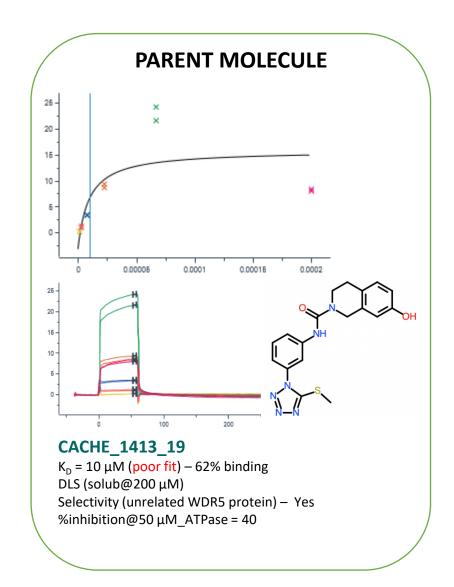
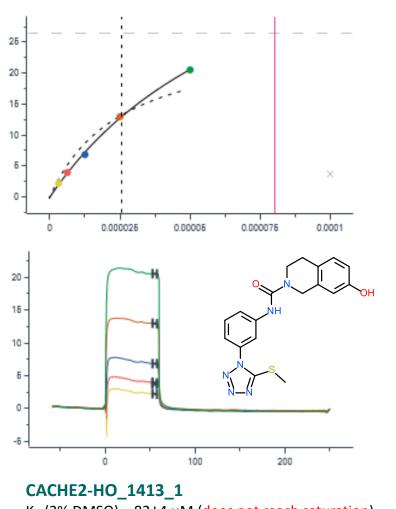
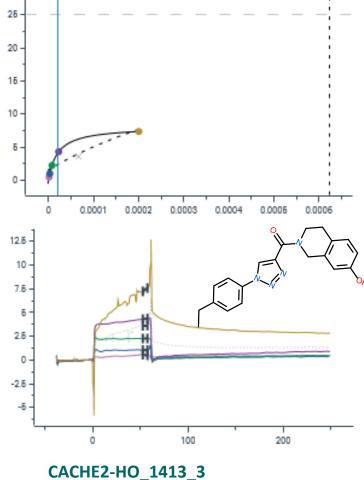
# CACHE#2 – NSP13 SARS2 Participant 1413



24 analogs of CACHE\_1413\_19 were submitted for round 2. 2 compounds, including re-supplied parent molecule showed weak binding profile.

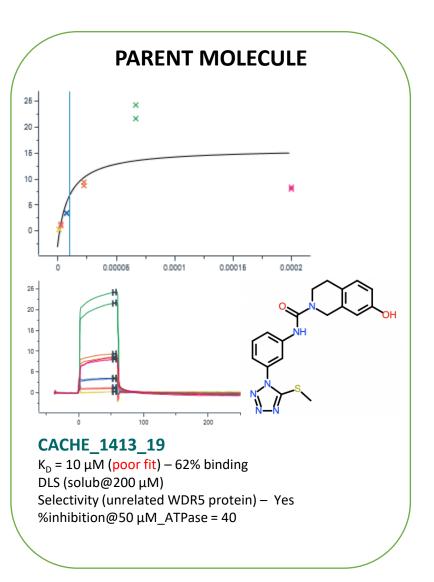


 $K_D$  (3% DMSO) = 82±4 μM (does not reach saturation) – 207±%7 binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@50 μM) %inhibition@50 μM\_ATPase = 20



 $K_D = 20 \mu M - 32\% \text{ binding}$ DLS (solub@200  $\mu$ M)

%inhibition@50 μM\_ATPase = 11

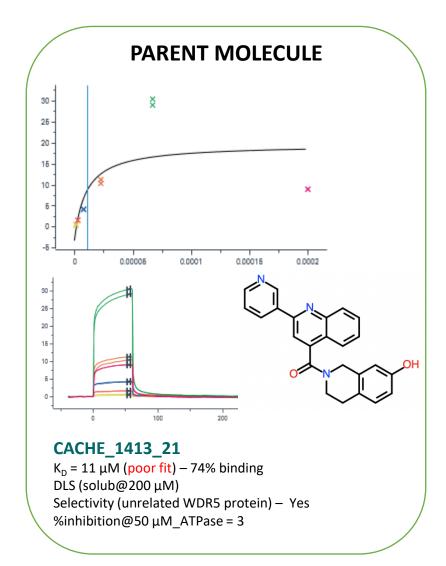


Dose response:		Dose response: Confirmed	ON NH	HO
CACHE_ID CACHE2-HO_1413_1 Parent CACHE_1413_19 distance_to_pai 0	CACHE_ID CACHE2-HO_1413_2 Parent CACHE_1413_19 distance_to_pai 0.1879	CACHE_ID CACHE2-HO_1413_3 Parent CACHE_1413_19 distance_to_pai 0.393	CACHE_ID CACHE2-HO_1413_4 Parent CACHE_1413_19 distance_to_pai 0.3956	CACHE_ID CACHE2-HO_1413_5 Parent CACHE_1413_19 distance_to_pai 0.407
O N OH	i HN O	NH OH		
CACHE_ID CACHE2-HO_1413_6 Parent CACHE_1413_19 distance_to_pai 0.42	CACHE_ID CACHE2-HO_1413_7 Parent CACHE_1413_19 distance_to_pai 0.4324	CACHE_ID CACHE2-HO_1413_8 Parent CACHE_1413_19 distance_to_pai 0.433	CACHE_ID CACHE2-HO_1413_9 Parent CACHE_1413_19 distance_to_pai 0.4691	CACHE_ID CACHE2-HO_1413_10 CACHE_1413_19 distance_to_pai 0.4702
	Q D (R)			F NH
CACHE_ID CACHE2-HO_1413_11 Parent CACHE_1413_19 distance_to_pai 0.495	CACHE_ID CACHE2-HO_1413_12 Parent CACHE_1413_19 distance_to_pai 0.4961	CACHE_ID CACHE2-HO_1413_13 Parent CACHE_1413_19 distance_to_pai 0.5098	CACHE_ID CACHE2-HO_1413_14 Parent CACHE_1413_19 distance_to_pai 0.5159	CACHE_ID CACHE2-HO_1413_15 Parent CACHE_1413_19 distance_to_pai 0.517

## **PARENT MOLECULE** 20 -16 -0.0002 0.00006 0.0001 0.00016 CACHE\_1413\_19 $K_D = 10 \mu M \text{ (poor fit)} - 62\% \text{ binding}$ DLS (solub@200 $\mu M$ ) Selectivity (unrelated WDR5 protein) – Yes %inhibition@50 µM\_ATPase = 40

## Tested analogs (cont...)

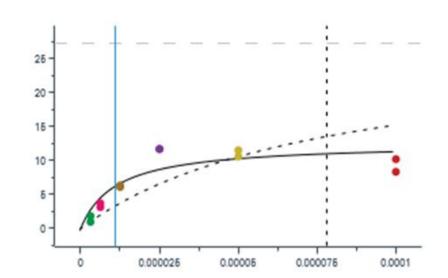
	4			
CACHE_ID	CACHE_ID CACHE2-HO_1413_22 Parent CACHE_1413_19 distance_to_pai 0.5517	CACHE_ID CACHE2-HO_1413_23 Parent CACHE_1413_19 distance_to_pai 0.5538	CACHE_ID CACHE2-HO_1413_24 Parent CACHE_1413_19 distance_to_pai 0.5635	CACHE_ID CACHE2-HO_1413_25 Parent CACHE_1413_19 distance_to_pai 0.5636
CACHE_ID	CACHE_ID CACHE2-HO_1413_27 Parent CACHE_1413_19 distance_to_pai 0.5656	CACHE_ID CACHE2-HO_1413_28 Parent CACHE_1413_19 distance_to_pai 0.5689	CACHE_ID CACHE2-HO_1413_29 Parent CACHE_1413_19 distance_to_pai 0.6069	

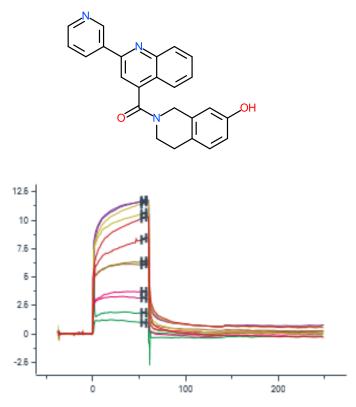


15 analogs of CACHE\_1413\_21 were submitted for round 2. 1 compound (re-supplied parent molecule) confirmed binding by SPR, showing the tendency to reach saturation.

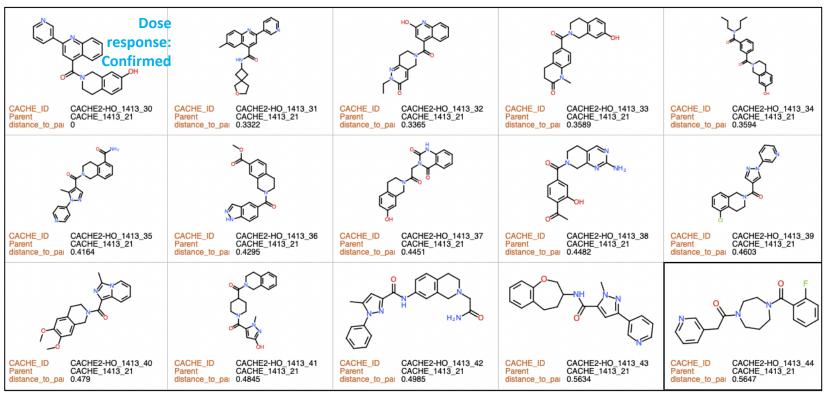
### CACHE2-HO\_1413\_30

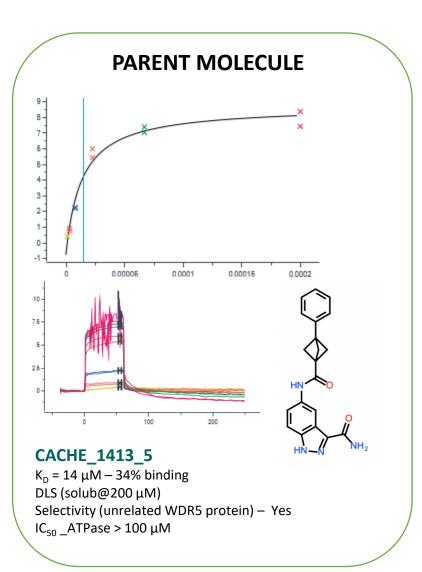
 $K_D$  = 11 μM (poor fit) – 46% binding DLS (solub@50 μM) Selectivity (unrelated WDR5 protein) – Yes %inhibition@50 μM ATPase = 3

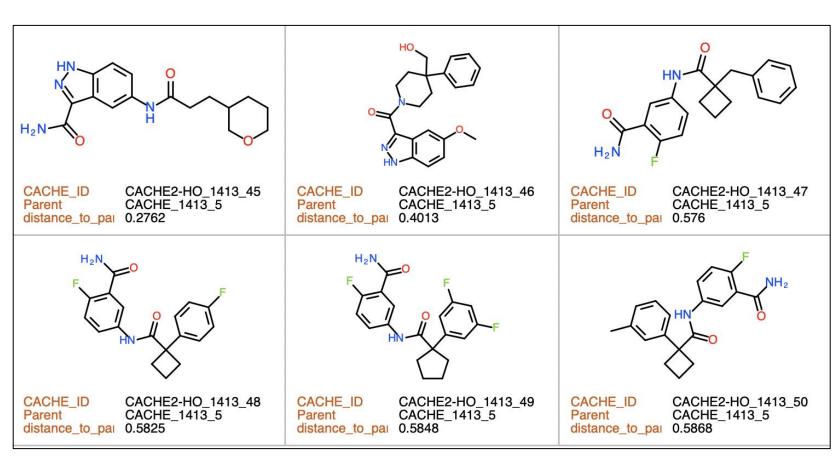


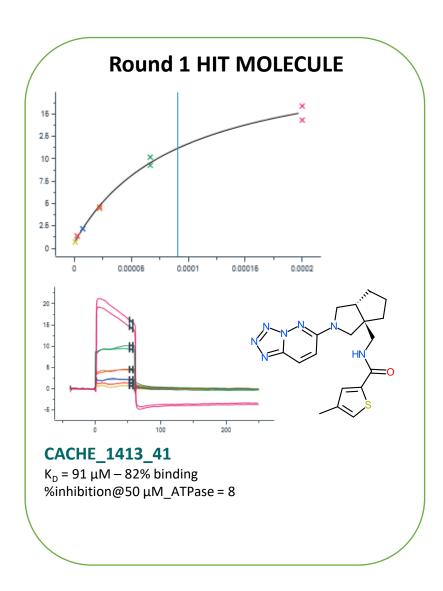


## **PARENT MOLECULE** 26 -20 -16 -0.00006 0.0001 0.0002 0.00016 CACHE\_1413\_21 $K_D = 11 \mu M \text{ (poor fit)} - 74\% \text{ binding}$ DLS (solub@200 µM) Selectivity (unrelated WDR5 protein) – Yes %inhibition@50 $\mu$ M ATPase = 3



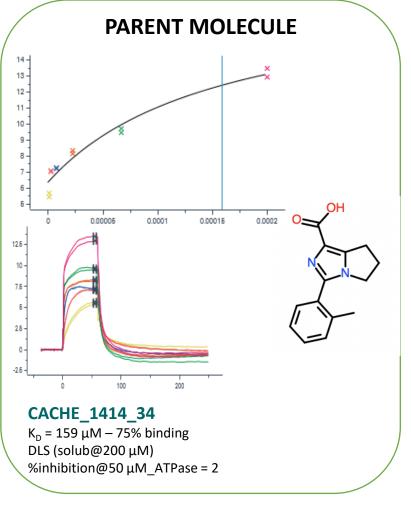






No compound in Round 2

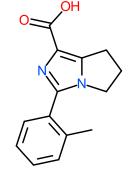
# CACHE#2 – NSP13 SARS2 Participant 1414

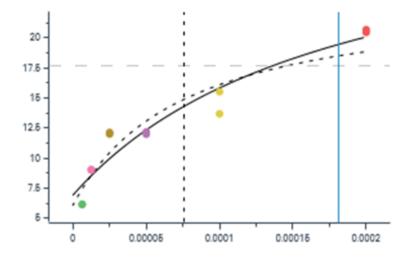


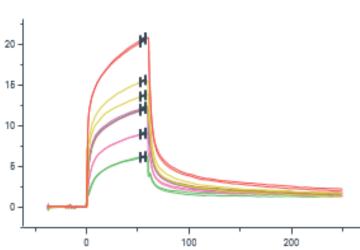
10 analogs of CACHE\_1414\_34 were submitted for round 2. 1 compound (re-supplied parent molecule) confirmed binding by SPR, showing a tendency to reach saturation.

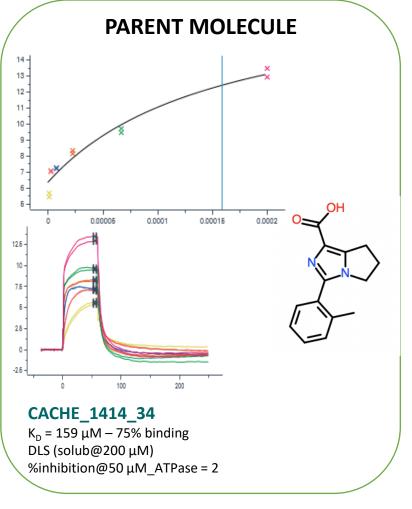
## CACHE2-HO\_1414\_1

 $K_D$  = 182  $\mu$ M – 142% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@50  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 11

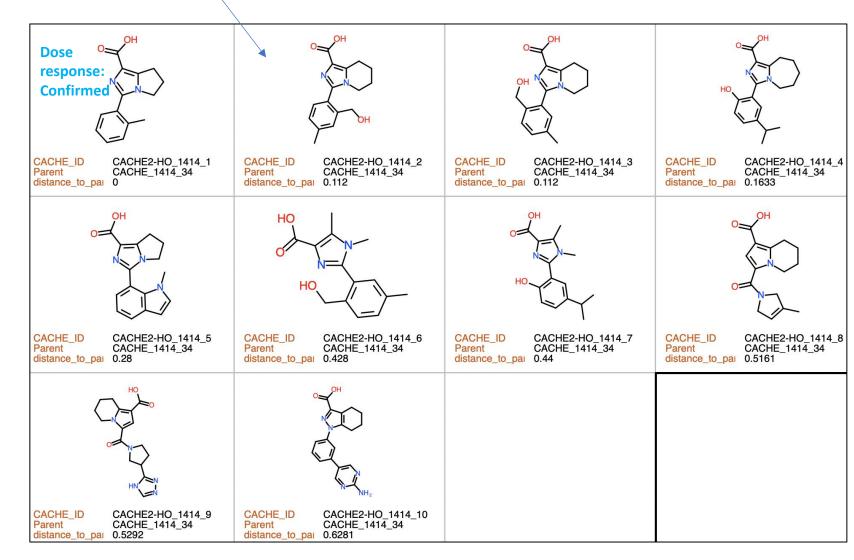


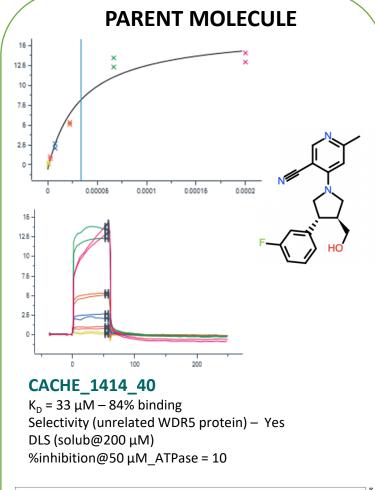


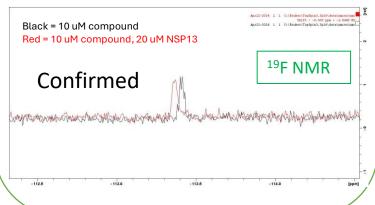




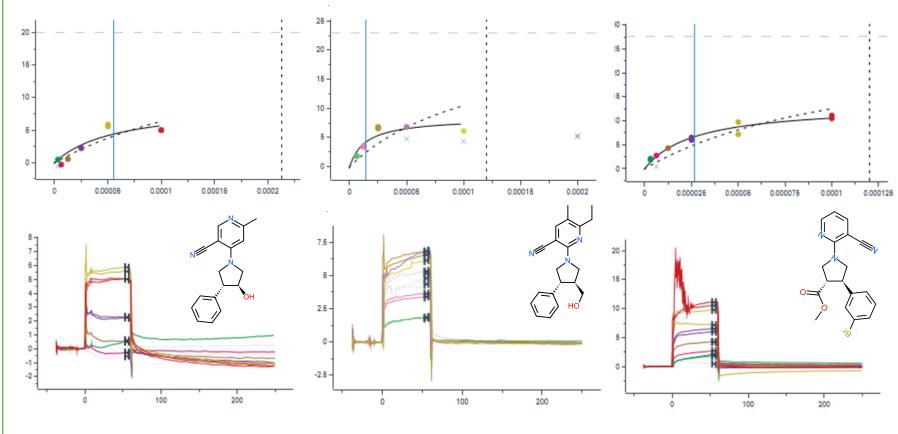
Blank means inactive Tested analogs







24 analogs of CACHE\_1414\_40 were submitted for round 2. 3 compounds showed a binding response by SPR.



## CACHE2-HO\_1414\_13

 $K_D_1$  = 41 μM – 66% binding  $K_D_2$  = 56 μM – 44% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 11

#### CACHE2-HO 1414 14

 $K_D$  (3% DMSO) = 14  $\mu$ M – 36% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 5

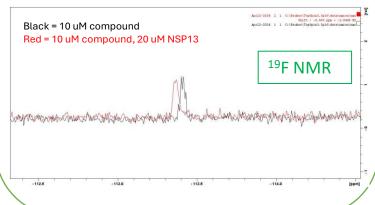
#### CACHE2-HO 1414 20

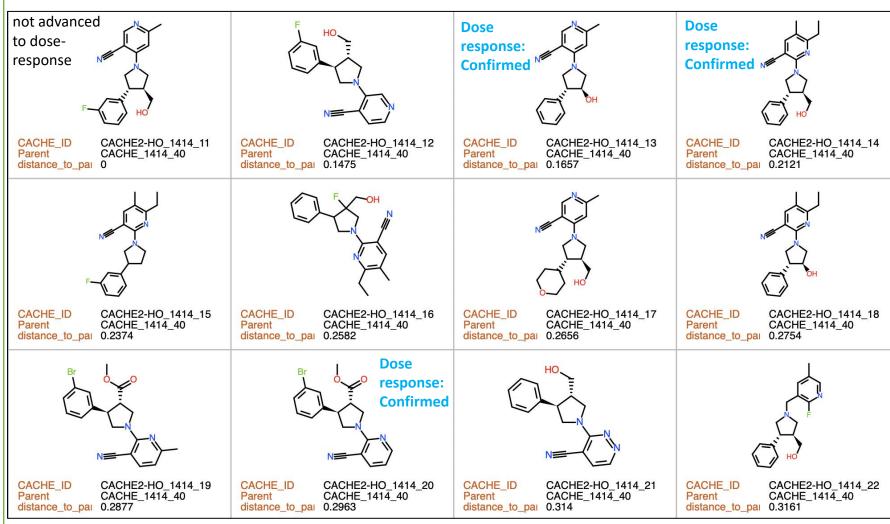
 $K_D = 1 = 36 \mu M - 62\%$  binding  $K_D = 27 \mu M - 49\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 8

## PARENT MOLECULE 12.5 0.00015 0.0002 0.00005 0.0001

## CACHE\_1414\_40

 $K_D = 33 \mu M - 84\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 10

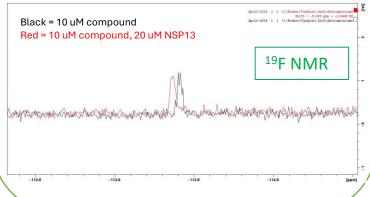




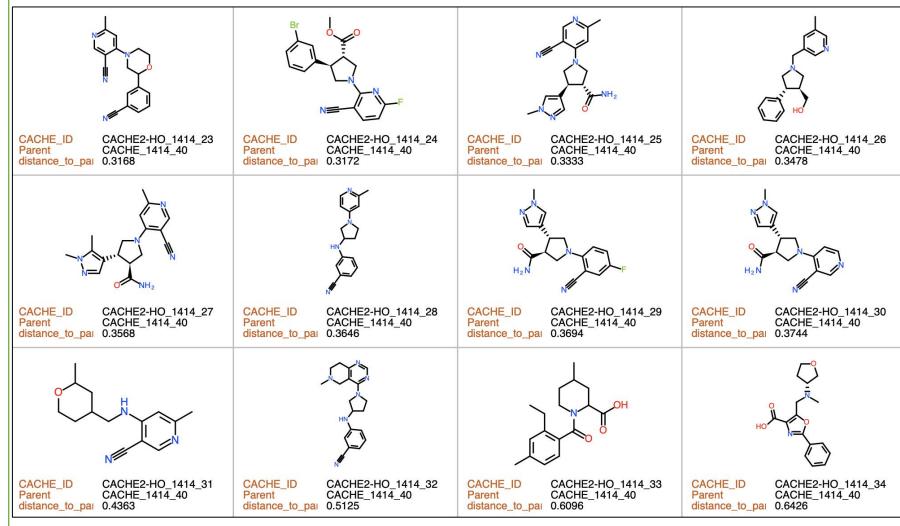
## PARENT MOLECULE 12.5 0.00015 0.0002 0.00005 0.0001

## CACHE\_1414\_40

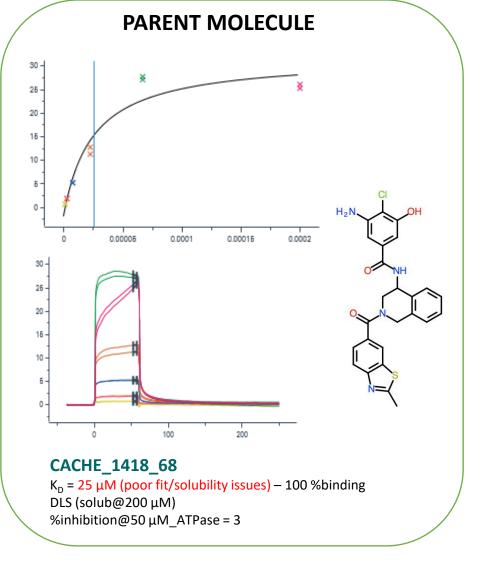
 $K_D = 33 \mu M - 84\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 10



## Tested analogs (cont...)



# CACHE#2 – NSP13 SARS2 Participant 1418



6 analogs of CACHE\_1418\_68 were submitted for round 2. 1 of them showed a super stoichiometric binding to NSP13 in addition to weak non-selective binding to the unrelated protein WDR5.

