

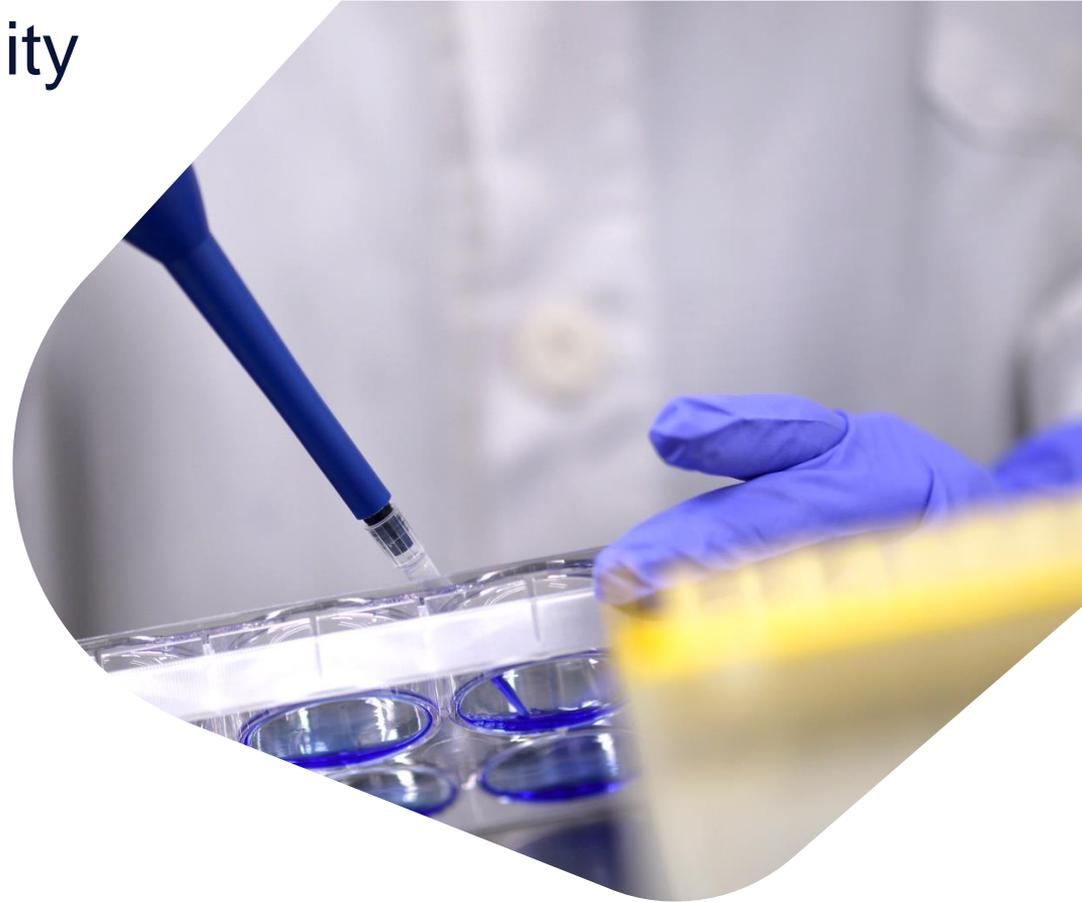
Emerging Modalities and Compound Developability Assessment in Small Molecule Early Development

MCERSI Co-Processed API and
Regulatory Requirements

Ahmad Sheikh

July 13, 2022

abbvie



VPAK-US-00002-E

Acknowledgements

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Last two decades

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

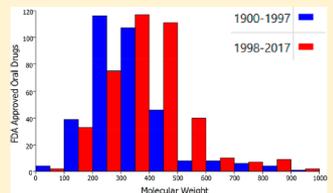
Miniperspective

Michael D. Shultz[#] 

Global Discovery Chemistry, Novartis Institutes for Biomedical Research, Inc., 181 Massachusetts Avenue, Cambridge, Massachusetts 02139, United States

Supporting Information

ABSTRACT: Two decades have passed since the rule of five ushered in the concept of “drug-like” properties. Attempts to quantify, correlate, and categorize molecules based on Ro5 parameters evolved into the introduction of efficiency metrics with far reaching consequences in decision making by industry leaders and scientists seeking to discover new medicines. Examination of oral drug parameters approved before and after the original Ro5 analysis demonstrates that some parameters such as clogP and HBD remained constant while the cutoffs for parameters such as molecular weight and HBA have increased substantially over the past 20 years. The time dependent increase in the molecular weight of oral drugs during the past 20 years provides compelling evidence to disprove the hypothesis that molecular weight is a “drug-like” property. This analysis does not validate parameters that have not changed as being “drug-like” but instead calls into question the entire hypothesis that “drug-like” properties exist.



J. Med. Chem. 2019, 62, 4, 1701–1714

Table 2. Analysis of FDA Approved Oral NCEs from 1998 to 2007^a

	clogP	MW	HBD	HBA	TPSA	RotB	Fsp ³	#ArRNG
1997 90 th percentile	4.7	470.3	4.0	10	139.8	10.0	0.83	3
90 th 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 [*]	0.3	0.0026	>0.99	0.69	0.97	0.039	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 [†]	0.35	0.28	0.94	0.55	0.94	0.066	0.56	0.025
p value [‡]	0.46	0.014	>0.99	0.044	0.43	0.0085	0.73	0.73
10 th 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 [*]	0.6	<0.0001	>0.99	>0.99	0.37	>0.99	<0.0001	>0.99

