

Supporting Information

Discovery of new quinoline-based diarylamides as potent B-RAF^{V600E}/C-RAF kinase inhibitors endowed with promising in vitro anticancer activity

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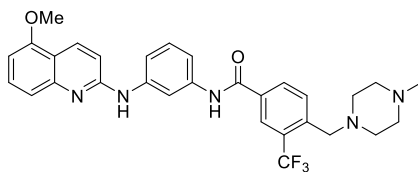
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⁵ Division of Bio-Medical Science & Technology, KIST School, Korea University of Science and Technology (UST), Seoul, 02792, Republic of Korea

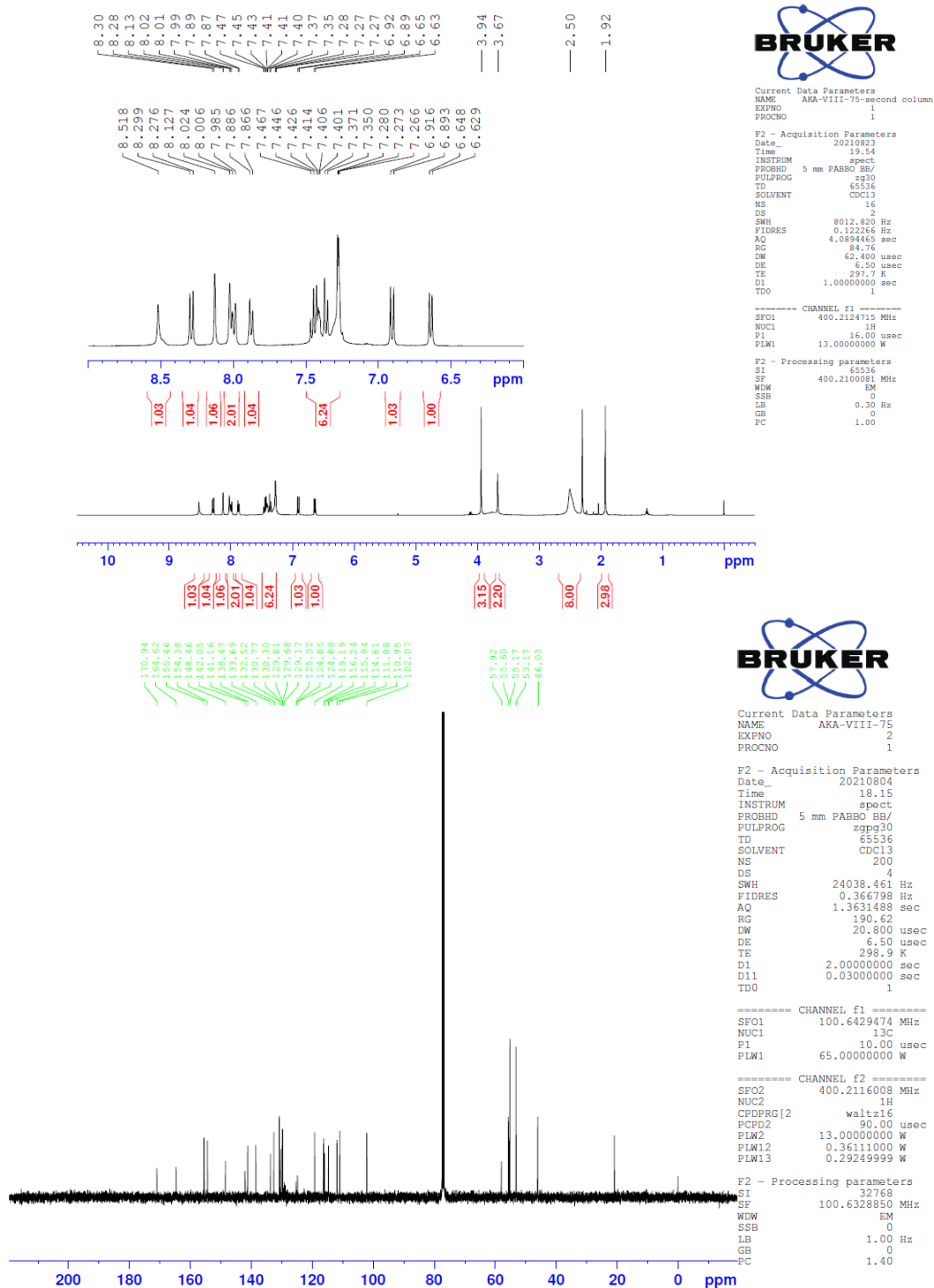
* Correspondence: gkeum@kist.re.kr (G. Keum), ashraf.el-damasy@kist.re.kr or phkarem2006@gmail.com (A.K.El-Damasy)

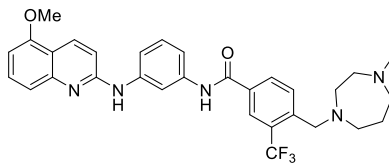
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1) ^1H NMR and ^{13}C NMR spectra

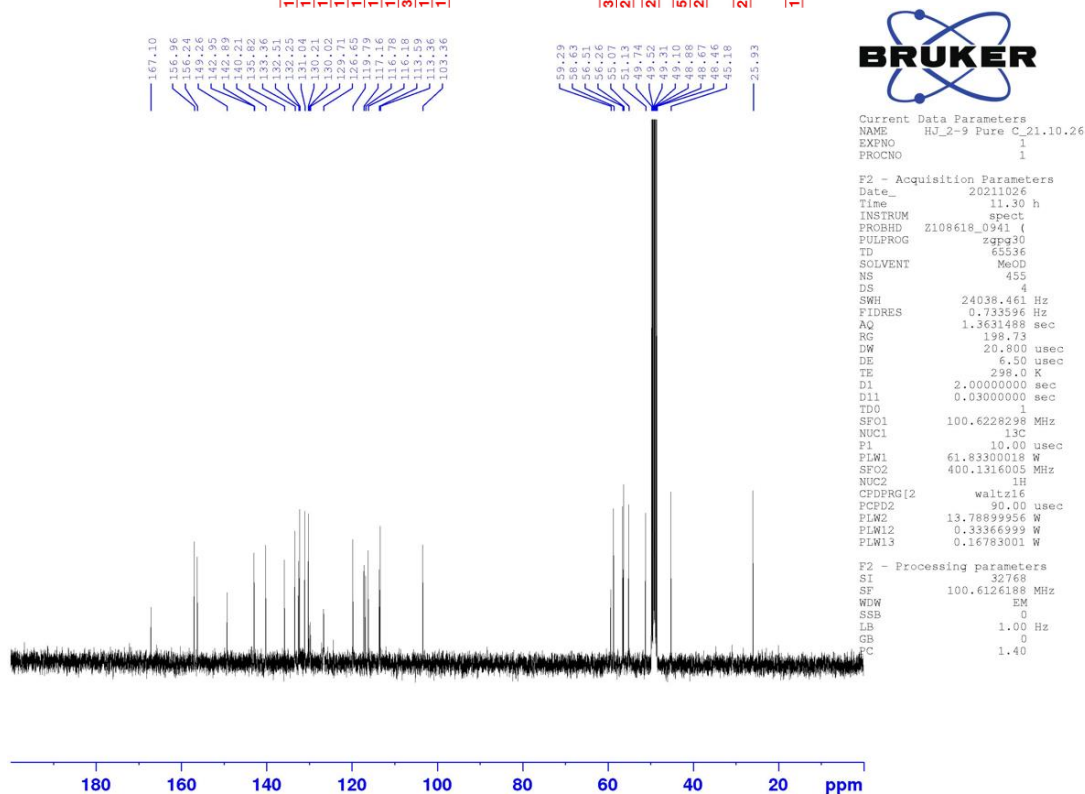
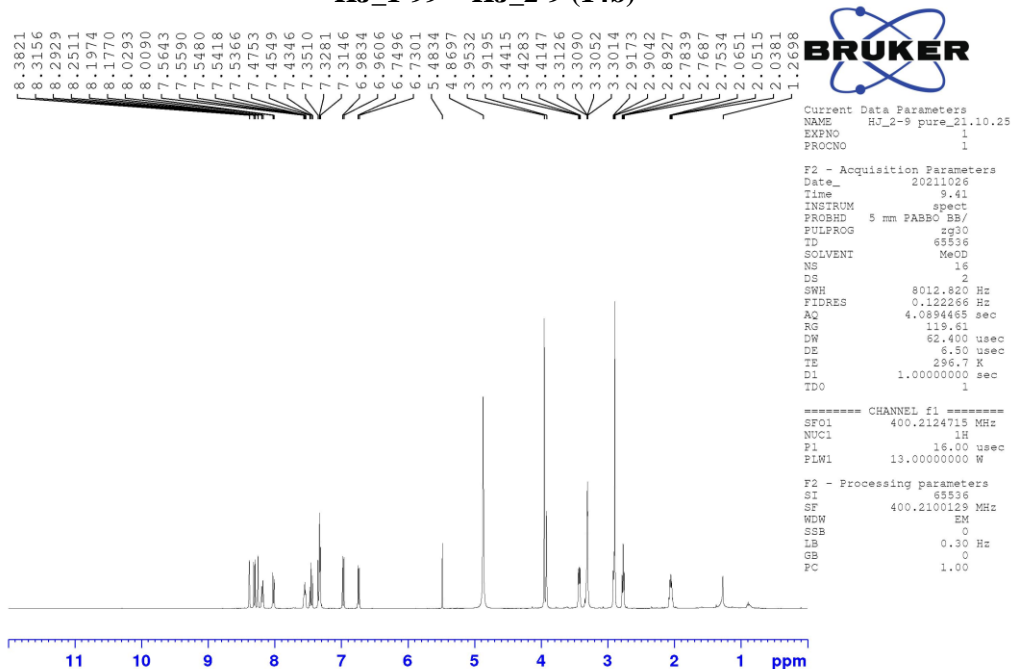


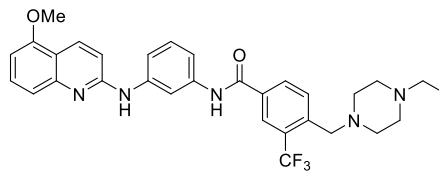
AKA-VIII-75 (14a)



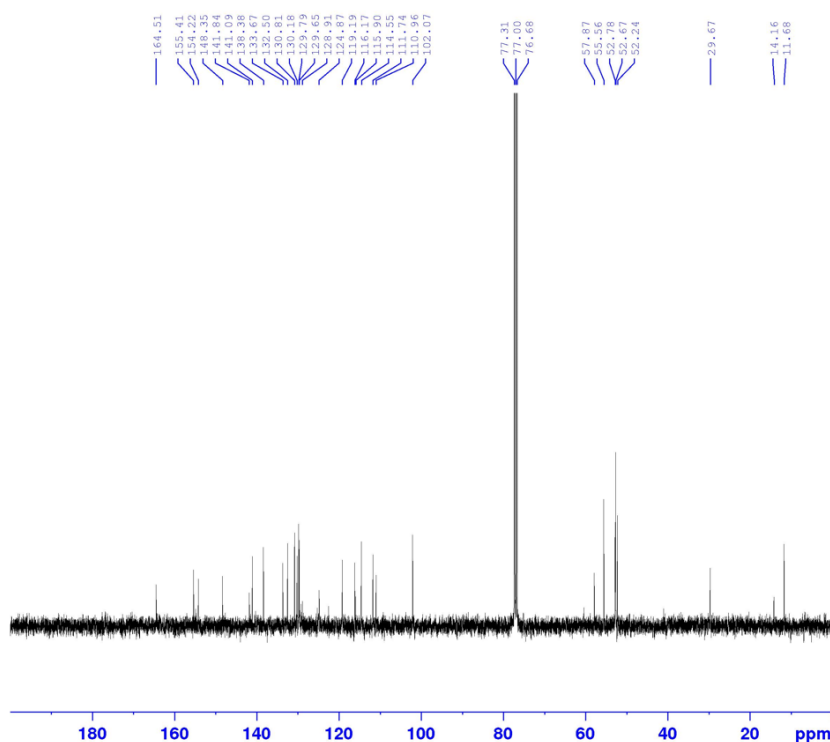
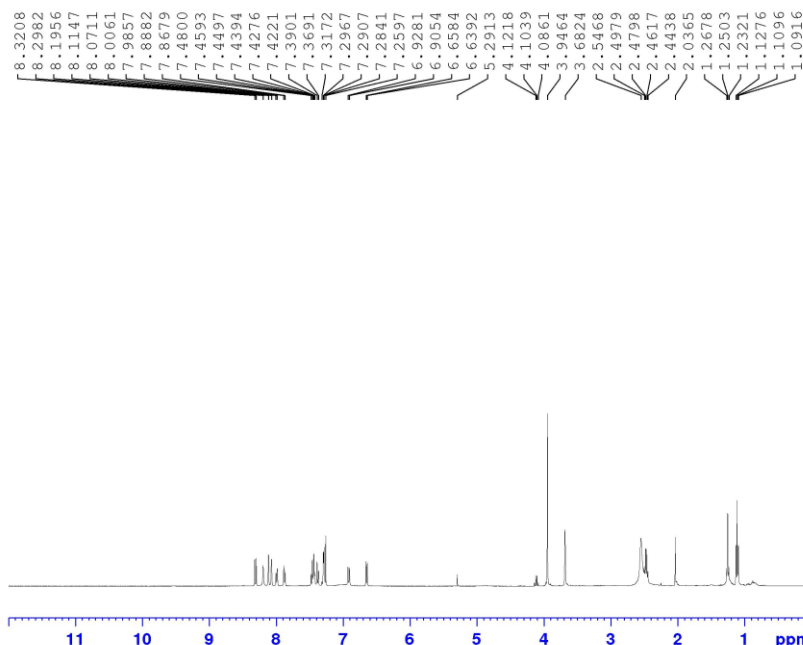


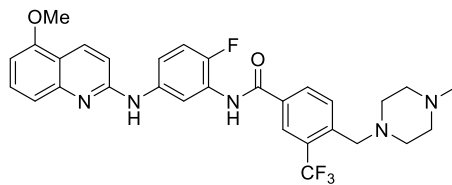
HJ_1-99 = HJ_2-9 (14b)



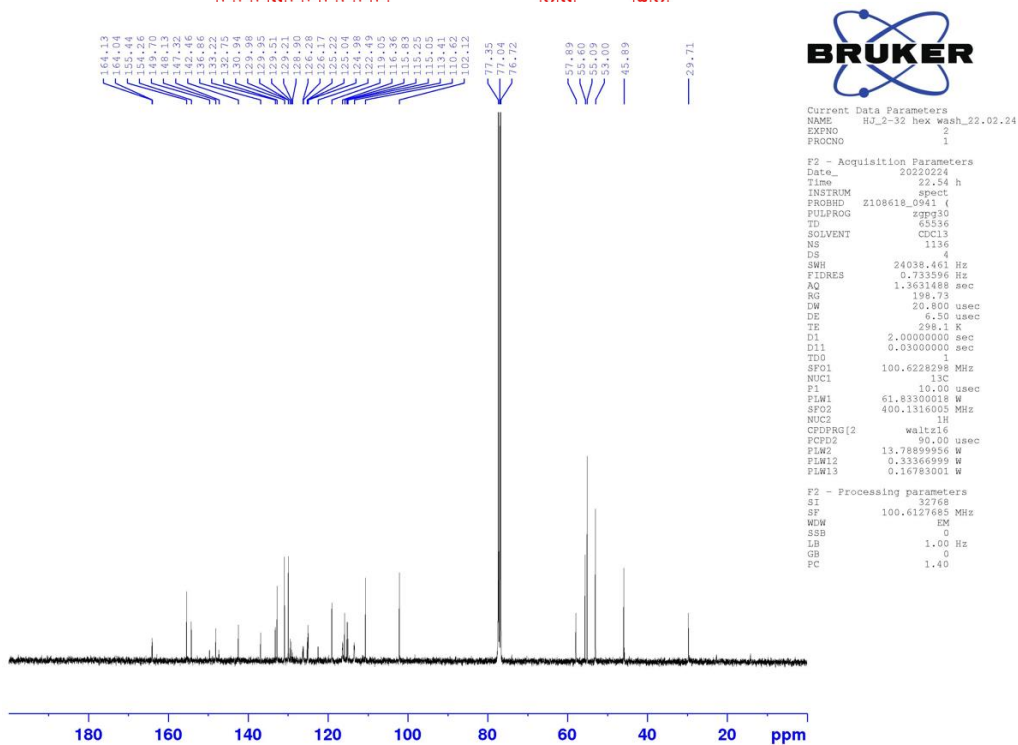
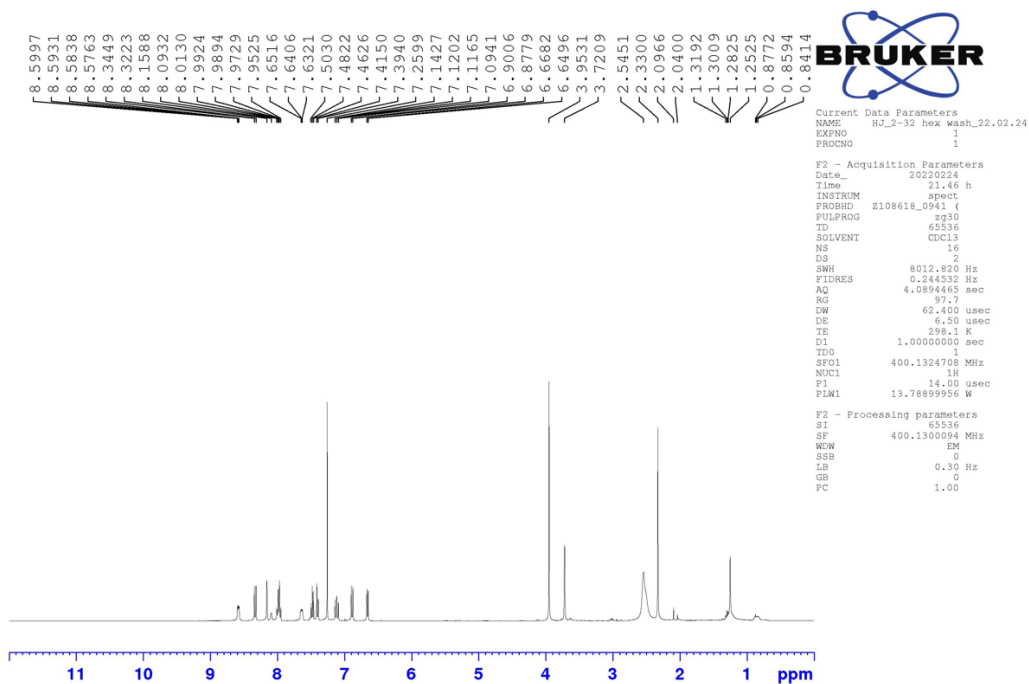


