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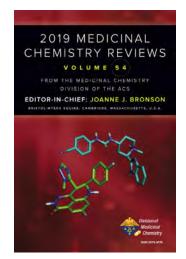
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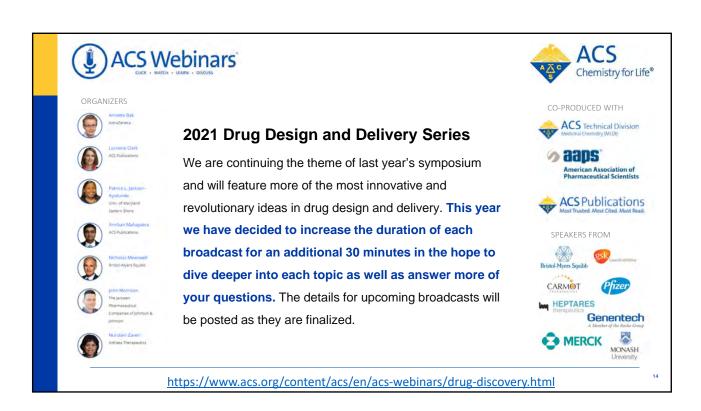




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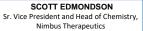






How Computational Chemistry is Accelerating Drug Discovery







Vice President, Bristol-Myers Squibb

Presentation slides available now! The edited recording will be made available as soon as possible.

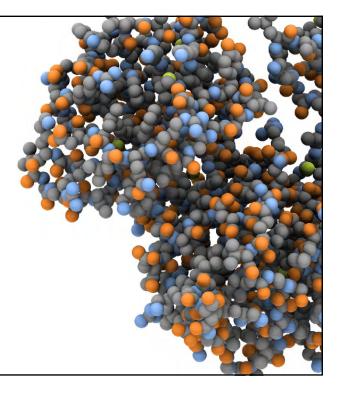
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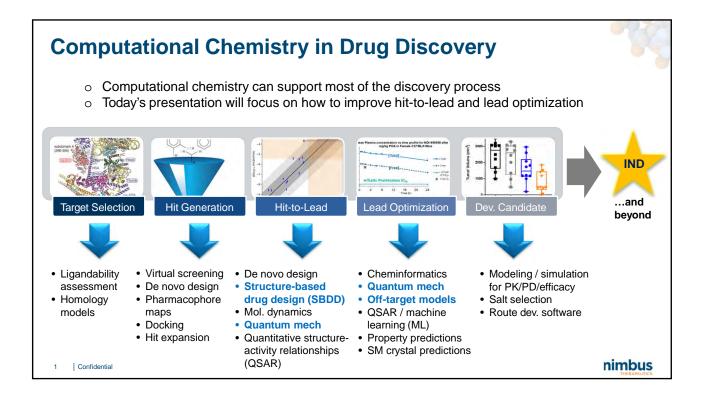
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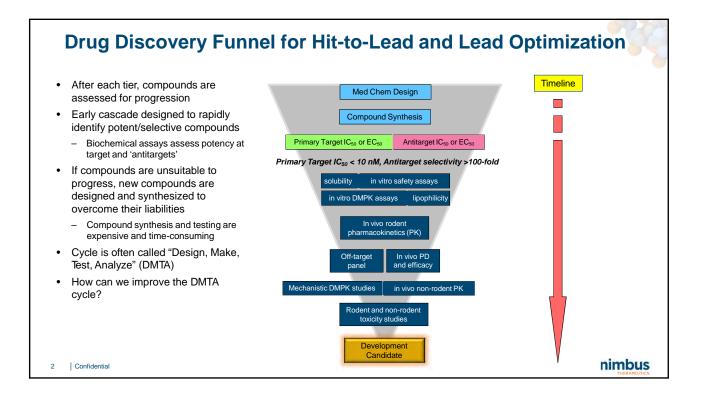
How Computational Chemistry with Structural Biology is Enabling Drug Discovery

Scott Edmondson, Nimbus Therapeutics ACS Webinar Series, 24-June-2021



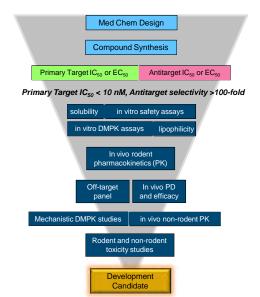
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How can we improve the DMTA Cycle?

- After each tier, compounds are assessed for progression
- Early cascade designed to rapidly identify potent/selective compounds
 - Biochemical assays assess potency at target and 'antitargets'
- If compounds are unsuitable to progress, new compounds are designed and synthesized to overcome their liabilities
 - Compound synthesis and testing are expensive and time-consuming
- Cycle is often called "Design, Make, Test, Analyze" (DMTA)
- How can we improve the DMTA cycle?



