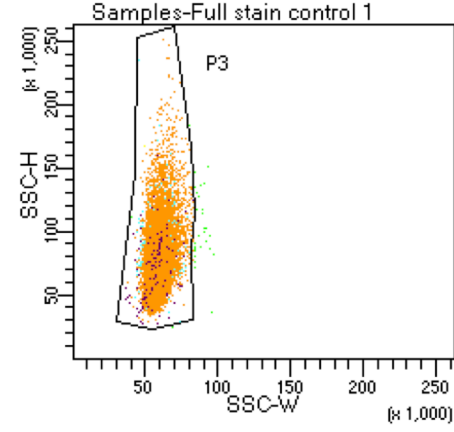
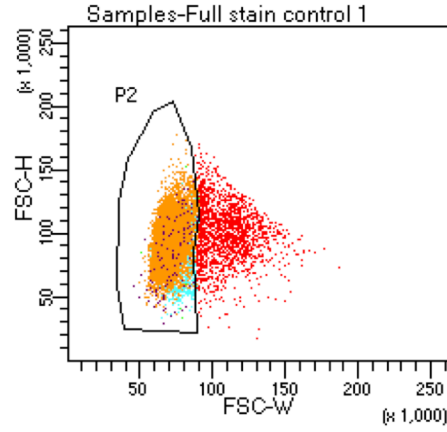
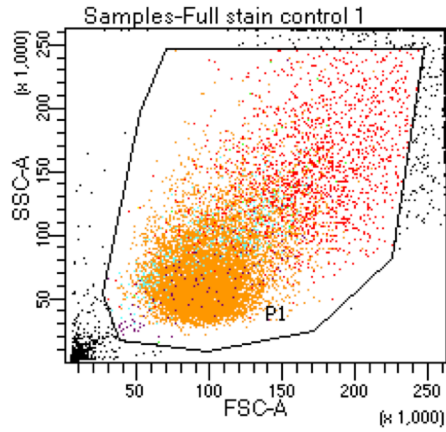


Tube: 40-1

Population	#Events	%Parent	%Total
All Events	40,946	####	100.0
P1	24,454	59.7	59.7
P2	23,221	95.0	56.7
P3	22,813	98.2	55.7
Q1	281	1.2	0.7
Q2	7,609	33.4	18.6
Q3	2,712	11.9	6.6
Q4	12,211	53.5	29.8

MCF10A vehicle control

BD FACSDiva 8.0.1

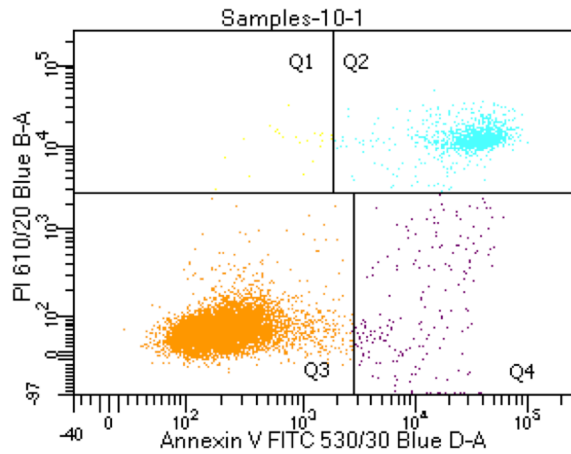
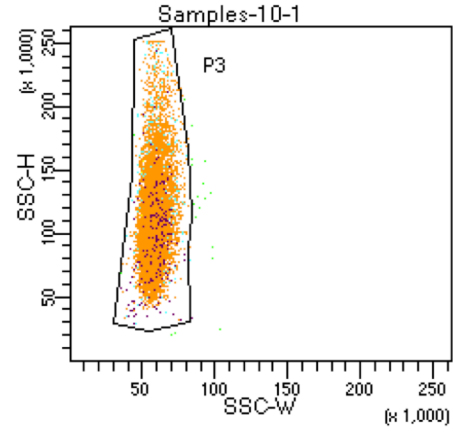
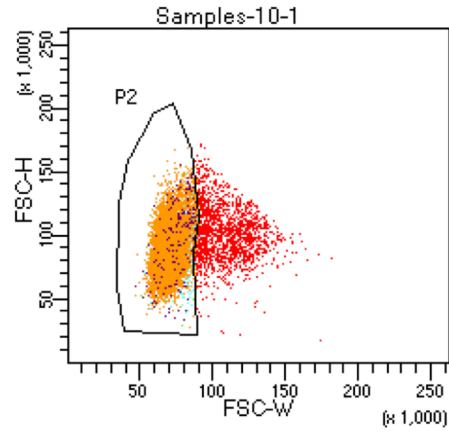
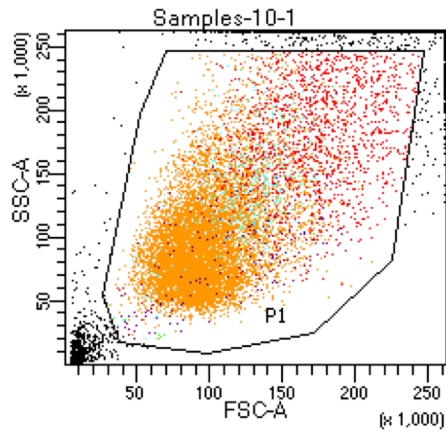