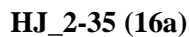
HJ-1-98 (14c)





HJ_2-32 (15a)





```
Current Data Parameters
NAME      HJ_2-35-2 3cc washed_22.01.17
EXPNO     1
PROCNO    1
```

```

F2 - Acquisition Parameters
Date_      20220117
Time       21:57
INSTRUM    spect
PROBHD     5 mm PABBO BB/
PULPROG    zg30
TD          65536
SOLVENT     CDCl3
NS          16
DS          2
SWH         8012.820 Hz
FIDRES      0.122266 Hz
AQ          4.089465 sec
RG          84.76
RG          62.400 usec
DE          6.50 usec
TE          298.1 K
D1          1.0000000 sec
TP0

```

```
===== CHANNEL f1 =====
SFO1      400.2124715 Mhz
NUC1      1H
P1         16.00 usec
PLW1      13.00000000 W
```

```
F2 - Processing parameters
SI                65536
SF                400.2100128 MHz
WDW               EM
SSB               0
LB                0.30 Hz
GB               0
PC               1.00
```



```
Current Data Parameters
NAME      HJ_2-35-2 3cc washed_22.01.17
EXPNO     2
PROCNO    1
```

```

F2 - Acquisition Parameters
Date_ 20220117
Time 22.39
INSTRUM spect
PROBHD 5 mm PABBO BB/
PULPROG zgpg30
TD 65536
SOLVENT CDCl3
NS 1500
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3631488 sec
RG 190.62
DW 20.800 usec
DE 6.50 usec
TE 298.1 K
D1 2.00000000 sec
D11 0.03000000 sec
TD0 1

```

```

===== CHANNEL f1 =====
SFO1      100.6429474 MHz
NUC1             13C
P1             10.00 usec
PLW1      65.000000000 W

```

```

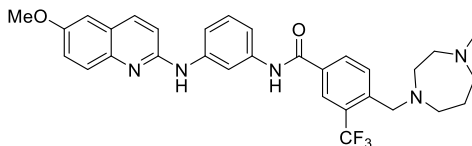
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SFO2      400.2116008 MHz
NUC2      1H
CPDPRG[2] waltz16
PCPD2      90.00 usec
PLW2      13.00000000 W
PLW12     0.36111000 W

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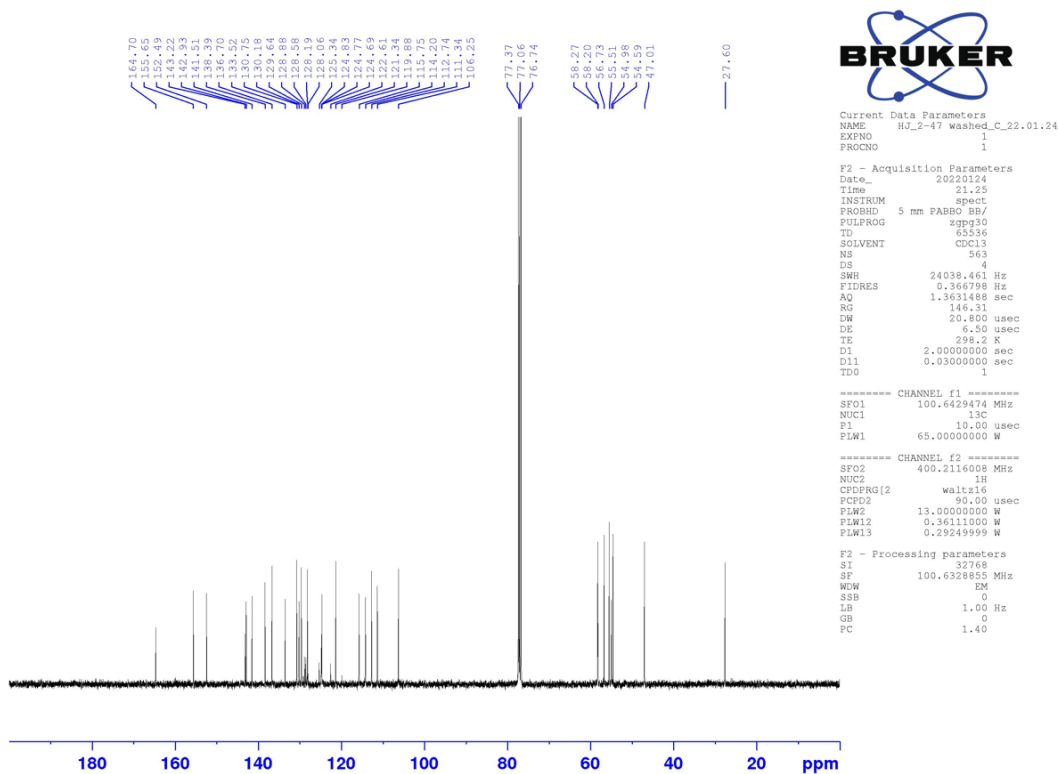
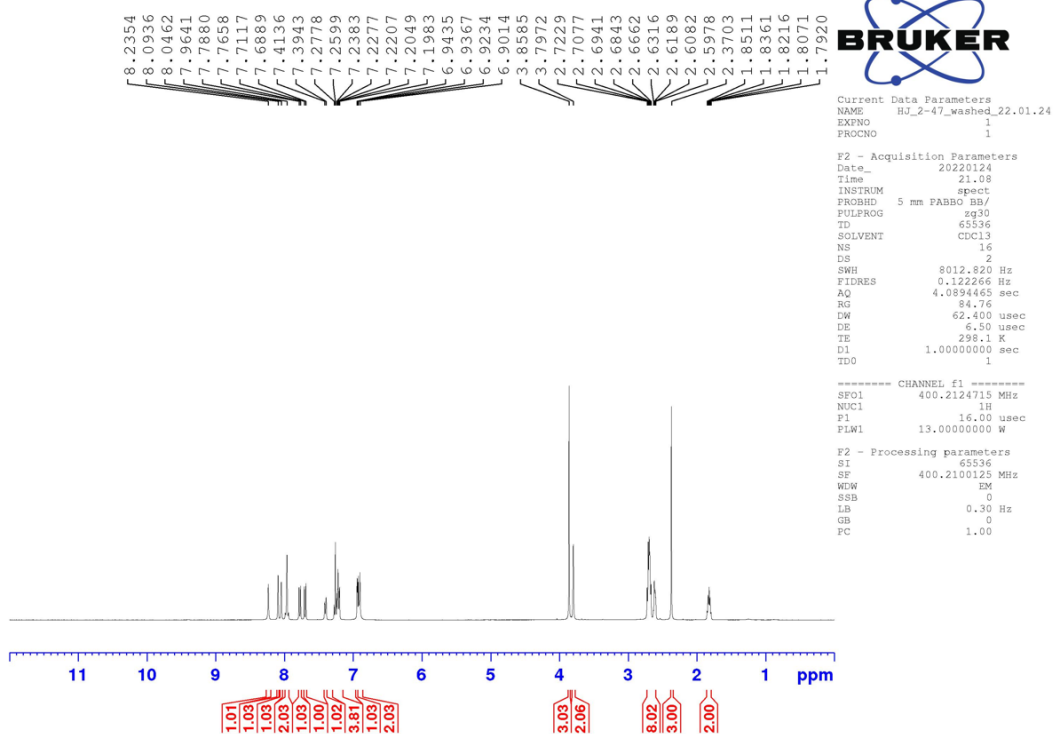
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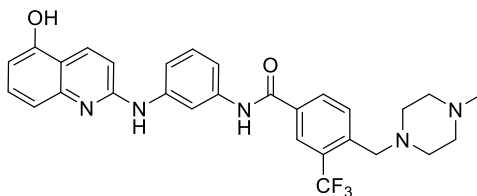
PLW13      0.29249999 W
F2 - Processing parameters
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SF          100.6328851 MHz
WDW         EM
SSB         0
LB          1.00 Hz
GB          0
PC          1.40

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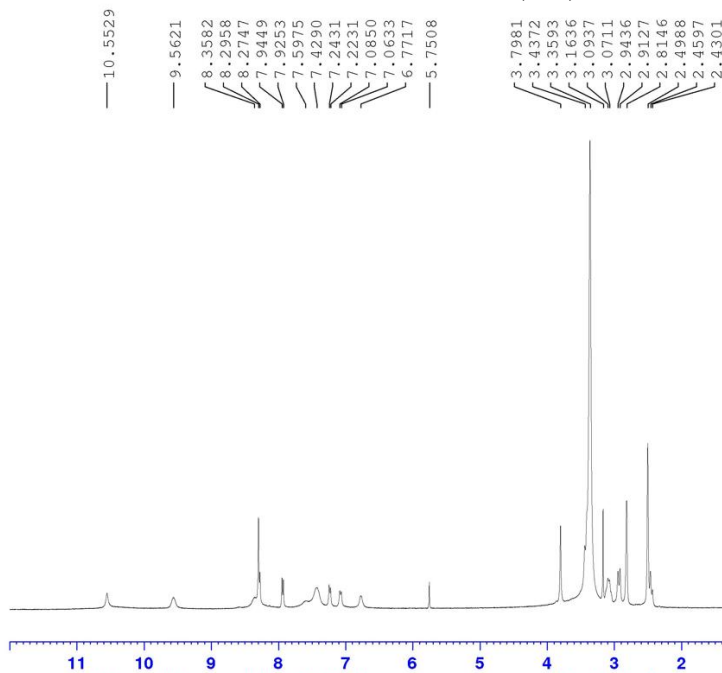


HJ_2-33 = 2-47 (16b)





HJ_2-14 (17a)

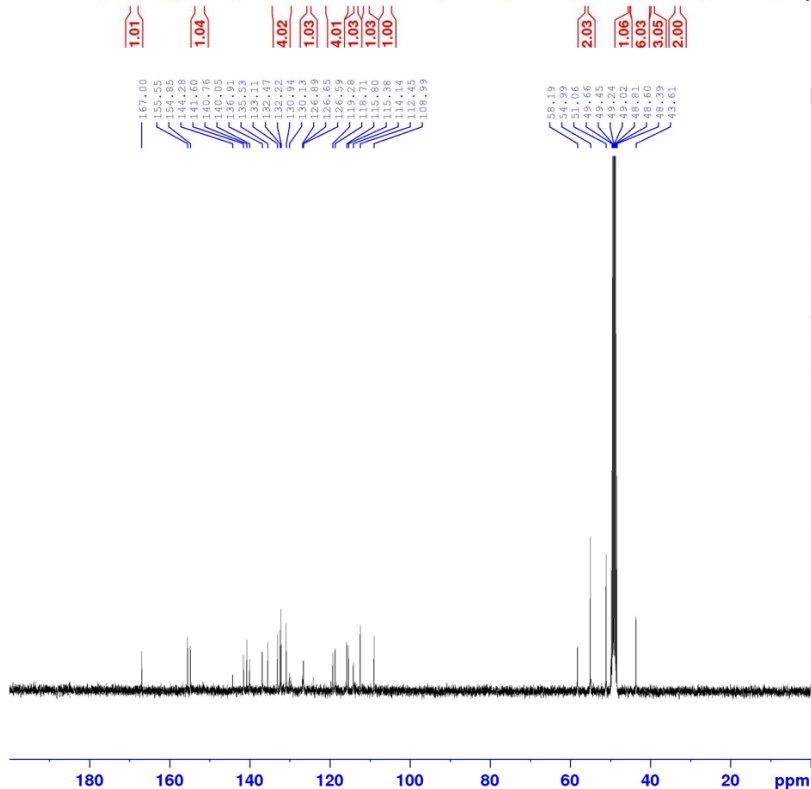


Current Data Parameters
NAME HJ_2-14 DMSO H_22.04.20
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20220420
Time 21.47
INSTRUM spect
PROBHD 5 mm PABBO BB/
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 171.62
DW 62.400 usec
DE 6.50 usec
TE 293.1 K
D1 1.00000000 sec
TDO 1

===== CHANNEL f1 =====
SFO1 400.2124715 MHz
NUC1 1H
P1 16.00 usec
PLW1 13.00000000 W

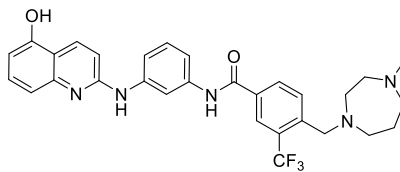
F2 - Processing parameters
SI 65536
SF 400.2100067 MHz
WVW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



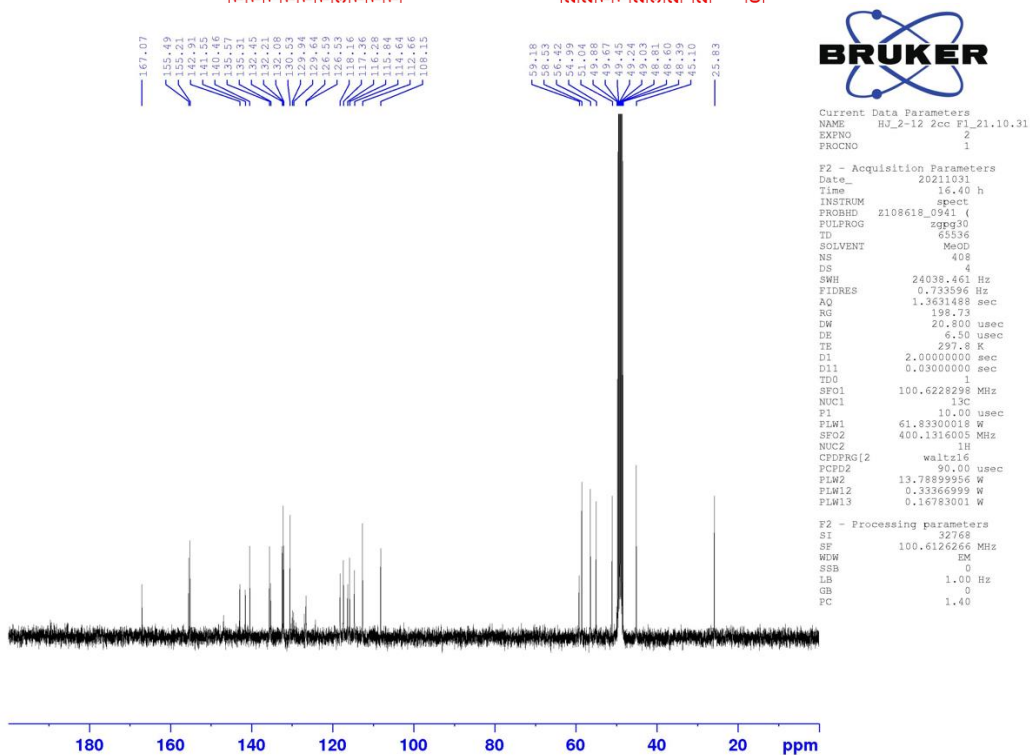
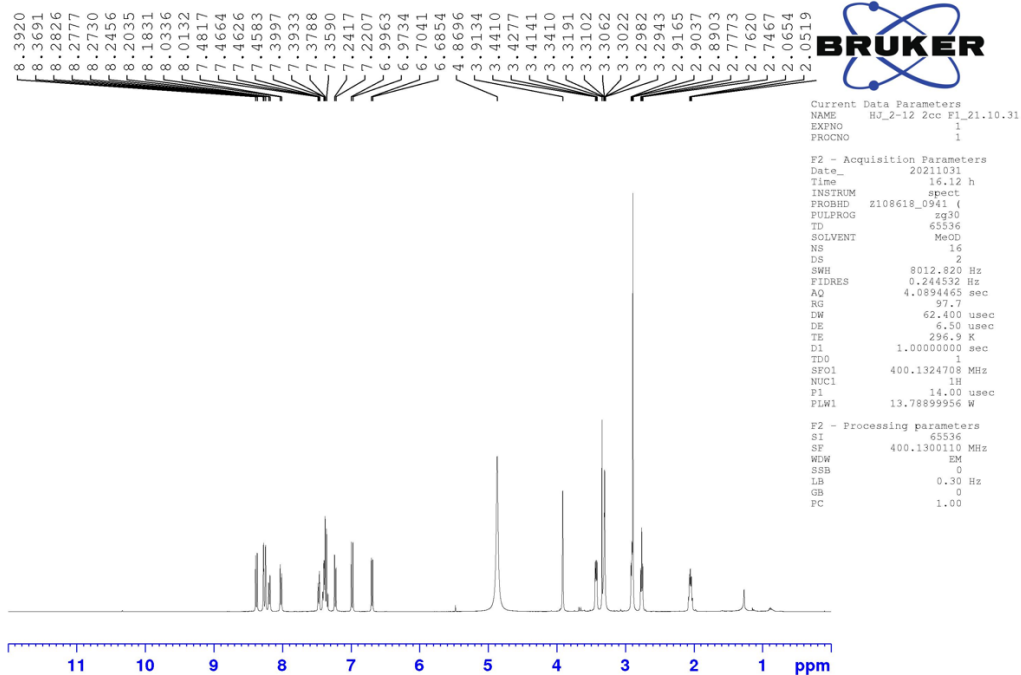
Current Data Parameters
NAME HJ_2-14_21.11.04
EXPNO 2
PROCNO 1

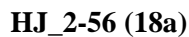
F2 - Acquisition Parameters
Date_ 20211104
Time 10.01 h
INSTRUM spect
PROBHD Z108618_0941 (
PULPROG zgpg30
TD 65536
SOLVENT MeOD
NS 310
DS 4
SWH 24038.461 Hz
FIDRES 0.733596 Hz
AQ 1.3631488 sec
RG 198.73
DW 20.800 usec
DE 6.50 usec
TE 297.9 K
D1 2.00000000 sec
D11 0.03000000 sec
TDO 1
SFO1 100.6228298 MHz
NUC1 13C
P1 10.00 usec
PLW1 61.83300018 W
SFO2 400.1316005 MHz
NUC2 1H
PCPD2 waltz16
PLW2 13.78899956 W
PLW12 0.33366999 W
PLW13 0.16783001 W

F2 - Processing parameters
SI 32768
SF 100.6126287 MHz
WVW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40



HJ_2-12 (17b)