## CACHE2-HO\_1418\_34

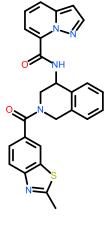
 $K_D _1 = 61 \mu M - 298 \% binding$ 

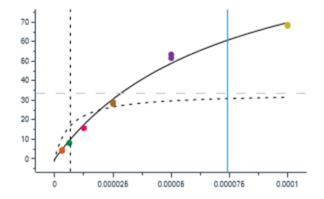
 $K_D = 2 = 74 \mu M - 362 \% binding$ 

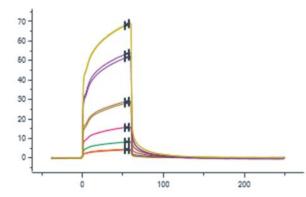
Selectivity (unrelated WDR5 protein) – weak binding

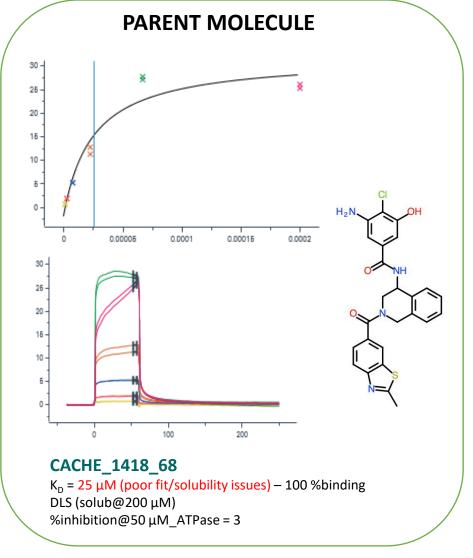
DLS (solub/no aggreg@200 µM)

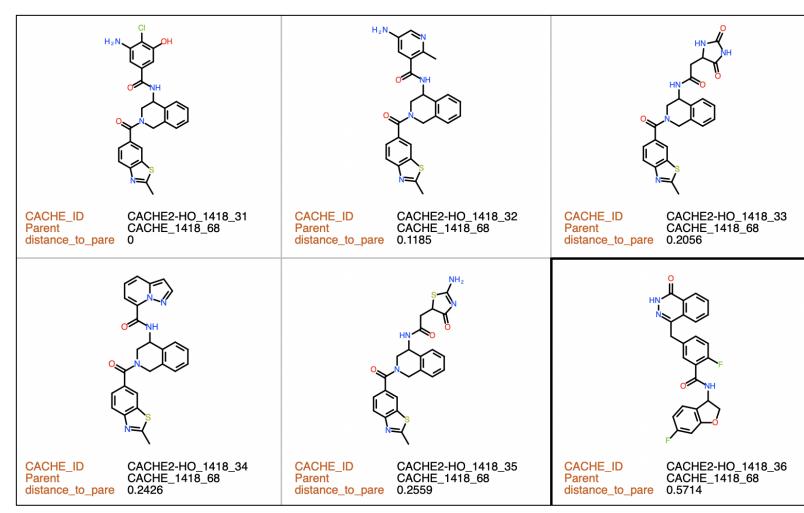
%inhibition@50  $\mu$ M\_ATPase = 6

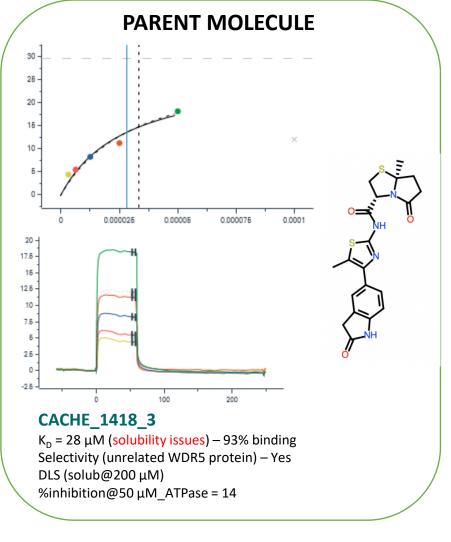








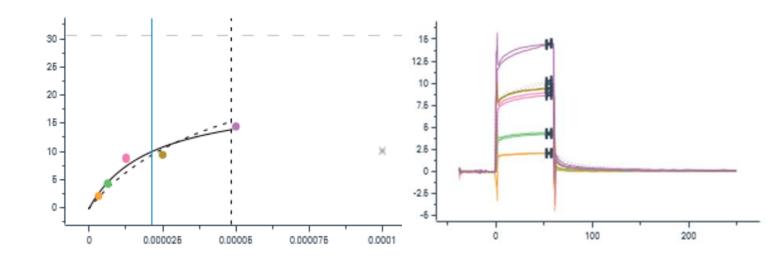




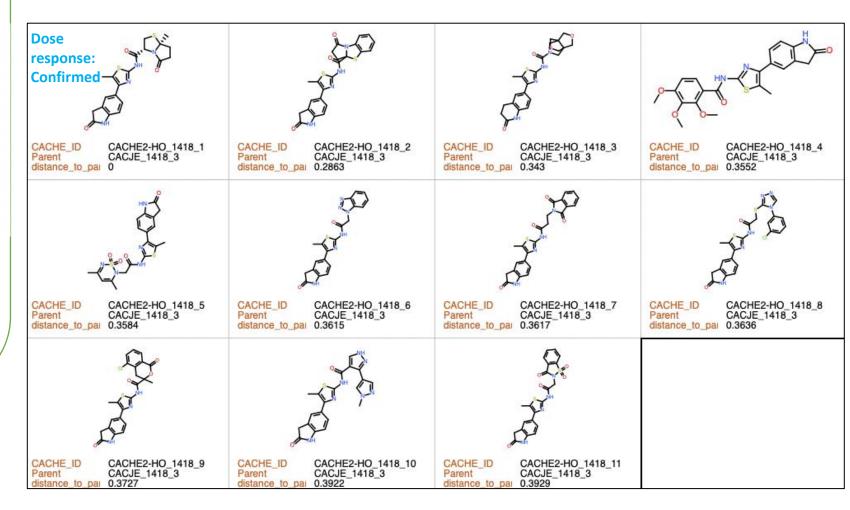
11 analogs of CACHE\_1418\_3 were submitted for round 2 including the re-supplied parent molecule. The Re-supplied hit confirmed dose dependent binding to NSP13.

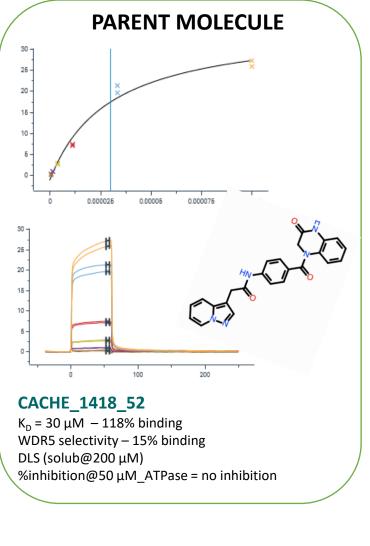
### CACHE2-HO\_1418\_1

 $K_D$  (3% DMSO) = 21 μM (solubility issues) – 65% binding Selectivity (unrelated WDR5 protein) – Yes %inhibition@50 μM ATPase = 13

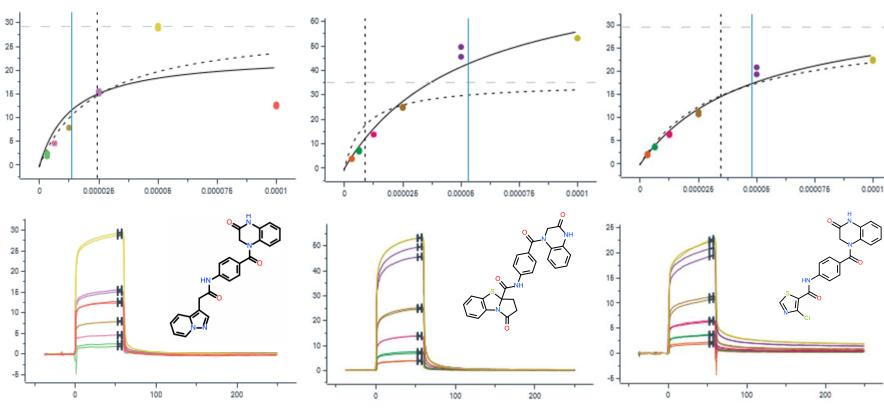


## PARENT MOLECULE 25 -0.000026 0.00006 0.000076 0.0001 17.5 15 -12.5 2.5 **CACHE 1418 3** $K_D = 28 \mu M$ (solubility issues) – 93% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 µM ATPase = 14





9 analogs of CACHE\_1418\_52 were submitted for round 2. 1 of them showed clear dose dependent binding response by SPR.



## CACHE2-HO\_1418\_22

 $K_D$  = 14 μM (poor fit) – 80% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 19

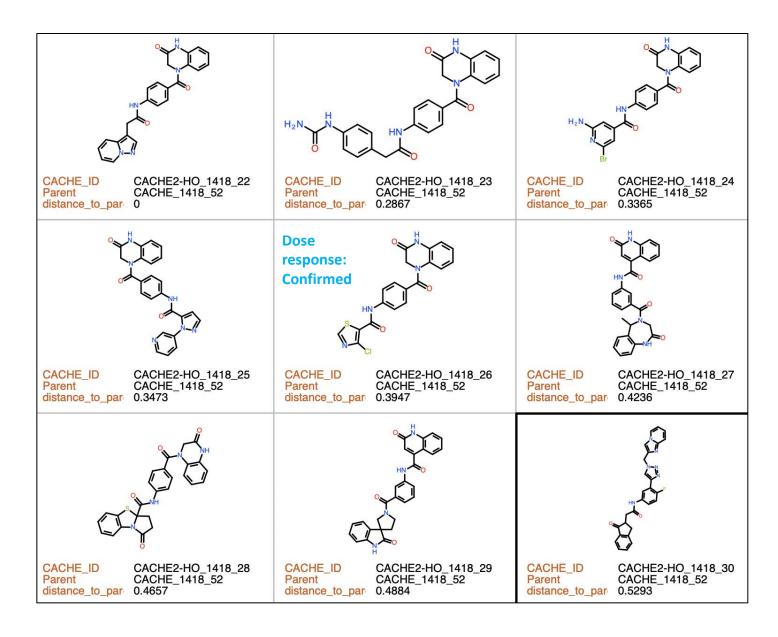
## CACHE2-HO\_1418\_28

 $K_D _1 = 56 \mu M - 781\%$  binding  $K_D _2 = 53 \mu M - 246\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 21

### CACHE2-HO\_1418\_26

 $K_D$  = 48  $\mu$ M – 118% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M\_ATPase = 15

## **PARENT MOLECULE** 0.000026 0.00006 0.000076 20 200 CACHE\_1418\_52 $K_D = 30 \,\mu\text{M} - 118\% \text{ binding}$ WDR5 selectivity - 15% binding DLS (solub@200 µM) %inhibition@50 µM ATPase = no inhibition

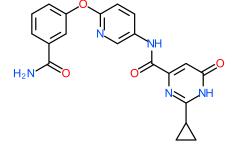


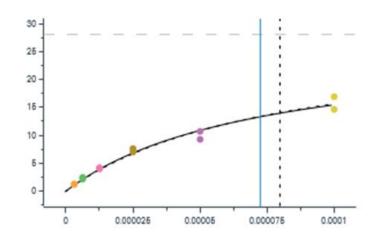
## **PARENT MOLECULE** 0.000026 0.0000B 0.000076 CACHE\_1418\_37 $K_D = 33\pm 1 \,\mu\text{M} - 89\pm 17\%$ binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 µM\_ATPase = 72% Not confirmed by dose response IC<sub>50</sub>>150

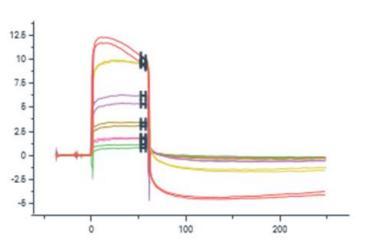
6 analogs of CACHE\_1418\_37 were submitted for round 2. 1 of them (re-supplied parent molecule) confirmed a selective dose dependent binding response.

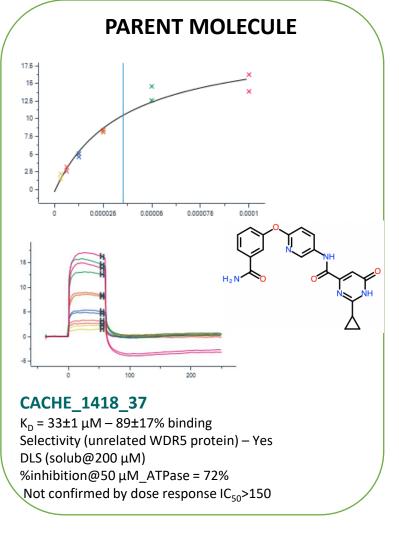
#### CACHE2-HO 1418 15

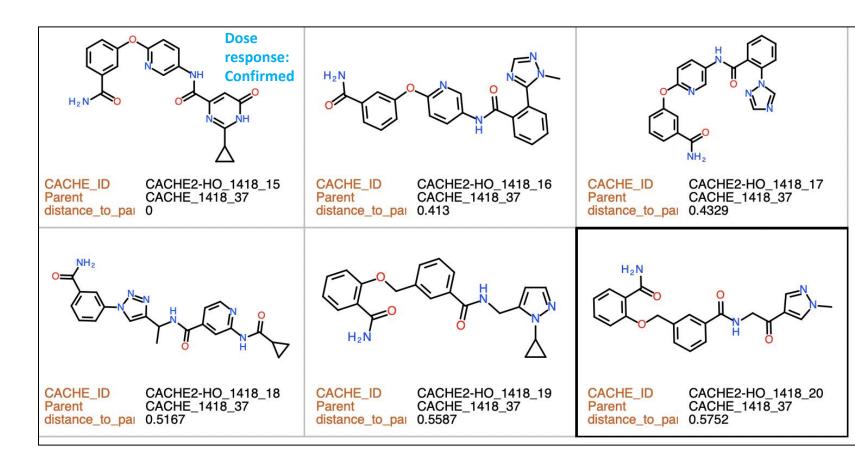
 $K_D$  (3% DMSO) = 72 μM – 95% binding At 100 uM concentration shows binding to the reference cell Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 16

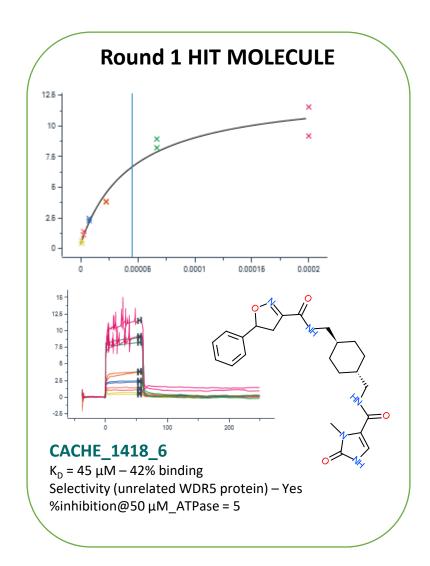


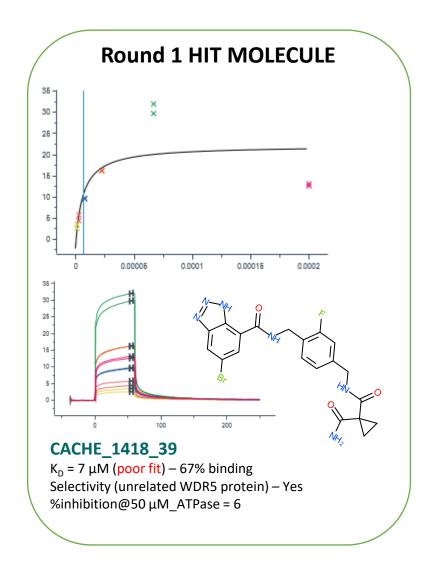




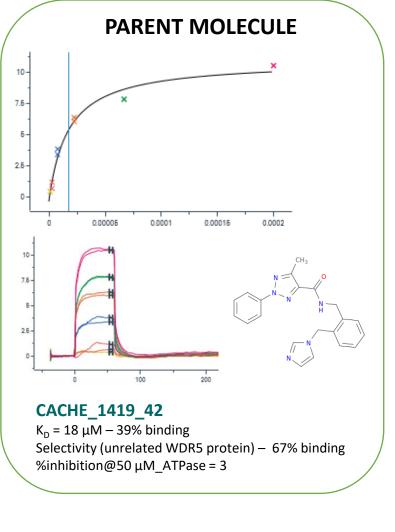




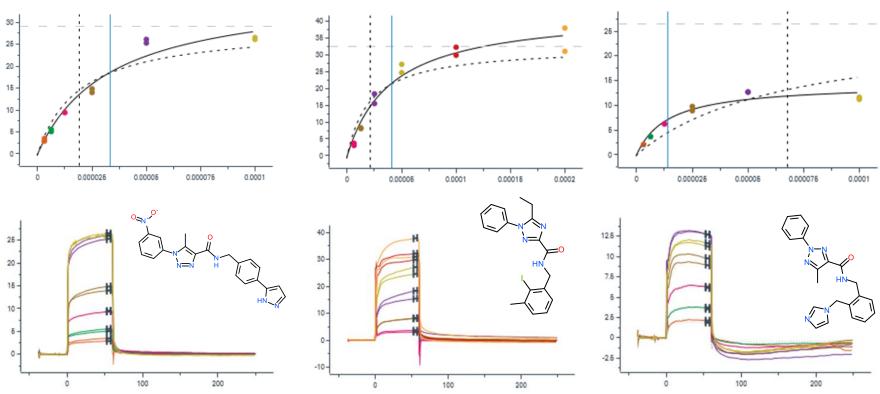




# CACHE#2 – NSP13 SARS2 Participant 1419



48 analogs of CACHE\_1419\_42 were submitted for round 2. 3 compounds including re-supplied parent molecule showed a binding response by SPR.