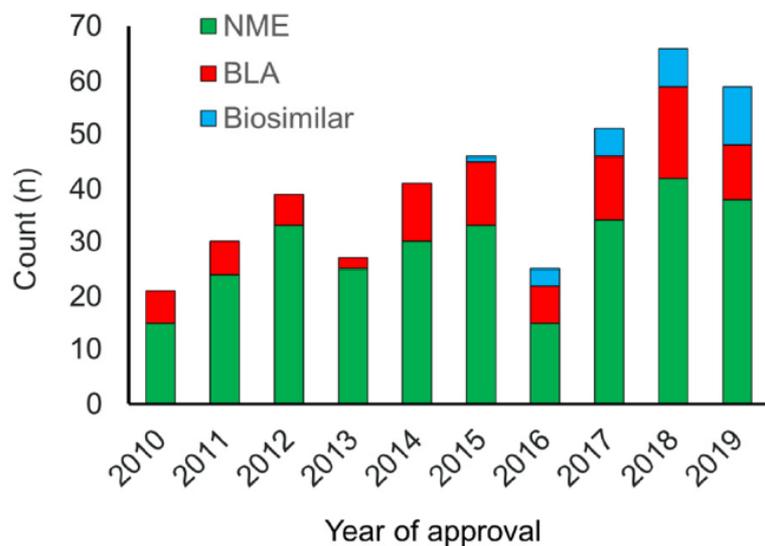
^a*n* = 195, or 26% of all FDA approved oral NCEs. The 90th percentile values determined for FDA approved oral drugs from 1900 to 1997 are included for reference. The change in values from FDA approved oral drugs from 1900 to 1997 values are in parentheses. ^{*}*p* value of the 10th and 90th percentile cutoffs based on one-way ANOVA analyses. [†]*p* values from one-way ANOVA (normal distribution). [‡]*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^a

	clogP	MW	HBD	HBA	TPSA	RotB	Fsp ³	#ArRNG
1997 90 th percentile	4.7	470.3	4.0	10	139.8	10.0	0.83	3
90 th 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 [*]	0.27	<0.0001	>0.99	<0.0001	0.20	<0.0001	<0.0001	<0.0001
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 [†]	<0.0001	<0.0001	0.57	0.0008	0.037	<0.0001	0.65	<0.0001
p value [‡]	<0.0001	<0.0001	0.35	<0.0001	0.005	<0.0001	0.99	>0.99
10 th 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 [*]	<0.0001	<0.0001	>0.99	0.024	0.0049	<0.0001	0.0005	>0.99

^a*n* = 214, or 29% of all FDA approved oral NCEs. The 90th 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. ^{*}*p* value of the 10th and 90th percentile cutoffs based on one-way ANOVA analyses. [†]*p* values from one-way ANOVA (normal distribution). [‡]*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.

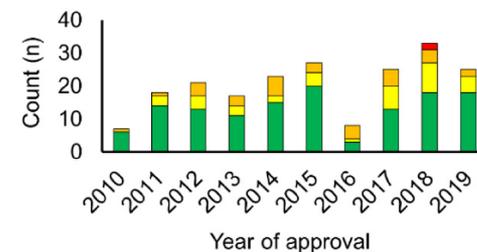
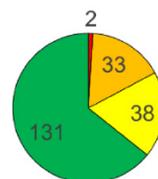
Last decade



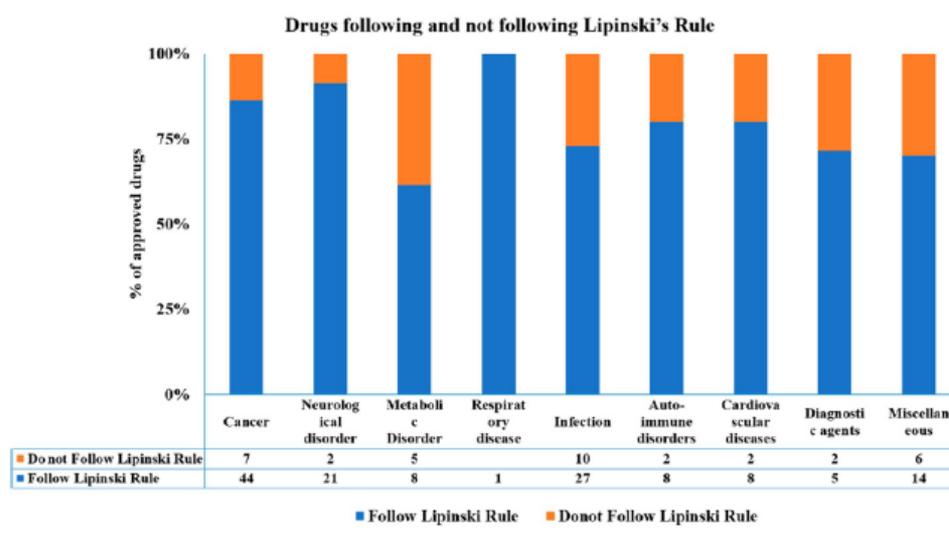
J. Med. Chem. 2021, 64, 5, 2312–2338

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 ₅₀ /D _{max}	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		MZF54	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