```
Current Data Parameters
NAME      HJ_2-56 3cc filter_22.02.21
EXPNO     1
PROCNO    1
```

```

F2 - Acquisition Parameters
Date_      20220221
Time       18.58
INSTRUM    spect
PROBHD     5 mm PABBO BB/
PULPROG    zg30
TD          65536
SOLVENT    MeOD
NS          16
DS          2
SWH         8012.820 Hz
FIDRES     0.122266 Hz
AQ         4.0849445 sec
RG          119.61
DW         62.400 usec
DE         6.50 usec
TE         296.7 K
D1         1.00000000 sec
TD0        .1

```

```
===== CHANNEL f1 =====
SFO1      400.2124715 MHz
NUC1              1H
P1              16.00 usec
PLW1      13.00000000 W
```

```
F2 - Processing parameters
SI                      65536
SF                      400.2100000 MHz
WDW                      EM
SSB                      0
LB                      0.30 Hz
GB                      0
PC                      1.00
```



```
Current Data Parameters
NAME      HJ_2-56 C_22.02.28
EXPNO      1
PROCNO     1
```

```

F2 - Acquisition Parameters
Date_      20220228
Time       22.43
INSTRUM    spect
PROBHD     5 mm PABBO BB/
PULPROG    zgpg30
ID         65536
SOLVENT    MeOD
NS         5200
DS         4
SWH        24038.461 Hz
FIDRES     0.366798 Hz
AQ         1.3631488 sec
RG         190.62
DE         20.800 use
TE         6.50 use
TD         298.1 K
D1         2.00000000 sec
D11        0.03000000 sec
TD0        1

```

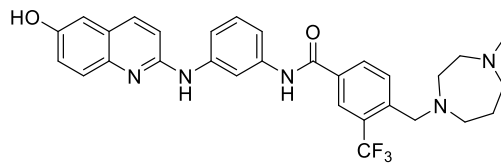
```
===== CHANNEL f1 =====
SF01      100.6429474 MHz
NUC1              13C
P1              10.00 usec
PLW1      65.00000000 W
```

```

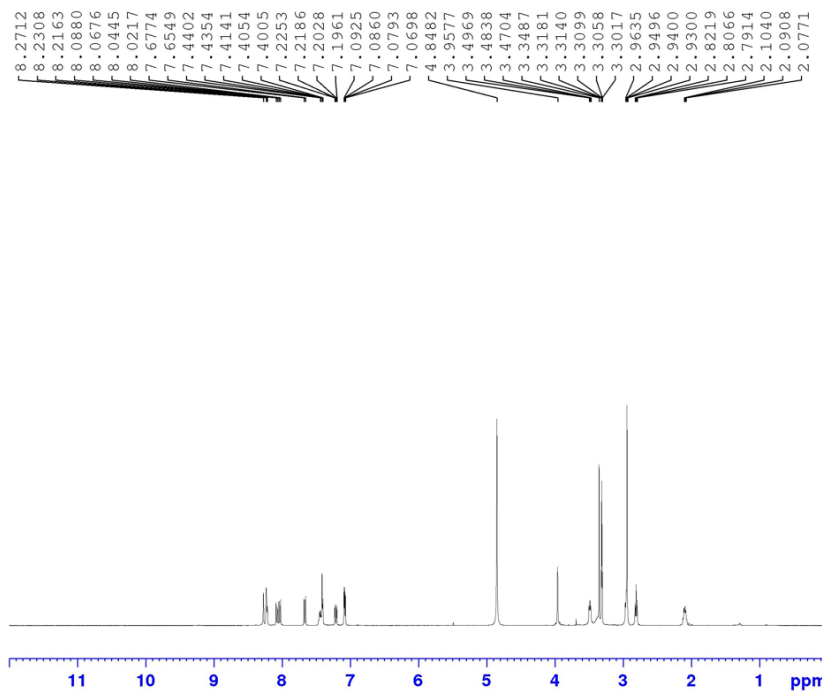
SFO2      400.2116008 MHz
NUC2      1H
CPDPRG2   waltz16
PCPD2     90.00 usec
PLW2      13.00000000 W
PLW12     0.36110000 W
PLW13     0.29249999 W

```