## CACHE2-HO\_1419\_18

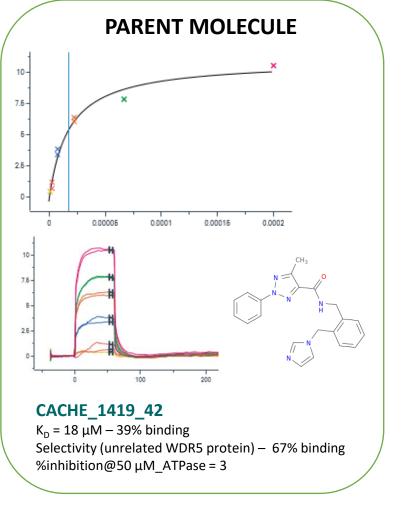
 $K_D$  = 33  $\mu$ M - 128% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub/@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 11

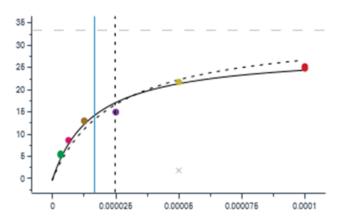
### CACHE2-HO 1419 30

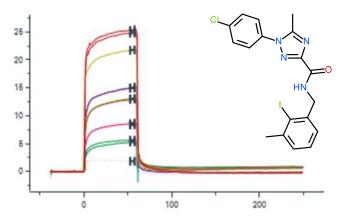
 $K_D$  = 41 μM – 133% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 14

#### **CACHE2-HO 1419 1**

 $K_D = 14 \mu M - 54\%$  binding Selectivity (unrelated WDR5 protein) – 16% binding DLS (solub@100 μM) %inhibition@50 μM ATPase = 8

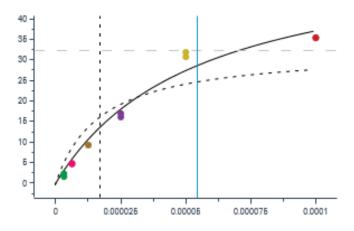


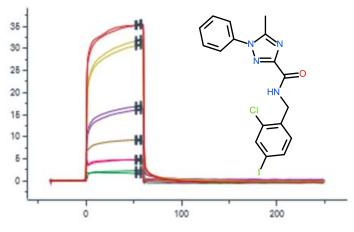




## CACHE2-HO\_1419\_28

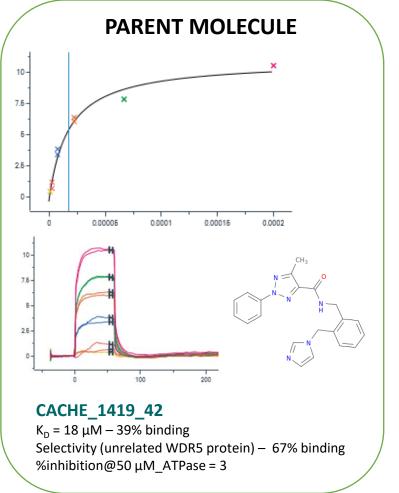
 $K_D _1 = 41 \ \mu M - 81\%$  binding  $K_D _2 = 17 \ \mu M - 85\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu M$ ) %inhibition@50  $\mu M$ \_ATPase = 14

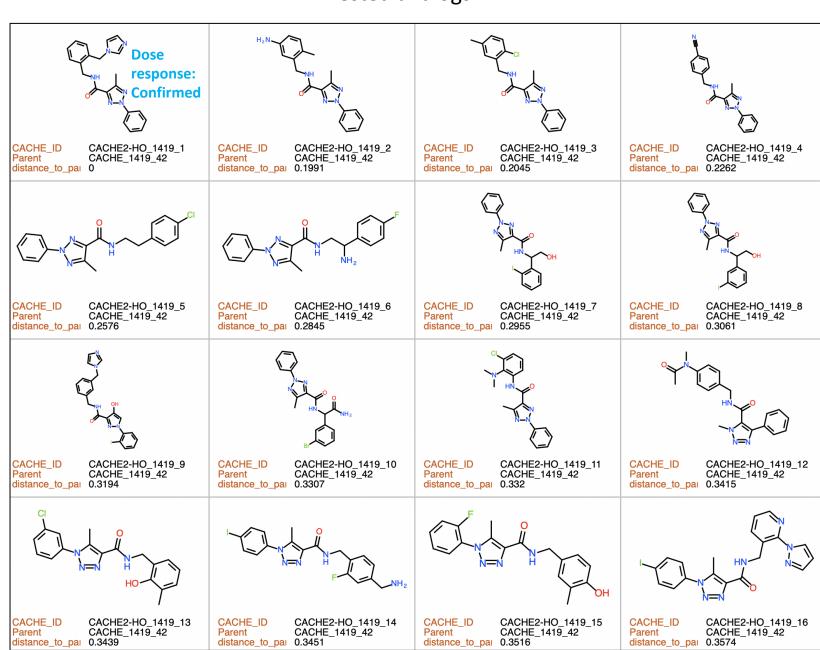


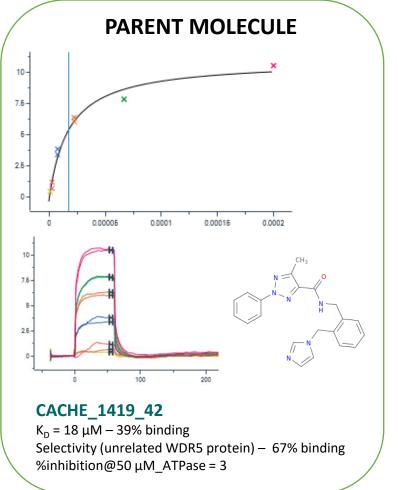


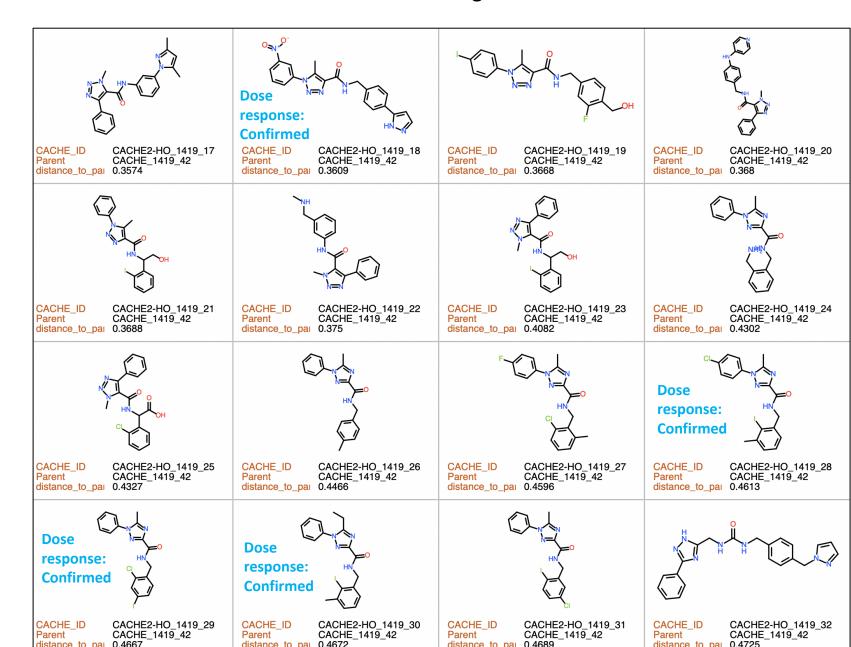
## CACHE2-HO\_1419\_29

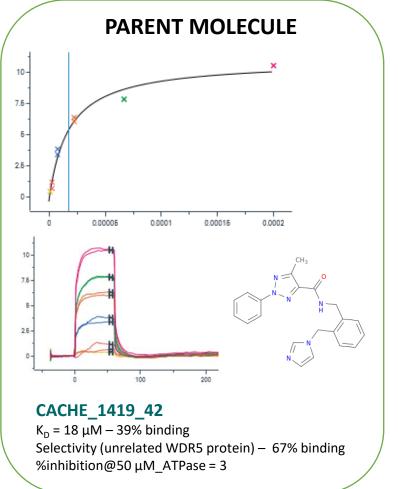
 $K_D _1 = 51 \mu M - 157\%$  binding  $K_D _2 = 55 \mu M - 177\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 13

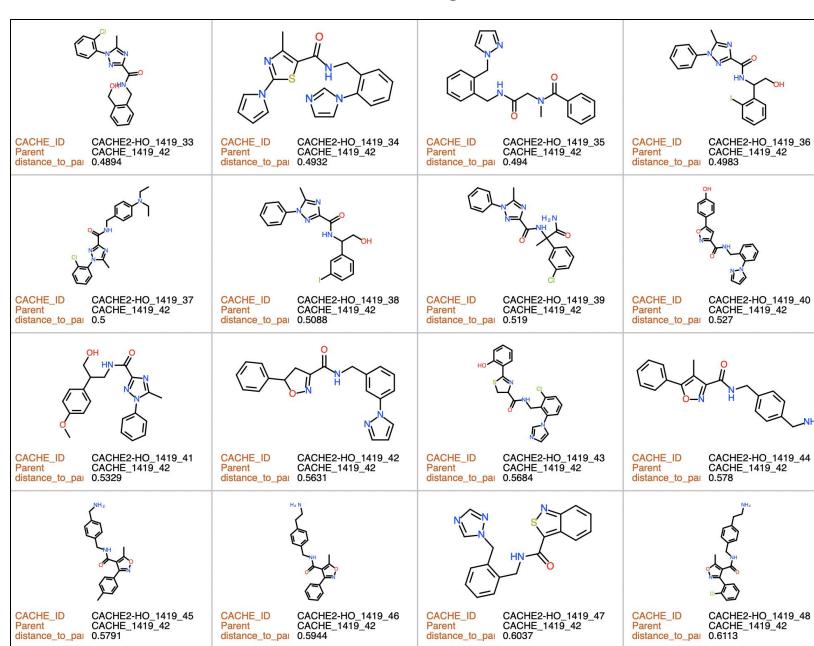








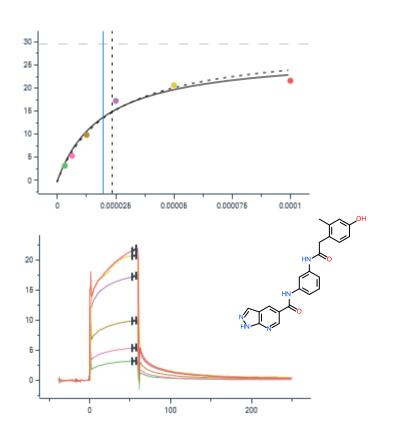




# CACHE#2 – NSP13 SARS2 Participant 1421

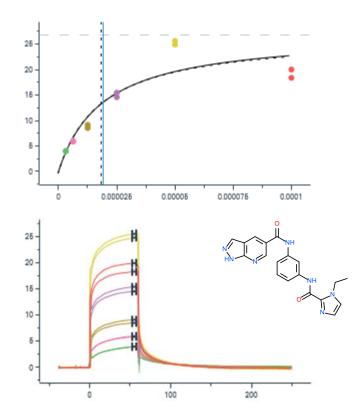
## **PARENT MOLECULE** 0.000026 0.00006 CACHE\_1421\_62 $K_D = 62 \mu M - 70\%$ binding %inhibition@50 µM ATPase = 13

11 analogs of CACHE\_1421\_62 were submitted for round 2. 2 compounds confirmed a selective dose dependent binding response by SPR.



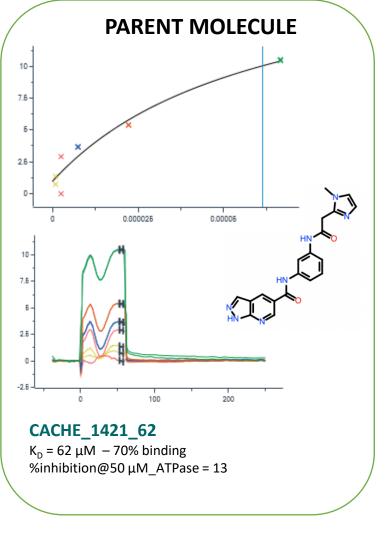


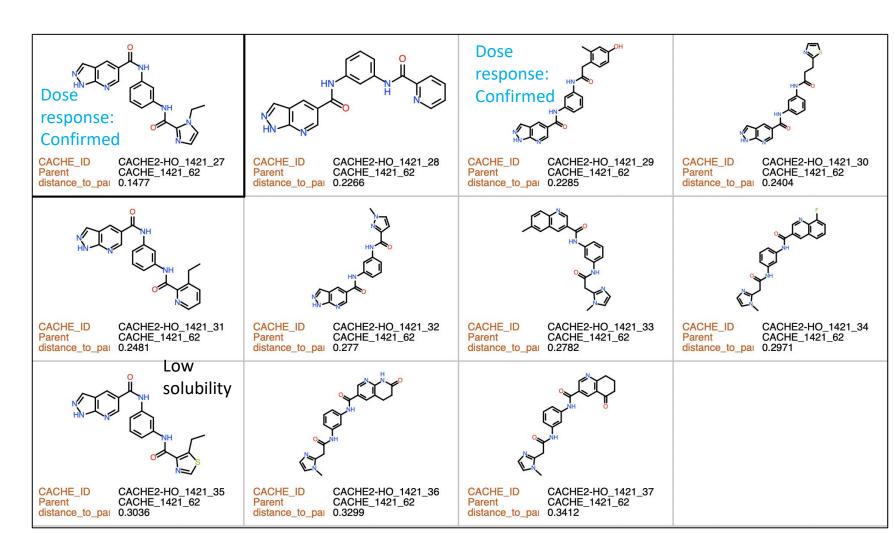
 $K_D$  (3%DMSO) = 18±2 μM – 94±1% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = -5

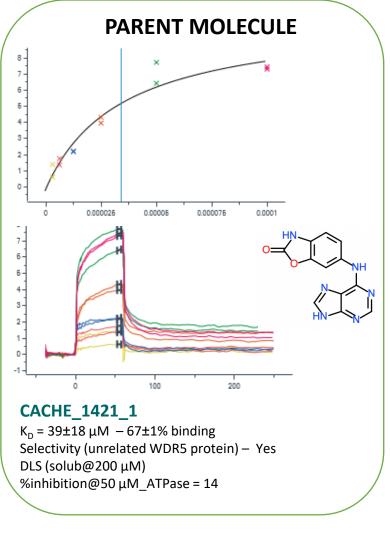


#### CACHE2-HO 1421 27

 $K_D$  (3% DMSO) = 19 μM (poor fit) – 101% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = -5



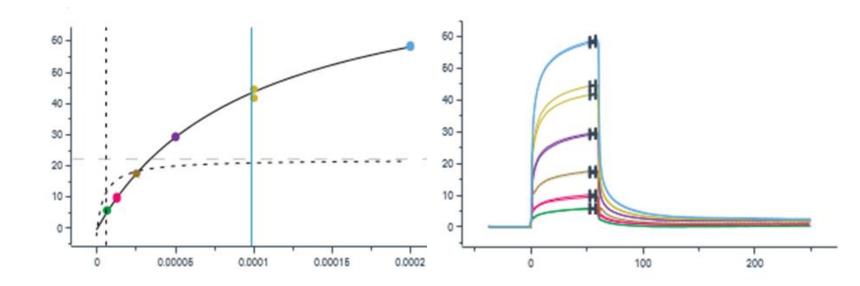




9 analogs of CACHE\_1421\_1 were submitted for round 2. 1 compound showed a superstoichiometric binding response with mild binding affinity to unrelated negative control protein.

## CACHE2-HO\_1421\_3

 $K_D = 104 \mu M - 395\%$  binding  $K_D = 200 \mu M - 392\%$  binding Selectivity (unrelated WDR5 protein) – mild binding DLS (solub@200 μM) %inhibition@50 μM ATPase = 40

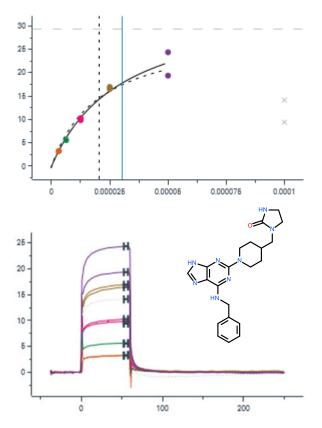


## **PARENT MOLECULE** 0.000026 0.00006 0.000076 0.0001 100 200 CACHE\_1421\_1 $K_D = 39\pm18 \ \mu M - 67\pm1\%$ binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 μM\_ATPase = 14

HN	HIN P	O NH	HN NH
CACHE_ID CACHE2-HO_1421_1 Parent CACHE_1421_1 distance_to_pai 0.1155	CACHE_ID CACHE2-HO_1421_2 Parent CACHE_1421_1 distance_to_pai 0.2174	CACHE_ID CACHE2-HO_1421_3 Parent CACHE_1421_1 distance_to_pai 0.2508	CACHE_ID CACHE2-HO_1421_4 Parent CACHE_1421_1 distance_to_pai 0.2593
F N N N N N N N N N N N N N N N N N N N	HIN O	O HN N N N N N N N N N N N N N N N N N N	HN CI
CACHE_ID CACHE2-HO_1421_5 Parent CACHE_1421_1 distance_to_pai 0.3831	CACHE_ID CACHE2-HO_1421_6 Parent CACHE_1421_1 distance_to_pai 0.3921	CACHE_ID CACHE2-HO_1421_7 Parent CACHE_1421_1 distance_to_pai 0.3968	CACHE_ID CACHE2-HO_1421_8 Parent CACHE_1421_1 distance_to_pai 0.4177
F HN N			
CACHE_ID CACHE2-HO_1421_9 Parent CACHE_1421_1 distance_to_pai 0.4244			

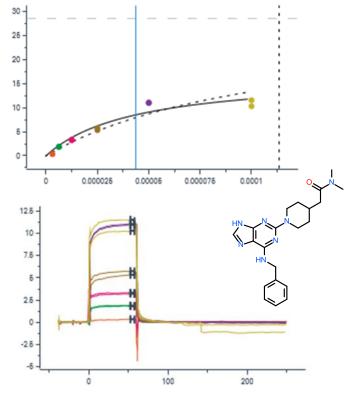
## **PARENT MOLECULE** 0.000026 0.00006 0.000076 2.5 -CACHE\_1421\_21 $K_D = 10 \pm 1 \mu M - 65 \pm 13\%$ binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@100 µM) %inhibition@50 µM\_ATPase = 12

14 analogs of CACHE\_1421\_21 were submitted for round 2. 2 compound showed a binding response.



### CACHE2-HO\_1421\_16

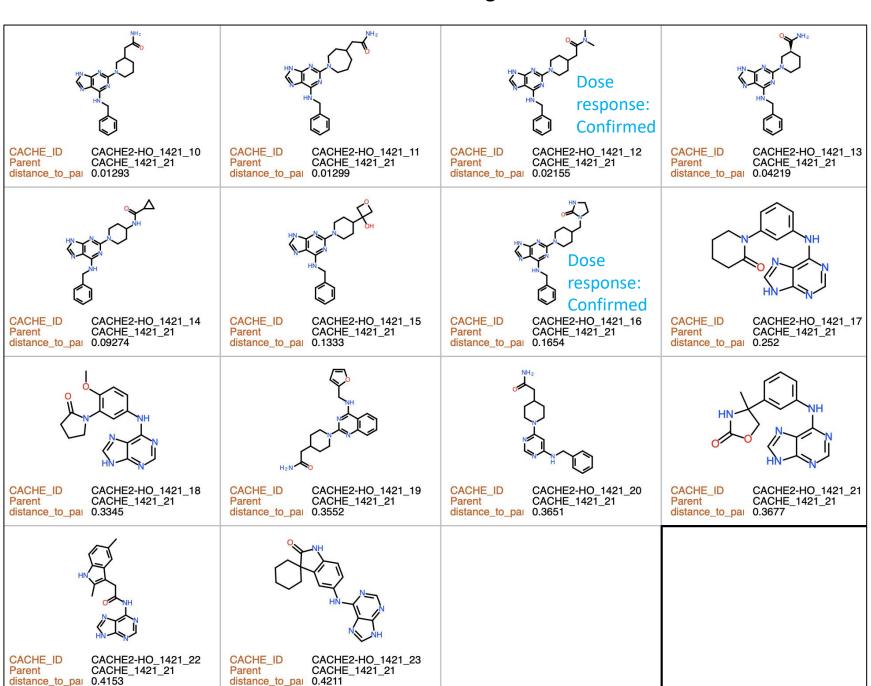
 $K_D$  = 30  $\mu$ M - 122% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M\_ATPase = 13



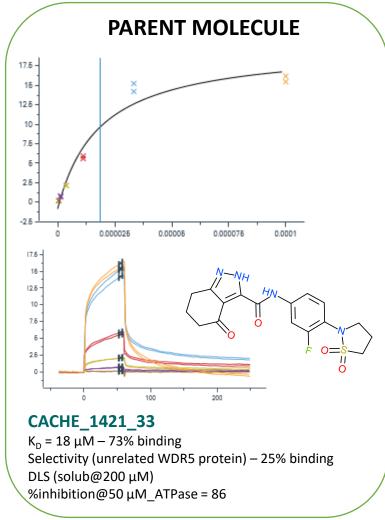
#### CACHE2-HO\_1421\_12

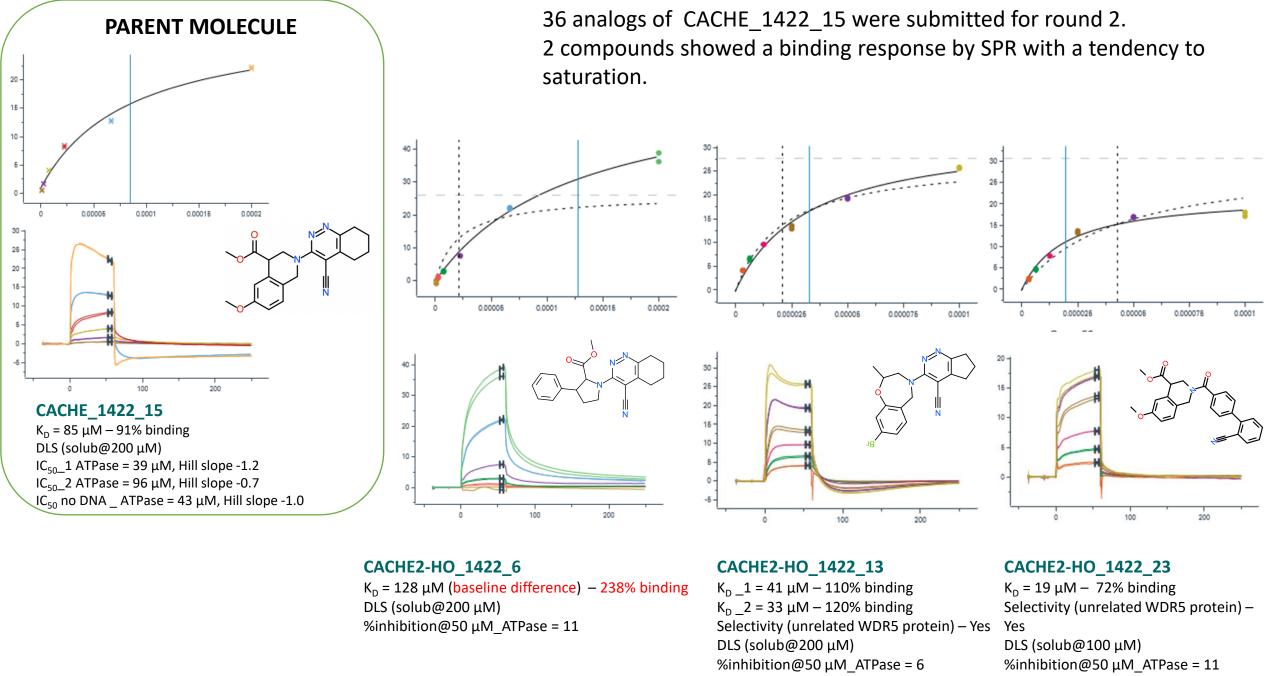
 $K_D$  (3% DMSO) = 43 μM - 60% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@100 μM) %inhibition@50 μM\_ATPase = 16

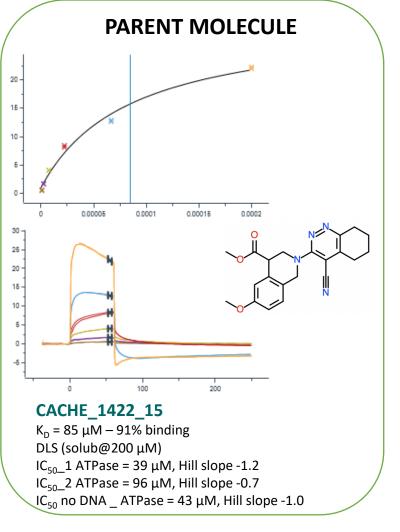
### PARENT MOLECULE 2.5 -0.000026 0.00006 0.000076 12.5 -2.5 -CACHE\_1421\_21 $K_D = 10 \pm 1 \mu M - 65 \pm 13\%$ binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@100 µM) %inhibition@50 µM ATPase = 12

