

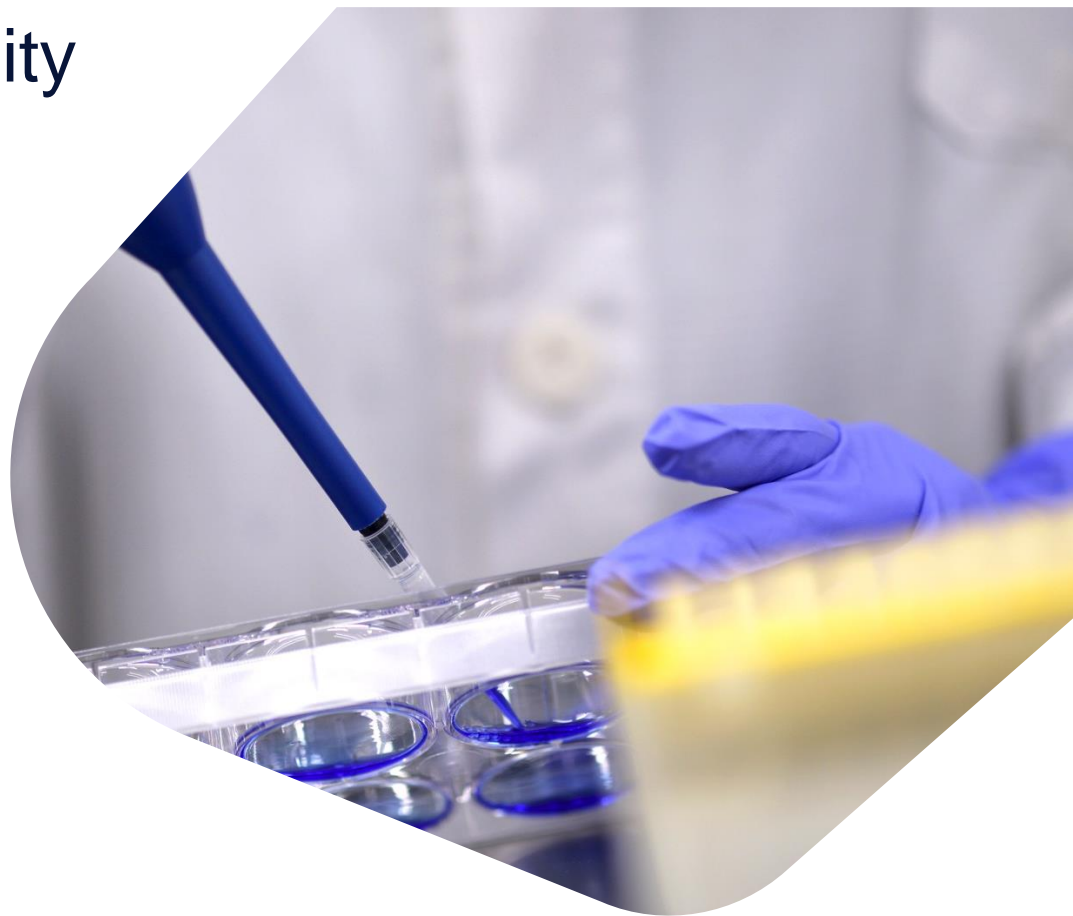
# Emerging Modalities and Compound Developability Assessment in Small Molecule Early Development

MCERSI Co-Processed API and  
Regulatory Requirements

Ahmad Sheikh

July 13, 2022

abbvie



VPAK-US-00002-E

# Acknowledgements

Rajni Miglani Bhardwaj

Richard Hong

Rodger Henry

Moiz Diwan

Vivian Suarez

David DeGoey

Paul Breckemeyer

Gabriela Schneider-Rauber

Alessandra Mattei

Nathan Abraham

Kenneth Engstrom

Yi Gao

Erin Jordan

Charles Hutchins

Yue Gui

Gerry Danzer

*Authors are employees of AbbVie and may own AbbVie stock. AbbVie sponsored and funded the study; contributed to the design; participated in the collection, analysis, and interpretation of data, and in writing, reviewing, and approval of the final publication.*

# Last two decades

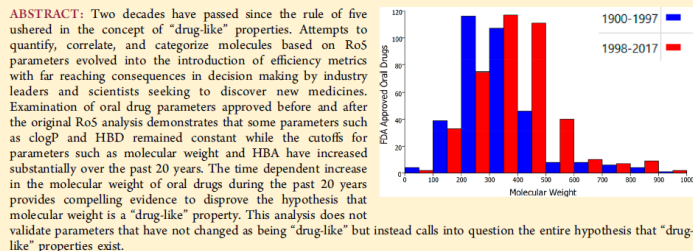
## Two Decades under the Influence of the Rule of Five and the Changing Properties of Approved Oral Drugs

Miniperspective

Michael D. Shultz<sup>\*,†</sup>

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<sup>‡</sup> Supporting Information



*J. Med. Chem.* 2019, 62, 4, 1701–1714

Table 2. Analysis of FDA Approved Oral NCEs from 1998 to 2007<sup>a</sup>

	clogP	MW	HBD	HBA	TPSA	RotB	Fsp <sup>3</sup>	#ArRNG
1997 90 <sup>th</sup> percentile	4.7	470.3	4.0	10	139.8	10.0	0.83	3
90 <sup>th</sup> percentile	4.7 (0)	525.5 (+ 55.2)	4.0 (0)	9.6 (-0.4)	142.3 (+2.5)	11 (+1)	0.78 (-0.05)	3.0 (0)
p value <sup>*</sup>	0.3	<b>0.0026</b>	>0.99	0.69	0.97	<b>0.039</b>	0.082	>0.99
Median	2.6 (+0.3)	348.4 (+40.1)	1 (0)	6 (+2)	74.7 (+7.2)	6 (+2)	0.43 (+0.03)	2 (+1)
Mean	2.4 (+0.3)	360.1 (+28.1)	1.8 (-0.1)	6.0 (+0.5)	82.0 (+3.2)	5.9 (+0.9)	0.46 (+0.03)	1.7 (+0.3)
p value <sup>†</sup>	0.35	0.28	0.94	0.55	0.94	<b>0.066</b>	0.56	<b>0.025</b>
p value <sup>‡</sup>	0.46	<b>0.014</b>	>0.99	<b>0.044</b>	0.43	<b>0.0085</b>	0.73	0.73
10 <sup>th</sup> percentile	-0.4 (+0.2)	201.3 (+30.1)	0.0 (0)	2.0 (0)	30.5 (+9.2)	1.0 (0)	0.08 (+0.1)	0 (0)
p value <sup>*</sup>	0.6	<b>&lt;0.0001</b>	>0.99	>0.99	0.37	>0.99	<b>&lt;0.0001</b>	>0.99

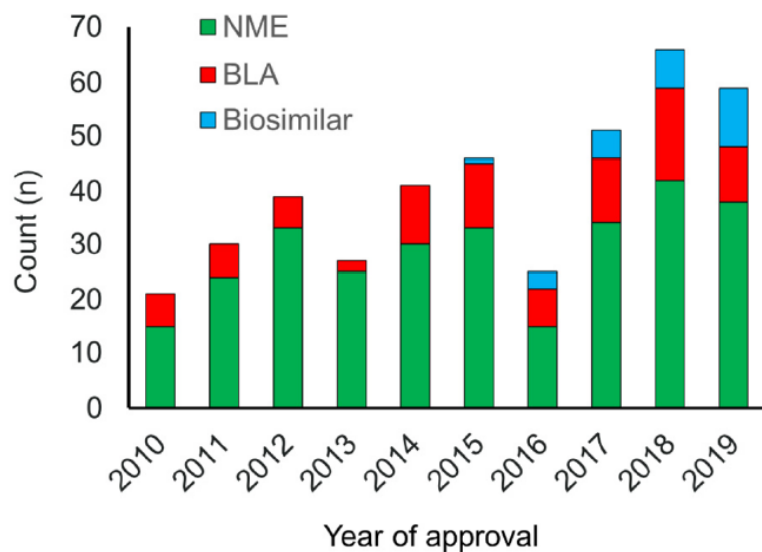
<sup>a</sup>*n* = 195, or 26% of all FDA approved oral NCEs. The 90<sup>th</sup> percentile values determined for FDA approved oral drugs from 1900 to 1997 are included for reference. The change in values from FDA approved oral drugs from 1900 to 1997 values are in parentheses. <sup>\*</sup>*p* value of the 10<sup>th</sup> and 90<sup>th</sup> percentile cutoffs based on one-way ANOVA analyses. <sup>†</sup>*p* values from one-way ANOVA (normal distribution). <sup>‡</sup>*p* values from Kurskal–Wallace (non-normal distribution). Differences that achieve statistical significance (*p* < 0.05) are in bold. Categories that have a statistically significant increase are shaded in red.