```
F2 - Processing parameters
SI              32768
SF              100.6327424 MHz
WDW             EM
SSB             0
LB              1.00 Hz
GB              0
PC              1.40
```



HJ_2-57 (18b)

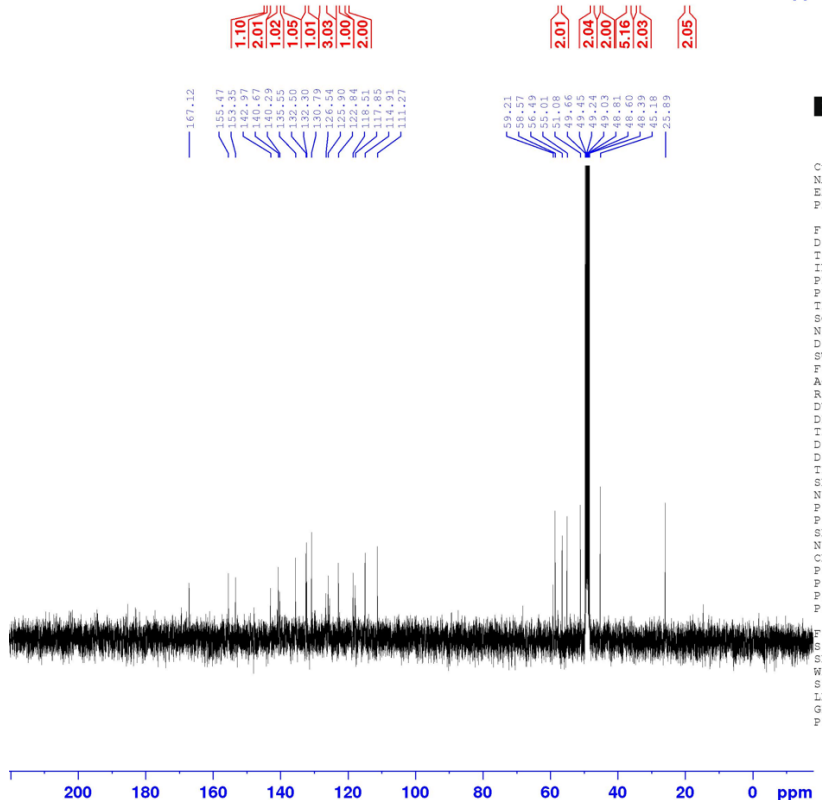


BRUKER

Current Data Parameters
NAME HJ_2-57_3cc v2.DCN wash_22.02.23
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20220301
Time 21.36 h
INSTRUM spect
PROBHD Z108618_0941 (
PULPROG zgpg30
TD 65536
SOLVENT MeOD
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.244532 Hz
AQ 4.088463 sec
RG 198.73
DW 65.400 usec
DE 6.50 usec
TE 298.1 K
D1 1.00000000 sec
TD0 1
SFO1 400.1324708 MHz
NUC1 1H
P1 14.00 usec
PLW1 13.78899956 W

F2 - Processing parameters
SI 65536
SF 400.1300078 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



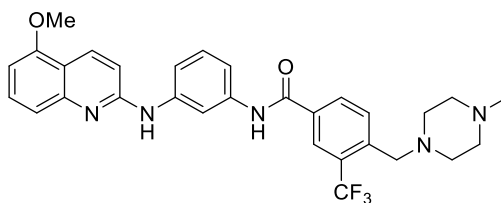
BRUKER

Current Data Parameters
NAME HJ_2-57 c_22.02.28
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20220301
Time 6.20 h
INSTRUM spect
PROBHD Z108618_0941 (
PULPROG zgpg30
TD 65536
SOLVENT MeOD
NS 8000
DS 4
SWH 24038.461 Hz
FIDRES 0.733596 Hz
AQ 1.3631488 sec
RG 198.73
DW 20.800 usec
DE 6.50 usec
TE 299.2 K
D1 2.00000000 sec
D11 0.03000000 sec
TD0 1
SFO1 100.6228298 MHz
NUC1 13C
P1 10.00 usec
PLW1 61.83300018 W
SFO2 400.1316005 MHz
NUC2 1H
CPDPRG[2] waltz16
PCPD2 90.00 usec
PLW2 13.78899956 W
PLW12 0.33366999 W
PLW13 0.16783001 W

F2 - Processing parameters
SI 32768
SF 100.6126259 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40

2) HRMS charts



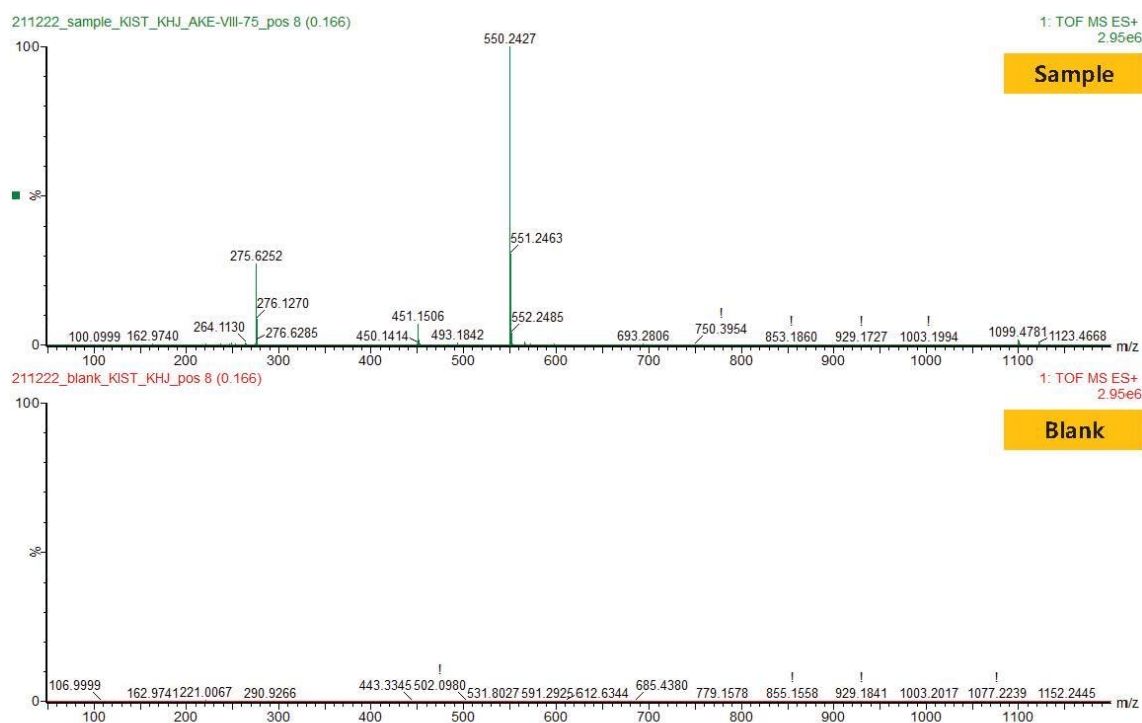
AKA-VIII-75 (14a)

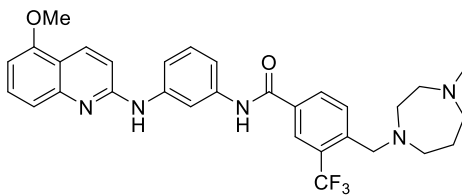
$C_{30}H_{31}F_3N_5O_2$ $[M+H]^+$:

Calculated: 550.2430

Found: 550.2427

Mass Spectrum: AKE-VIII-75 (Positive mode)





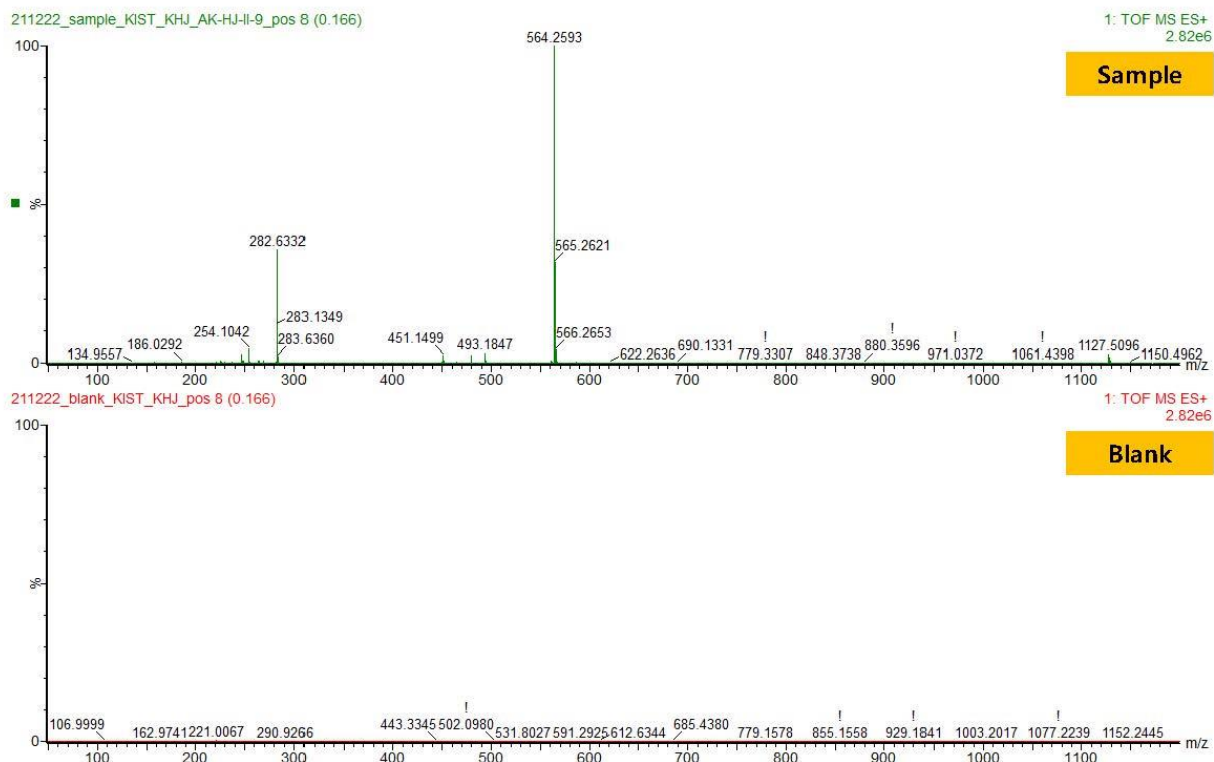
HJ_1-99 = HJ_2-9 (14b)

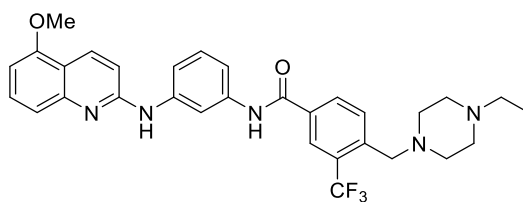
$\text{C}_{31}\text{H}_{33}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 564.2586

Found: 564.2593

Mass Spectrum: AK-HJ-II-9 (Positive mode)





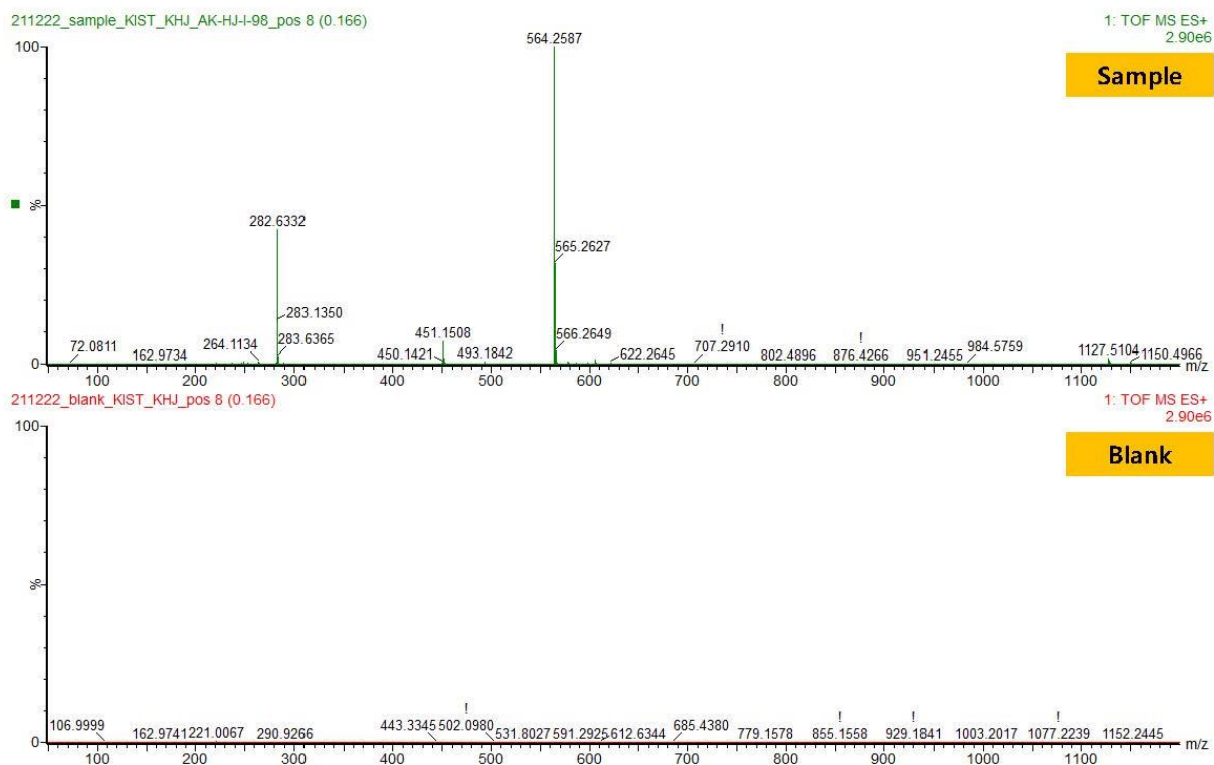
HJ-1-98 (14c)

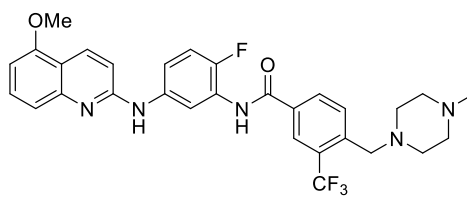
$\text{C}_{31}\text{H}_{33}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 564.2586

Found: 564.2587

Mass Spectrum: AK-HJ-I-98 (Positive mode)





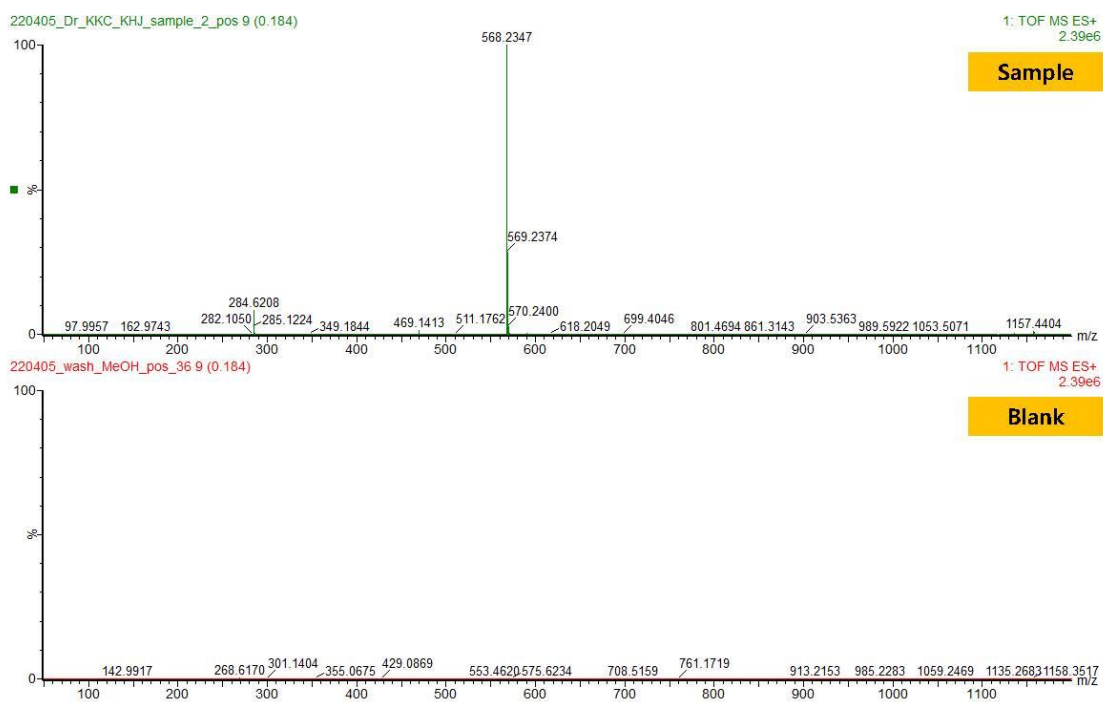
HJ_2-32 (15a)

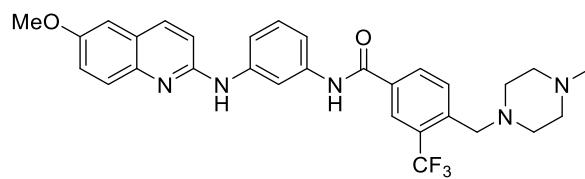
$C_{30}H_{30}F_4N_5O_2$ $[M+H]^+$:

Calculated: 568.2336

Found: 568.2347

Mass Spectrum: AK-HJ-II-32 (Positive mode)





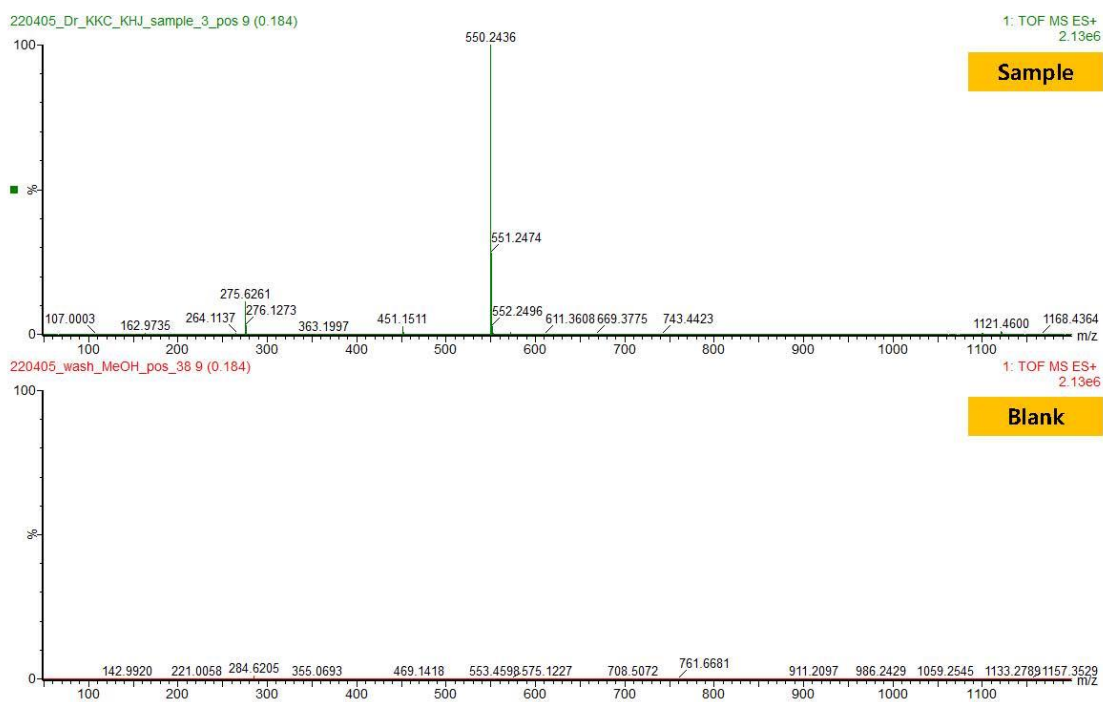
HJ_2-35 (16a)

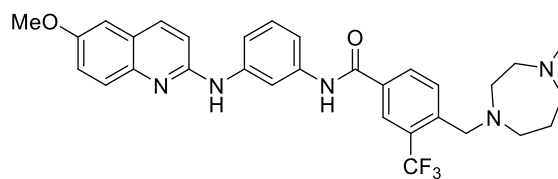
$\text{C}_{30}\text{H}_{31}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 550.2430

Found: 550.2436

Mass Spectrum: AK-HJ-II-35 (Positive mode)





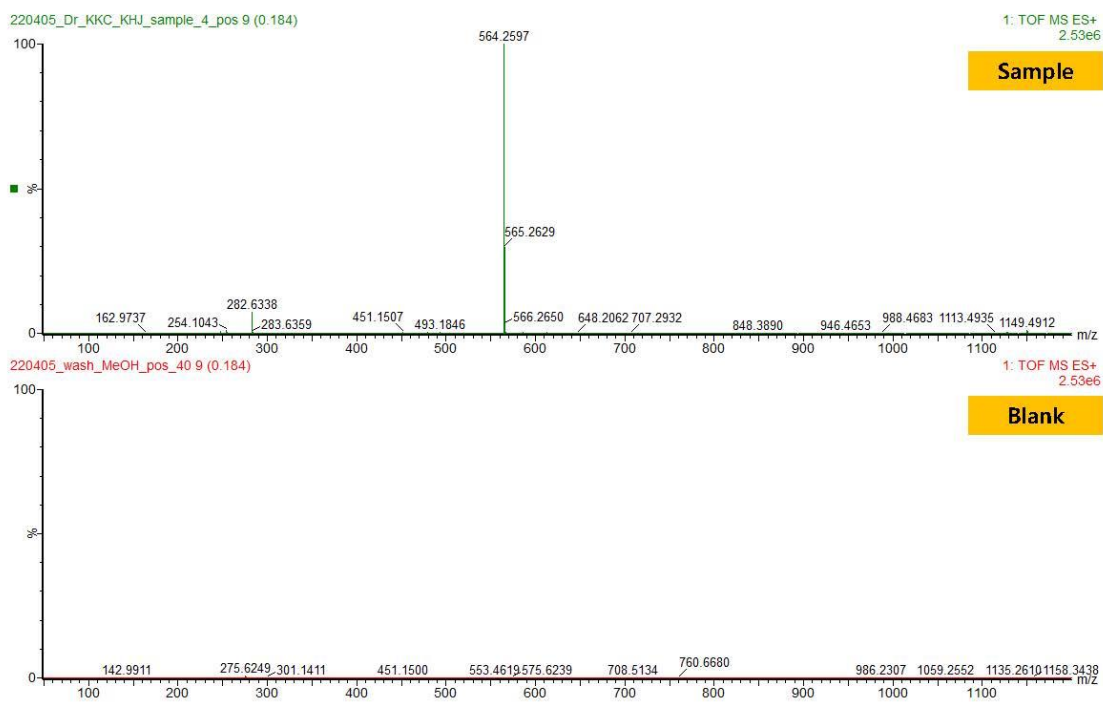
HJ_2-33 = 2-47 (16b)

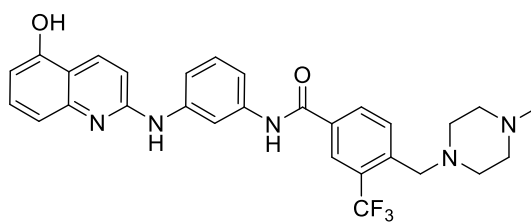
$\text{C}_{31}\text{H}_{33}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 564.2586

Found: 564.2597

Mass Spectrum: AK-HJ-II-47 (Positive mode)





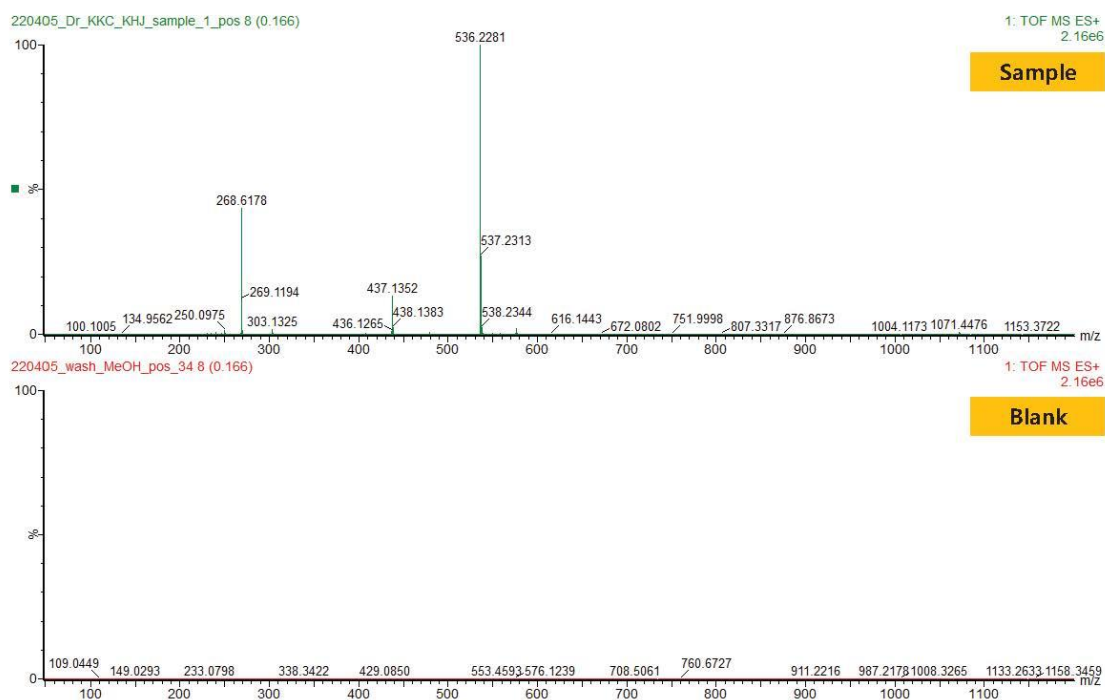
HJ_2-14 (17a)

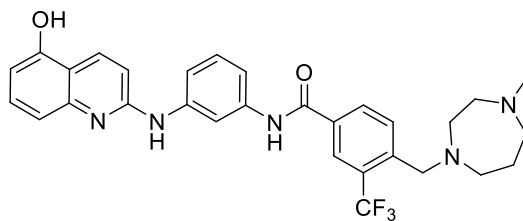
$\text{C}_{29}\text{H}_{29}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 536.2274

Found: 536.2281

Mass Spectrum: AK-HJ-II-14 (Positive mode)





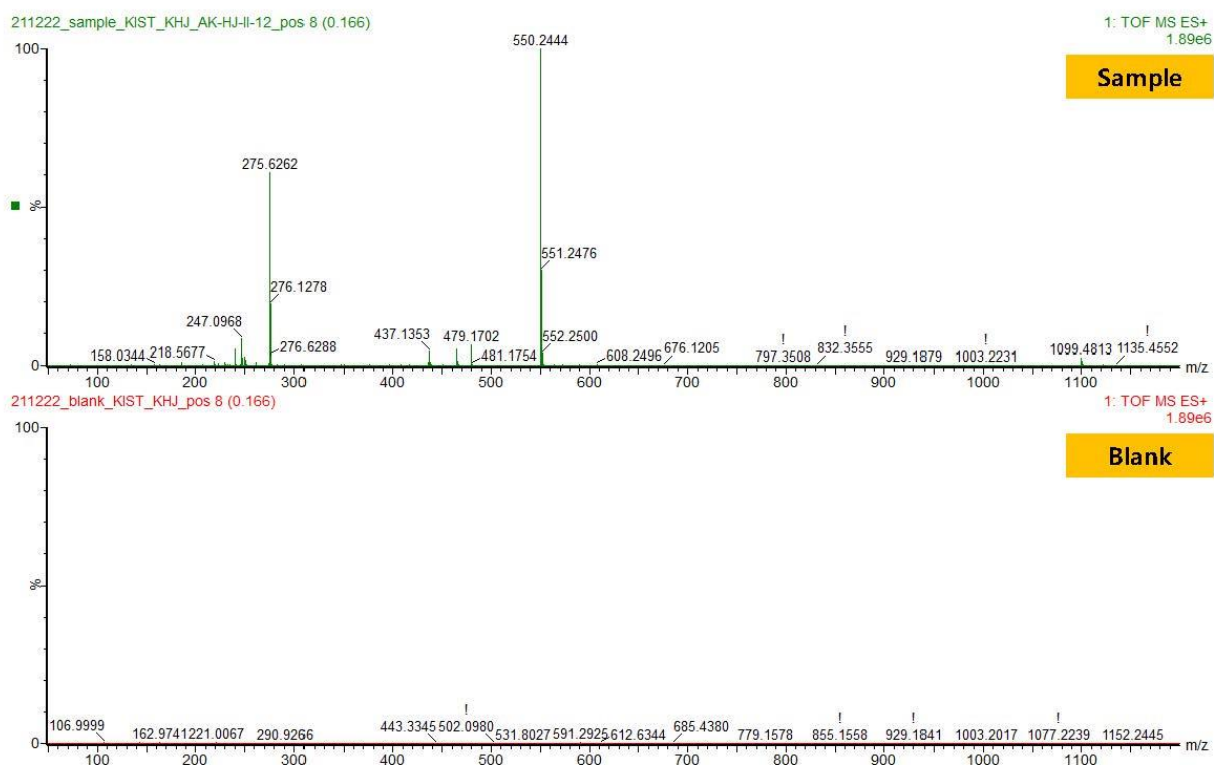
HJ_2-12 (17b)