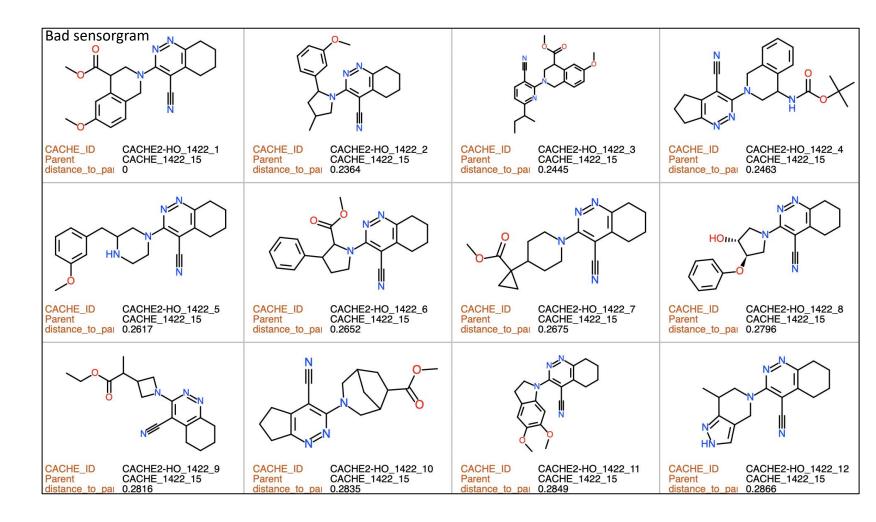


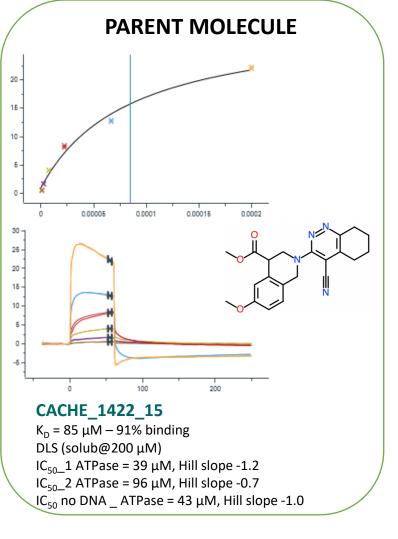
## ARENT MOLECULE Tested analogs

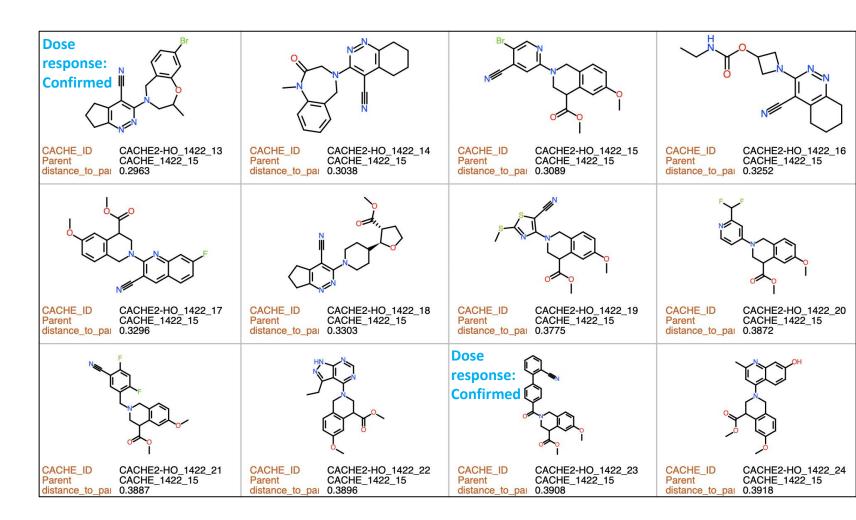


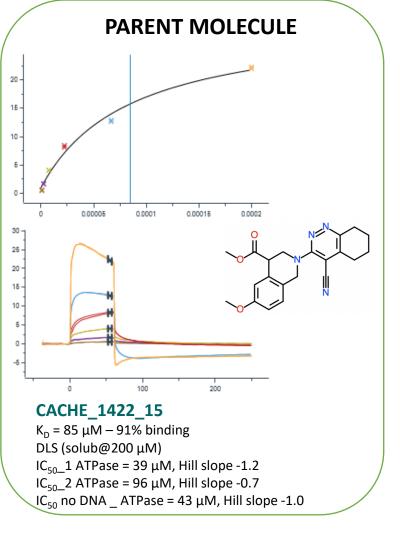








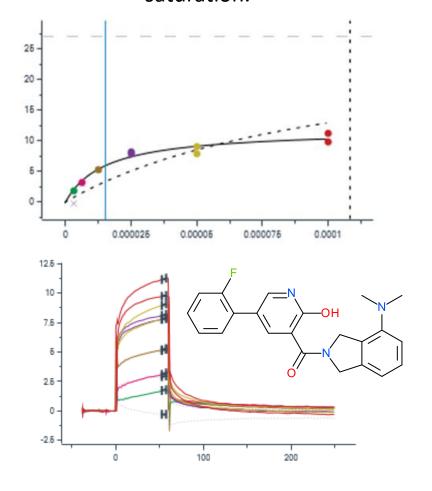




C   N			
CACHE_ID CACHE2-HO_1422_25 Parent CACHE_1422_15 distance_to_pai 0.3982	CACHE_ID CACHE2-HO_1422_26 Parent CACHE_1422_15 distance_to_pai 0.4074	CACHE_ID CACHE2-HO_1422_27 Parent CACHE_1422_15 distance_to_pai 0.4142	CACHE_ID CACHE2-HO_1422_28 Parent CACHE_1422_15 distance_to_pai 0.419
CACHE_ID CACHE2-HO_1422_29 Parent CACHE_1422_15 distance_to_pai 0.4294	CACHE_ID CACHE2-HO_1422_30 CACHE_1422_15 distance_to_pai	CACHE_ID CACHE2-HO_1422_31 Parent CACHE_1422_15 distance_to_pai 0.4368	CACHE_ID CACHE2-HO_1422_32 Parent CACHE_1422_15 distance_to_pai  CACHE2-HO_1422_32 CACHE_1422_15 0.4378
	S-N-N-Q-O	N N N N N N N N N N N N N N N N N N N	
CACHE_ID CACHE2-HO_1422_33 Parent CACHE_1422_15 distance_to_pai 0.446	CACHE_ID CACHE2-HO_1422_34 Parent CACHE_1422_15 distance_to_pai 0.4583	CACHE_ID CACHE2-HO_1422_35 Parent CACHE_1422_15 distance_to_pai 0.4697	CACHE_ID CACHE2-HO_1422_36 Parent CACHE_1422_15 distance_to_pai 0.478

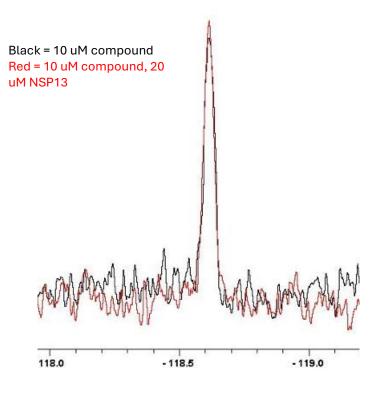
## **PARENT MOLECULE** 0.00005 0.0001 0.00016 200 CACHE\_1425\_80 $K_D = 19 \mu M$ (baseline difference/poor fit) -95% binding Selectivity (unrelated WDR5 protein) - Yes %inhibition@50 μM\_ATPase = no inhibition

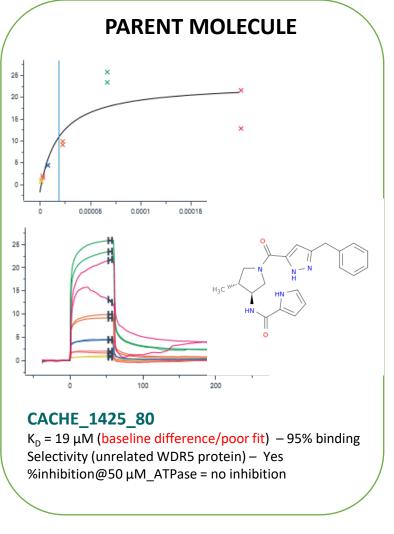
12 follow-ups of CACHE\_1425\_80 were submitted for round 2. 1 compound showed a mild binding response reaching saturation.

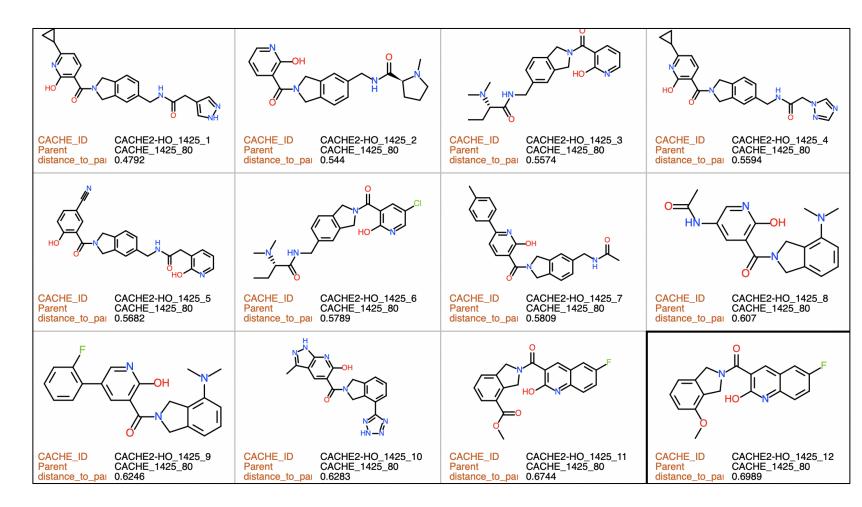


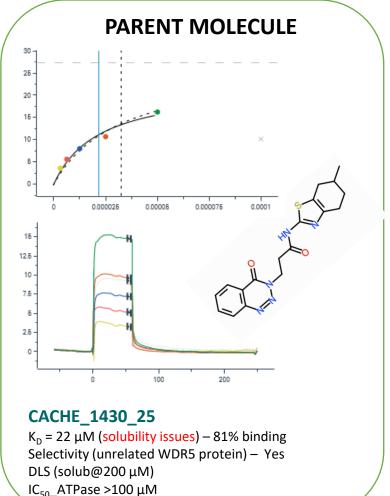
#### CACHE2-HO\_1425\_9

 $K_D$  = 15  $\mu$ M - 44% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 10 <sup>19</sup>F NMR Did not confirm binding

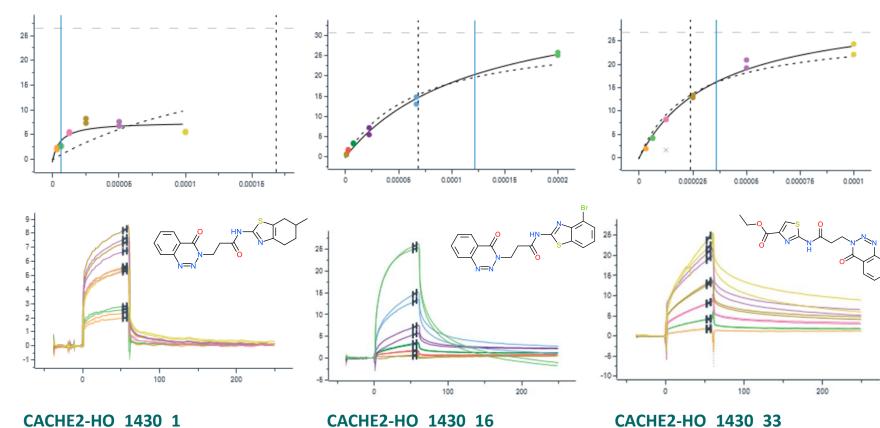








48 analogs of CACHE\_1430\_25 were submitted for round 2. 2 compounds showed a weak binding response by SPR.



 $K_D = 122 \,\mu\text{M} - 133\%$  binding

%inhibition@50 µM ATPase = 25

DLS (solub@200 µM)

 $K_D$  (3% DMSO) = 36  $\mu$ M (baseline

%inhibition@50 µM ATPase = 10

Selectivity (unrelated WDR5 protein) - Yes

difference) - 121% binding

DLS (solub@200 µM)

 $K_D$  (3%DMSO) = 6  $\mu$ M (poor fit) – 29%

%inhibition@50 µM ATPase = 11

Selectivity (unrelated WDR5 protein) – Yes

binding

DLS (solub@100 µM)

## **PARENT MOLECULE** 0.000026 0.00006 0.000076 12.5 -CACHE\_1430\_25 $K_D = 22 \mu M$ (solubility issues) – 81% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM)

IC<sub>50</sub>\_ATPase >100 μM

Dose response: Confirmed		HN NH2	HN N
CACHE_ID CACHE2-HO_1430_1 Parent CACHE_1430_25 distance_to_pai 0	CACHE_ID CACHE2-HO_1430_2 Parent CACHE_1430_25 distance_to_pai 0.01749	CACHE_ID CACHE2-HO_1430_3 Parent CACHE_1430_25 distance_to_pai 0.04274	CACHE_ID CACHE2-HO_1430_4 Parent CACHE_1430_25 distance_to_pai 0.04651
HN N	HN S	HIN N	HN S
CACHE_ID CACHE2-HO_1430_5 Parent CACHE_1430_25 distance_to_pai 0.07042	CACHE_ID CACHE2-HO_1430_6 Parent CACHE_1430_25 distance_to_pai 0.1444	CACHE_ID CACHE2-HO_1430_7 Parent CACHE_1430_25 distance_to_pai 0.1519	CACHE_ID CACHE2-HO_1430_8 Parent CACHE_1430_25 distance_to_pai  CACHE_1738
HN N	HN N	HN STA	HN ST Br
CACHE_ID CACHE2-HO_1430_9 Parent CACHE_1430_25 distance_to_pai 0.1771	CACHE_ID CACHE2-HO_1430_10 Parent CACHE_1430_25 distance_to_pai 0.1782	CACHE_ID CACHE2-HO_1430_11 Parent CACHE_1430_25 distance_to_pai 0.1933	CACHE_ID CACHE2-HO_1430_12 Parent CACHE_1430_25 distance_to_pai 0.2039
HN N CI	HN WHO	HN STORY	Dose response: Confirmed
CACHE_ID CACHE2-HO_1430_13 Parent CACHE_1430_25 distance_to_pai 0.2082	CACHE_ID CACHE2-HO_1430_14 Parent CACHE_1430_25 distance_to_pai 0.2087	CACHE_ID CACHE2-HO_1430_15 Parent CACHE_1430_25 distance_to_pai  CACHE2-HO_1430_15 CACHE3-HO_1430_15 C	CACHE_ID CACHE2-HO_1430_16 Parent CACHE_1430_25 distance_to_pai  CACHE_1430_25 0.2105

## **PARENT MOLECULE** 0.000026 0.00006 0.000076 12.5 -CACHE\_1430\_25 $K_D = 22 \mu M$ (solubility issues) – 81% binding Selectivity (unrelated WDR5 protein) – Yes

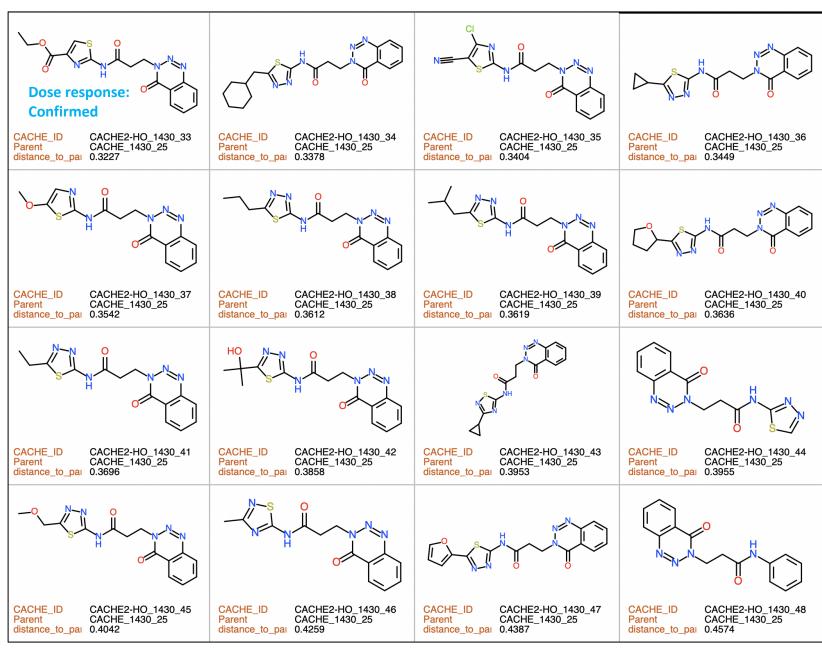
DLS (solub@200 μM)

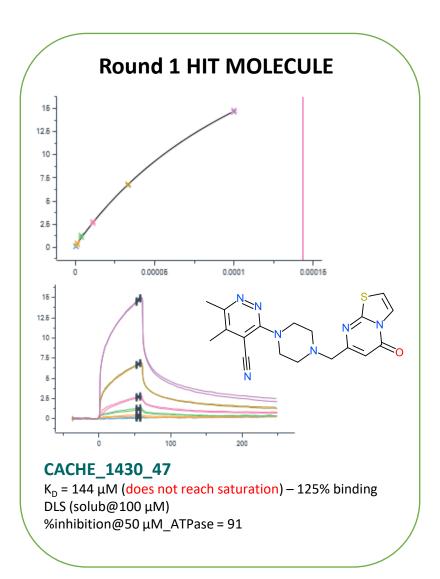
IC<sub>50</sub>\_ATPase >100 μM

CACHE_ID CAGHE2HO_1430_17 Parent_CACHE_ID CAGHE2HO_1430_18 Parent_CACHE_ID CAGHE2HO_1430_29 Parent_CACHE_ID CACHE2HO_1430_29 Parent_CACHE_ID_CACHE2HO_1430_29				
Parent   CACHE_1430_25   Parent   CACHE_1430	HN N	HN N	HN N	HN N
Parent	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25
Parent	N N N N N N N N N N N N N N N N N N N	N N N N N N N N N N N N N N N N N N N		S O N N N
Parent distance_to_pai 0.2693  Parent distance_to_pai 0.2727  Parent distance_to_pai 0.2727  Parent distance_to_pai 0.2727  Parent distance_to_pai 0.279  Parent distance_to_pai 0.279  Parent distance_to_pai 0.279  Parent distance_to_pai 0.279  Parent distance_to_pai 0.2923  Parent distance_to_pai 0.2923  Parent distance_to_pai 0.279  CACHE_1430_25  CACHE_1430_25  CACHE_1430_25  CACHE_1D CACHE2-HO_1430_30  Parent CACHE_1D CACHE2-HO_1430_31  CACHE_1D CACHE2-HO_1430_32  Parent CACHE_1430_25  Parent CACHE_1430_25  Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25
Parent distance_to_pai 0.2693  Parent distance_to_pai 0.2727  Parent distance_to_pai 0.2727  Parent distance_to_pai 0.2727  Parent distance_to_pai 0.279  Parent distance_to_pai 0.279  Parent distance_to_pai 0.279  Parent distance_to_pai 0.279  Parent distance_to_pai 0.2923  Parent distance_to_pai 0.2923  Parent distance_to_pai 0.279  CACHE_1430_25  CACHE_1430_25  CACHE_1430_25  CACHE_1D CACHE2-HO_1430_30  Parent CACHE_1D CACHE2-HO_1430_31  CACHE_1D CACHE2-HO_1430_32  Parent CACHE_1430_25  Parent CACHE_1430_25  Parent CACHE_1430_25	N O N N N N N N N N N N N N N N N N N N	Br WH	F S N N N N	N N N N N N N N N N N N N N N N N N N
CACHE_ID CACHE2-HO_1430_29 CACHE_ID CACHE2-HO_1430_30 CACHE_1H30_25 CACH	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25
Parent CACHE_1430_25 Parent CACHE_1430_25 Parent CACHE_1430_25 Parent CACHE_1430_25		N N N N N N N N N N N N N N N N N N N		H <sub>2</sub> N S O N N N
	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25	Parent CACHE_1430_25

## PARENT MOLECULE 0.000025 0.00006 0.000076 12.5 -CACHE\_1430\_25 $K_D = 22 \mu M$ (solubility issues) – 81% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM)

 $IC_{50}$ ATPase >100  $\mu$ M

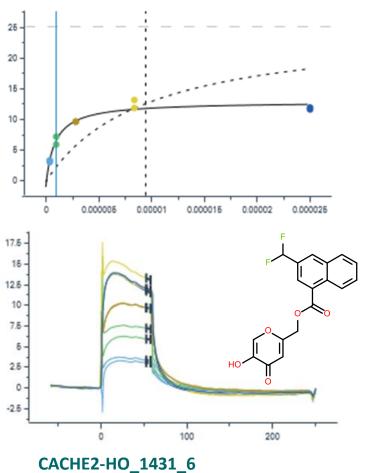




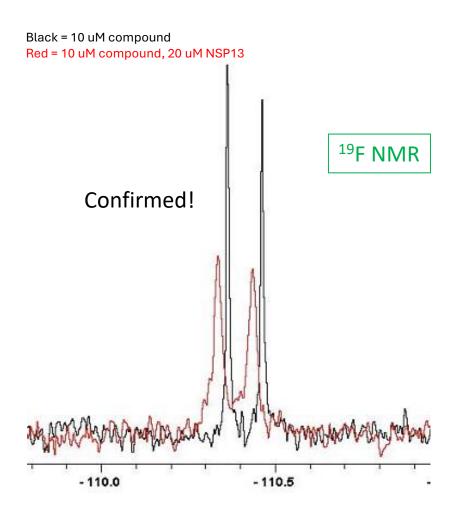
No analog in Round 2

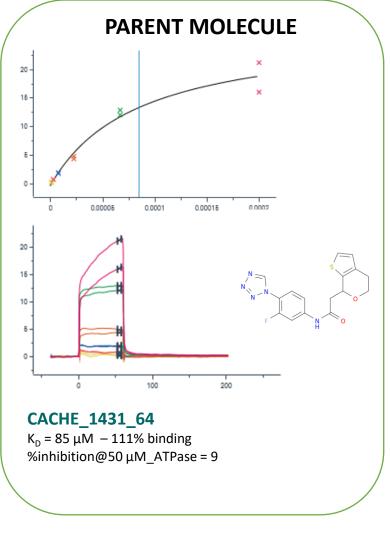
## **PARENT MOLECULE** 0.00005 0.00016 200 CACHE\_1431\_64 $K_D = 85 \mu M - 111\%$ binding %inhibition@50 µM\_ATPase = 9

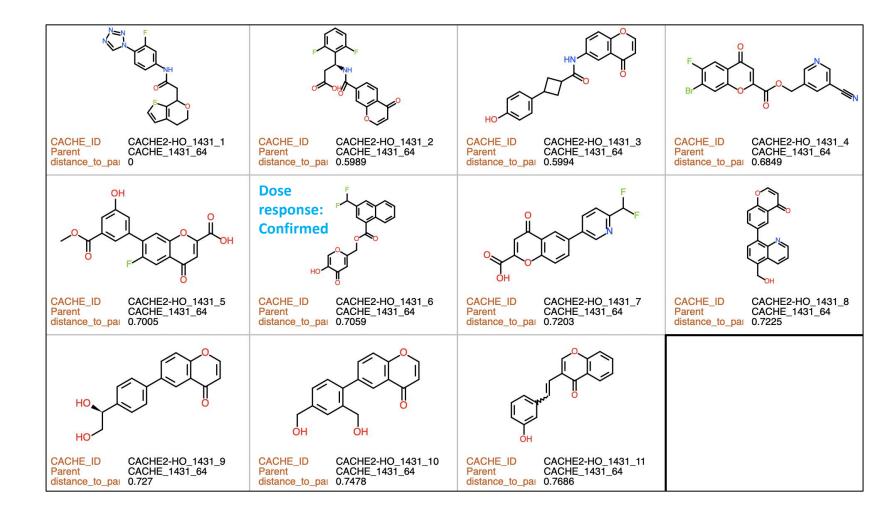
11 non-analog follow-ups of CACHE\_1431\_64 were submitted for round 2. One compound showed a high binding affinity by SPR that was confirmed by <sup>19</sup>F NMR.