Table 3. Analysis of FDA Approved Oral NCEs from 2008 to 2017<sup>a</sup>

	clogP	MW	HBD	HBA	TPSA	RotB	Fsp <sup>3</sup>	#ArRNG
1997 90 <sup>th</sup> percentile	4.7	470.3	4.0	10	139.8	10.0	0.83	3
90 <sup>th</sup> percentile	5.0 (+0.3)	607.3 (+137.0)	4.0 (0)	11.7 (+2.0)	150.7 (+10.9)	13.7 (+3.7)	0.69 (-0.14)	4.0 (+1.0)
p value <sup>*</sup>	0.27	<b>&lt;0.0001</b>	>0.99	<b>&lt;0.0001</b>	0.20	<b>&lt;0.0001</b>	<b>&lt;0.0001</b>	<b>&lt;0.0001</b>
Median	3.3 (+1.0)	420.0 (+111.7)	2 (+1)	6 (+2)	80.8 (+13.3)	6 (+2)	0.38 (-0.02)	2 (+1)
Mean	2.9 (+0.8)	436.5 (+104.5)	2.1 (+0.2)	6.8 (+1.4)	93.6 (+14.8)	7.0 (+2.0)	0.40 (-0.03)	2.2 (+0.8)
p value <sup>†</sup>	<b>&lt;0.0001</b>	<b>&lt;0.0001</b>	0.57	<b>0.0008</b>	<b>0.037</b>	<b>&lt;0.0001</b>	0.65	<b>&lt;0.0001</b>
p value <sup>‡</sup>	<b>&lt;0.0001</b>	<b>&lt;0.0001</b>	0.35	<b>&lt;0.0001</b>	<b>0.005</b>	<b>&lt;0.0001</b>	0.99	<b>&gt;0.99</b>
10 <sup>th</sup> percentile	0.3 (+0.9)	235.5 (+64.3)	0.0 (0)	3.0 (+1)	39.7 (+14.8)	3 (+2)	0.16 (+0.07)	0 (0)
p value <sup>*</sup>	<b>&lt;0.0001</b>	<b>&lt;0.0001</b>	>0.99	<b>0.024</b>	<b>0.0049</b>	<b>&lt;0.0001</b>	<b>0.0005</b>	>0.99

<sup>a</sup>*n* = 214, or 29% of all FDA approved oral NCEs. The 90<sup>th</sup> percentile values determined for FDA approved oral drugs from 1900 to 1997 are included for reference. The change in values from FDA approved oral drugs from 1900 to 1997 values are in parentheses. <sup>\*</sup>*p* value of the 10<sup>th</sup> and 90<sup>th</sup> percentile cutoffs based on one-way ANOVA analyses. <sup>†</sup>*p* values from one-way ANOVA (normal distribution). <sup>‡</sup>*p* values from Kurskal–Wallace (non-normal distribution). Differences that achieve statistical significance (*p* < 0.05) are in bold. Categories that have a statistically significant increase are shaded in red, and those that have a statistically significant decrease are shaded in green.

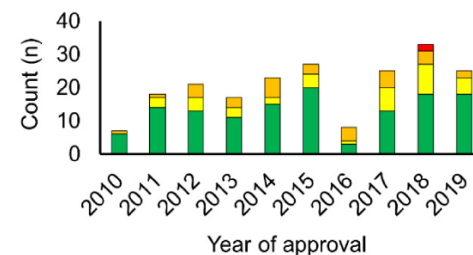
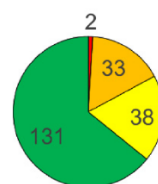
# Last decade



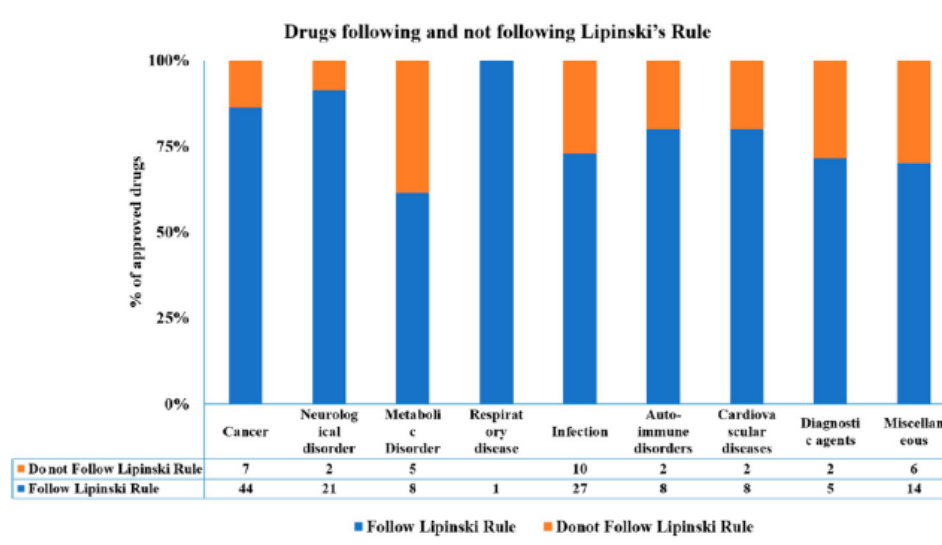
*J. Med. Chem.* 2021, 64, 5, 2312–2338

*J. Med. Chem.* 2021, 64, 5, 2339–2381

**a** Ro5 violations 2010-2019 (oral drugs) **b** Ro5 violations by year 2010-2019 (oral drugs)



(a) Total analysis of Ro5 violations for 204 approved oral drugs and (b) Ro5 violations per year on 204 approved oral drugs.



Bar graph represents percent of approved FDA drugs from the year 2015 until June 2020, following and not following Lipinski's Rule.