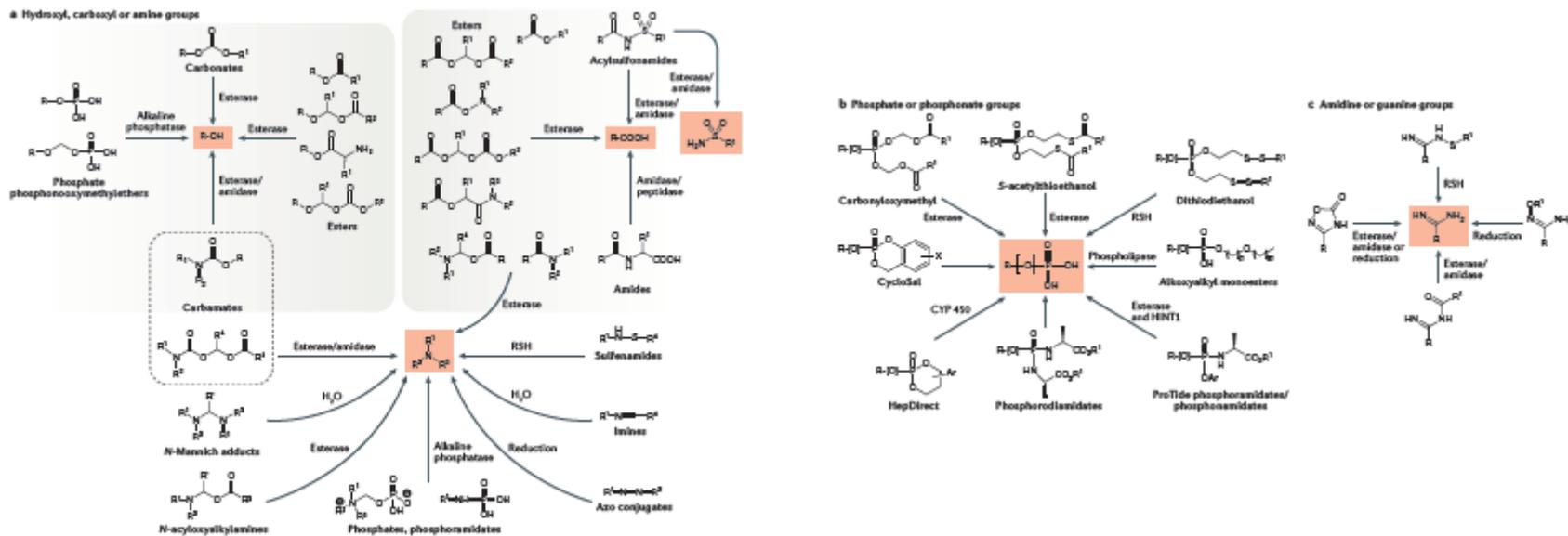
Table 2
In silico metrics for selected PROTACs.

Compound	E3 Ligase	MW	cLogP	HBD	HBA	PSA	nRotB	N _{rule-of-5}	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

Biorganic & Medicinal Chemistry Letters 29 (2019) 1555–1564

Prodrugs- chemically labile by design

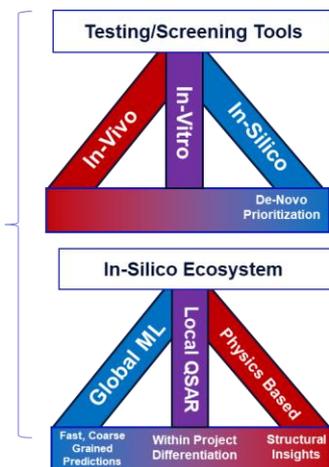
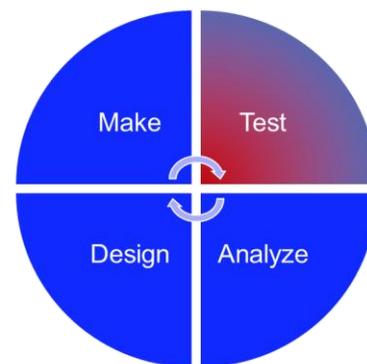
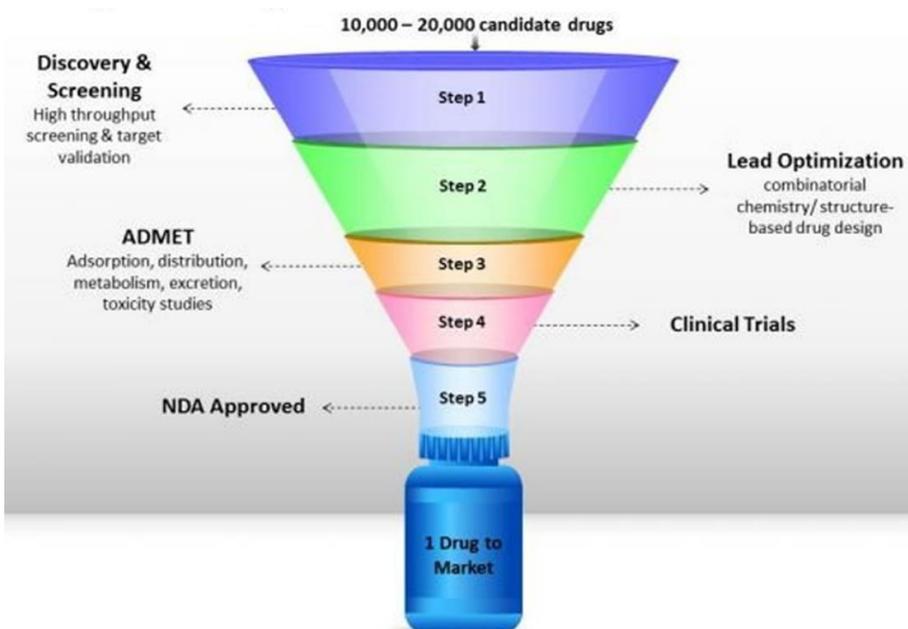
- Over 30 approved in the last decade ('08-17), representing ~ 12% of all approved small molecules