 $K_D _1 = 0.65 \mu M - 72\%$  binding  $K_D _2 (3\% DMSO) = 0.9 \mu M - 51\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@100 μM) %inhibition@50 μM ATPase = 11

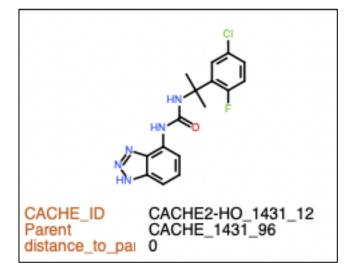


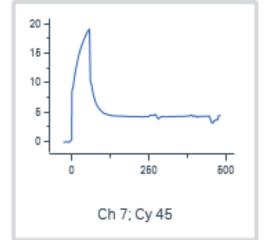


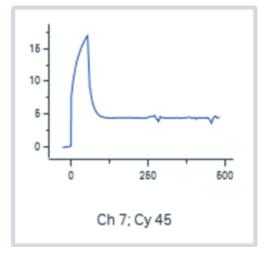


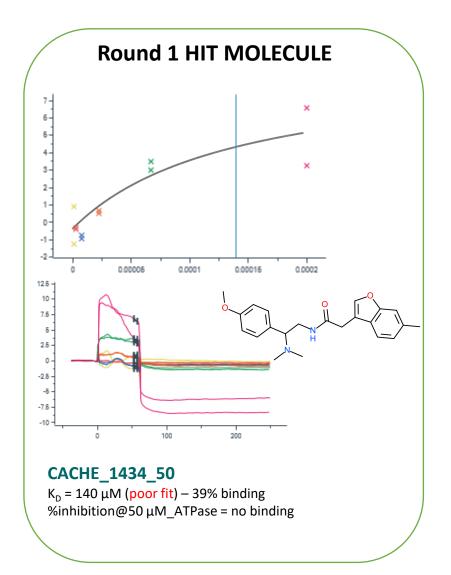
#### PARENT MOLECULE 26 20 16 -10 -0.00006 0.0001 0.00016 0.0002 20 16 CACHE\_1431\_96 $K_D = 46 \mu M$ (slow on/off)– 124% binding Selectivity (unrelated WDR5 protein) - mild binding to WDR\$ %inhibition@50 µM\_ATPase = 18

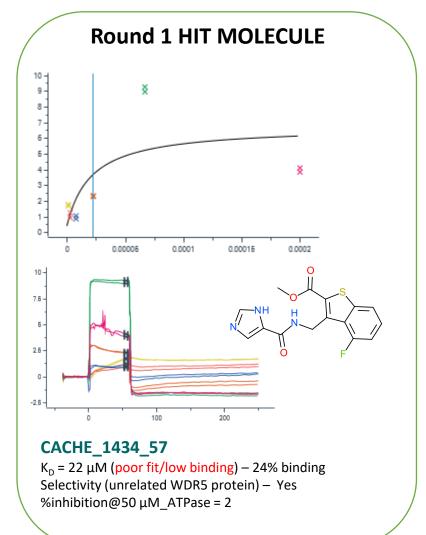
## Resupply – showed baseline difference and was not included for a dose - response



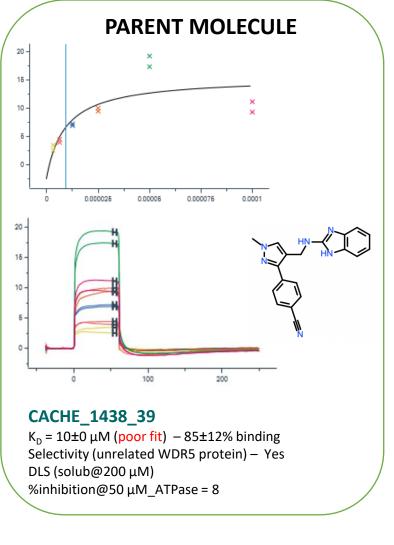






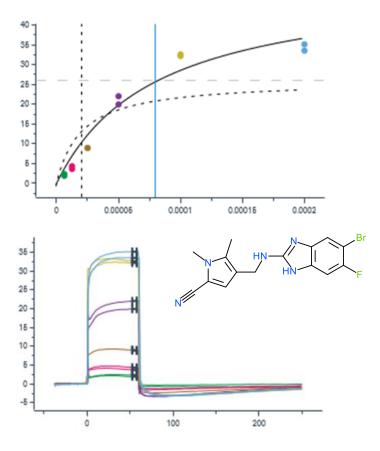


No analogs in Round 2



45 analogs of CACHE\_1438\_39 were submitted for round 2. Three compounds (including re-supplied parent molecule) showed a binding response by SPR with a tendency to reach saturation. 1 compound confirmed binding affinity by <sup>19</sup>F NMR.

Black = 10 uM compound



Red = 10 uM compound, 20 uM NSP13

19 F NMR

Confirmed!

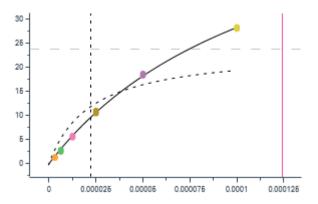
-117

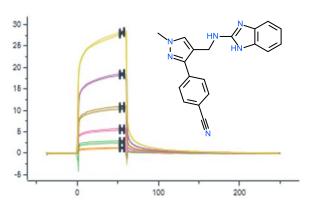
-118

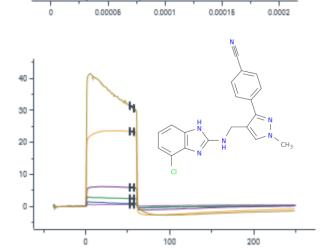
#### CACHE2-HO\_1438\_19

 $K_D _1 = 90 \mu M - 166\%$  binding  $K_D _2 = 80 \mu M - 198\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 11

#### **PARENT MOLECULE** 20 -0.000026 0.00006 0.000076 0.0001 20 -15 -10 -CACHE\_1438\_39 $K_D = 10\pm0 \,\mu\text{M} \,(\text{poor fit}) - 85\pm12\% \,\text{binding}$ Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 $\mu$ M\_ATPase = 8







20 -

#### CACHE2-HO\_1438\_1

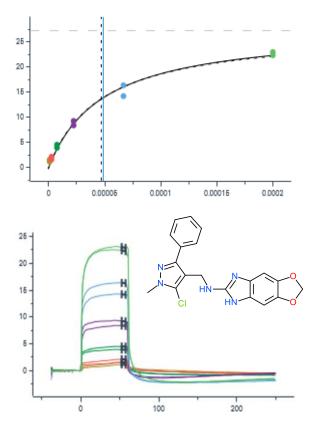
 $K_D$  (3% DMSO) = 124 μM (does not reach saturation) – 268% binding Selectivity (unrelated WDR5 protein) – 31% binding (linear response) DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 11

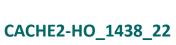
#### CACHE2-HO\_1438\_3

 $K_D$  = 78 μM - 178% binding Suspicious sensorgram Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = -3

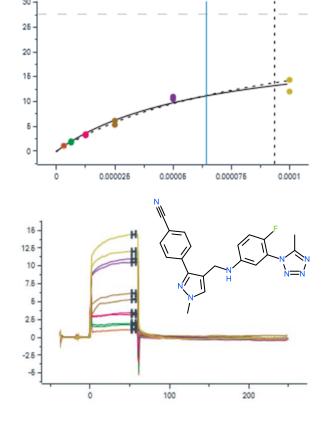
## **PARENT MOLECULE** 20 -0.000026 0.00006 0.000076 0.0001 20 -CACHE\_1438\_39 $K_D = 10\pm0 \,\mu\text{M} \,(\text{poor fit}) - 85\pm12\% \,\text{binding}$ Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM)

%inhibition@50  $\mu$ M\_ATPase = 8





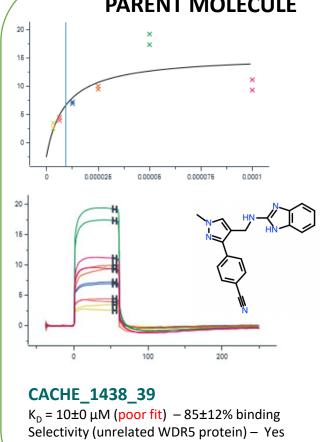
 $K_D$  = 49 μM - 102% binding Selectivity (unrelated WDR5 protein) - weak binding (linear response) DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 10



#### CACHE2-HO\_1438\_41

 $K_D$  = 64 μM – 81% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 8

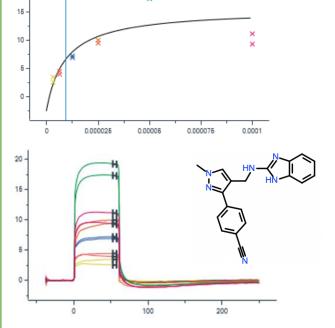
#### Tested analogs **PARENT MOLECULE**



DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 8

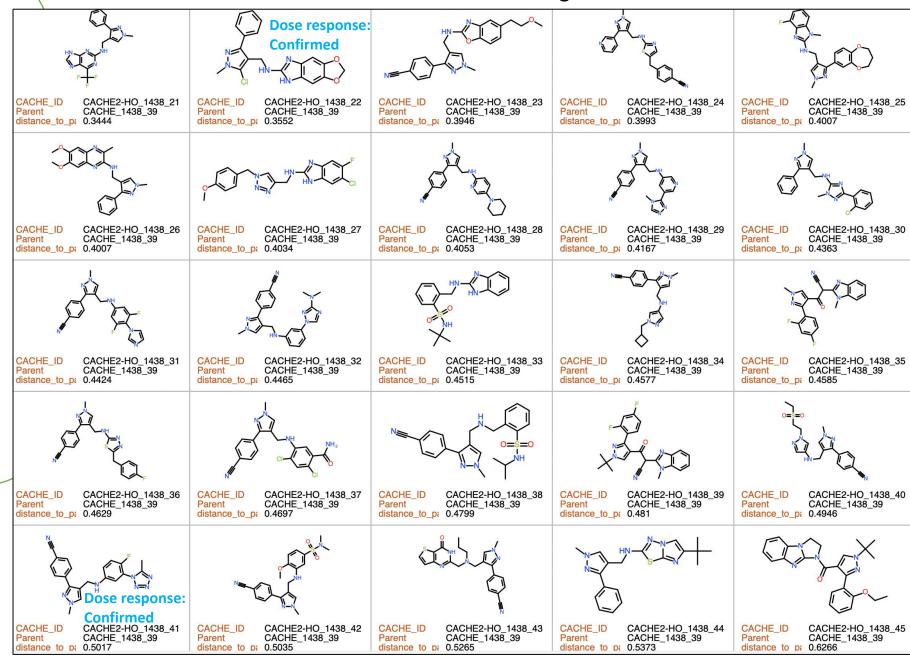
	HN-HN-HN-HN-HN-HN-HN-HN-HN-HN-HN-HN-HN-H	HN HN HN TO	HN HN HN	N HN HN	
CACHE_ Parent distance	D CACHE2-HO_1438_1 CACHE_1438_39 to_p; 0	CACHE_ID CACHE2-HO_1438_2 Parent CACHE_1438_39 distance_to_pa 0.0793	CACHE_ID CACHE2-HO_1438_3 Parent CACHE_1438_39 distance_to_pa 0.0793	CACHE_ID CACHE2-HO_1438_4 Parent CACHE_1438_39 distance_to_pa 0.09589	CACHE_ID CACHE2-HO_1438_5 Parent CACHE_1438_39 distance_to_pa 0.1292
	N HN N	N HN HN	HN HN F	HN HN F	HN HN CI
CACHE_ Parent distance	ID CACHE2-HO_1438_6 CACHE_1438_39 _to_p; 0.1429	CACHE_ID CACHE2-HO_1438_7 Parent CACHE_1438_39 distance_to_p: 0.1502	CACHE_ID CACHE2-HO_1438_8 Parent CACHE_1438_39 distance_to_p; 0.1646	CACHE_ID CACHE2-HO_1438_9 Parent CACHE_1438_39 distance_to_pa 0.1674	CACHE_ID CACHE2-HO_1438_10 Parent CACHE_1438_39 distance_to_pa 0.214
7-2	HN HN NH <sub>2</sub>	N HN HN F	N HN HN HN	N HN C	N HN N
CACHE_ Parent distance	D CACHE2-HO_1438_11 CACHE_1438_39 to_p; 0.2205	CACHE_ID CACHE2-HO_1438_12 Parent CACHE_1438_39 distance_to_pa 0.2205	CACHE_ID CACHE2-HO_1438_13 Parent CACHE_1438_39 distance_to_pa 0.2302	CACHE_ID CACHE2-HO_1438_14 Parent CACHE_1438_39 distance_to_pa 0.251	CACHE_ID CACHE2-HO_1438_15 Parent CACHE_1438_39 distance_to_pa 0.2734
			N HN HN TC	Dose response:	HN HN HN
CACHE_ Parent distance	D CACHE2-HO_1438_16 CACHE_1438_39 _to_p; 0.2915	CACHE_ID CACHE2-HO_1438_17 Parent CACHE_1438_39 distance_to_pa 0.3101	CACHE_ID CACHE2-HO_1438_18 Parent CACHE_1438_39 distance_to_p; 0.3288	CACHE_ID CACHE2-HO_1438_19 Parent CACHE_1438_39 distance_to_p; 0.3385	CACHE_ID CACHE2-HO_1438_20 Parent CACHE_1438_39 distance_to_pa 0.3404

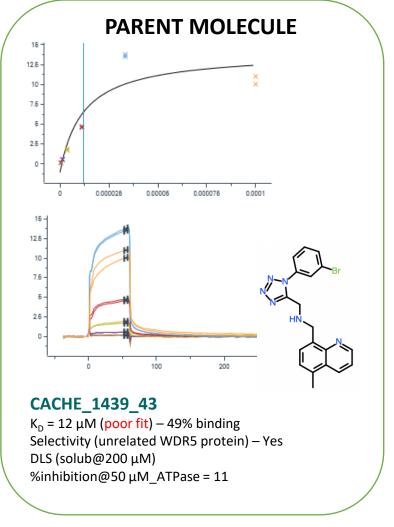
#### **PARENT MOLECULE**



#### CACHE\_1438\_39

 $K_D = 10\pm0~\mu\text{M}$  (poor fit)  $-85\pm12\%$  binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200  $\mu\text{M}$ ) %inhibition@50  $\mu\text{M}$  ATPase = 8





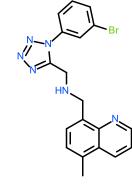
25 analogues of CACHE\_1439\_43, including re-supplied parent molecule, were submitted for round 2.

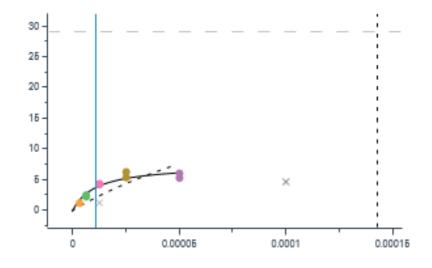
Re-supplied hit showed a very weak dose dependent binding response.

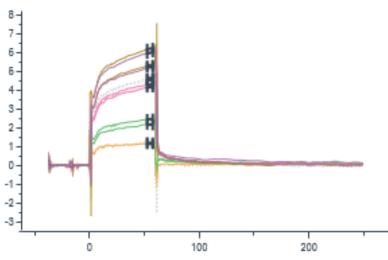
#### CACHE2-HO\_1439\_21

 $K_D$  (3% DMSO) = 11  $\mu$ M (low binding) – 25% binding

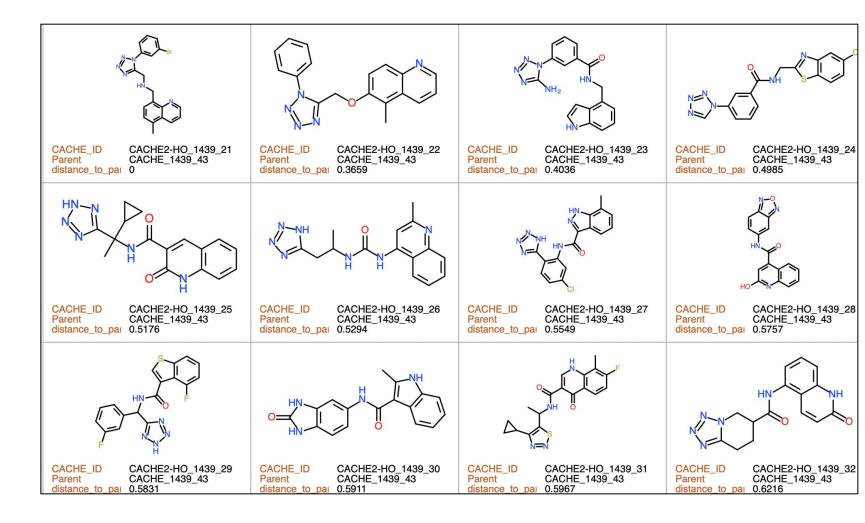
Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 32







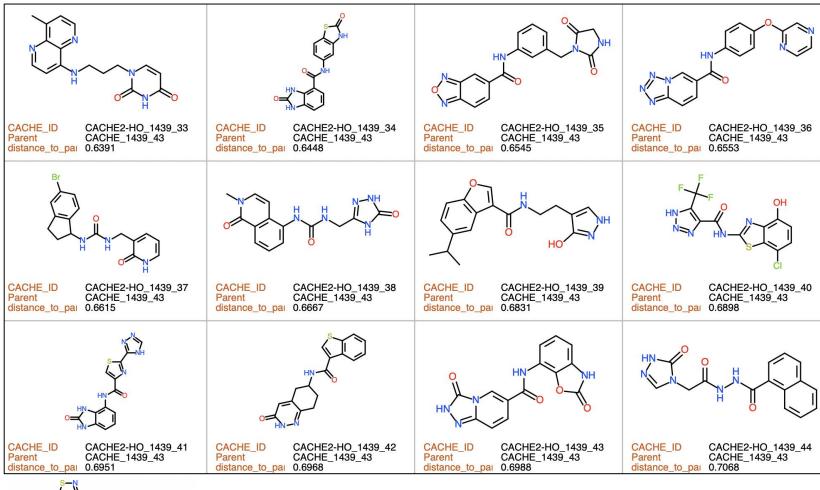
#### PARENT MOLECULE 12.5 2.5 -0.000026 0.00006 0.000076 0.0001 12.5 -10 2.5 CACHE\_1439\_43 $K_D = 12 \mu M \text{ (poor fit)} - 49\% \text{ binding}$ Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 µM\_ATPase = 11

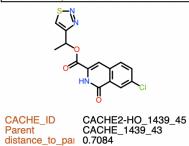


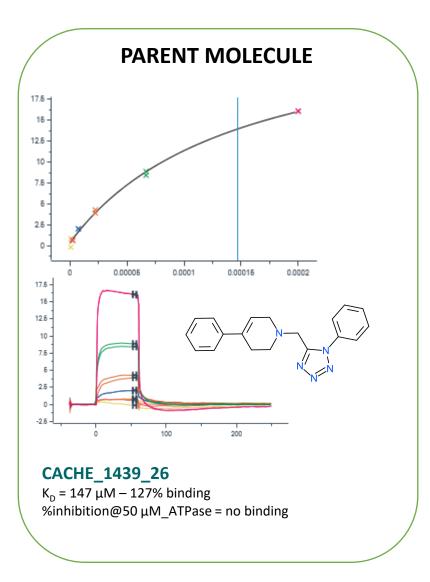
### PARENT MOLECULE 12.5 2.5 -0.000026 0.00006 0.000076 0.0001 12.5 2.5 CACHE\_1439\_43 $K_D = 12 \mu M \text{ (poor fit)} - 49\% \text{ binding}$ Selectivity (unrelated WDR5 protein) – Yes

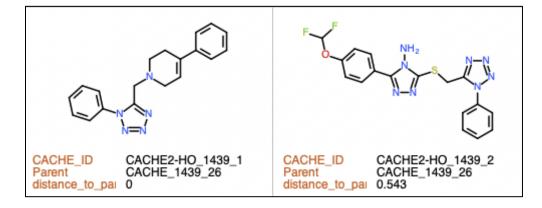
DLS (solub@200 µM)

%inhibition@50 µM\_ATPase = 11

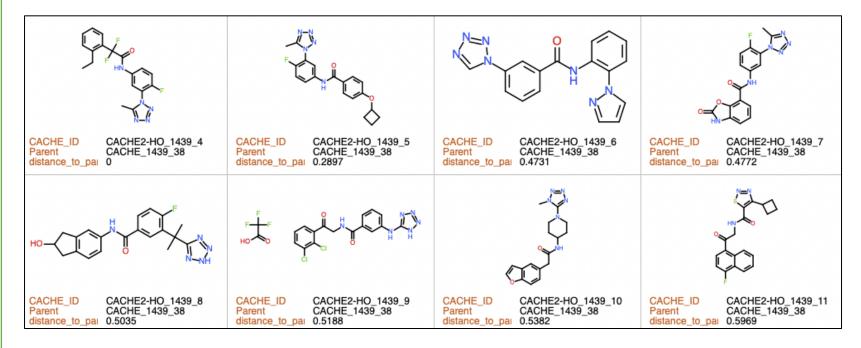


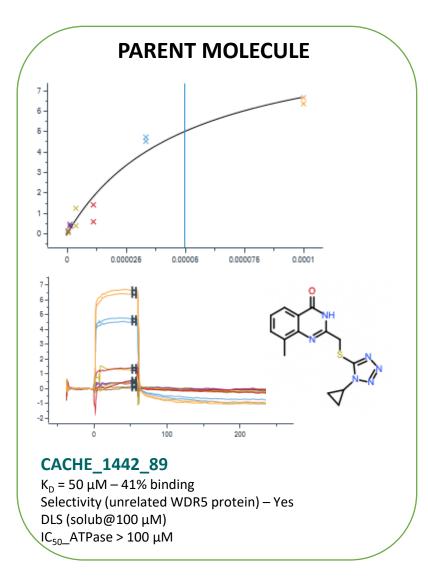






## **PARENT MOLECULE** 0.00006 0.0001 0.00016 0.0002 200 CACHE\_1439\_38 $K_D = 114 \, \mu M - 124\% \, binding$ DLS (solub@200 µM) %inhibition@50 µM\_ATPase = 11



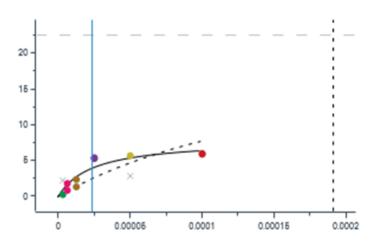


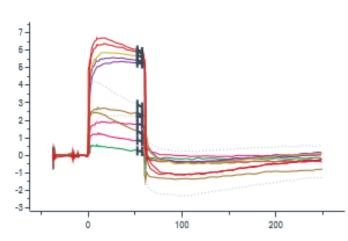
36 analogues of CACHE\_1442\_89, including re-supplied parent molecule, were submitted for round 2.