# PROTAC's

## way beyond rule of 5

**Table 1**  
Representative recent PROTACs organized by E3 warhead: MDM2, IAP, VHL, and CRBN binders.

PROTAC	Structure	Name	Target (POI)	E3 Ligase	DC <sub>50</sub> /D <sub>max</sub>	Cellular/ <i>in vitro</i>	Ref
1		Compound 14	AR	MDM2	10 µM/-	<i>In vitro</i>	18
2		A1874	BRD4	MDM2	32 nM/98% at 100 nM	<i>In vitro</i>	19
3		SNIPER(BRD4)-1	BRD4	IAP	> 3 nM & < 10 nM/70% at 10 nM	<i>In vitro</i> probe	20
4		SNIPER(ABL)-39	ABL	IAP	> 3 nM & < 10 nM/ > 90% at 100 nM	<i>In vitro</i> probe	21
5		SNIPER(ER)-87	ERα	IAP	> 1 nM & < 3 nM/70% at 10 nM	<i>In vitro</i> efficacy (IP injection)	22
6		SNIPER(ER)-110	ERα	IAP	< 3 nM/80% at 100 nM	<i>In vitro</i> probe	23
7		MZ1	BRD4	VHL	< 100 nM (BRD4)/ > 96% at 50 nM	<i>In vitro</i> cellular probe	26
8		12b	BRD4	VHL	0.083 µM/-	<i>In vitro</i>	27
9		ARV-771	BRD4	VHL	< 5 nM for BRD2/3/4/ > 99%	<i>In vitro</i> (SC) efficacy	28
10		AT1	BET	VHL	> 10 nM & < 100 nM for BRD4 short/ > 90%	<i>In vitro</i> prob	29
11		MZP54	BET	VHL	10 nM- < 100 nM/87% at 50 nM	<i>In vitro</i> cellular prob	30
12		PROTAC_ERRα (1)	ERRα	VHL	100 nM/86% at 1 µM	<i>In vitro</i> probe (IP injection)	31

(continued on next page)

Will these be orally bioavailable  
Would we crystallize them  
What would the formulations be

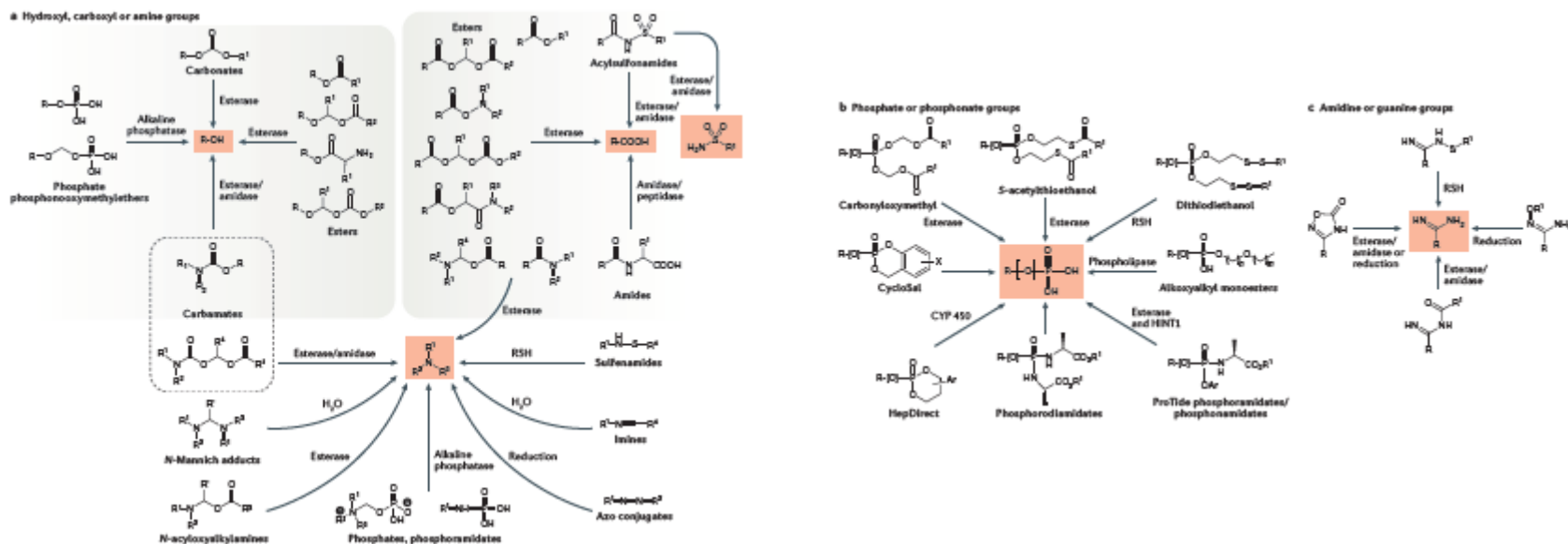
**Table 2**  
*In silico* metrics for selected PROTACs.

Compound	E3 Ligase	MW	cLogP	HBD	HBA	PSA	nRotB	N <sub>rule-of-5</sub>	Ar Rings	cLogD	AB-MPS
1	MDM2	1210	11	4	22	268	31	3	5	7.3	40.3
2	MDM2	1174	9.1	4	16	200	27	3	6	7.9	37.9
Average	MDM2	1192	10.1	4.0	19	234	29	3.0	5.5	7.6	39.1
3	IAP	1057	6.1	3	17	197	27	3	5	4.7	33.7
4	IAP	1115	5.7	4	20	228	29	3	5	5.9	36.9
5	IAP	1044	8.9	3	15	182	31	3	6	6.5	40.5
6	IAP	1122	11.4	4	16	196	33	3	5	7.5	42.5
Average	IAP	1085	8.0	3.5	17	201	30	3.0	5.3	6.2	38.4
7	VHL	1003	4.9	4	17	210	25	2	5	3.5	30.5
8	VHL	1040	4.2	4	19	229	24	2	6	3.4	30.4
9	VHL	987	5.9	4	16	202	23	3	5	3.2	28.2
10	VHL	973	6.1	4	14	184	22	3	5	4.4	28.4
11	VHL	1037	6.3	5	16	211	27	3	5	4.6	33.6
12	VHL	949	6.9	3	15	202	21	3	4	4.5	26.5

*Bioorganic & Medicinal Chemistry Letters* 29 (2019) 1555–1564

# Prodrugs- chemically labile by design

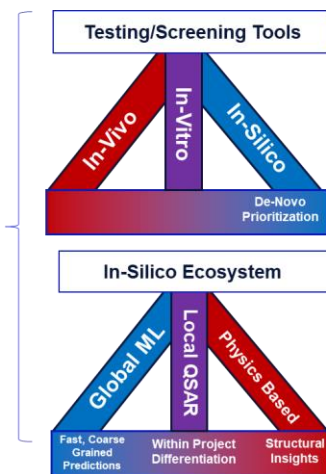
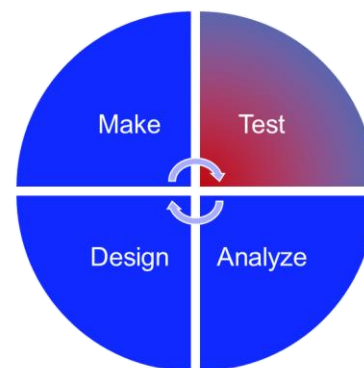
- Over 30 approved in the last decade ('08-17), representing ~ 12% of all approved small molecules



- Limited stability across physiologically relevant pH range, limited understanding of bio-conversion and metabolism
- Complex synthesis, complex solid state, analytical (CMC and bio-analytical)
- Regulatory challenges in setting specs for impurities such as “parent” and other pro-drug type impurities