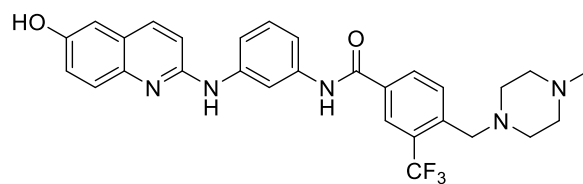
$\text{C}_{30}\text{H}_{31}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 550.2430

Found: 550.2444

Mass Spectrum: AK-HJ-II-12 (Positive mode)





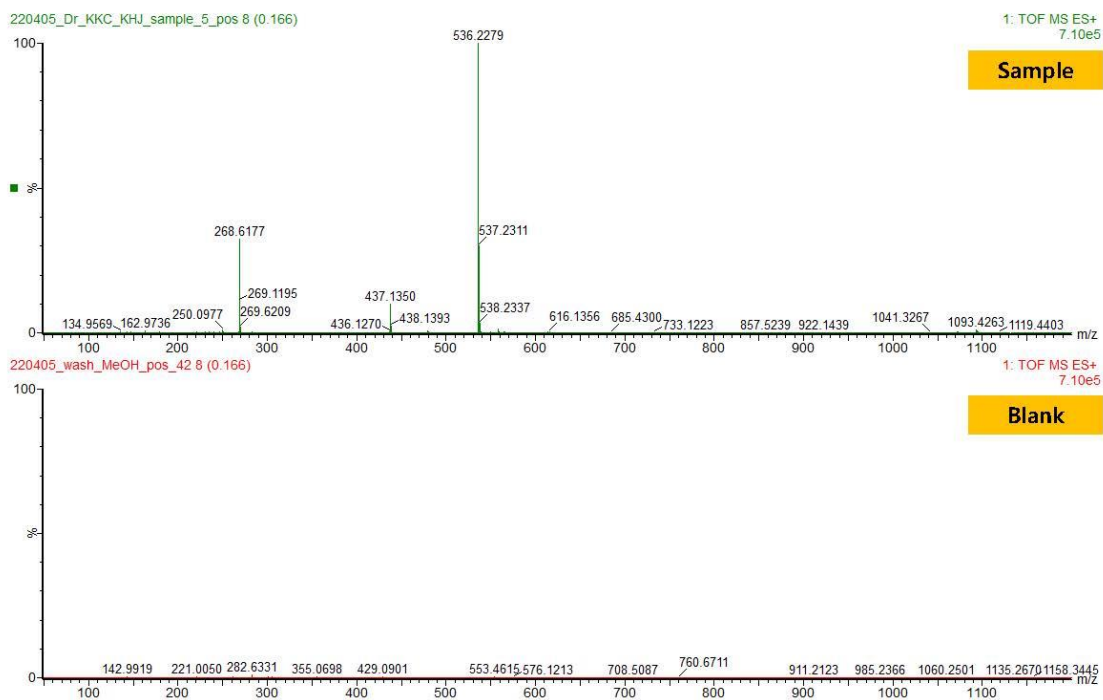
HJ_2-56 (18a)

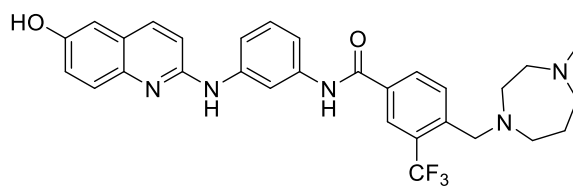
$\text{C}_{29}\text{H}_{29}\text{F}_3\text{N}_5\text{O}_2$ $[\text{M}+\text{H}]^+$:

Calculated: 536.2274

Found: 536.2279

Mass Spectrum: AK-HJ-II-56 (Positive mode)





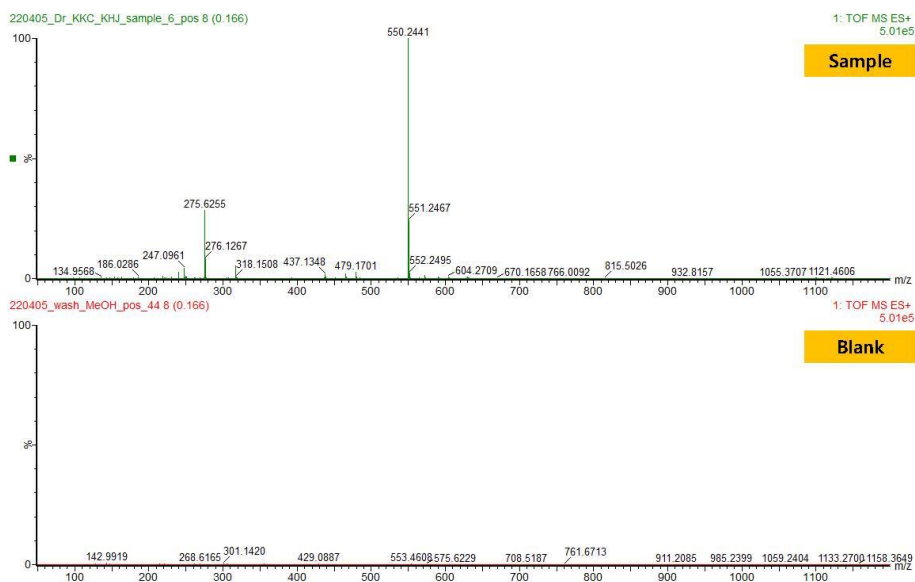
HJ_2-57 (18b)

$C_{30}H_{31}F_3N_5O_2$ $[M+H]^+$:

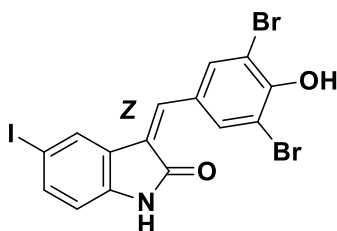
Calculated: 550.2430

Found: 550.2441

Mass Spectrum: AK-HJ-II-57 (Positive mode)

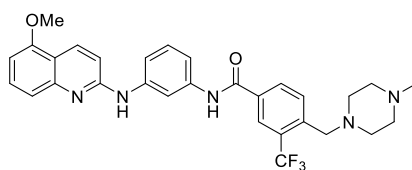


3) Chemical structure of GW5074

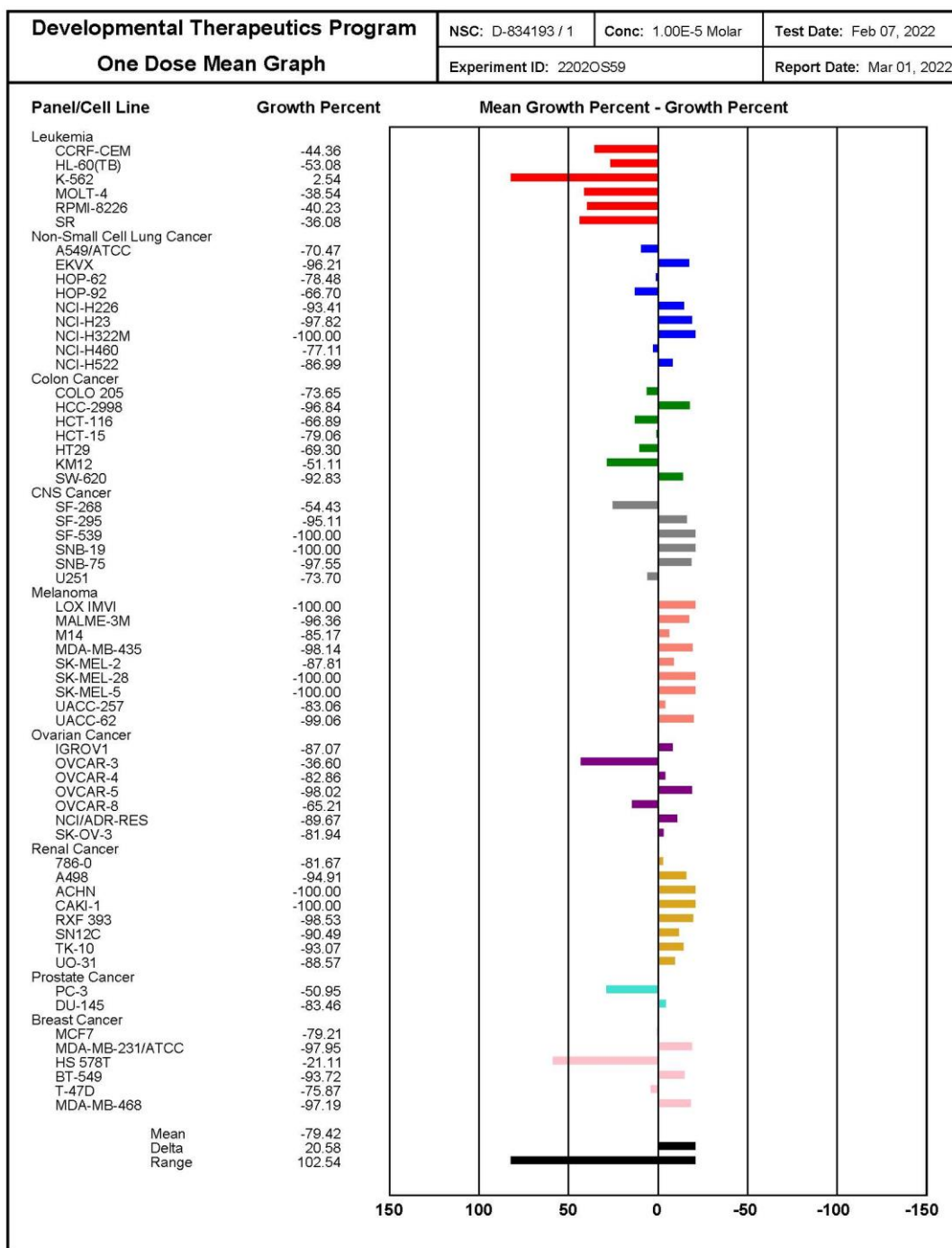


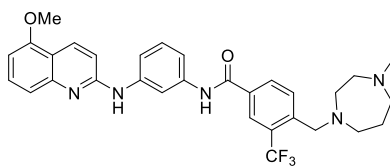
GW5074

4) NCI data

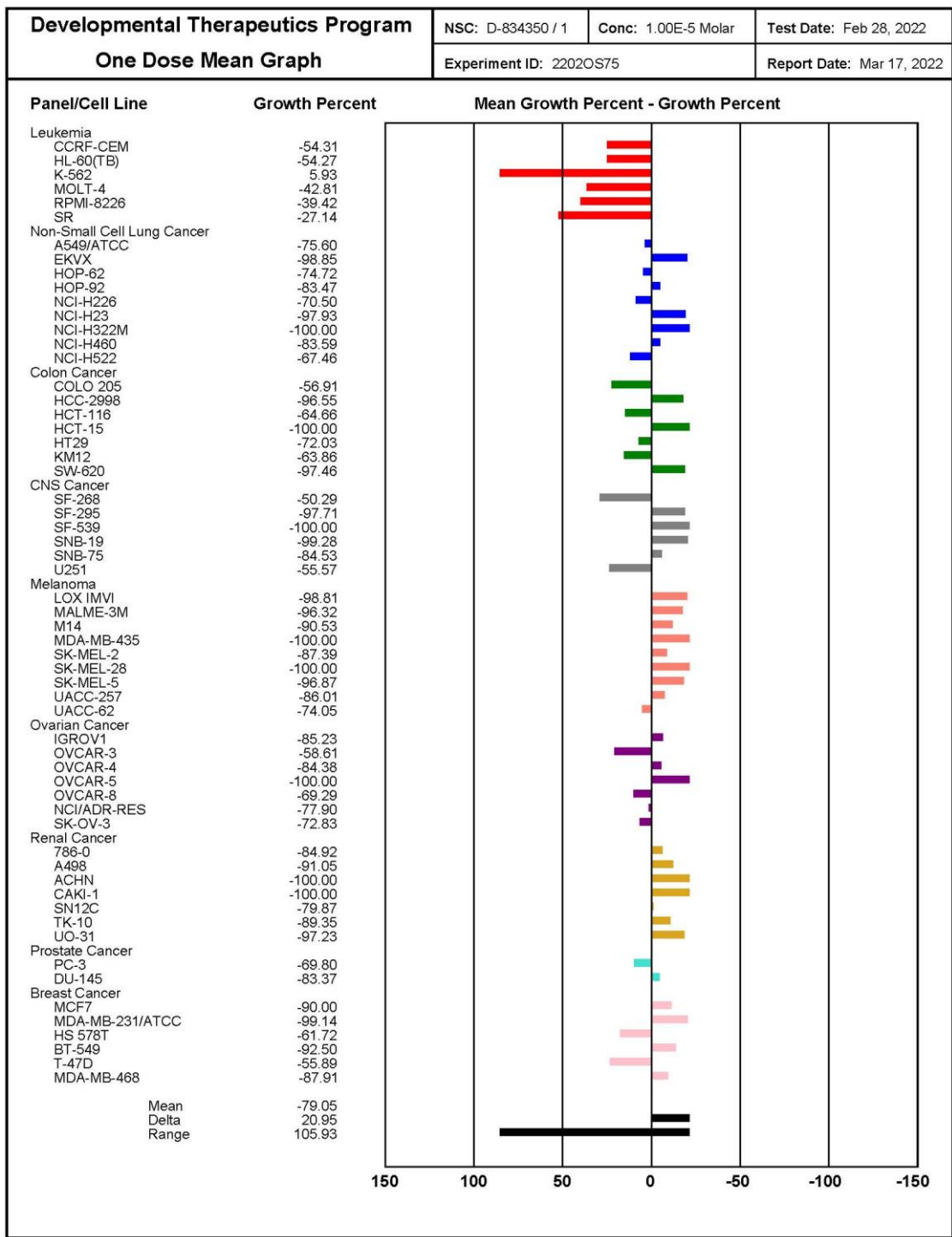


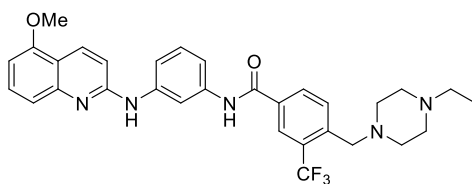
AKA-VIII-75 (14a)



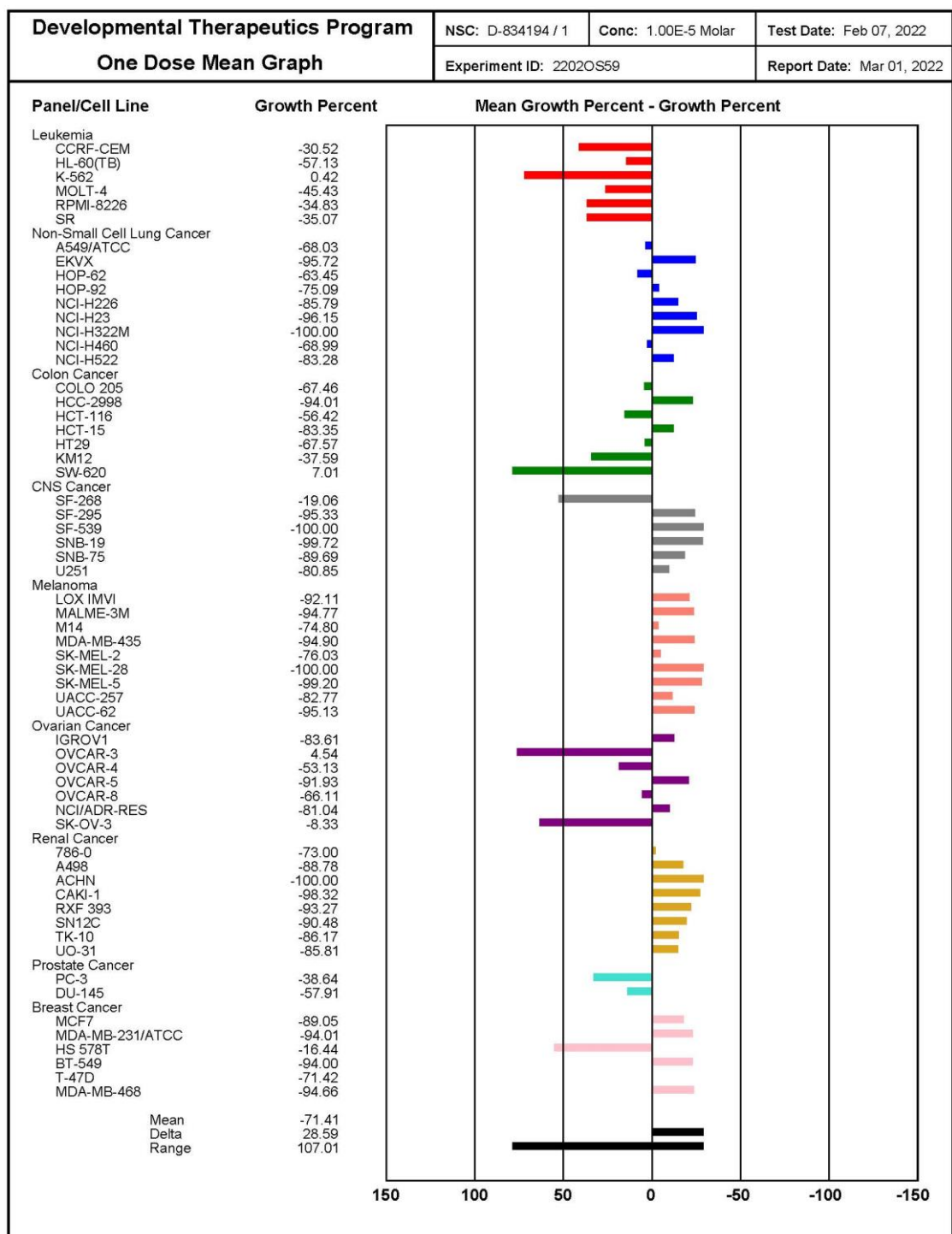


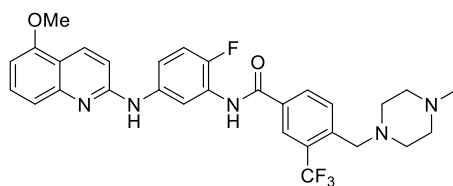
HJ_1-99 = HJ_2-9 (14b)



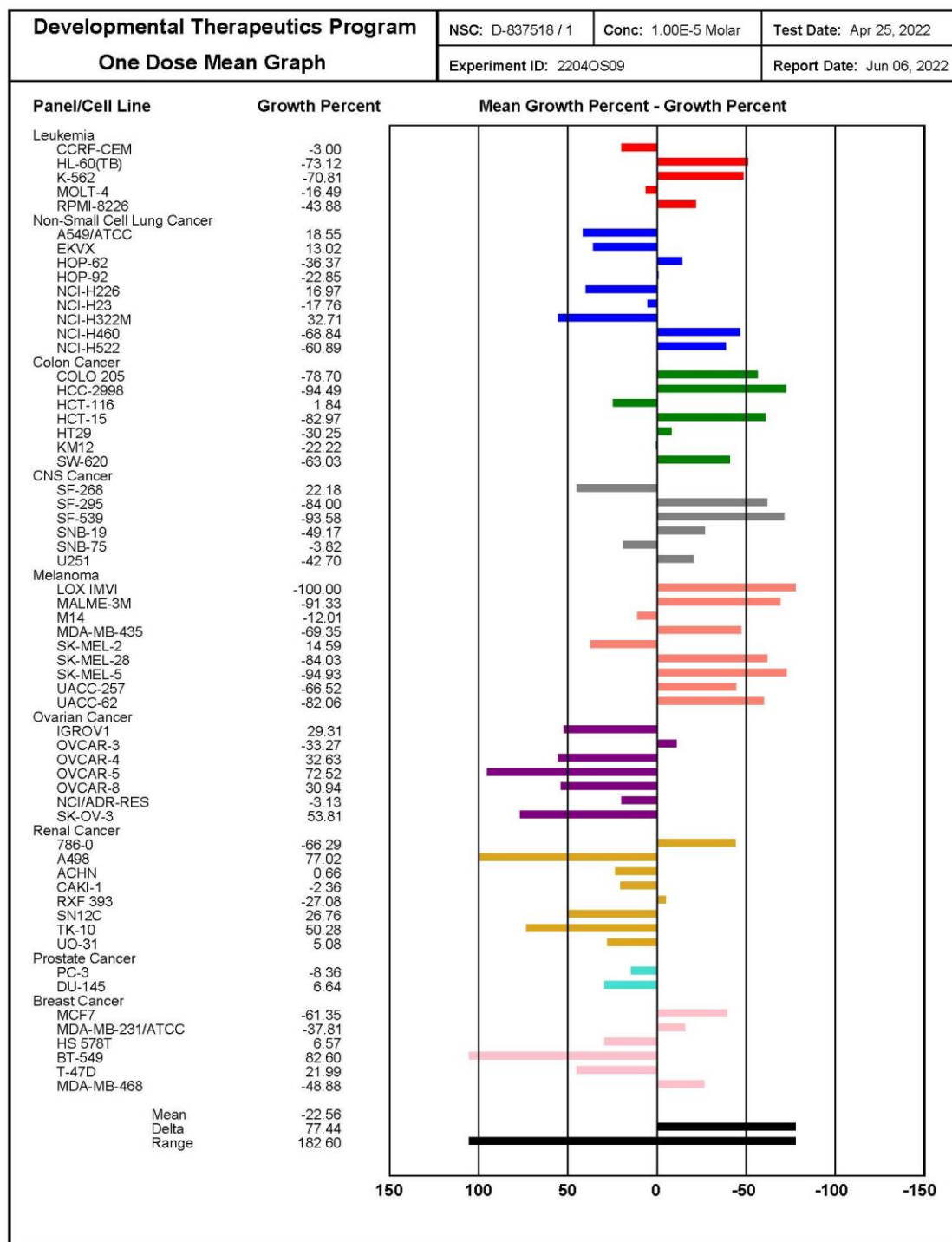


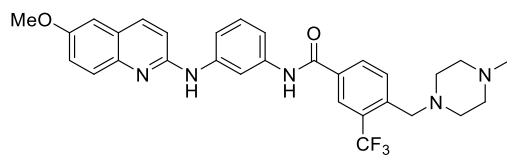
HJ-1-98 (14c)



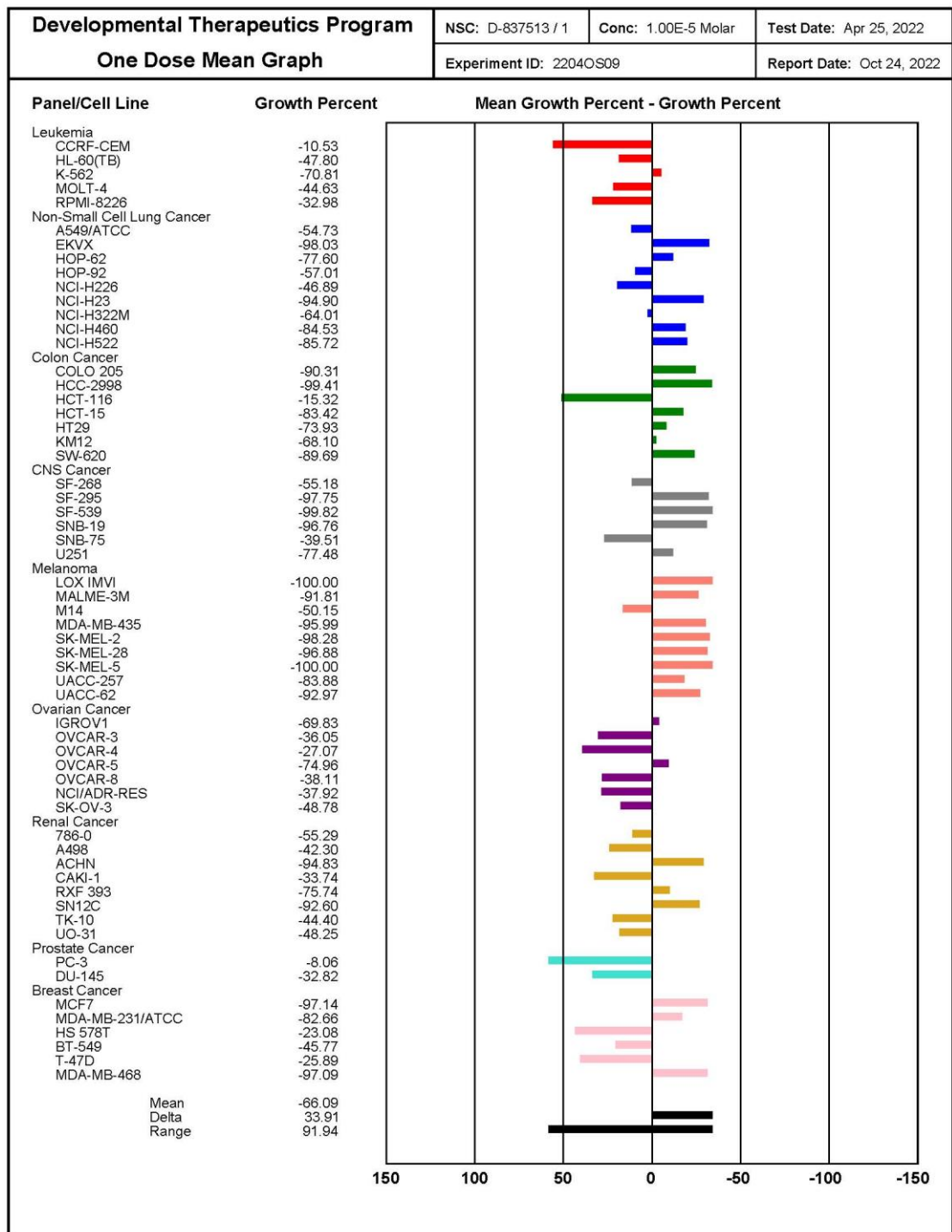


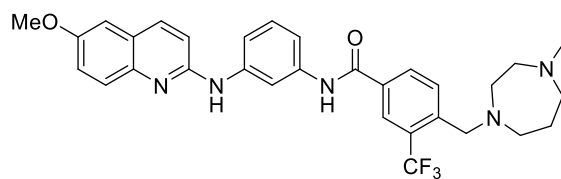
HJ_2-32 (15a)



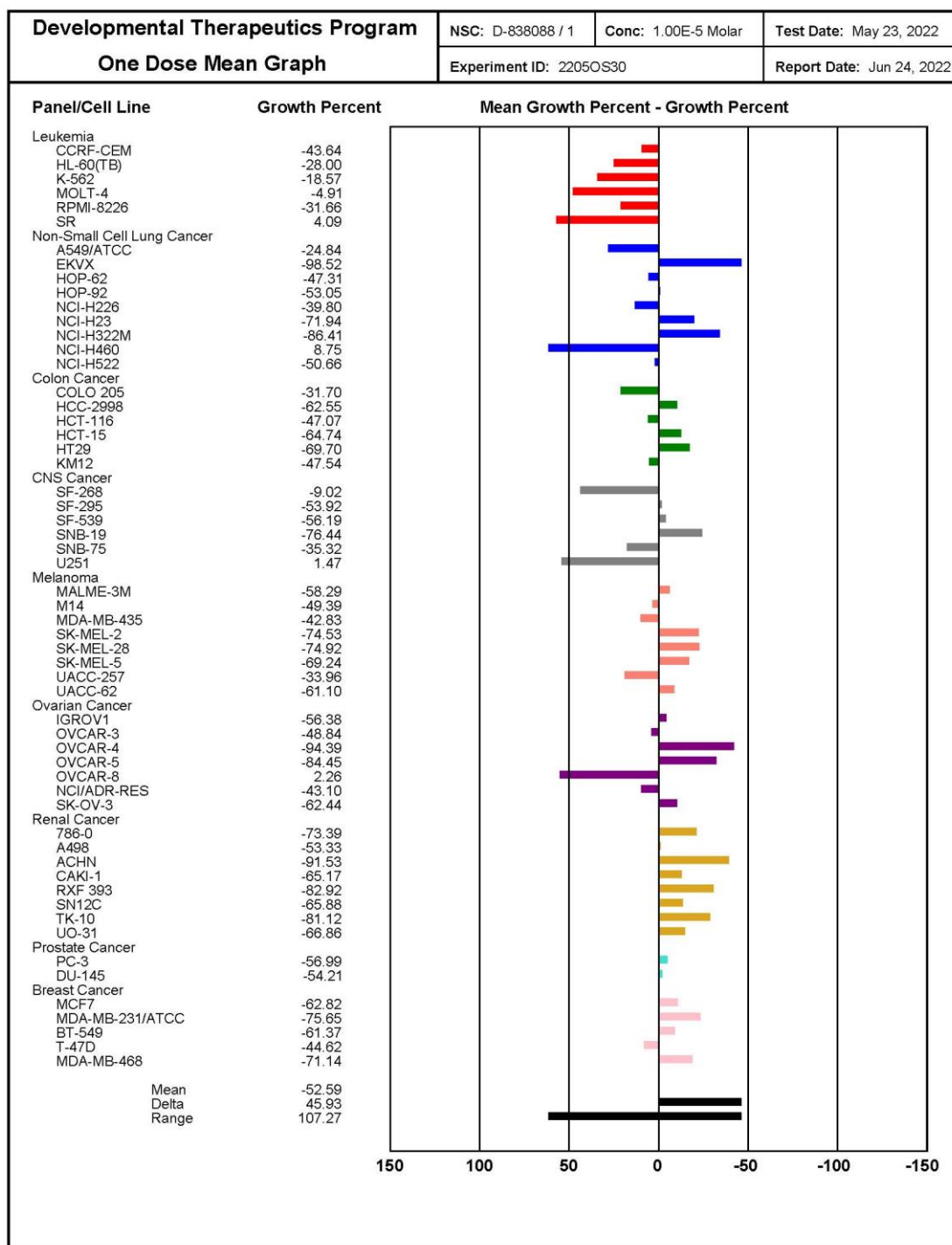


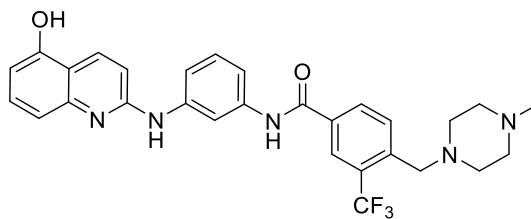
HJ_2-35 (16a)



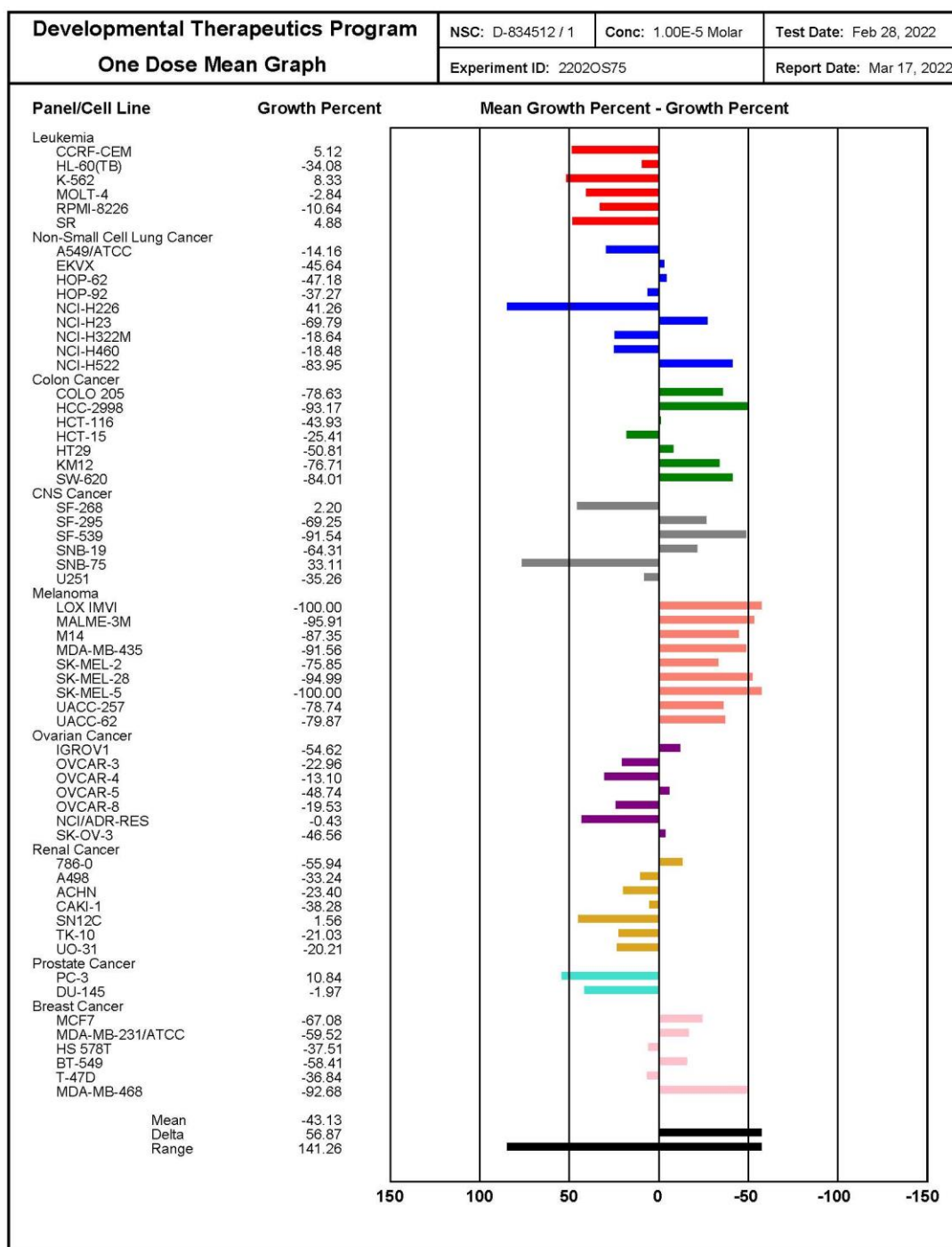


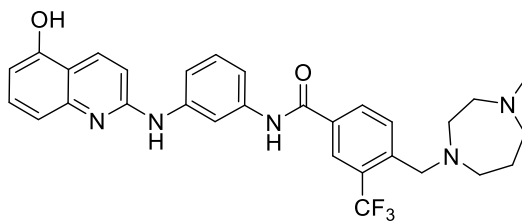
HJ_2-33 = 2-47 (16b)



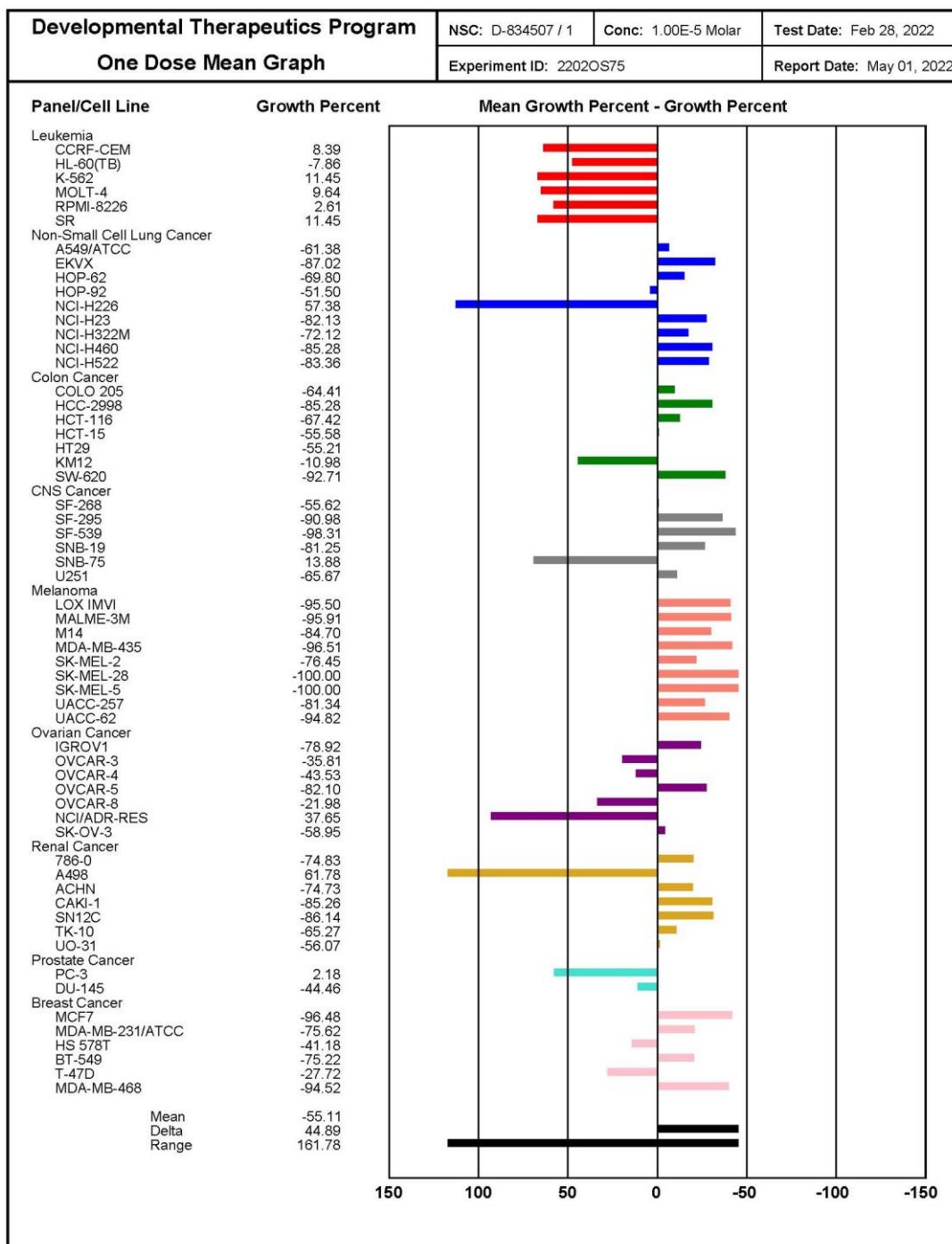


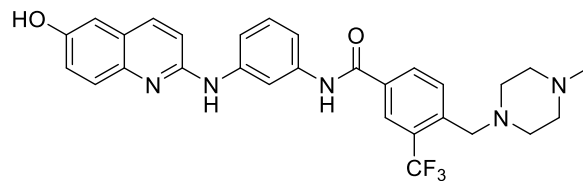
HJ_2-14 (17a)



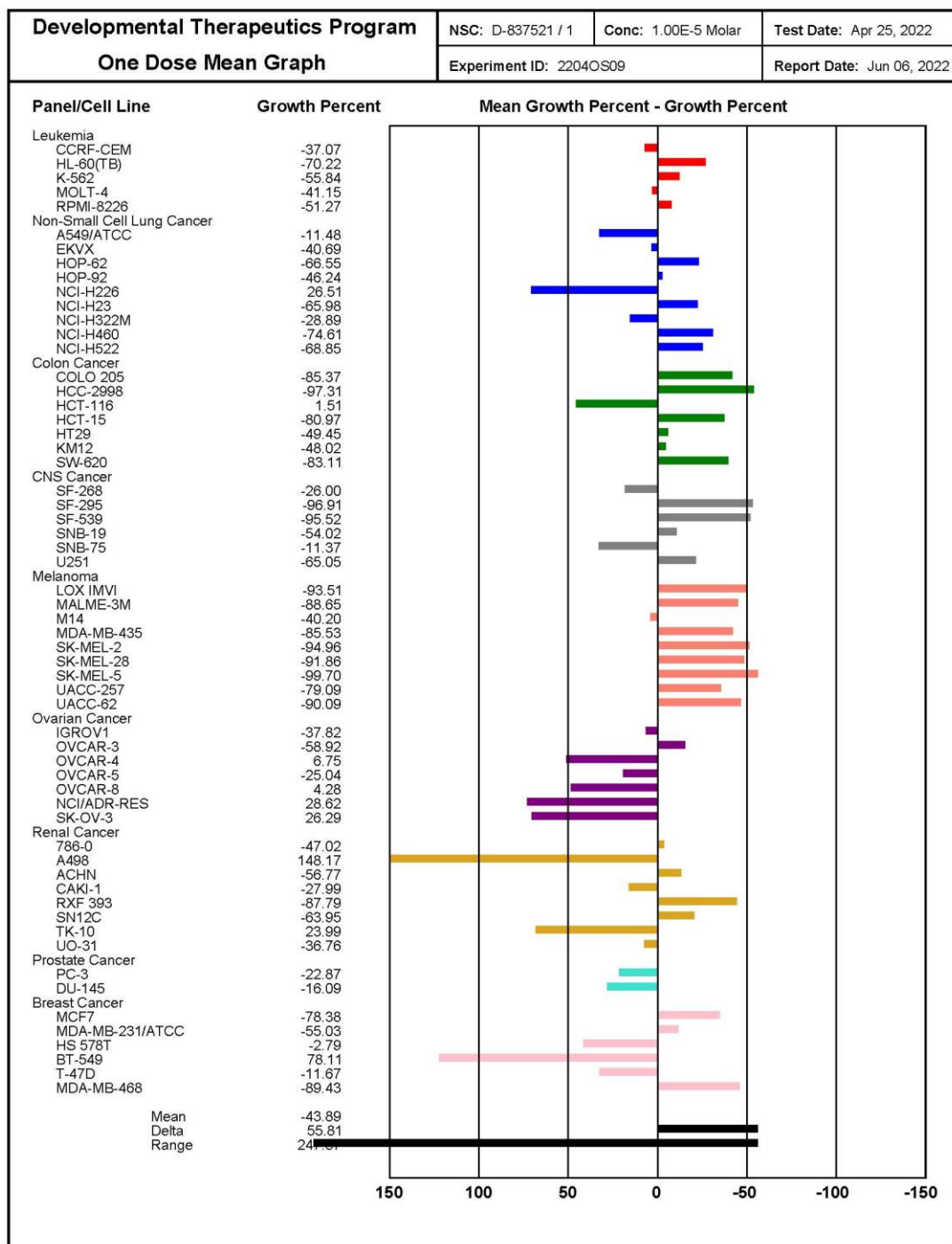


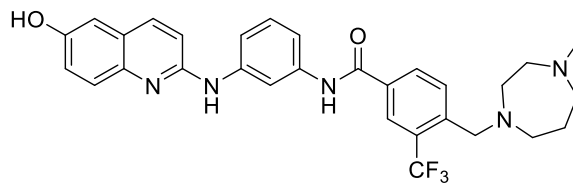