- 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

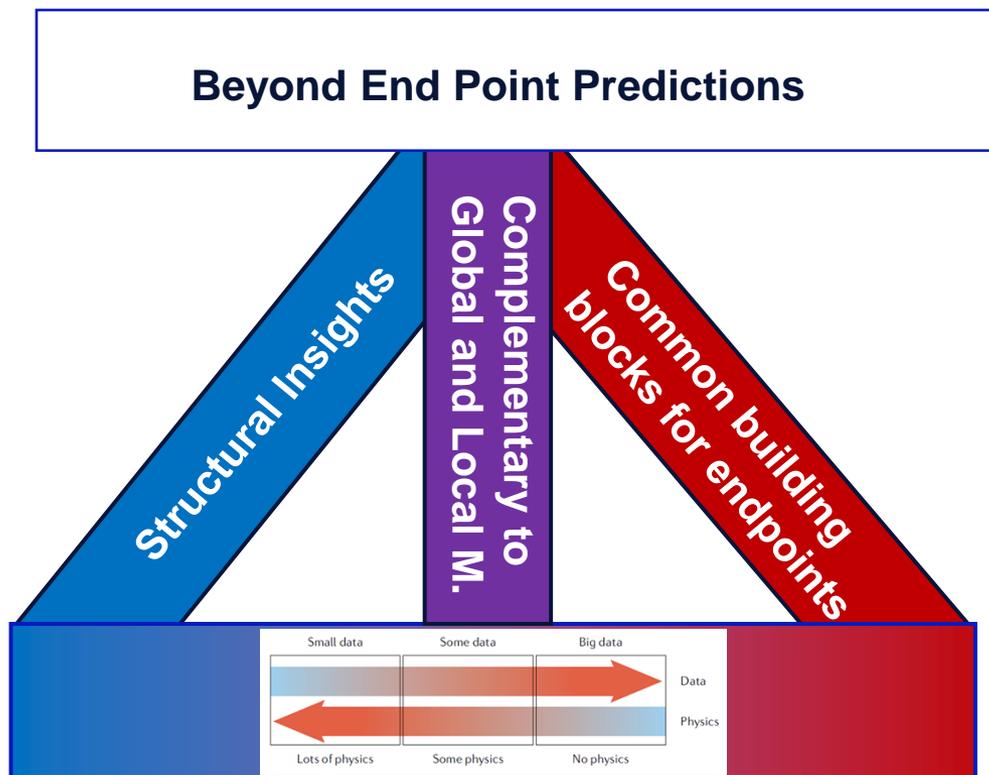
Positioning of Developability earlier in the LO Funnel



Absorption	Distribution	Metabolism	Excretion	Toxicity	PhysChem
Caco-2 permeability	Human serum albumin binding	Microsomal stability	CL _{int}	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)	

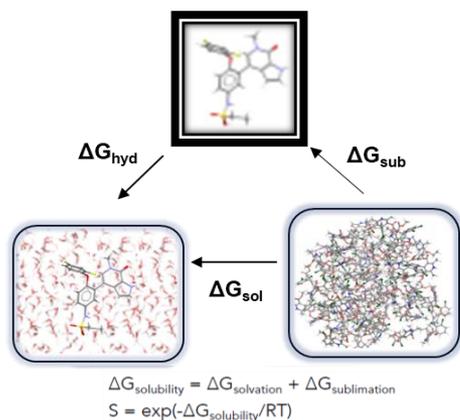
Drug Disc Today, Volume 27, Issue 4, 2022, 967-984

Physics Based Models

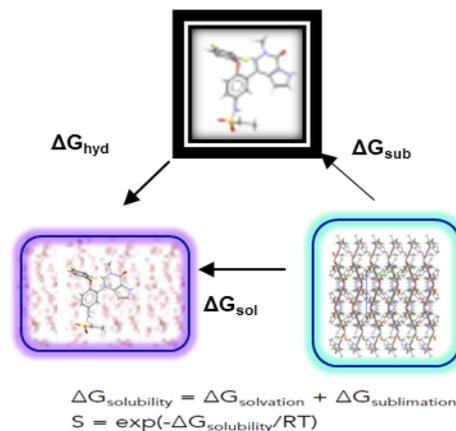
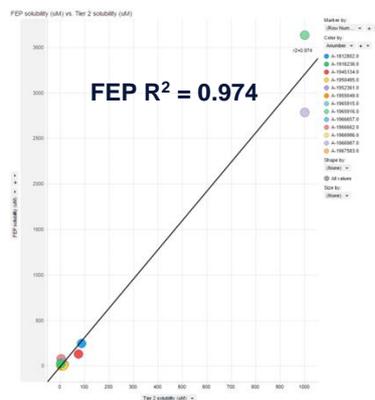


Nat Rev Phys 3, 422–440 (2021)

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

JCIM 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, Guangru Sun, Sirzhu Li, Lingle Wang, Sayan Mondal, Jianguo Ji, and Thomas B. Borchardt

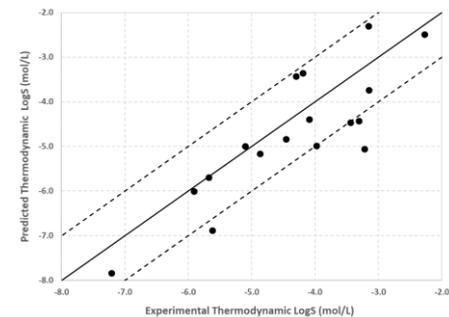
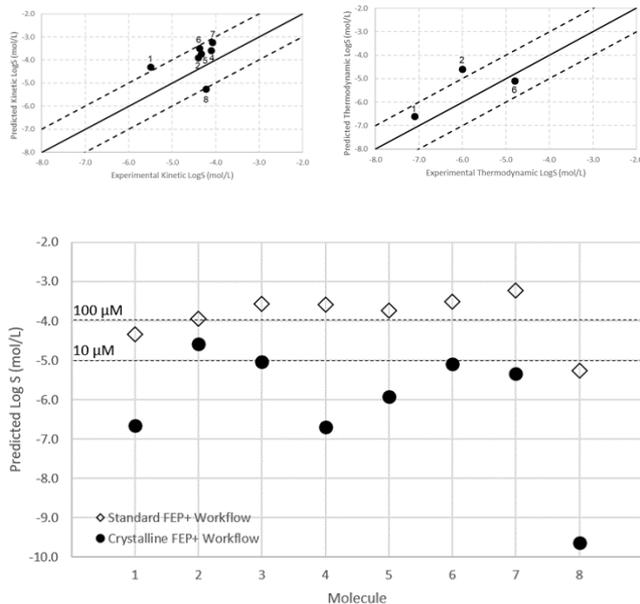
Cite This: *J. Chem. Inf. Model.* 2021, 61, 1412–1426