Nature Reviews Drug Discovery, (2018) 559-587

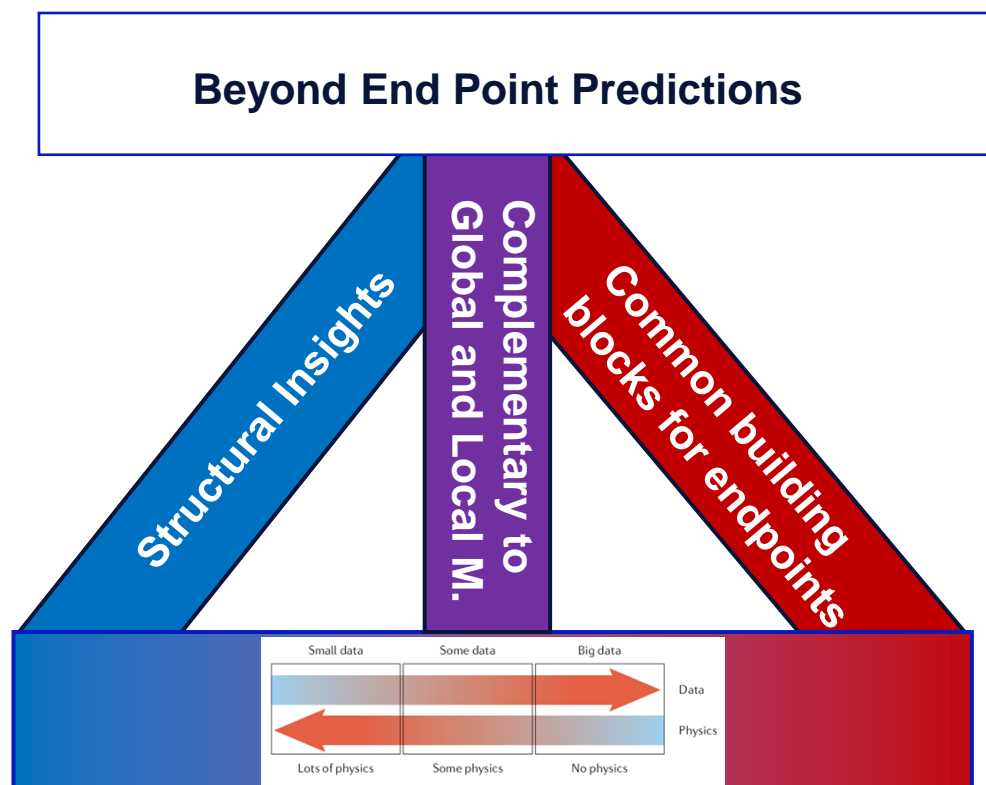
# Positioning of Developability earlier in the LO Funnel



Absorption	Distribution	Metabolism	Excretion	Toxicity	PhysChem
Caco-2 permeability	Human serum albumin binding	Microsomal stability	CL <sub>int</sub>	Ames mutagenicity	Aq. solubility
Caco-2 efflux ratio	Plasma-protein binding	Hepatocyte stability	Terminal half-life	CYP inhibition	LogD
PAMPA permeability	Fraction unbound (fu)	Site of metabolism		Phaspholipidosis	Membrane affinity
Blood-brain barrier penetration	Volume of distribution (vdss)			hERG inhibition	pKa
	Brain/plasma ratio (Kp)			Drug-induced liver injury (DILI)	

*Drug Disc Today*, Volume 27, Issue 4, 2022, 967-984

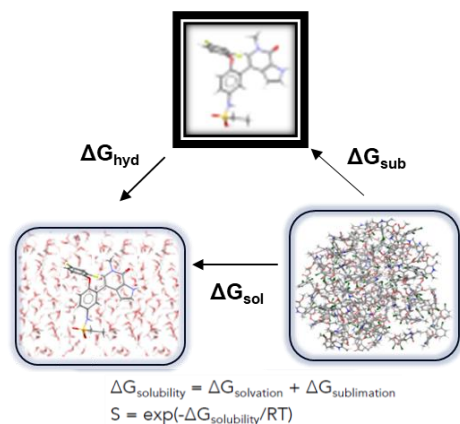
# Physics Based Models



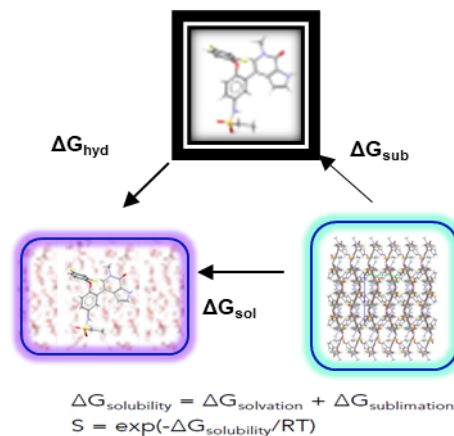
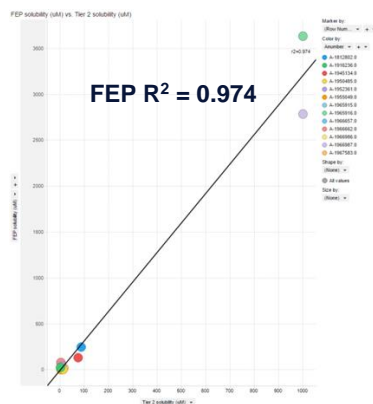
*Nat Rev Phys* **3**, 422–440 (2021)



# Solubility Prediction: Trending and Quantitatively differentiating Structurally Related Molecules



Utilizing a thermodynamic cycle, solvation & sublimation energies are calculated



- Ability to qualitatively predict solubility within a series
- Help remove low soluble compounds with ~30 % false negative

- Crystal structure prediction (CSP) to identify energetically favorable 3-D crystal packing using 2-D molecular structure
- Global minimum from CSP as input for thermodynamic cycle  
Quick version of CSP to bring efficiency for LO stage

# Thermodynamic Solubility Prediction before Synthesis and Crystallization

JMIM  
JOURNAL OF  
CHEMICAL INFORMATION  
AND MODELING

pubs.acs.org/jcim

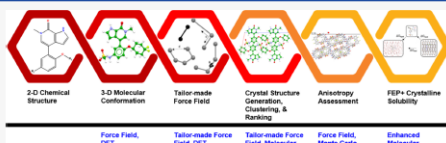
## Novel Physics-Based Ensemble Modeling Approach That Utilizes 3D Molecular Conformation and Packing to Access Aqueous Thermodynamic Solubility: A Case Study of Orally Available Bromodomain and Extraterminal Domain Inhibitor Lead Optimization Series

Richard S. Hong, Alessandra Mattei, Ahmad Y. Sheikh,\* Rajni Miglani Bhardwaj, Michael A. Bellucci, Keith F. McDaniel, M. Olivia Pierce, Guangxu Sun, Sirzhu Li, Lingle Wang, Sayan Mondal, Jianguo Ji, and Thomas B. Borchardt

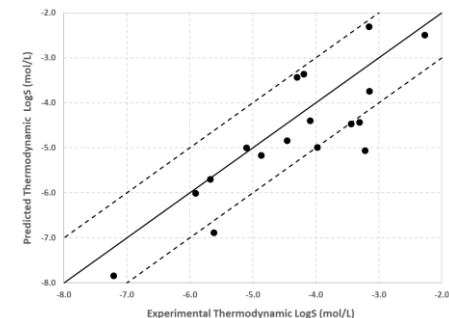
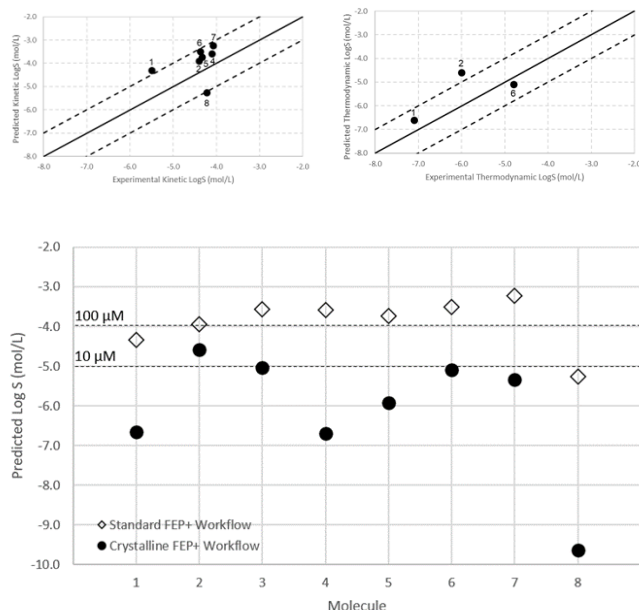
Cite This: *J. Chem. Inf. Model.* 2021, 61, 1412–1426

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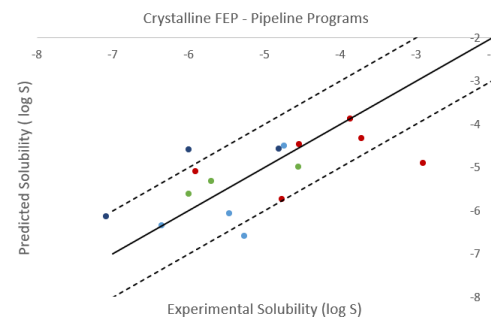
ACCESS | Metrics & More | Article Recommendations | Supporting Information