HJ_2-12 (17b)



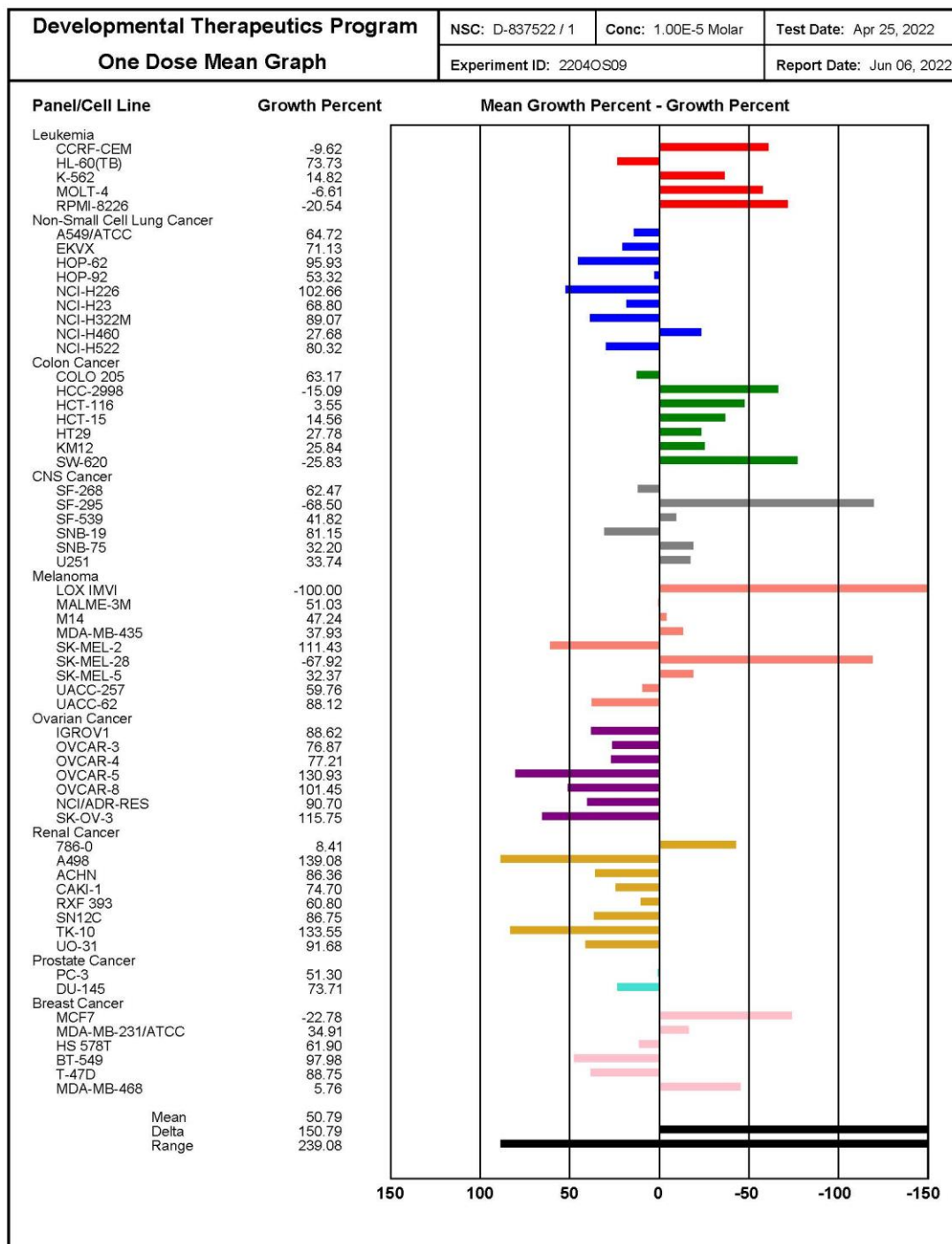


HJ_2-56 (18a)





HJ_2-57 (18b)



5) Molecular docking

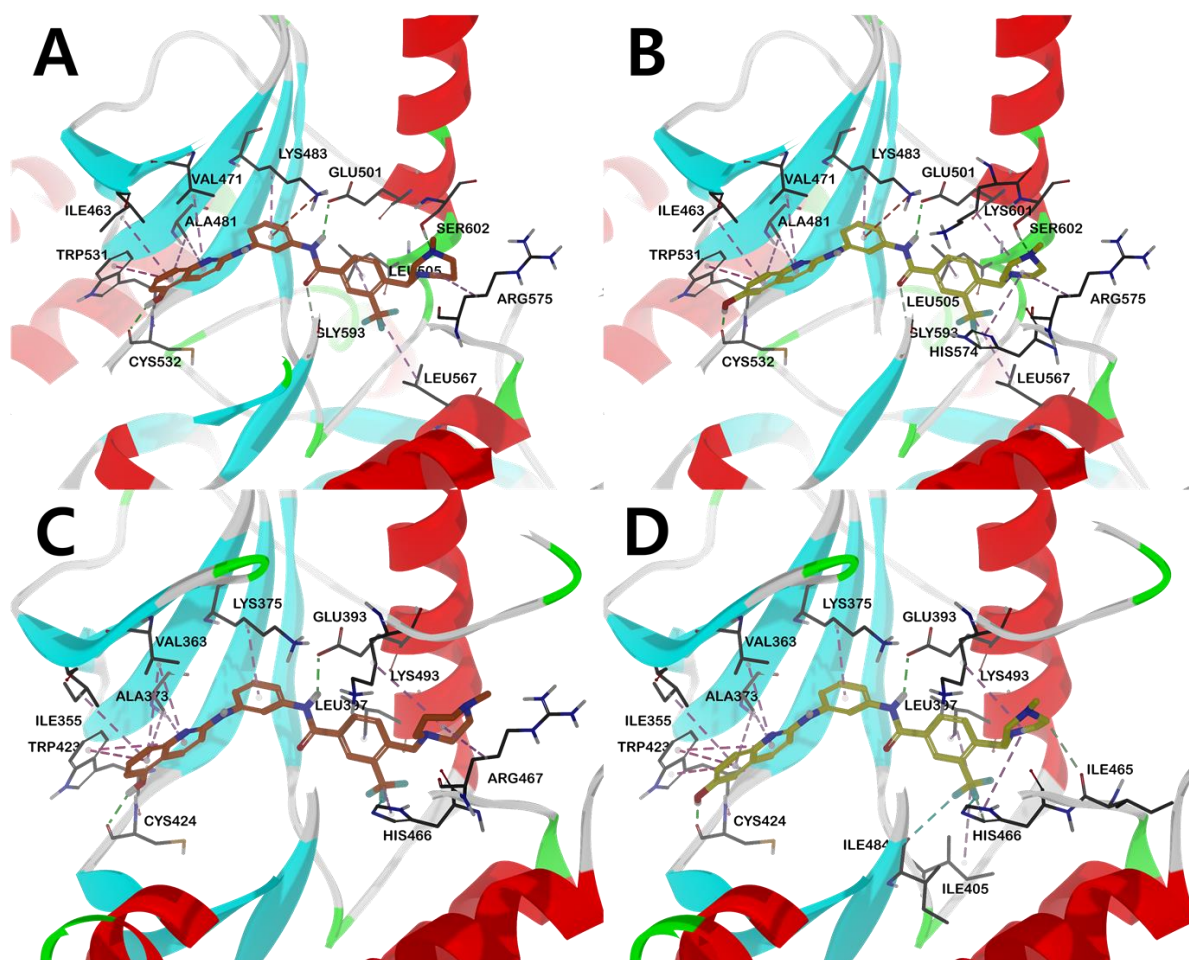


Figure S1. The binding mode of compounds (A) **17b** and (B) **18a** in the catalytic kinase domain of B-RAF^{V600E} model. The binding mode of compound (C) **17b** and (D) **18a** in the C-RAF homology model. For clarity purpose, the residues showing interactions were shown. Compounds **17b** and **18a** are shown in orange and yellow colored stick model. Binding interactions are shown in various colored dashes lines, crucial hydrogen bonding shown in green dashes lines.

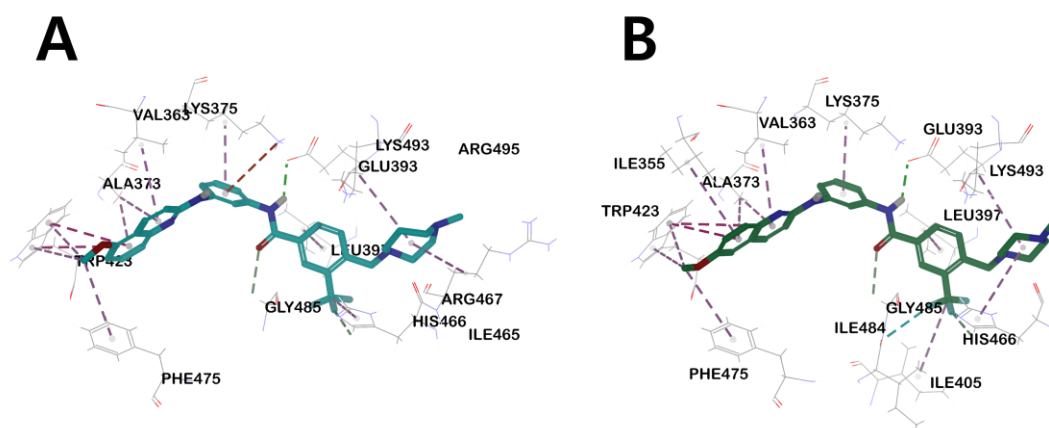


Figure S2. Binding model of compounds (A) **14b** and (B) **16a** in the C-Raf homology model. Compounds **14b** and **16a** shown by sky-blue and green colored stick model, respectively. The surrounding residues were shown as line model and various interactions shown as dashes line.

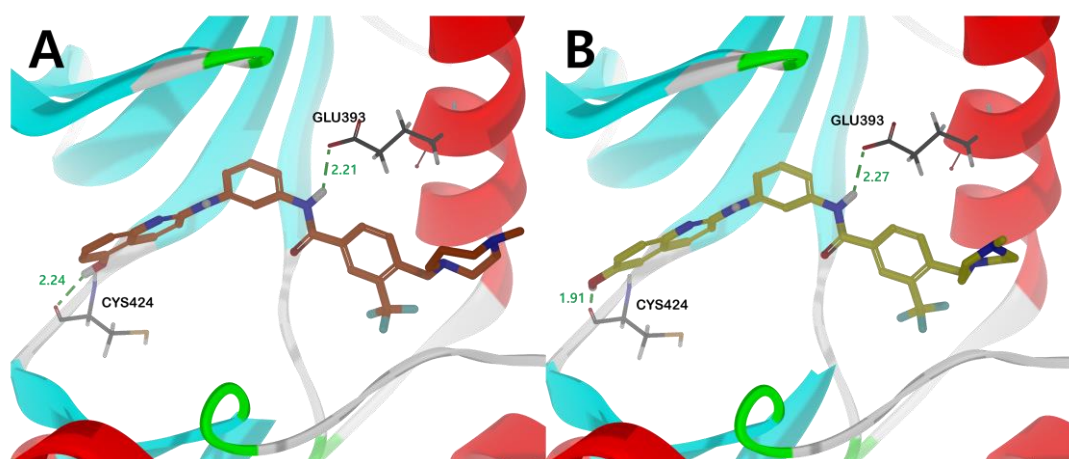


Figure S3. Binding model of compounds (A) **17b** and (B) **18a** in the C-Raf homology model. For clarity purpose, only key residues showing interactions were shown. Compounds **17b** and **18a** are shown in orange and yellow colored stick model. Key hydrogen bonding interactions are shown in green dashes. The number near the dashes indicates the bonding distance, in units of Å.

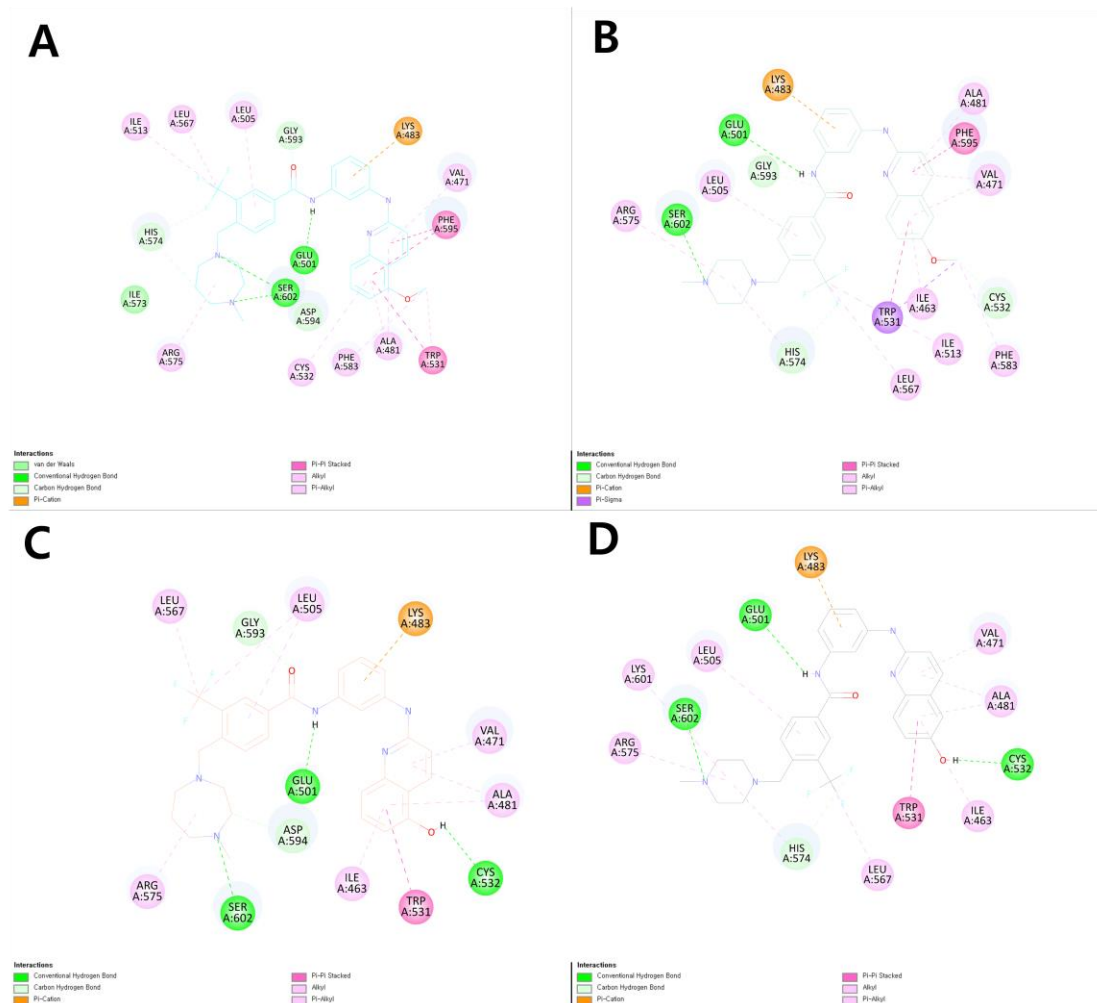


Figure S4. 2D binding mode of compound (A) **14b**, (B) **16a**, (C) **17b** and (D) **18a** with B-Raf^{V600E}. Various interactions are depicted by different color legends. The inhibitors are shown by line, interacting residues by colored sphere, and interactions by dash lines.