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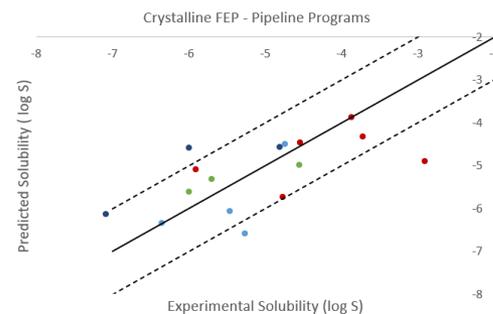
ACCESS | Metrics & More | Article Recommendations | Supporting Information



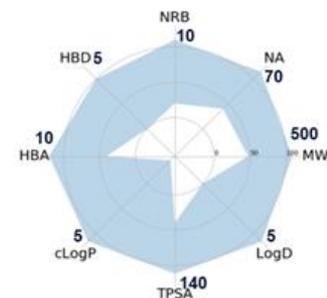
ABSTRACT: Drug design with patient centrality for ease of administration and pill burden requires robust understanding of the impact of chemical modifications on relevant physicochemical properties early in lead optimization. To this end, we have developed a physics-based ensemble approach to predict aqueous thermodynamic crystalline solubility, with a 2D chemical structure as the input. Predictions for the bromodomain and extraterminal domain (BET) 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)



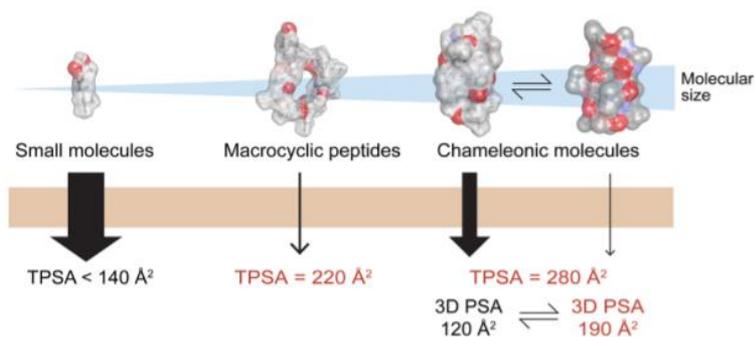
J. Chem. Inf. Model. 2021, 61, 3, 1412–1426

Incorporation of Physical Properties considerations in the design cycle

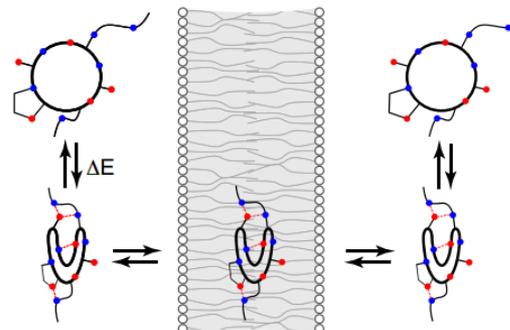
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Company Confidential © 2021