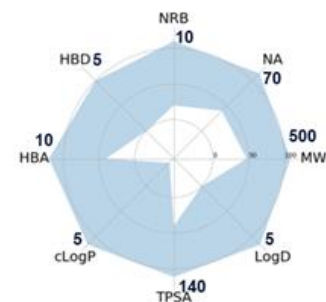
**ABSTRACT:** Drug design with patient-centricity for ease of administration and pill burden requires robust understanding of the impact of chemical modifications on relevant physicochemical properties early in lead optimization. To this end, we have developed a physics-based ensemble approach to predict aqueous thermodynamic crystalline solubility, with a 2D chemical structure as the input. Predictions for the bromodomain and extraterminal domain (RET) inhibitor series show very close match (0.5 log unit) with measured thermodynamic solubility for cases with low crystal anisotropy and good match (1 log unit) for high anisotropy structures. The importance of thermodynamic solubility is clearly demonstrated by up to a 4 log unit drop in solubility compared to kinetic (amorphous) solubility in some cases and implications thereof for instance on human dose. We have also demonstrated that incorporating predicted crystal structures in thermodynamic solubility prediction is necessary to differentiate (up to 4 log unit) between solubility of molecules within the series. Finally, our physics-based ensemble approach provides valuable structural insights into the origins of 3-D conformational landscapes, crystal polymorphism, and anisotropy that can be leveraged for both drug design and development.



## Structurally distinct AbbVie Dataset (17)



## Structurally related AbbVie Dataset (4 series)



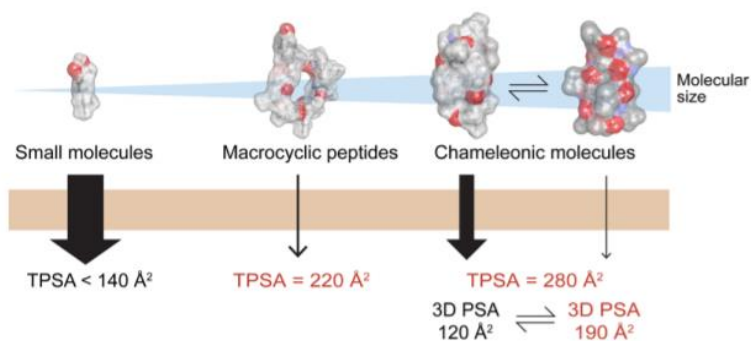
Incorporation of Physical Properties considerations in the design cycle

abbvie

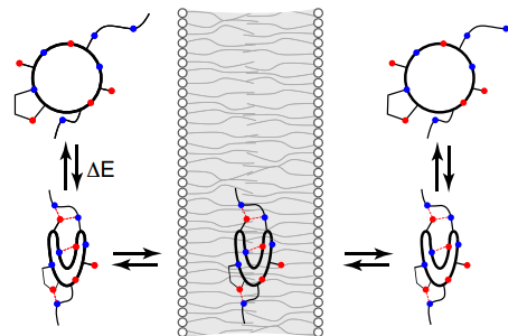
Company Confidential © 2021

*J. Chem. Inf. Model.* 2021, 61, 3, 1412–1426

# bR05 Chameleonicity: Towards Improved Passive Permeability

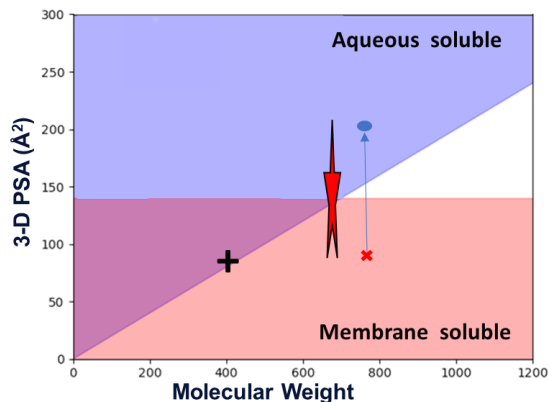


Size boundaries for membrane-permeable molecules



Conformational changes due to intramolecular hydrogen bonding (IMHB) + alkyl/ aryl shielding of polar groups facilitate permeation

A Framework to Quantify Chameleonic Behavior



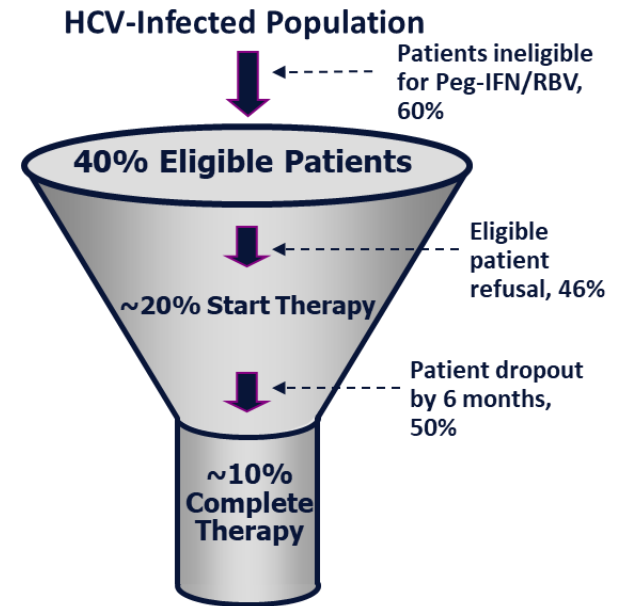
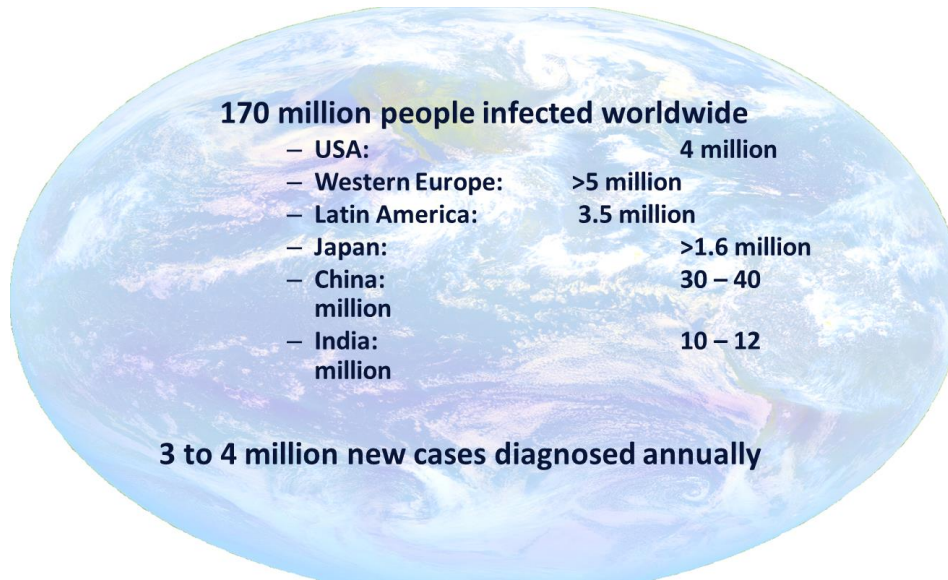
+ mean value reported for 1193 drugs  
★ MW ≥ 700, no single PSA value can satisfy both conditions

Generation of conformations in polar (aqueous) and non-polar (membrane) media and calculation of their BW SA 3D-PSA  
Provide information on IMHB, shielding of polarity in non-polar media which help improving the permeability



*J Med Chem*, 2017, 60, 1662-1664  
*Drug Disc Today*, 21, 2016, 713-717

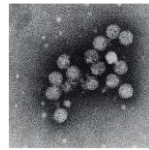
# Hepatitis C Infection- where were we a decade ago