Re-supplied hit showed a weak dose dependent binding response.

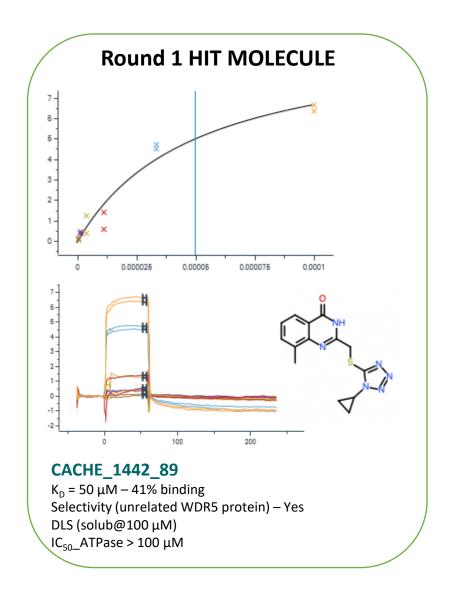
#### CACHE2-HO\_1442\_1

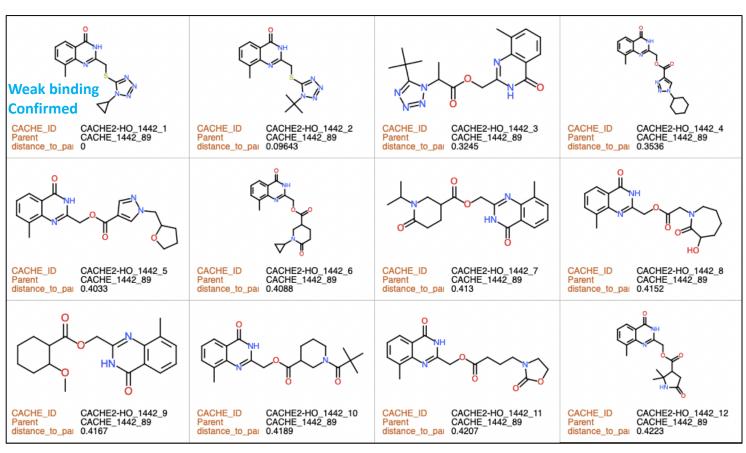
K<sub>D</sub> = 24 μM (poor fit) – 35% binding Selectivity (unrelated WDR5 protein) – Yes %inhibition@50 μM\_ATPase = 18



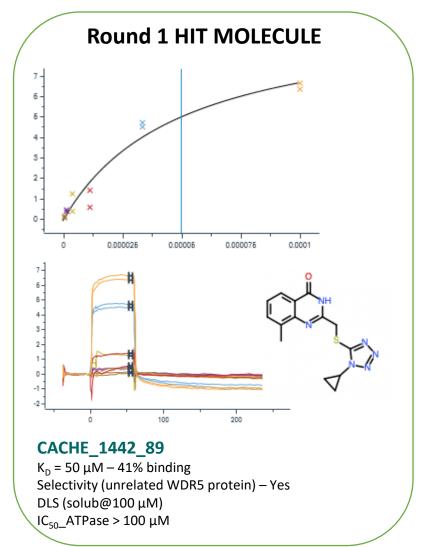


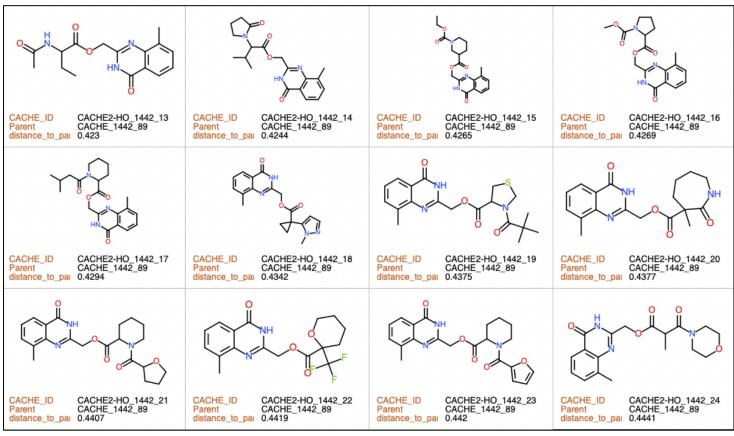
#### Tested follow-ups



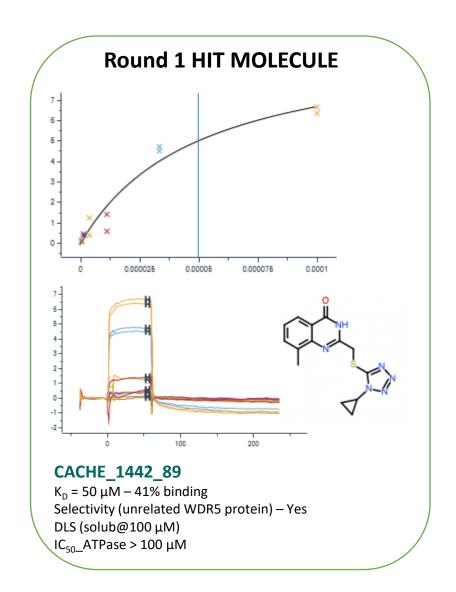


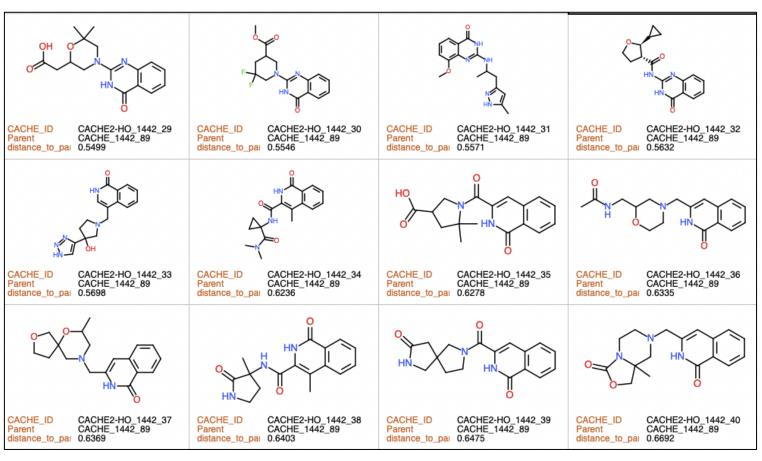
#### Tested follow-ups



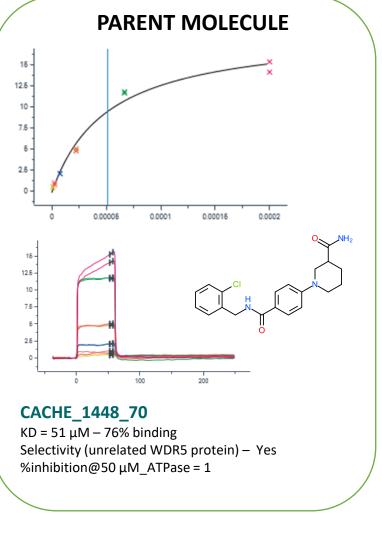


#### Tested follow-ups

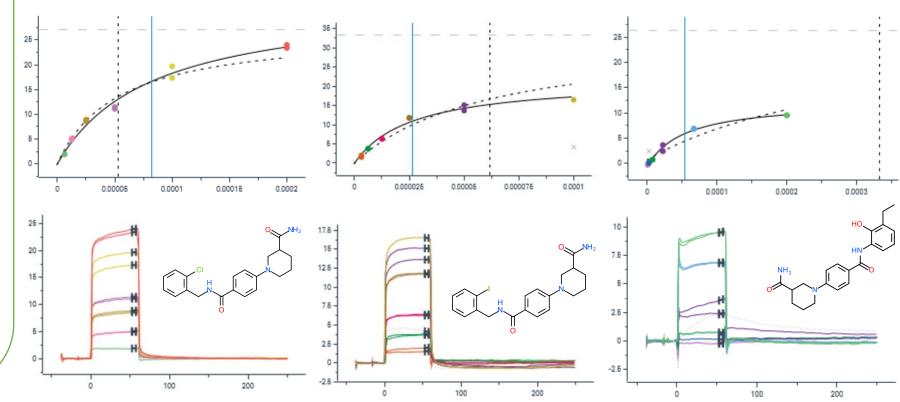




# CACHE#2 – NSP13 SARS2 Participant 1448



44 analogs of CACHE\_1448\_70 were submitted for round 2. 3 compounds (including re-supplied parent molecule) showed a dose dependent binding response by SPR.



#### **CACHE2-HO 1448 1**

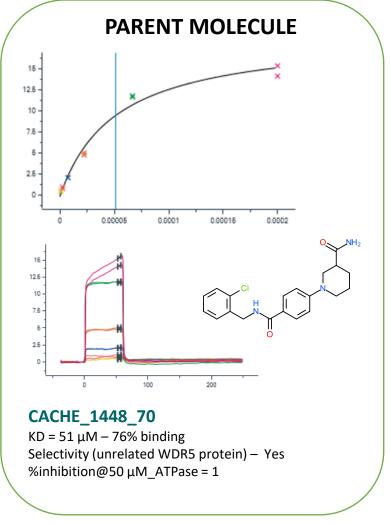
 $K_D$  (3% DMSO) = 82  $\mu$ M – 123% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M\_ATPase = 9

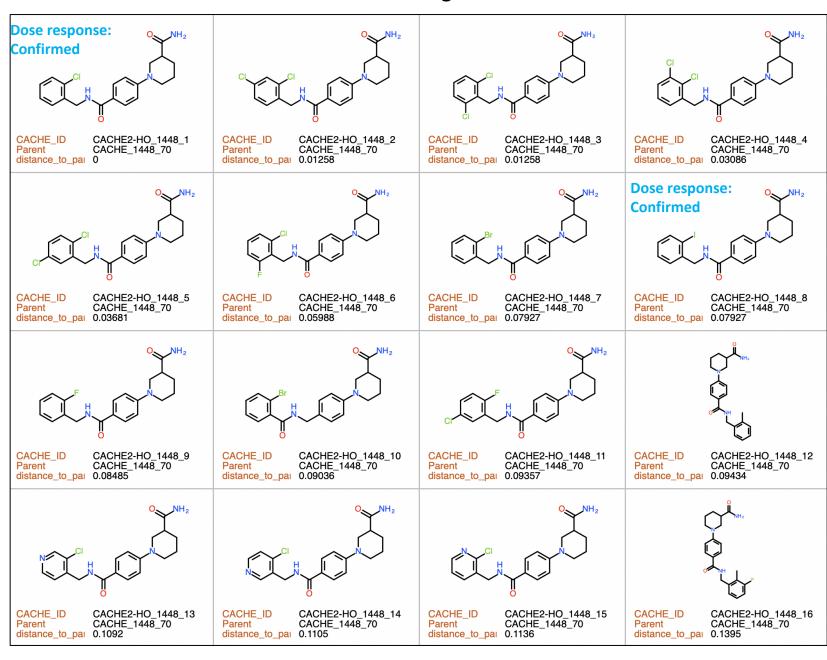
#### **CACHE2-HO 1448 8**

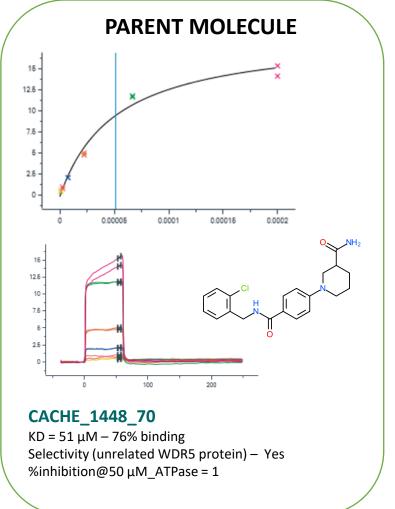
 $K_D = 26 \mu M - 66\%$  binding DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 6

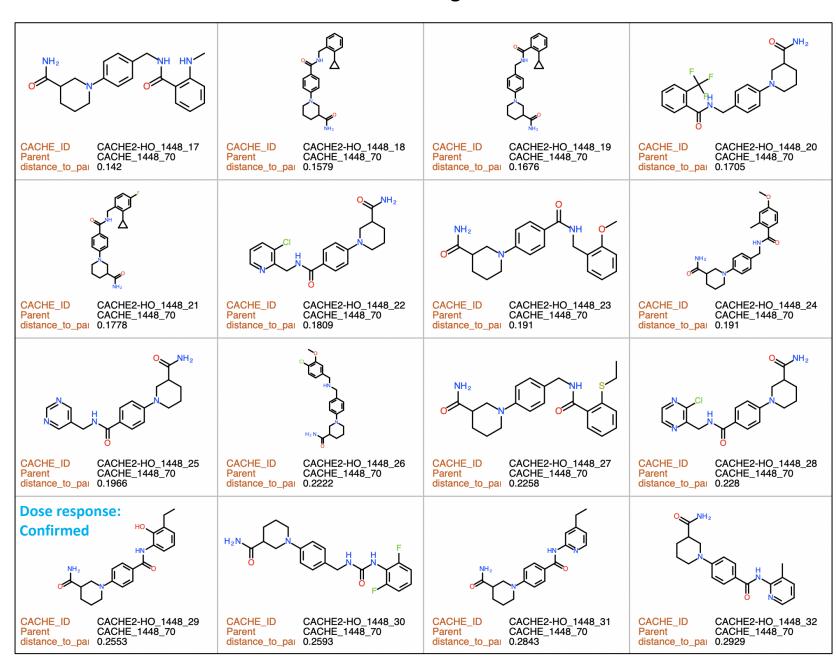
#### CACHE2-HO 1448 29

 $K_D$  = 54 μM - 49% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 6

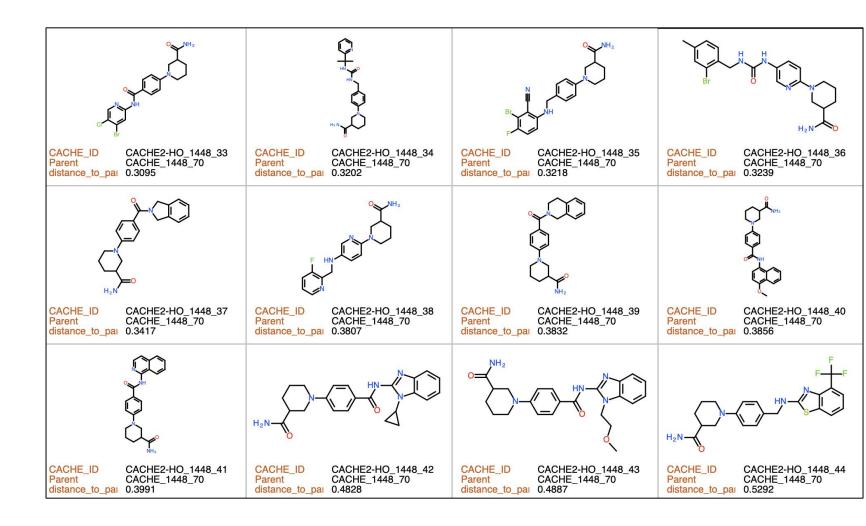






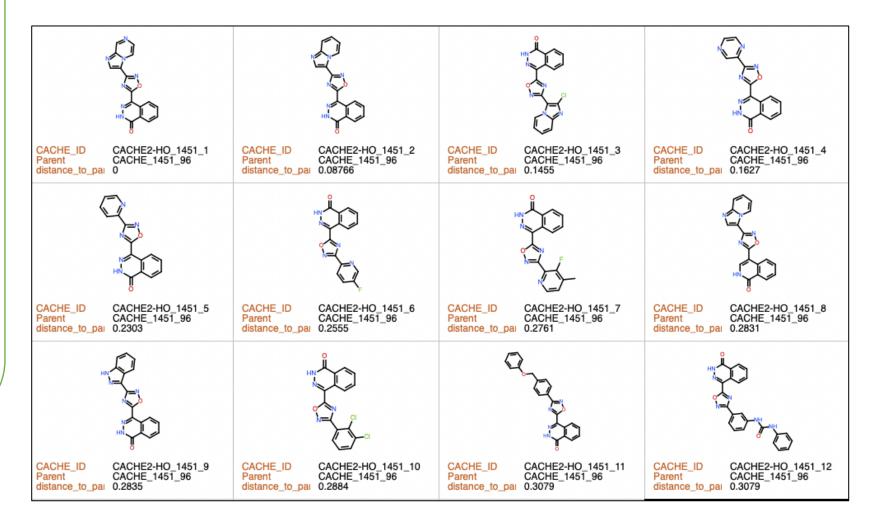


### PARENT MOLECULE 12.5 -2.5 -0.0002 0.00005 0.0001 0.00015 12.5 10 -CACHE\_1448\_70 $KD = 51 \mu M - 76\%$ binding Selectivity (unrelated WDR5 protein) - Yes %inhibition@50 µM\_ATPase = 1

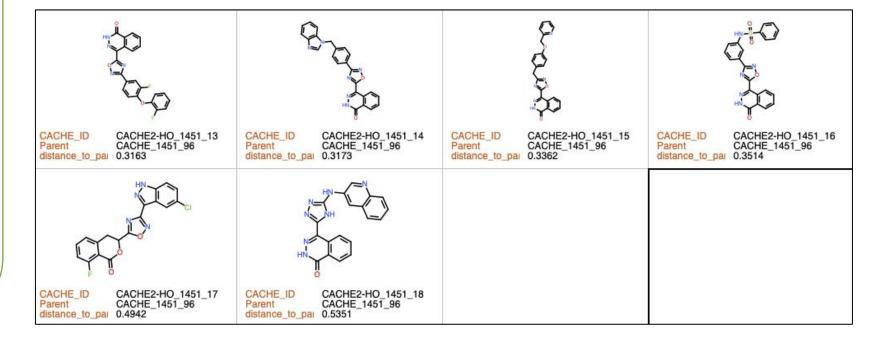


# CACHE#2 – NSP13 SARS2 Participant 1451

## PARENT MOLECULE 0.0001 0.0002 0.00005 0.00016 CACHE\_1451\_96 $K_D = 36 \mu M - 42\%$ binding DLS (solub@200 µM) $IC_{50}$ ATPase = 24 $\mu$ M



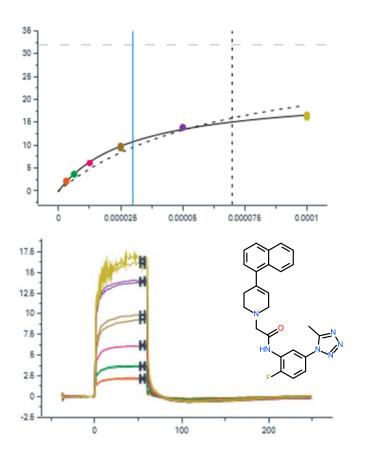
## **PARENT MOLECULE** 0.00005 0.0001 0.00016 0.0002 CACHE\_1451\_96 $K_D = 36 \mu M - 42\%$ binding DLS (solub@200 µM) $IC_{50}$ ATPase = 24 $\mu$ M

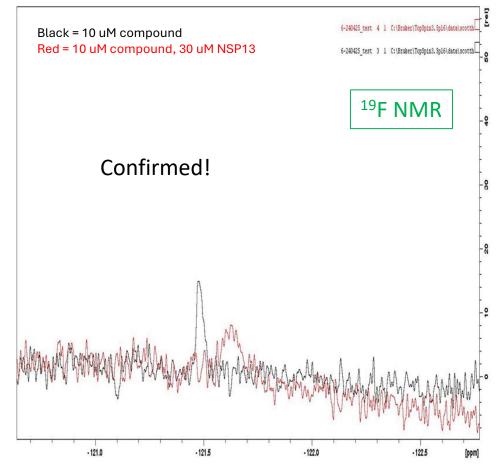


# CACHE#2 – NSP13 SARS2 Participant 1454

# **PARENT MOLECULE** 0.000026 0.00006 **CACHE 1454 20** $K_D = 64 \mu M$ (does not reach saturation) – 85% binding Selectivity (unrelated WDR5 protein) – Yes $IC_{50}$ ATPase >160 $\mu$ M

8 analogs of CACHE\_1454\_20 were submitted for round 2. The re-supplied parent molecule confirmed a selective binding response by SPR and by <sup>19</sup>F NMR.