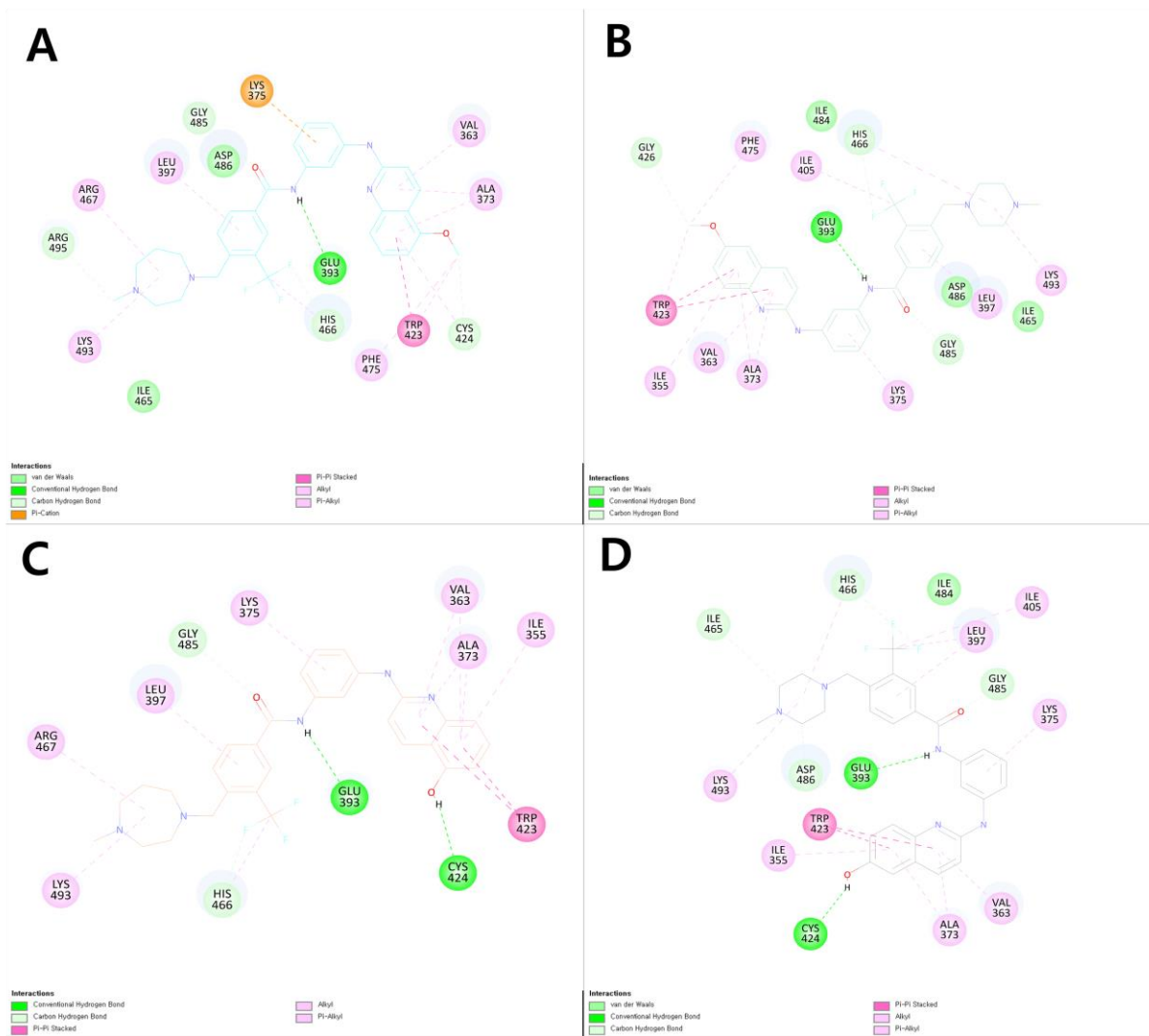


Figure S5. 2D binding mode of compounds (A) **14b**, (B) **16a**, (C) **17b** and (D) **18a** with C-RAF homology model. Various interactions are depicted by different color legends. Inhibitors are shown by line, interacting residues by colored sphere, and interactions by dash lines.

6) Molecular dynamics

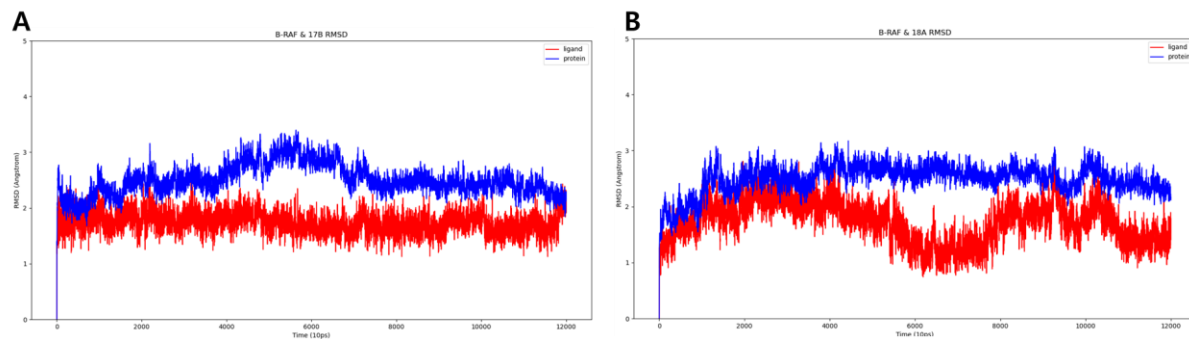


Figure S6. RMSD values of (A) **17b** and (B) **18a** compounds bounded B-RAF^{V600E} protein during 120 ns of molecular dynamics simulation.

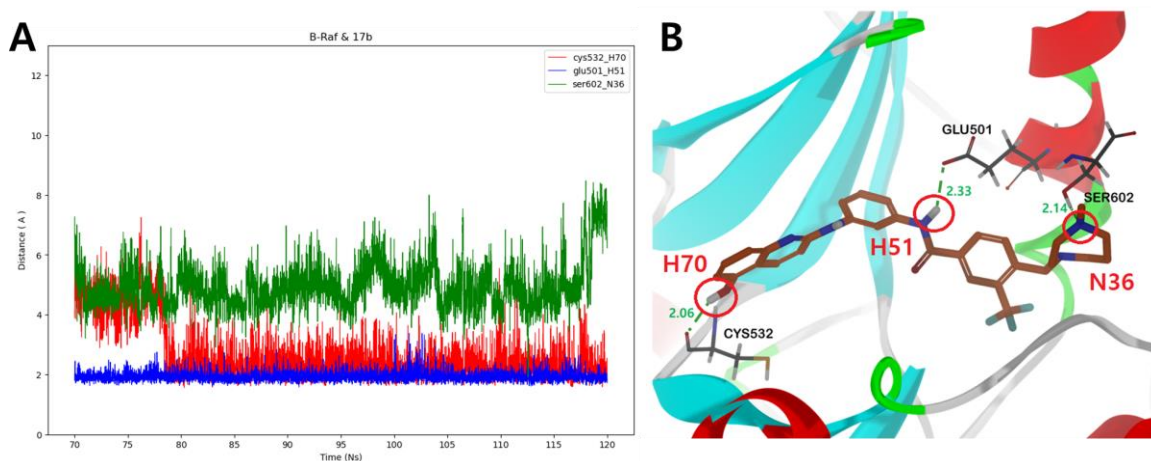


Figure S7. (A) Distance of three key hydrogen bonding interaction between B-RAF^{V600E} protein and **17b** during 70-120 ns dynamics simulation. The distances between Cys532 and H70, Glu501 and H51, and Ser602 and N36 are marked in red, blue, and green, respectively. (B) Initial binding structure of **17b** and B-RAF^{V600E} used in dynamics simulation. Atoms participating in three hydrogen bonding bonds whose distances are monitored are marked with red circles.

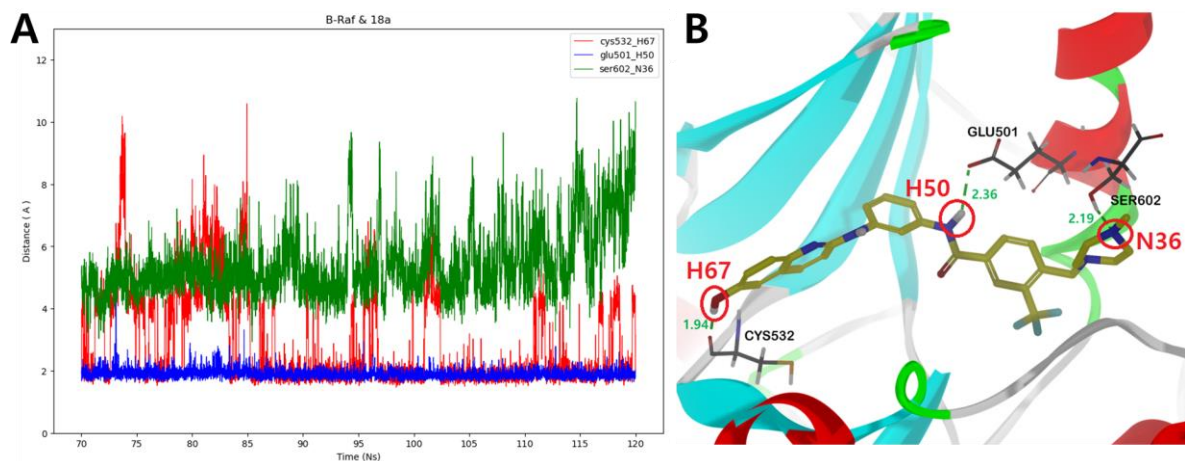


Figure S8. (A) Distance of three key hydrogen bonding interaction between B-Raf^{V600E} protein and **18a** during 70-120 ns dynamics simulation. The distances between Cys532 and H67, Glu501 and H50, and Ser602 and N36 are marked in red, blue, and green, respectively. (B) Initial binding structure of **18a** and B-Raf^{V600E} used in dynamics simulation. Atoms participating in three hydrogen bonding bonds whose distances are monitored are marked with red circle.