**SVR\* or Relapse?**

First treatment:  
PEG-IFN/Ribavirin

- PEG-IFN: fatigue, flu-like symptoms, headache, dizziness, anorexia, depression, abdominal pain, irritability, insomnia, injection-site reaction, partial hair loss
- Ribavirin: hemolytic anemia, insomnia, teratogenic effects



Sustained viral  
response (SVR)

- Genotype 1 (48 wks, ≤50% SVR)
- Genotype 2 (24 wks, 80% SVR)



Poor response in  
special populations

- Cirrhosis patients (SVR 30%)
- HIV co-infected patients (SVR < 30%)
- African American patients (SVR < 20%)

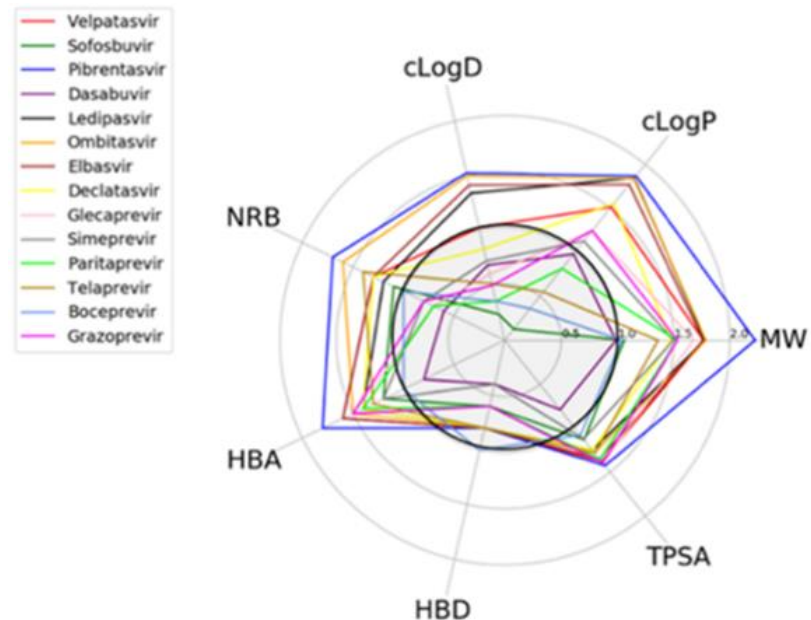
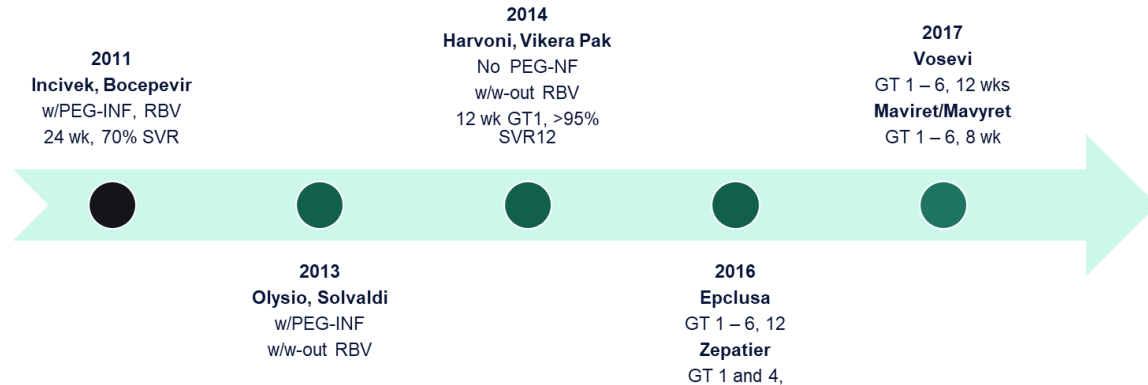
# and Since..

7 approved all-oral  
interferon free direct-  
acting antiviral (DAA)  
combination regimens.

Target nonstructural  
proteins responsible for  
replication and infection of  
HCV

Reach SVR12 > 95%  
across all prevalent  
genotypes

SVR12 also reduces  
adverse liver outcomes,  
such as cirrhosis, hepatic  
decompensation, and  
mortality



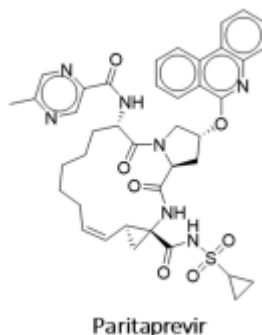
# Paritaprevir- the molecule

First generation NS3/4A protease inhibitor- discovered by AbbVie and Enanta Pharmaceuticals

Approved in 2014 as part of 1<sup>st</sup> gen DAA treatment regimen

Among 25 highest molecular weight approved oral drugs

Large triaryl phenanthridine group required for high potency in flat and featureless active binding site



## Implications of the Conformationally Flexible, Macrocyclic Structure of the First-Generation, Direct-Acting Anti-Viral Paritaprevir on Its Solid Form Complexity and Chameleonic Behavior

Ahmad Y. Sheikh,\* Alessandra Mattei, Rajni Miglani Bhardwaj, Richard S. Hong, Nathan S. Abraham, Gabriela Schneider-Rauber, Kenneth M. Engstrom, Moiz Diwan, Rodger F. Henry, Yi Gao, Vivian Juarez, Erin Jordan, David A. DeGoey, and Charles W. Hutchins

Cite This: *J. Am. Chem. Soc.* 2021, 143, 17479–17491

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## Distinct Hybrid Hydrates of Paritaprevir: Combined Experimental and Computational Assessment of their Hydration–Dehydration Behavior and Implications for Regulatory Controls

Published as part of a *Crystal Growth and Design* virtual special issue in Celebration of the Career of Roger Davey

Richard S. Hong, Rajni Miglani Bhardwaj, Rodger Henry, Alessandra Mattei, Moiz Diwan, Albert Thomas, Gerald D. Danzer, and Ahmad Y. Sheikh\*

Cite This: *Cryst. Growth Des.* 2022, 22, 726–737

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## Origins and Implications of Extraordinarily Soft Crystals in a Fixed-Dose Combination Hepatitis C Regimen

Published as part of a *Crystal Growth and Design* joint virtual special issue on Crystallizing the Role of Solid-State Form in Drug Delivery

Rajni Miglani Bhardwaj, Raimundo Ho, Yue Gui, Paul Brackemeyer, Gabriela Schneider-Rauber, Fredrik L. Nordstrom, and Ahmad Y. Sheikh\*

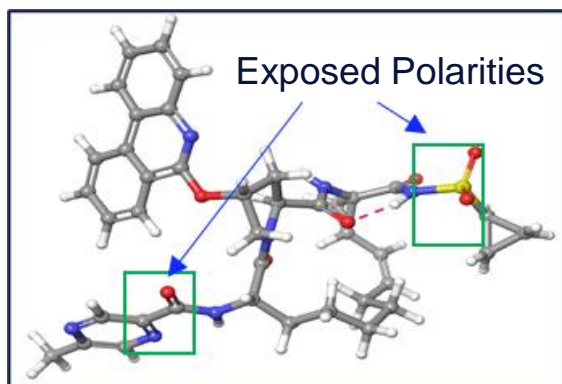
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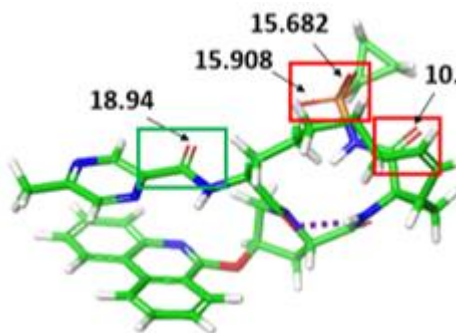
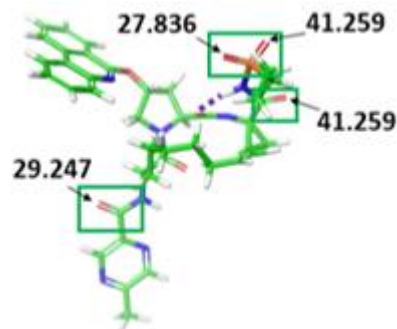
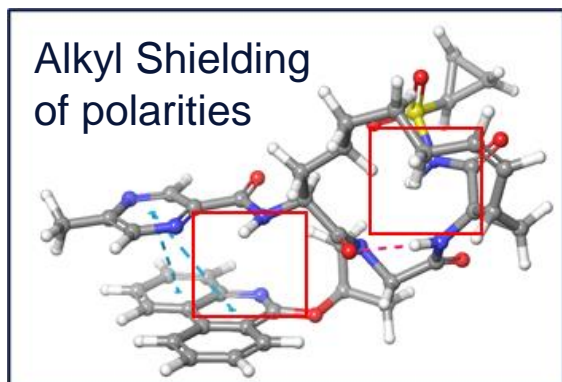