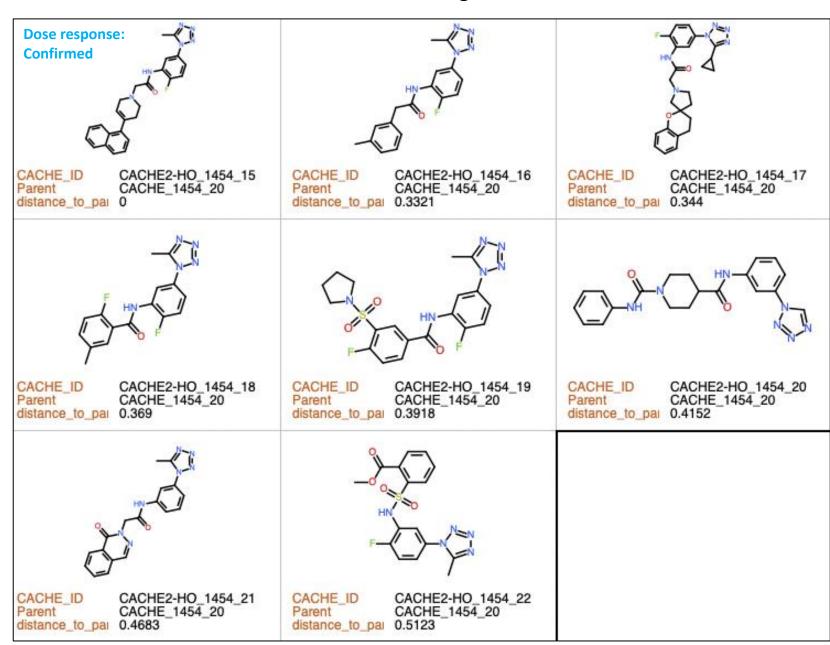


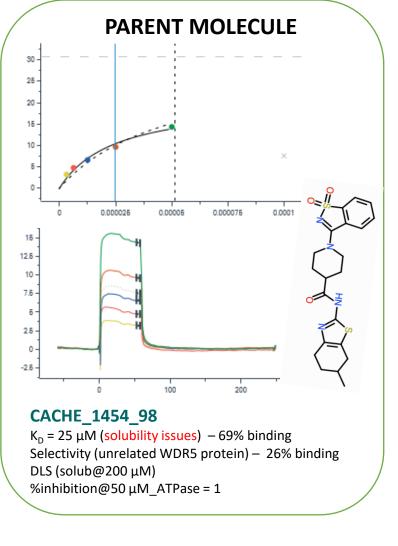
#### CACHE2-HO\_1454\_15

 $K_D_1 = 31 \mu M - 53\%$  binding  $K_D_2 = 30 \mu M - 68 \%$ binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M ATPase = 21

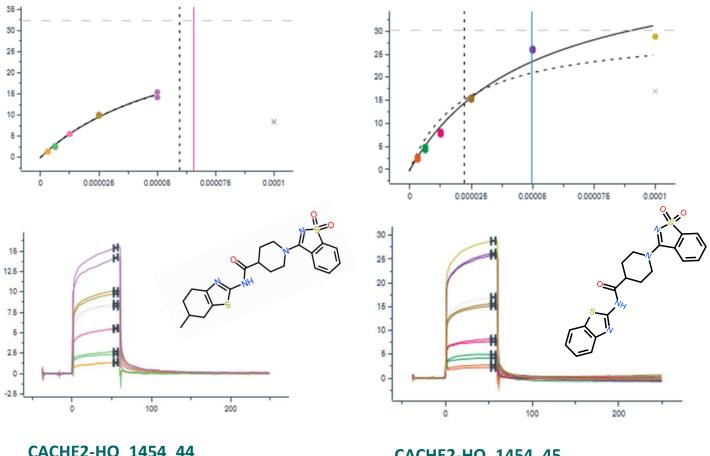
# PARENT MOLECULE 0.000026 0.00005 12.5 -**CACHE 1454 20** $K_D = 64 \mu M$ (does not reach saturation) – 85% binding

 $K_D$  = 64  $\mu$ M (does not reach saturation) – 85% bindi Selectivity (unrelated WDR5 protein) – Yes  $IC_{50}$ \_ATPase >160  $\mu$ M





7 analogs of CACHE\_1454\_98 were submitted for round 2. 2 compound (including re-supplied parent molecule) showed a dose dependent binding response by SPR with a tendency to reach saturation.

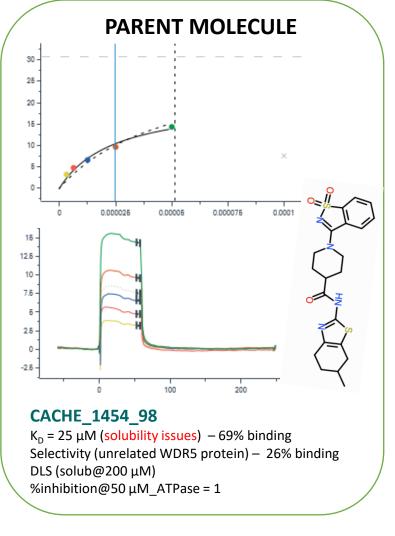


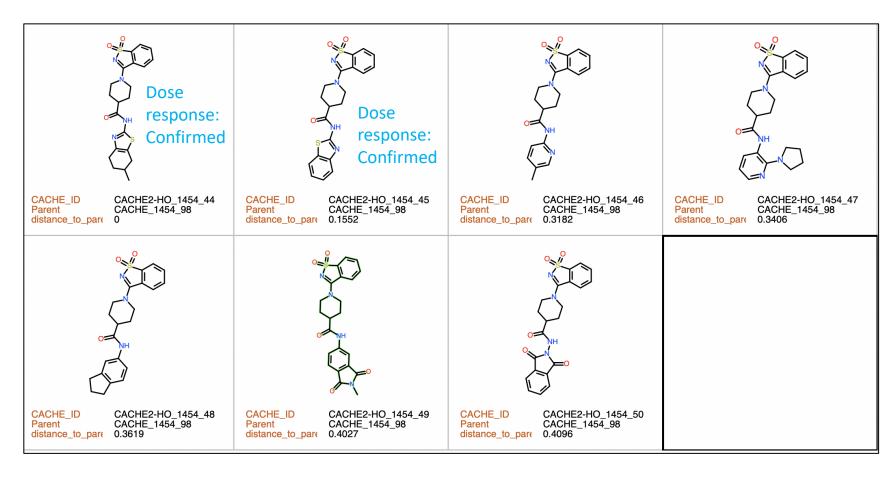
#### CACHE2-HO\_1454\_44

 $K_D = 65 \mu M$  (solubility issues) -107%binding %inhibition@50 µM ATPase = 8

#### CACHE2-HO\_1454\_45

 $K_D$  1 = 44  $\mu$ M – 135% binding  $K_D = 2 = 50 \,\mu\text{M} - 155\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 µM ATPase = 11



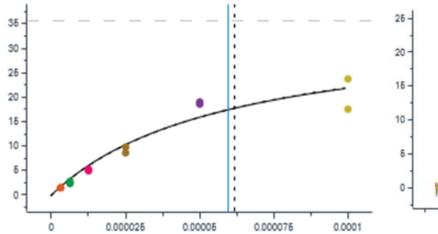


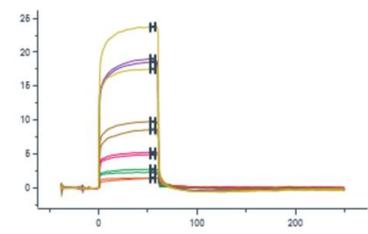
# **PARENT MOLECULE** 0.000076 CACHE\_1454\_13 $K_D = 86\pm 4 \,\mu\text{M} - 142\pm 11\% \text{ binding}$ Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM) %inhibition@50 $\mu$ M\_ATPase = 13

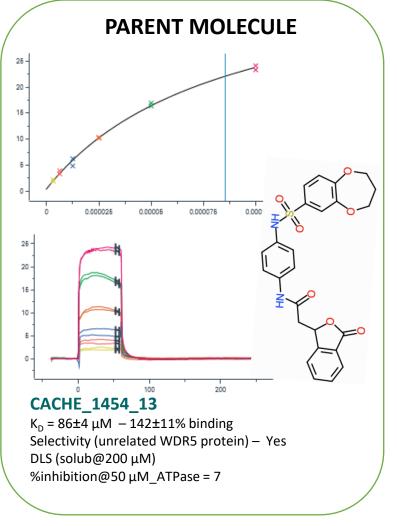
6 analogs of CACHE\_1454\_13 were submitted for round 2. The resupplied parent molecule confirmed a selective dose dependent binding response with a tendency to reach saturation.

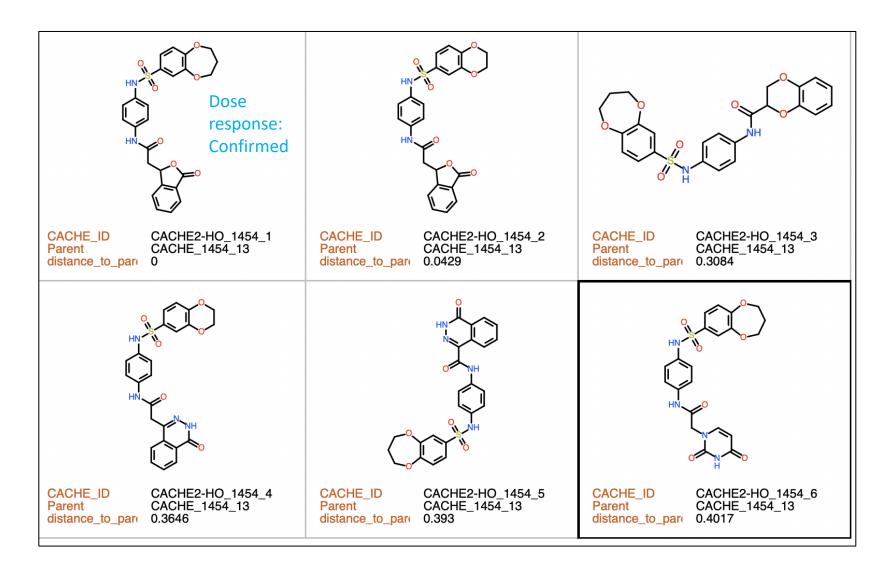
#### CACHE2-HO\_1454\_1

 $K_D _1 = 92 \mu M - 111\%$  binding  $K_D _2 = 60 \mu M - 98\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 uM) %inhibition@50  $\mu M_ATPase = 5$ 



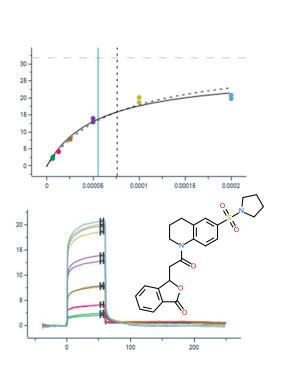




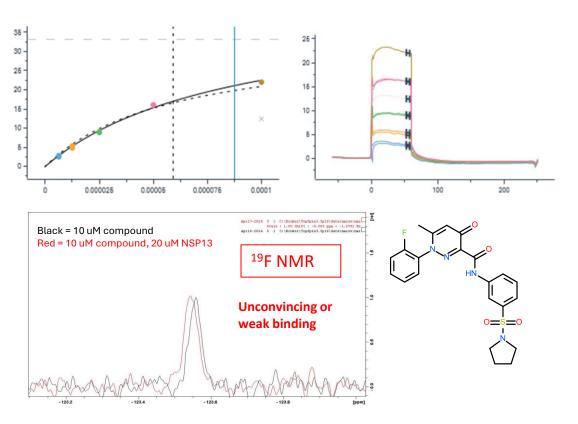


# PARENT MOLECULE 0.000026 **CACHE 1454 91** $K_D = 35\pm42 \,\mu\text{M} - 71\pm34\% \,\text{binding}$ Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM) %inhibition@50 $\mu$ M\_ATPase = 1

23 analogs of CACHE\_1454\_91 were submitted for round 2. 2 compound (including re-supplied parent molecule) confirmed a selective dose dependent binding response with a tendency to reach saturation.



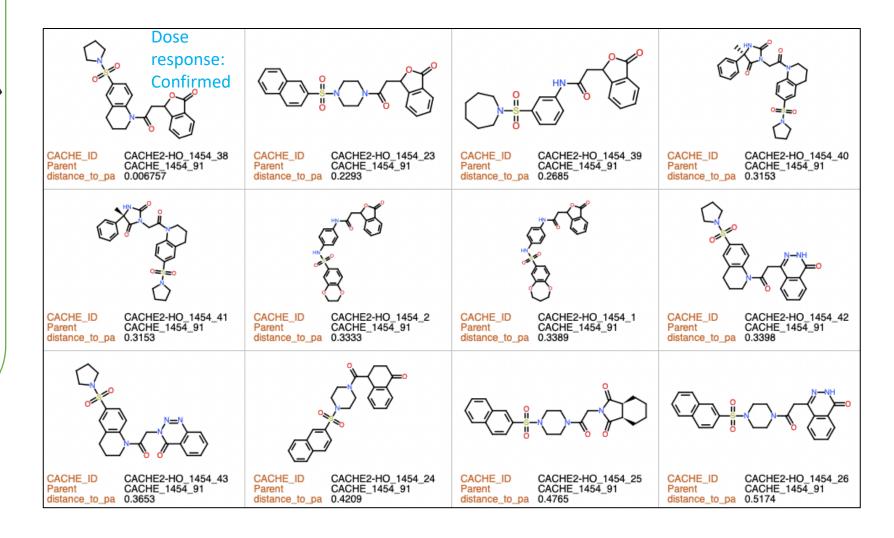
# CACHE2-HO\_1454\_38 $K_D = 55 \mu M - 86\% \text{ binding}$ Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 $\mu M$ ) %inhibition@50 $\mu M$ ATPase = 9



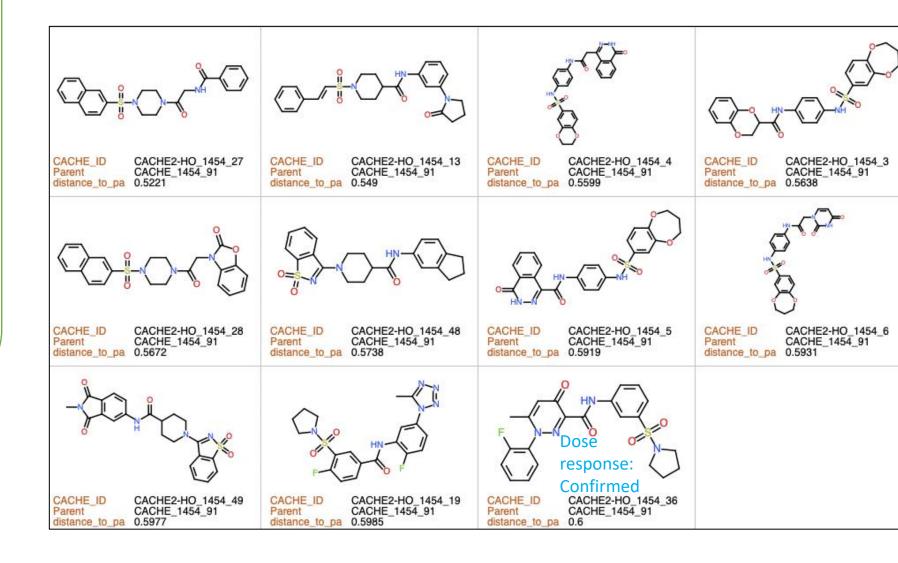
#### CACHE2-HO\_1454\_36

 $K_D$  (3% DMSO) = 88 μM - 127% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 10

### PARENT MOLECULE 12.5 -2.5 -0.000026 0.00005 0.000078 15 -12.5 2.5 200 **CACHE 1454 91** $K_D = 35\pm42 \,\mu\text{M} - 71\pm34\% \,\text{binding}$ Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM) %inhibition@50 µM\_ATPase = 1

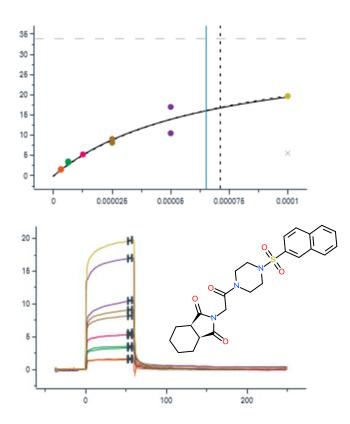


### PARENT MOLECULE 12.5 -2.5 -0.000026 0.00005 0.000078 15 -12.5 2.5 200 **CACHE 1454 91** $K_D = 35\pm42 \,\mu\text{M} - 71\pm34\% \,\text{binding}$ Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM) %inhibition@50 µM\_ATPase = 1



# **PARENT MOLECULE** 12.5 -0.000025 0.000076 200 CACHE\_1454\_46

 $K_D$  = 17 μM -58% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = no inhibition 6 analogs of CACHE\_1454\_46 were submitted for round 2. One compounds confirmed a selective dose dependent binding response by SPR.



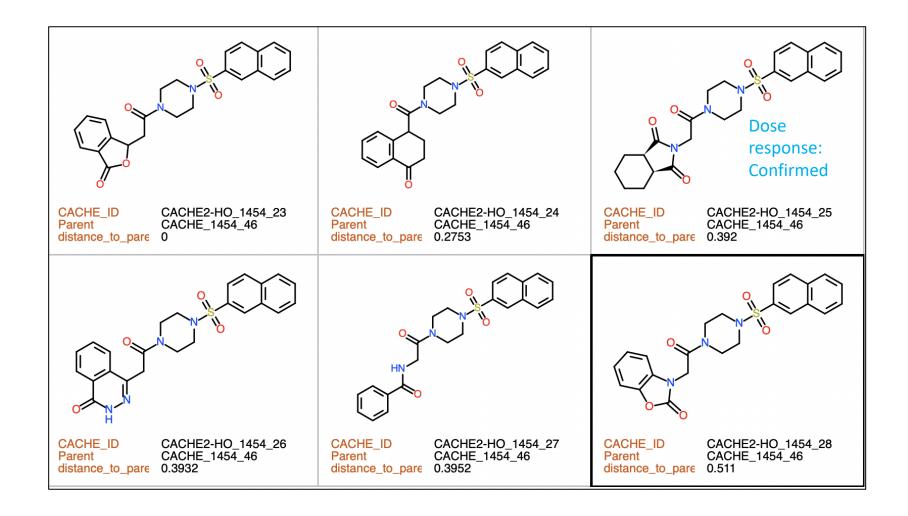
#### CACHE2-HO\_1454\_25

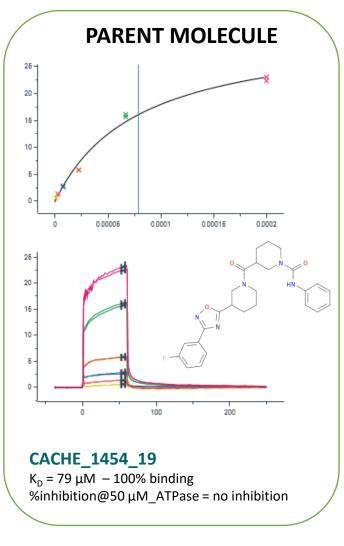
 $K_D _1 = 43 \mu M - 69\%$  binding  $K_D _2 = 65 \mu M - 95\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 14

### **PARENT MOLECULE** 12.5 -10 -7.5 -0.000026 0.00005 0.000076 12.5 -2.5 -100 200

#### **CACHE 1454 46**

 $K_D = 17 \mu M - 58\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 µM) %inhibition@50 µM\_ATPase = no inhibition



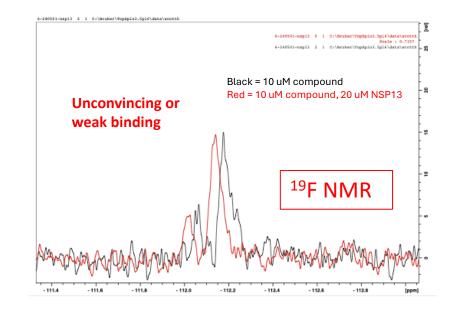


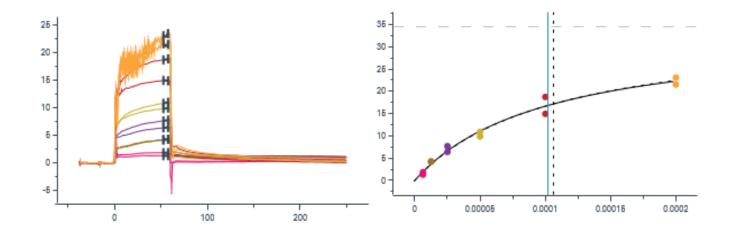
5 analogs of CACHE\_1454\_19 were submitted for round 2. Re-supplied parent molecule confirmed a selective dose dependent binding

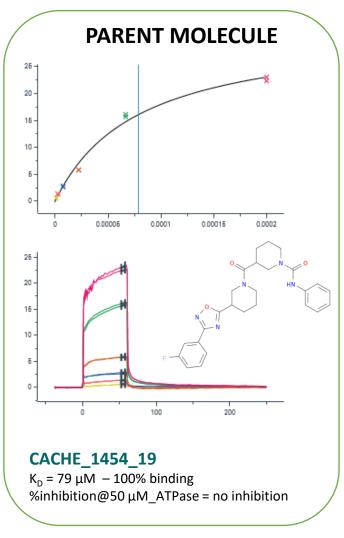
response by SPR.

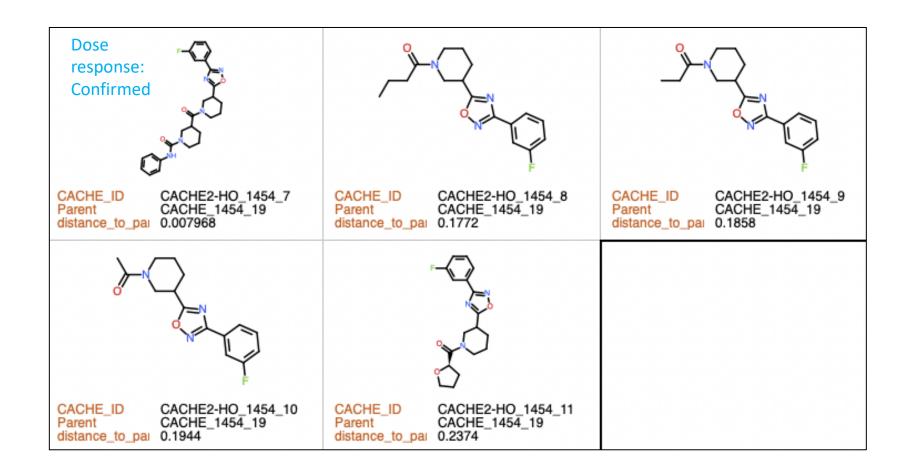
#### CACHE2-HO\_1454\_7

 $K_D = 102 \mu M - 98\%$  binding Selectivity (unrelated WDR5 protein) – Yes %inhibition@50  $\mu M$  ATPase = 11

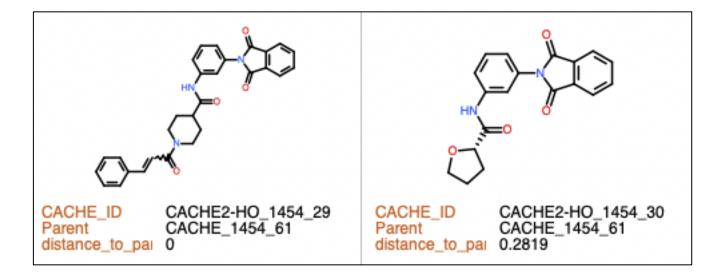






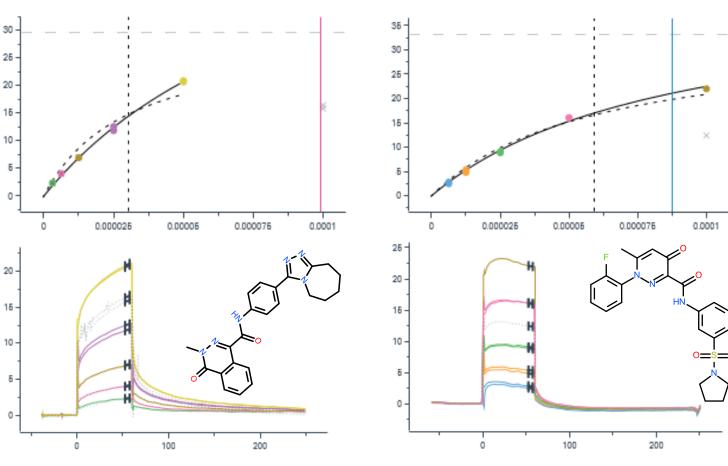


## **PARENT MOLECULE** 7.5 -2.5-0.00001 0.00002 0.00003 7.5 -2.6-CACHE\_1454\_61 $K_D = 5 \mu M - 36\%$ binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@100 µM) $IC_{50}$ ATPase = 29 $\mu$ M



## **PARENT MOLECULE** 26 -20 0.00005 0.0001 0.00016 26 20 **CACHE 1454 75** $K_D = 47 \mu M (slow on/off) - 106\% binding$ Selectivity (unrelated WDR5 protein) – Yes %inhibition@50 µM\_ATPase = 1

7 analogs of CACHE\_1454\_75 were submitted for round 2. 1 compounds showed a selective dose dependent binding response by SPR.