bR05 Chameleonicity: Towards Improved Passive Permeability

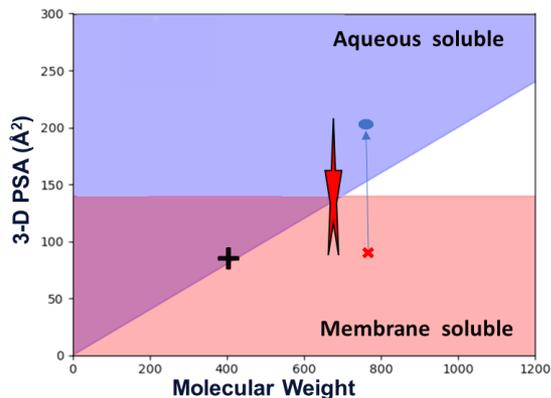


Size boundaries for membrane-permeable molecules



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

A Framework to Quantify Chameleonic Behavior



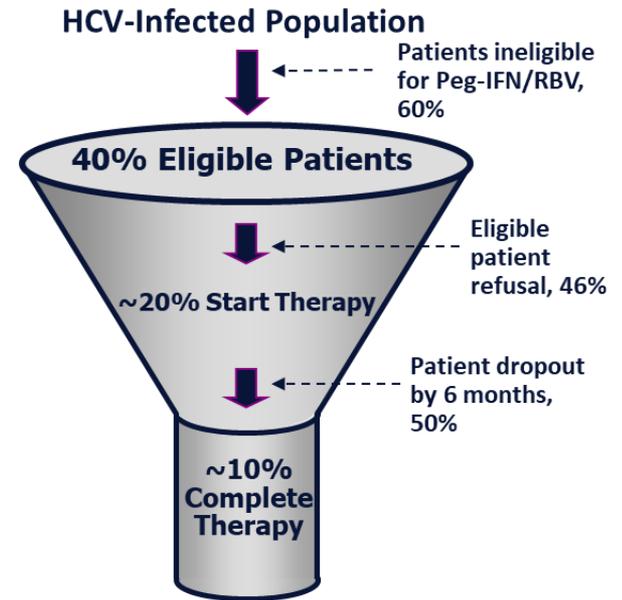
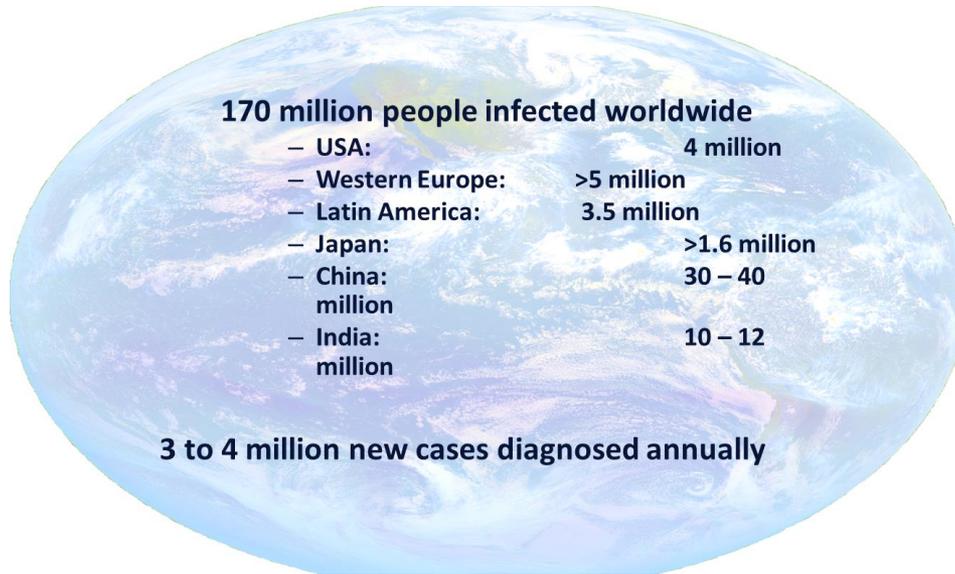
- + mean value reported for 1193 drugs
- ★ MW ≥ 700, no single PSA value can satisfy both conditions

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

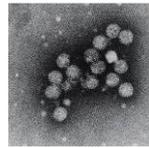
Hepatitis C Infection- where were we a decade ago



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%)

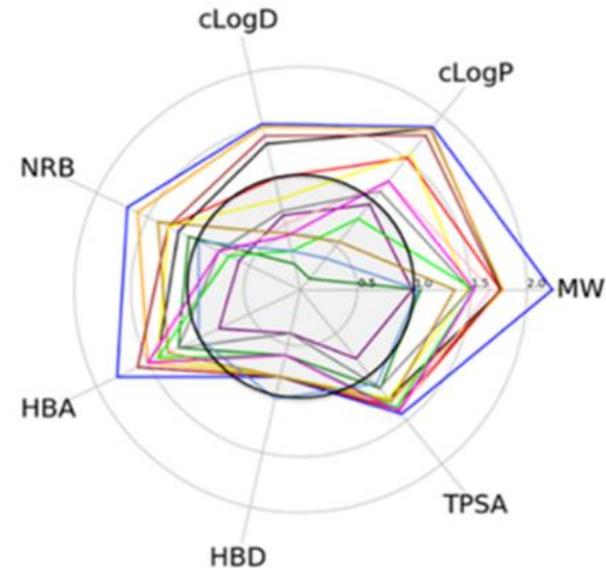
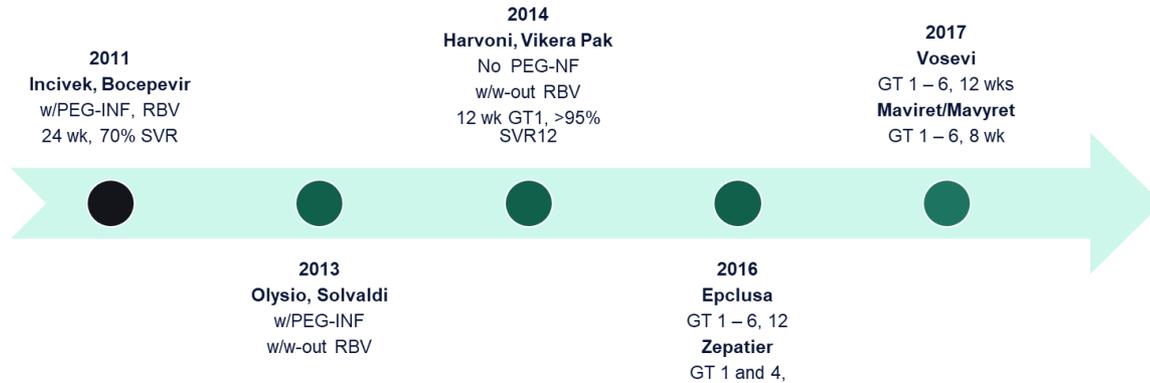
and Since..

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



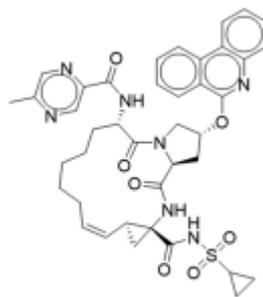
Paritaprevir- the molecule

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

Approved in 2014 as part of 1st 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



Paritaprevir

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

Read Online

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*

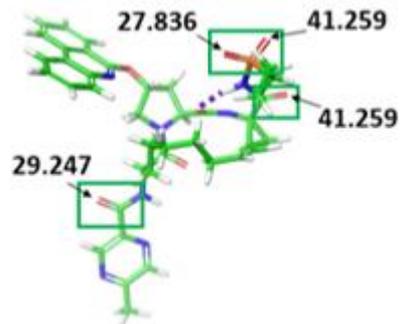
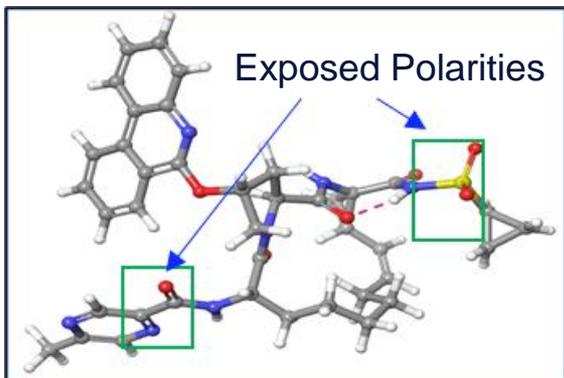
Rajni Miglani Bhardwaj, Raimundo Ho, Yue Gui, Paul Brackemeyer, Gabriela Schneider-Rauber, Fredrik L. Nordstrom, and Ahmad Y. Sheikh*