# Computational Assessment of Chameleonicity

Water Conformation\*  
3D PSA: 201.2 Å<sup>2</sup>



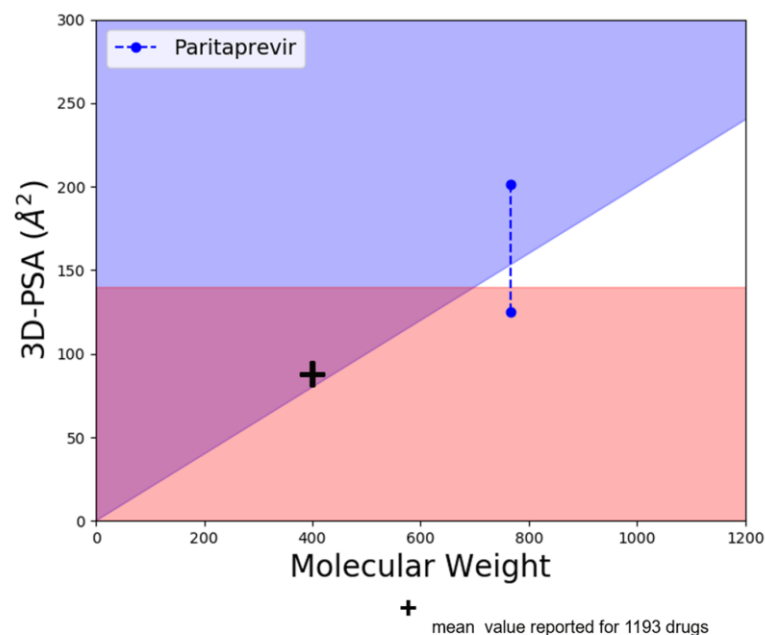
Chloroform Conformation  
3D PSA: 125.1 Å<sup>2</sup>



Measured logD<sub>1-octano/pH 6.8</sub> 3.1

Caco-2 permeability of > 60 × 10<sup>-6</sup> cm/second

EPSA 132 Å<sup>2</sup>



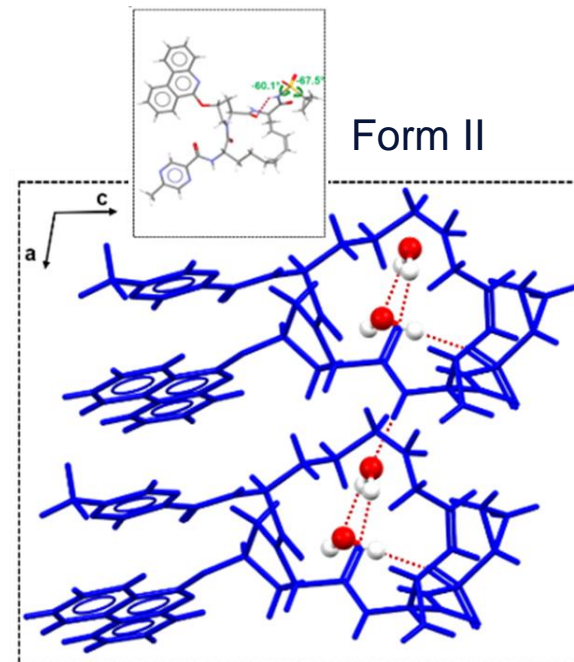
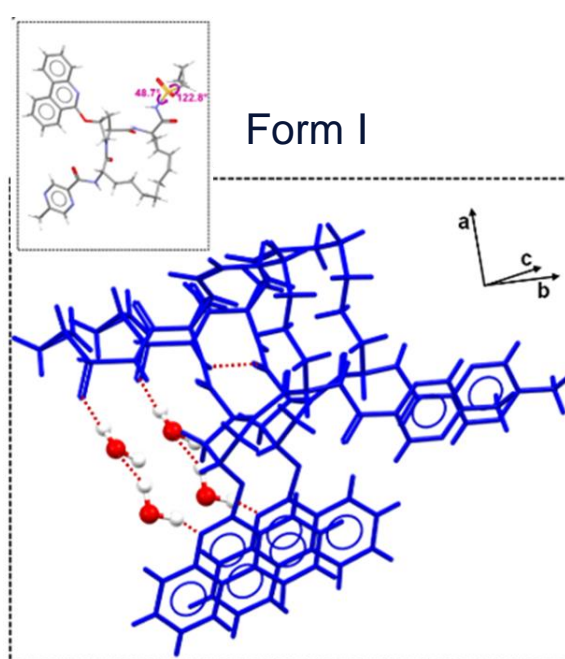
Whitty, et.al, DDT, 21, 2016

\*open conformations ensemble account for 52% compared to 10% in gas phase

abbvie J. Am. Chem. Soc. 2021, 143, 42, 17479–17491

# Form I and II- Conformations and interactions

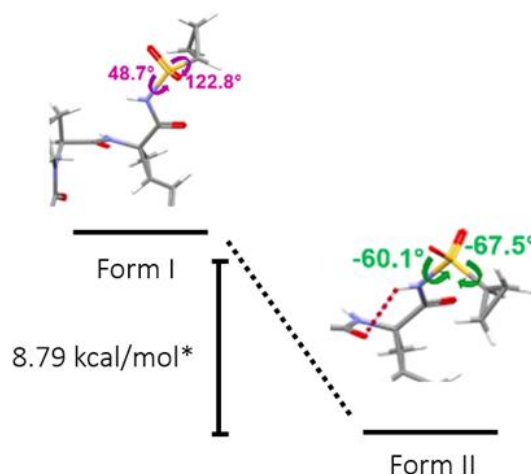
Form I- No IMHB. Two intermolecular Hydrogen bonded chain motifs, leading to H-bonded planes parallel to *bc* plane. Weak  $\pi$ - $\pi$  interactions along *c* axis between Interdigitated phenanthridine rings



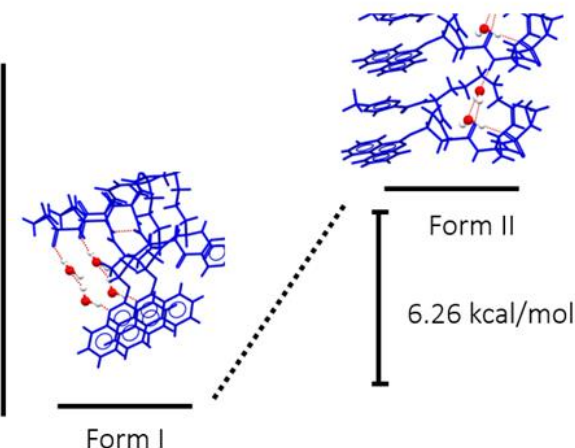
Form II- IMHB between the carbonyl oxygen and the sulfonamide N-H. Stacking related layers parallel to the *bc* plane related, stabilized by weak  $O\cdots H-C$  intermolecular interactions\*