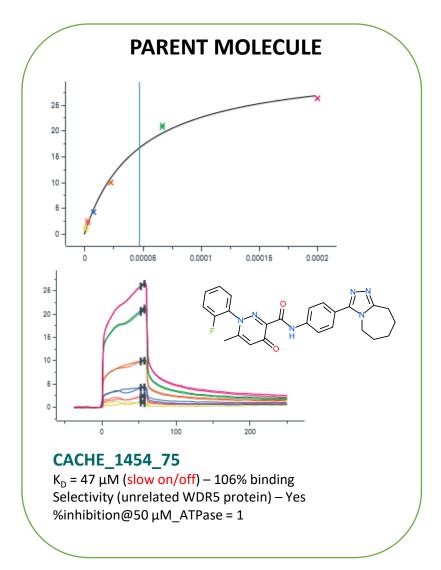


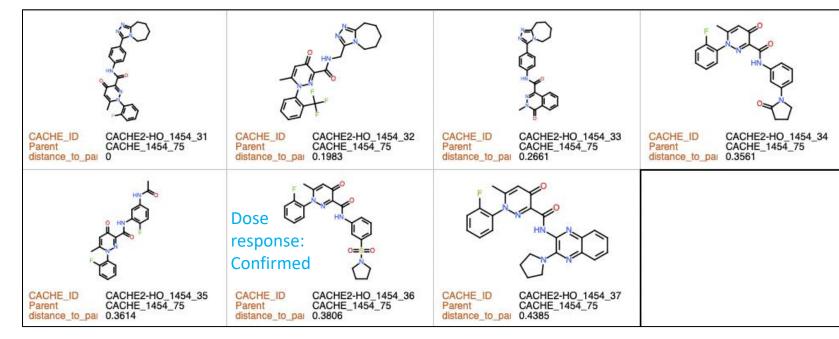
### **CACHE2-HO\_1454\_33** $K_D$ (3% DMSO) = 99 $\mu$ M (does not reach

saturation/solubility issue) – 209% binding DLS(solub@200 μM) %inhibition@50 μM ATPase = 14

#### CACHE2-HO\_1454\_36

 $K_D$  (3% DMSO) = 88  $\mu$ M – 127% binding Selectivity (unrelated WDR5 protein) – Yes DLS(solub@200  $\mu$ M) %inhibition@50  $\mu$ M\_ATPase = 10

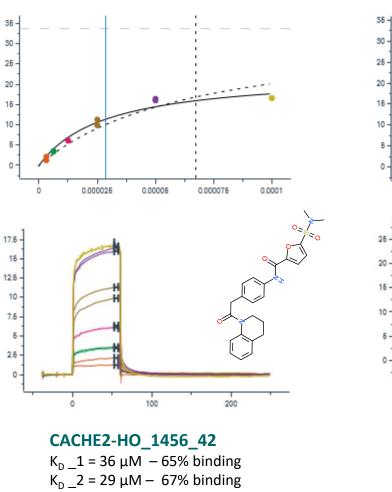




# CACHE#2 – NSP13 SARS2 Participant 1456

## **PARENT MOLECULE** 0.00006 0.0001 0.00016 0.0002 CACHE\_1456\_73 $K_D = 45 \mu M - 88\%$ binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM) %inhibition@50 µM\_ATPase = no inhibition

16 analogs of CACHE\_1456\_73 were submitted for round 2. 2 compound showed a dose dependent binding response by SPR.



 $K_D _1 = 36 \mu M - 65\%$  binding  $K_D _2 = 29 \mu M - 67\%$  binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM\_ATPase = 9

#### CACHE2-HO\_1456\_26

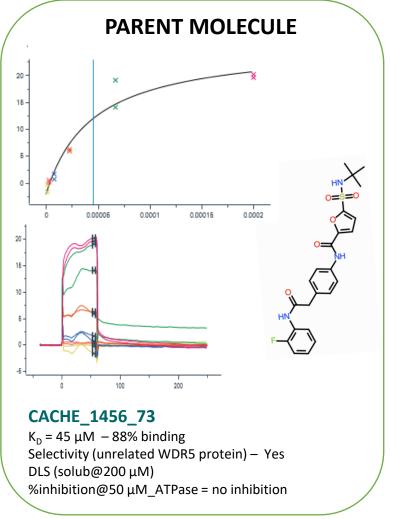
0.00005

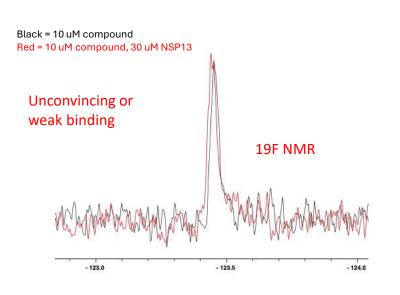
0.0001

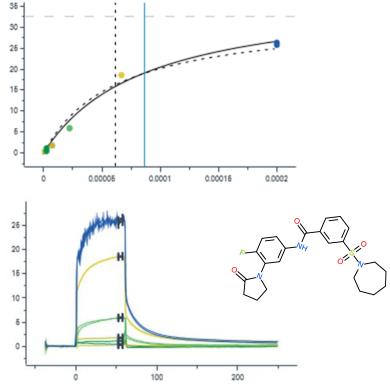
0.00015

0.0002

 $K_D$  = 86 μM – 116% binding Selectivity (unrelated WDR5 protein) – Yes DLS (solub@200 μM) %inhibition@50 μM ATPase = 19





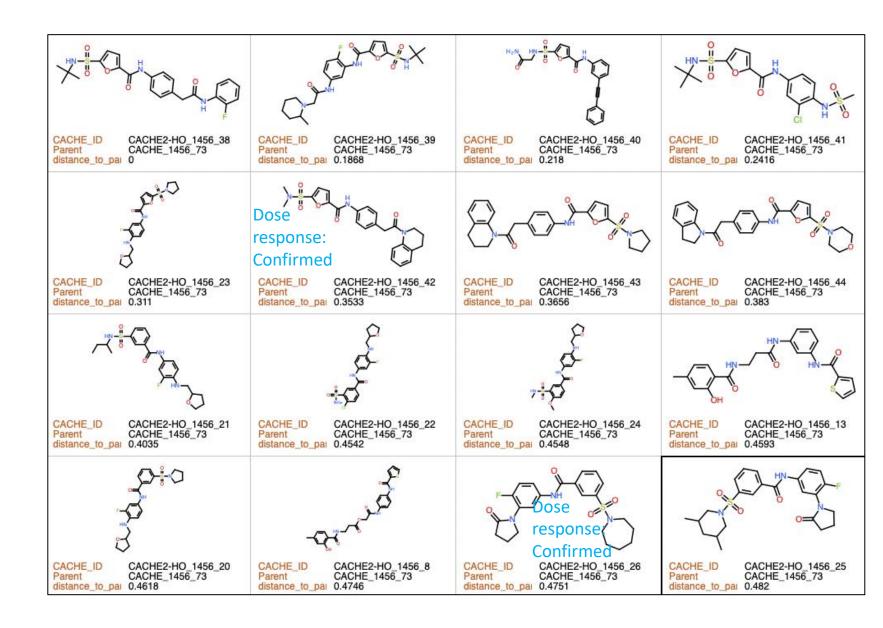


#### CACHE2-HO\_1456\_26

 $K_D$  = 86  $\mu$ M - 117% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M\_ATPase = 19

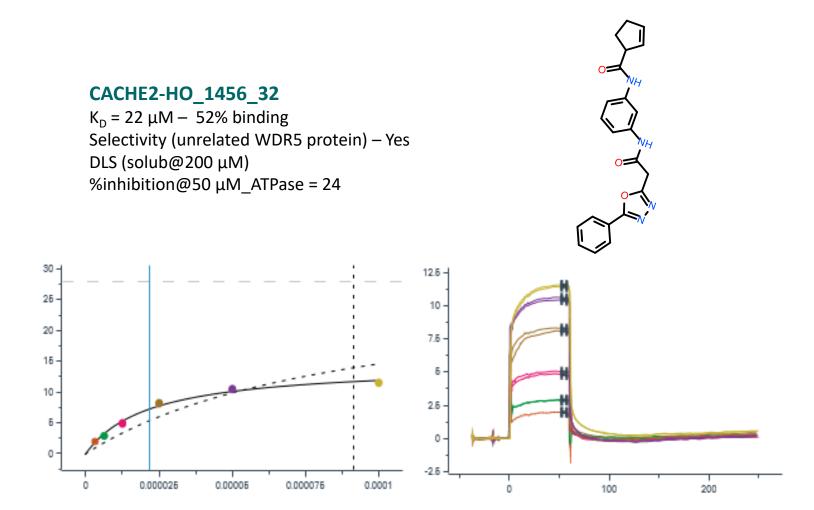
## PARENT MOLECULE 0.00006 0.0001 0.00016 0.0002 CACHE\_1456\_73 $K_D = 45 \mu M - 88\%$ binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200 µM)

%inhibition@50 μM ATPase = no inhibition

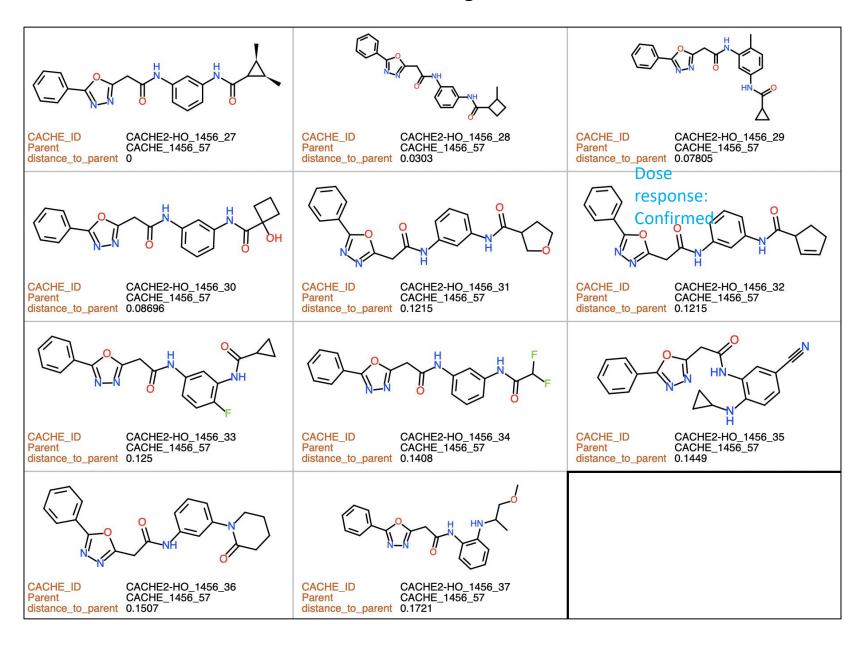


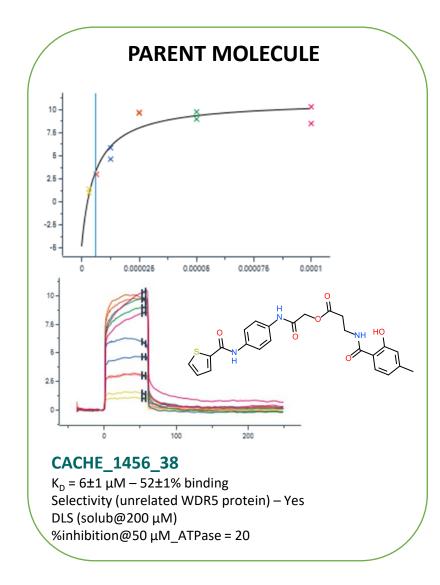
# **PARENT MOLECULE** 0.0001 CACHE\_1456\_57 $K_D$ (3% DMSO) = 12 $\mu$ M - 43% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@100 µM) %inhibition@50 μM\_ATPase = 59%, upon retesting was 31%

11 analogs of CACHE\_1456\_57 were submitted for round 2. 1 compound showed a binding response reaching saturation.



#### PARENT MOLECULE 25 20 -16 -0.000026 0.00006 0.000076 0.0001 0.000125 12.5 -10 -2.5 -200 CACHE\_1456\_57 $K_D$ (3% DMSO) = 12 $\mu$ M - 43% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@100 µM) %inhibition@50 µM ATPase = 59%, upon retesting was 31%



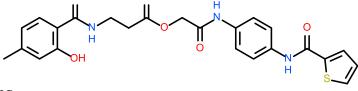


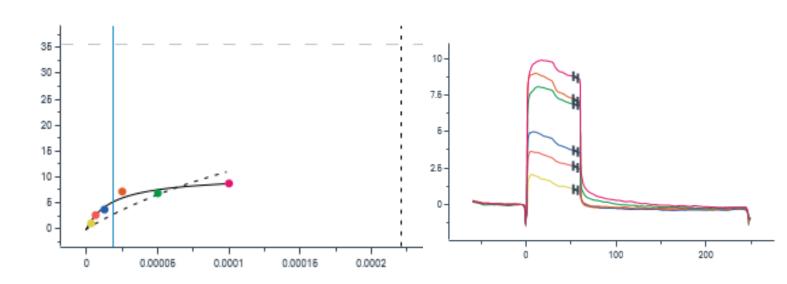
12 analogs of CACHE\_1456\_38, including the re-supplied hit molecule, were submitted for round 2.

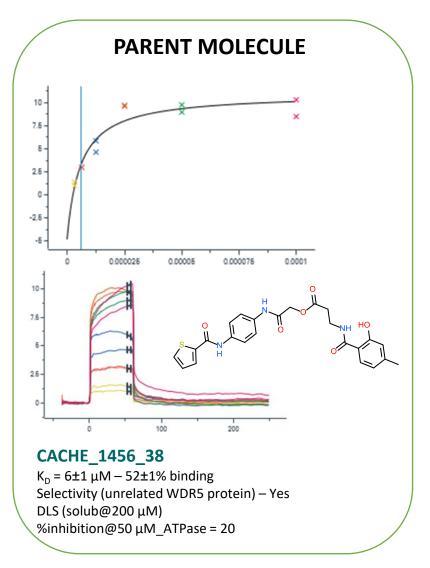
1 compound (re-supplied parent molecule) confirmed weak binding by SPR, reaching saturation.

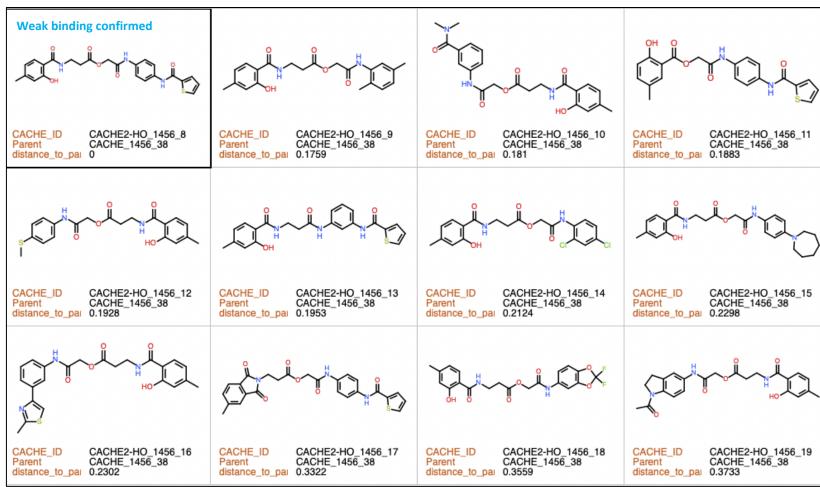
#### CACHE2-HO\_1456\_8

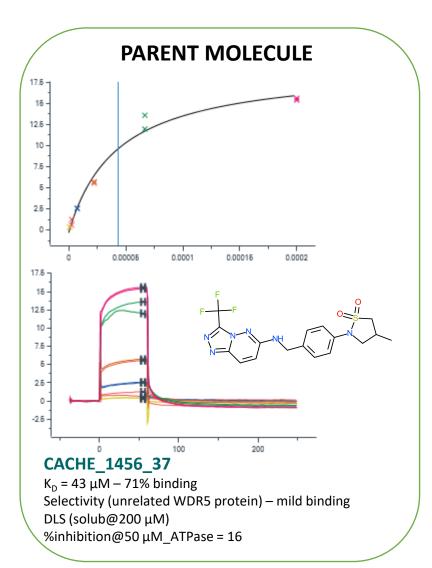
 $K_D$  = 19±0.4 μM – 30±1% binding Selectivity (unrelated WDR5 protein) – 10% binding  $IC_{50}$  ATPase > 100 μM

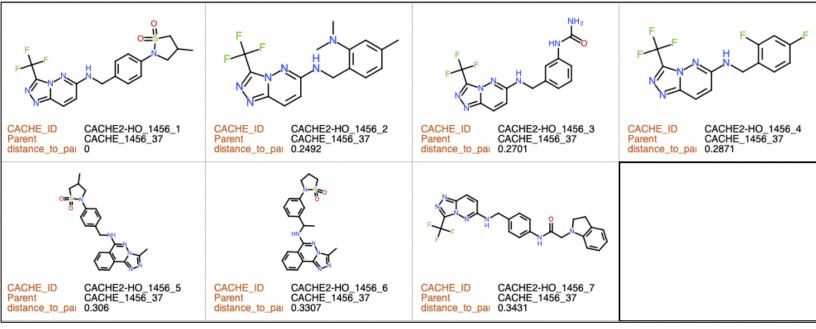






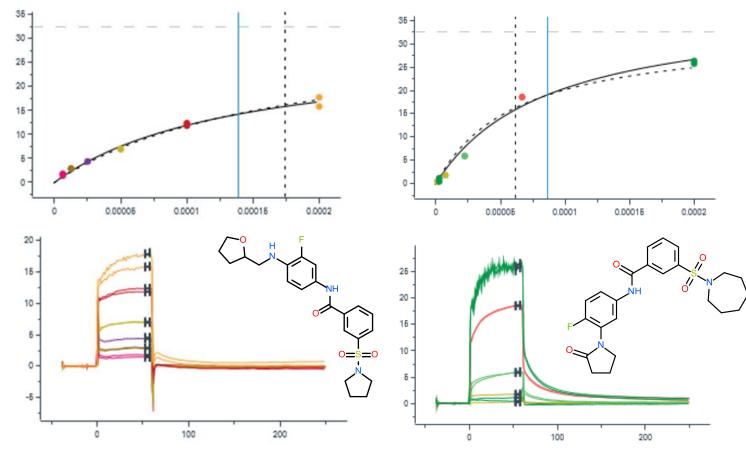






## **PARENT MOLECULE** 0.000026 0.000076 0.00006 0.0001 2.5-**CACHE 1456 42** $K_D = 95 \mu M$ (does not reach saturation) – 59% binding aggreg@200 μM %inhibition@50 µM ATPase = 12

7 analogs of CACHE\_1456\_42 were submitted for round 2. 2 compounds (including re-supplied parent molecule) showed a dose dependent binding response with a tendency to reach saturation.



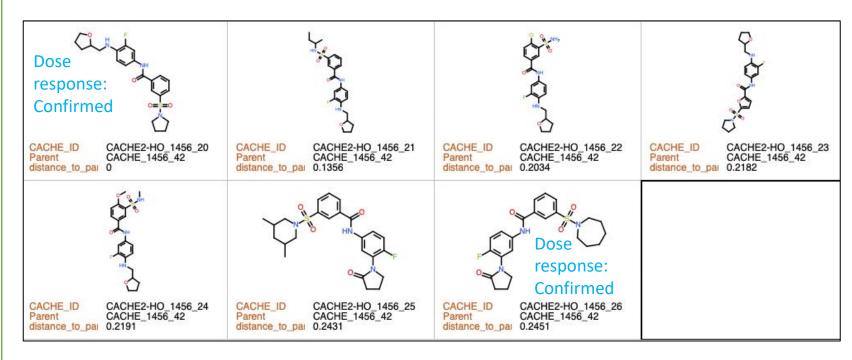
#### CACHE2-HO\_1456\_20

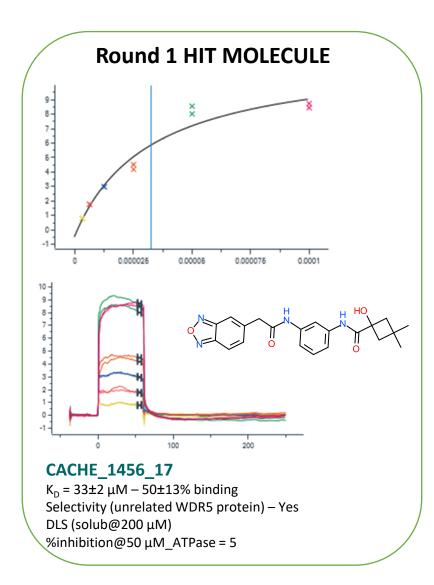
 $K_D$  = 139  $\mu$ M - 88% binding Selectivity (unrelated WDR5 protein) - Yes %inhibition@50  $\mu$ M\_ATPase = 13

#### CACHE2-HO 1456 26

 $K_D$  = 86  $\mu$ M - 117% binding Selectivity (unrelated WDR5 protein) - Yes DLS (solub@200  $\mu$ M) %inhibition@50  $\mu$ M\_ATPase = 19

## **PARENT MOLECULE** 0.000026 0.00006 0.000076 0.0001 2.5-CACHE\_1456\_42 $K_D = 95 \mu M$ (does not reach saturation) – 59% binding aggreg@200 μM %inhibition@50 µM\_ATPase = 12





No analog in Round 2