Cite This: <https://doi.org/10.1021/acs.cgd.2c00264>

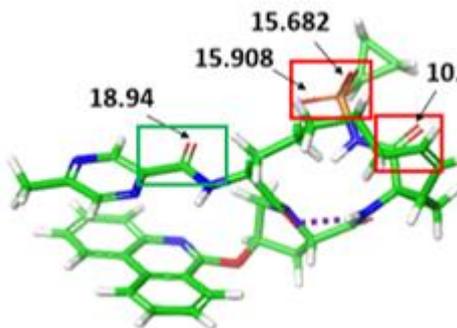
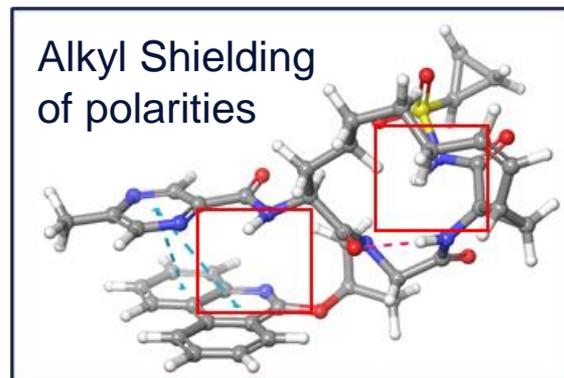
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Computational Assessment of Chameleonicity

Water Conformation*
3D PSA: 201.2 Å²



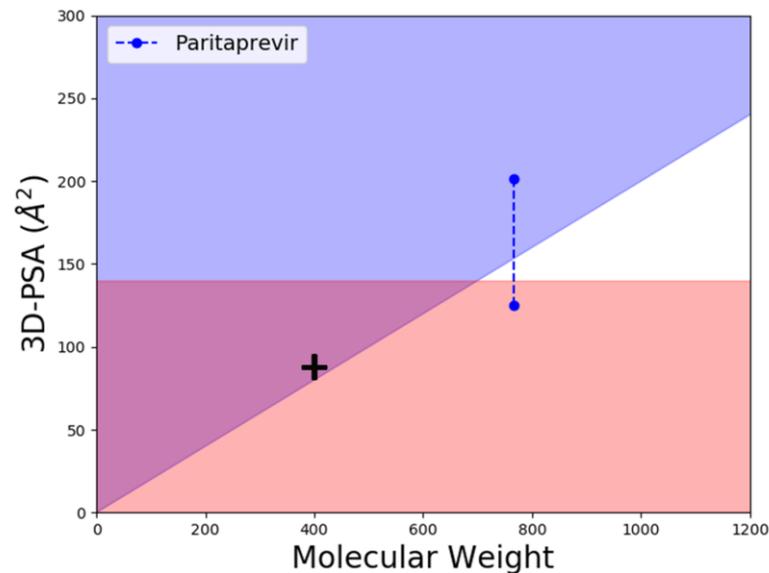
Chloroform Conformation
3D PSA: 125.1 Å²



Measured logD^{1-octano/pH 6.8} 3.1

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

EPSA 132 Å²



+ mean value reported for 1193 drugs

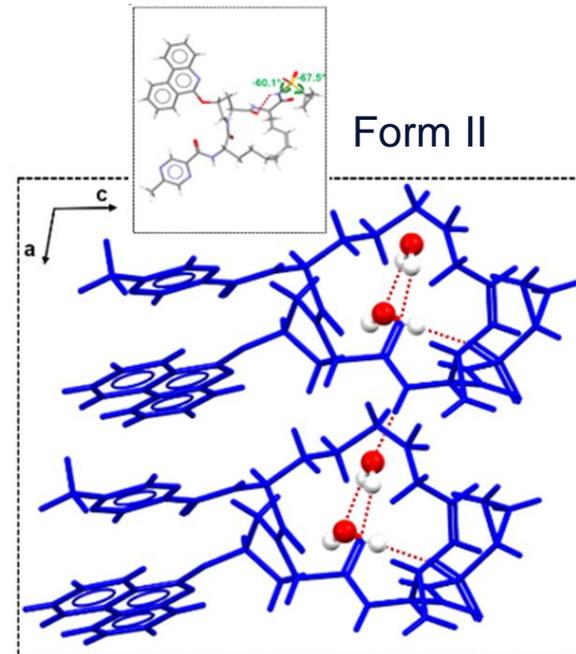
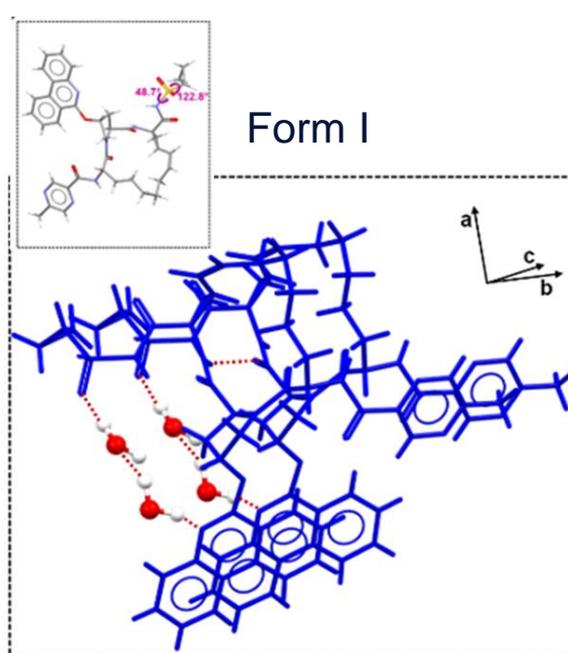
Whitty, et.al, DDT, 21, 2016