Tube: Full stain control 1

Population	#Events	%Parent	%Total
All Events	13,566	####	100.0
P1	11,809	87.0	87.0
P2	10,036	85.0	74.0
P3	10,000	99.6	73.7
Q1	6	0.1	0.0
Q2	1,263	12.6	9.3
Q3	8,620	86.2	63.5
Q4	111	1.1	0.8

MCF10A 10 uM ZW4864

BD FACSDiva 8.0.1

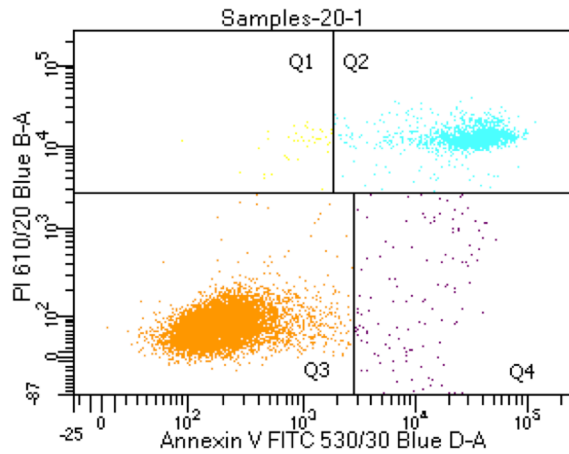
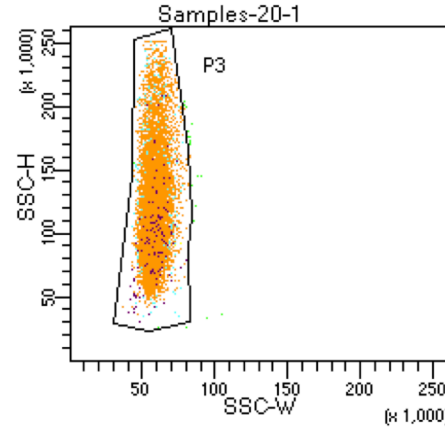
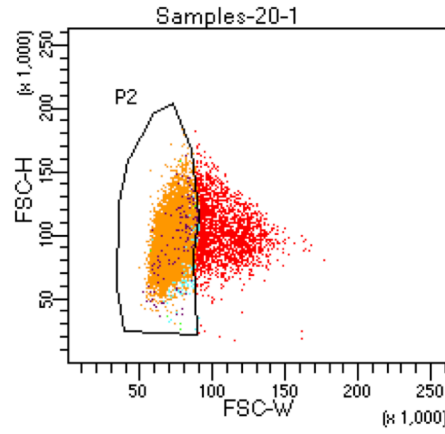
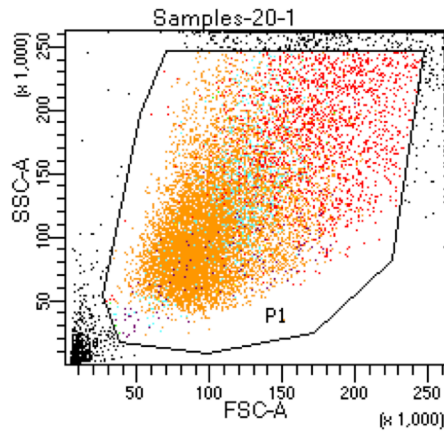