\* May result in offset between layers, higher void volume and lower long-range periodicity

Intramolecular Energy



Intermolecular Energy

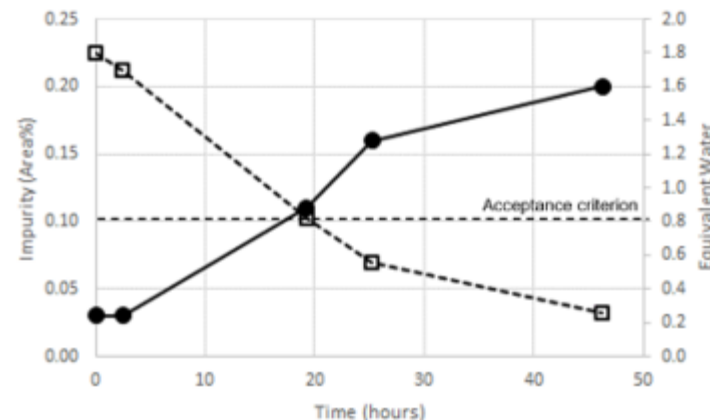
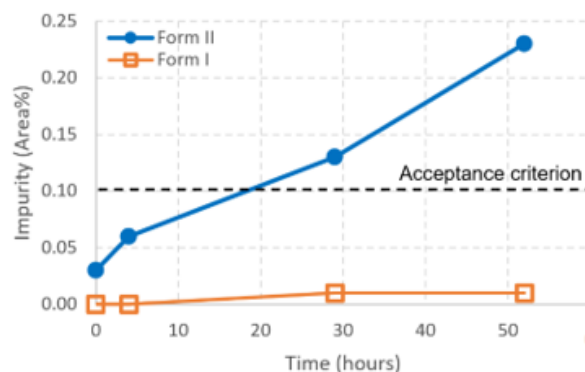
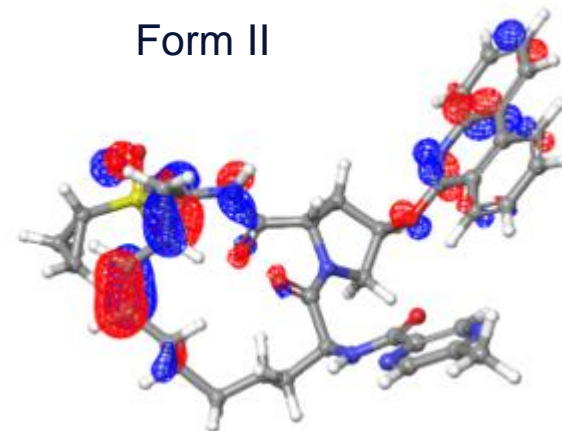
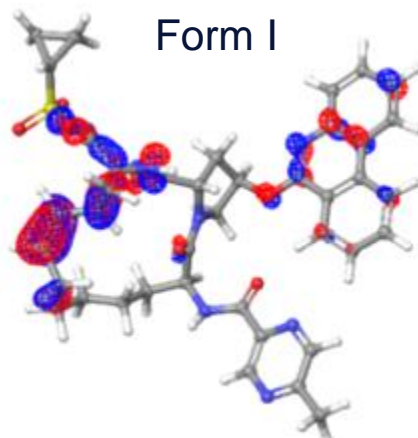




# Form I and II- Differing solid state chemical reactivity

In Form I conformation, HOMO distribution spreads over double bond of macrocycle ring and not delocalized on the sulfonamide group. Distance between the centers of mass of the cyclopropyl-sulfonamide moiety and the olefin group is 1.3 Å larger than for Form II

In Form II conformation, HOMO distribution mainly delocalized on double bond of macrocycle ring and sulfonamide group. Electron density map indicates cyclopropyl-sulfonamide moiety and olefin groups can react to form the observed oxidative impurity. Removal of water adversely affects the reaction



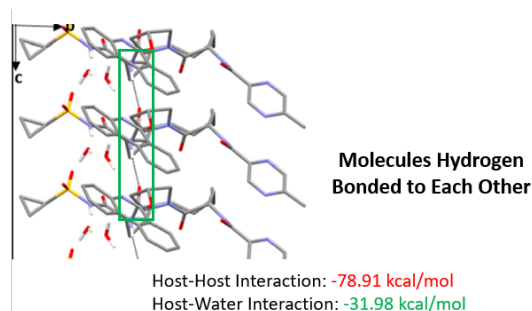
# Dehydration

Form I exhibits strong host-host intermolecular interactions due to its lack of IMHB while Form II exhibits weak host-host interactions but strong host-water interactions

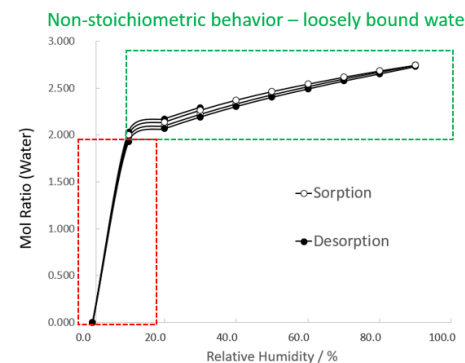
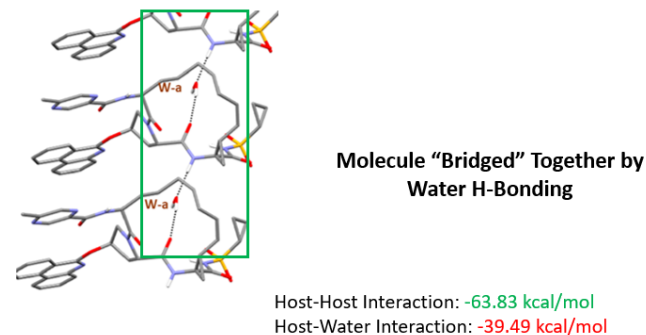
Due to weaker host-host interactions, Form II more amenable for hot melt extrusion processing

Form II exhibits hybrid stoichiometric and non-stoichiometric behavior with an enthalpically favored “stoichiometric regime and entropically favored non-stoichiometric regime consisting of ensembles of water bonding networks

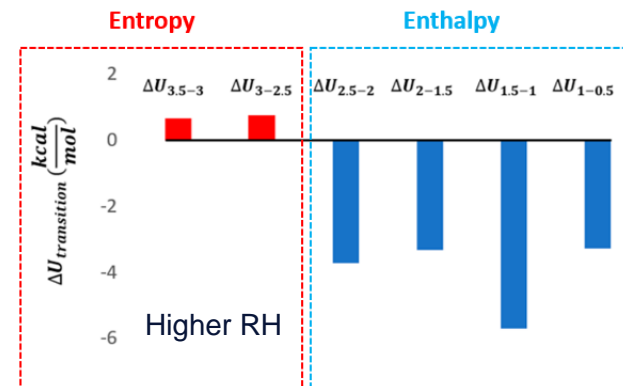
## Form I



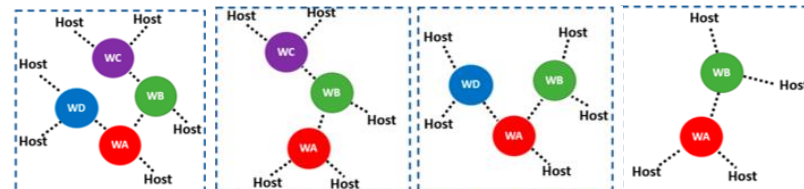
## Form II



Stoichiometric behavior – tightly bound water Di-Hydrate



## Lower RH



# Concluding Summary



## Reducing or Managing complexity



Screen out developability challenges to the extent possible



Some targets can only be engaged with complex matter



Early identification of the liabilities



Broader toolkit

Co-processed  
API



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