*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

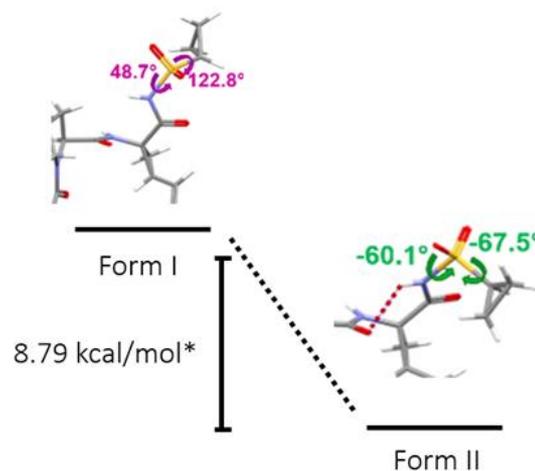
Form I- No IMHB. Two intermolecular Hydrogen bonded chain motifs, leading to H-bonded planes parallel to *bc* plane. Weak π - π interactions along *c* axis between Interdigitated phenanthridine rings



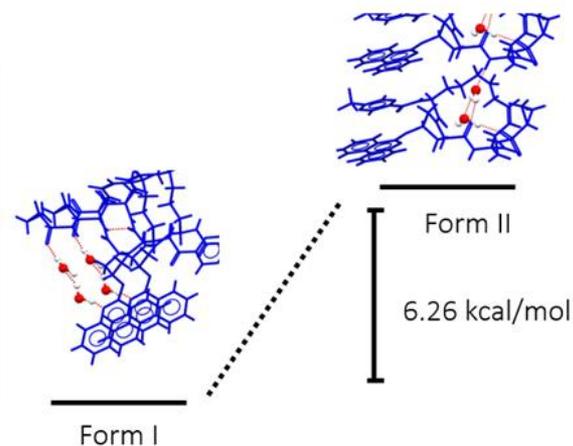
Form II- IMHB between the carbonyl oxygen and the sulfonamide N-H. Stacking related layers parallel to the *bc* plane related, stabilized by weak $O\cdots H-C$ intermolecular interactions*

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

Intramolecular Energy



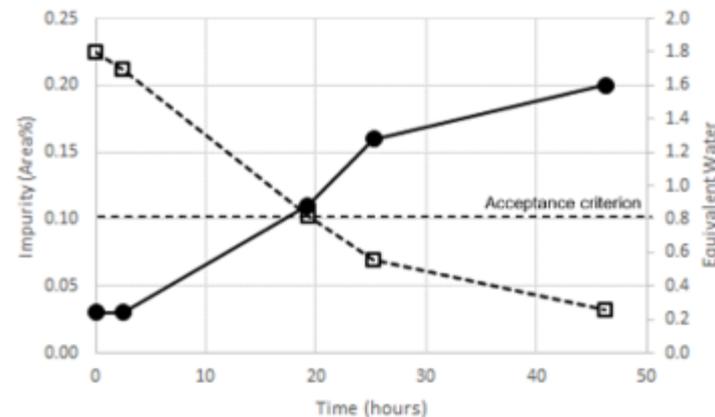
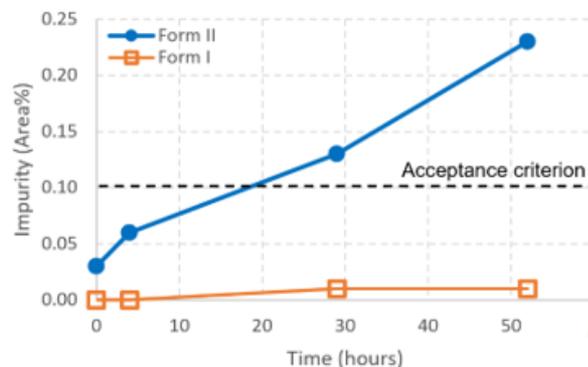
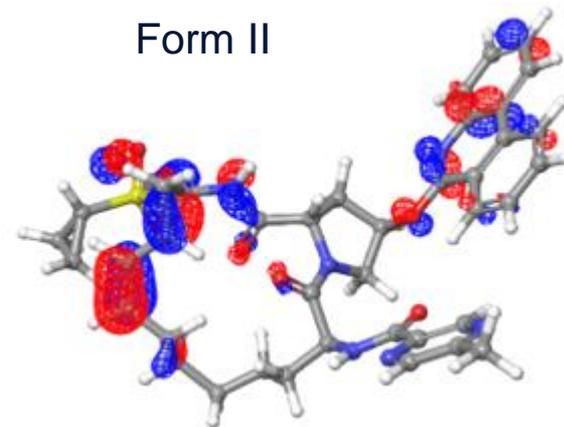
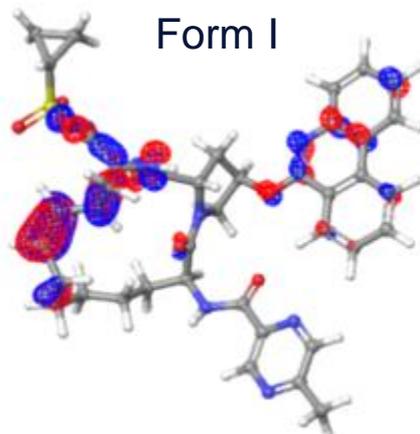
Intermolecular Energy



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



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

abbvie