Tube: 10-1

Population	#Events	%Parent	%Total
All Events	14,805	####	100.0
P1	11,754	79.4	79.4
P2	10,102	85.9	68.2
P3	10,082	99.8	68.1
Q1	24	0.2	0.2
Q2	1,238	12.3	8.4
Q3	8,628	85.6	58.3
Q4	192	1.9	1.3

MCF10A 20 uM ZW4864

BD FACSDiva 8.0.1

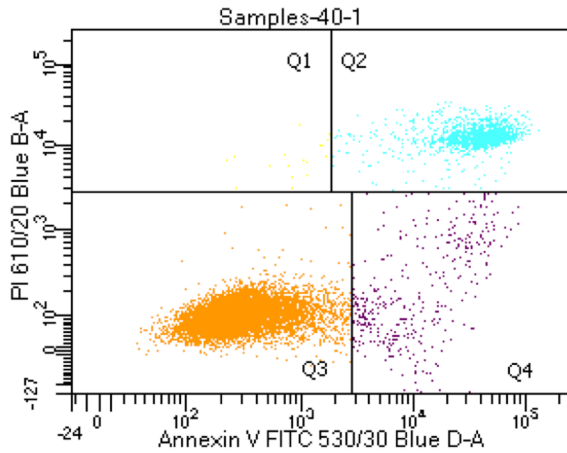
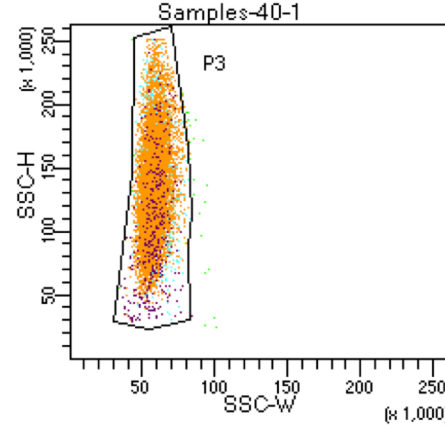
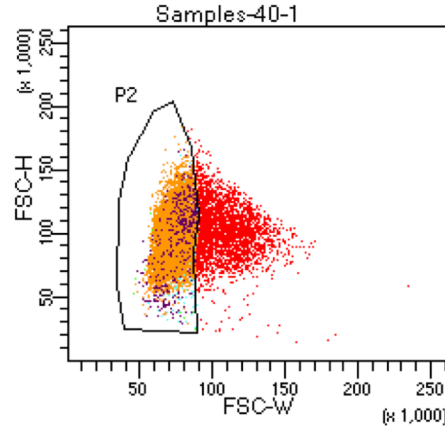
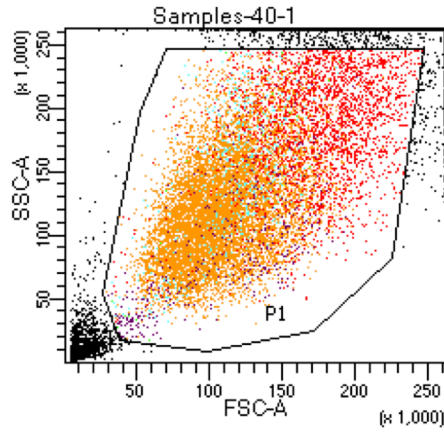


Tube: 20-1

Population	#Events	%Parent	%Total
All Events	15,847	####	100.0
P1	12,108	76.4	76.4
P2	10,022	82.8	63.2
P3	10,000	99.8	63.1
Q1	40	0.4	0.3
Q2	1,726	17.3	10.9
Q3	8,107	81.1	51.2
Q4	127	1.3	0.8

MCF10A 40 uM ZW4864

BD FACSDiva 8.0.1



Tube: 40-1

Population	#Events	%Parent	%Total
All Events	18,450	####	100.0
P1	13,209	71.6	71.6
P2	10,034	76.0	54.4
P3	10,000	99.7	54.2
Q1	17	0.2	0.1
Q2	1,528	15.3	8.3
Q3	8,068	80.7	43.7
Q4	387	3.9	2.1