- A. Assess compound efficacy in vivo in parallel to primary potency
- B. Synthesize more compounds
- C. Use computational methods to improve designs
- D. Skip the assay cascade and only make the development candidate

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How can we improve the DMTA Cycle?

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- How can we improve the DMTA cycle?
 - Better designs that leverage target and antitarget potency predictions

SBDD / Computational Methods – Evaluate Potency/Selectivity

Compound Synthesis

Primary Target IC₅₀ or EC₅₀

Antitarget selectivity >100-fold

solubility in vitro safety assays
in vitro DMPK assays lipophilicity

In vivo rodent pharmacokinetics (PK)

Off-target panel In vivo PD and efficacy

Mechanistic DMPK studies in vivo non-rodent PK

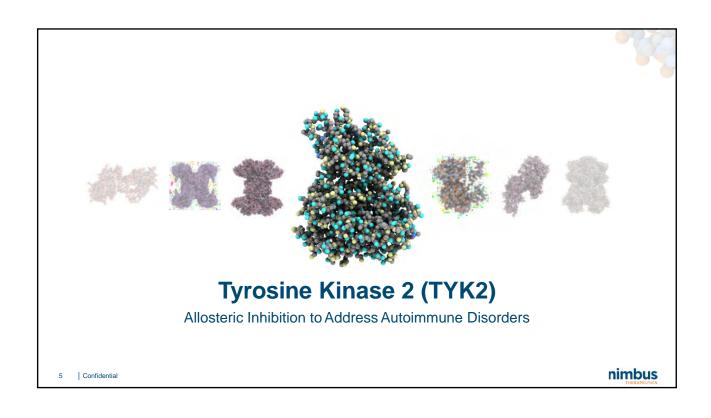
Rodent and non-rodent toxicity studies

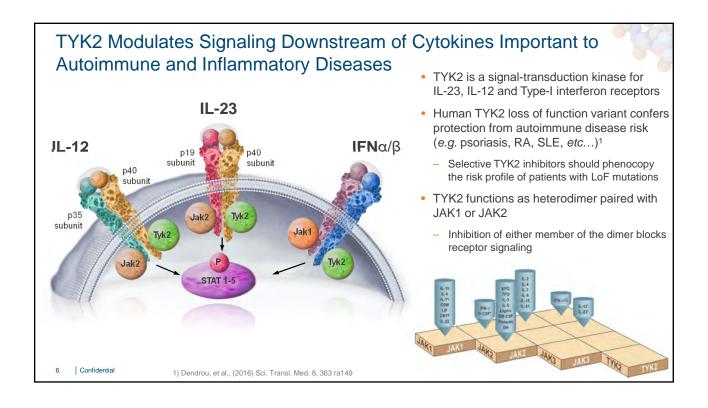
Development

- A. Assess compound efficacy in vivo in parallel to primary potency
- B. Synthesize more compounds
- C. Use computational methods to improve designs
- D. Skip the assay cascade and only make the development candidate

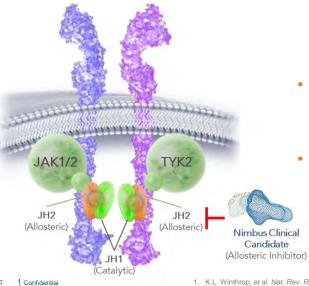
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Targeting TYK2 May be Safer than JAK Inhibitors... but Challenging to **Achieve Selectivity**



- · JAK inhibitor medicines such as tofacitinib carry black box warnings
 - Enhanced risk of venous thromboembolism (VTE), serious infections, malignancy, cytopenias, lipid
- High binding site homology between orthosteric (catalytic) binding sites of TYK2 and JAK1/2/3 kinases
 - Important to achieve high selectivity vs JAK family
- Allosteric TYK2 inhibition at the JH2 site may achieve high selectivity vs other JAK members

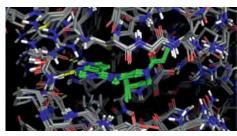
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1. K.L. Winthrop, et al. Nat. Rev. Rheumatol. 2017; 13; 234-243.

Poor Prospects of Achieving Selectivity vs JAKs in Catalytic Binding Pocket

Biochemical IC₅₀s (nM) of Inhibitors of the Catalytic (JH1) Sites of TYK2 and JAK1-31

Inhibitor	TYK2 catalytic domain	JAK1 catalytic domain	JAK2 catalytic domain	JAK3 catalytic domain
Tofacitinib	489	15	77	55
Baricitinib	61	4	7	787
Filgotinib	2,600	363	2,400	>10,000
Upadacitinib	4,690	47	120	2,304
PF-06700841	23	17	77	6,494
PF-06826647	17	383	74	>10,000



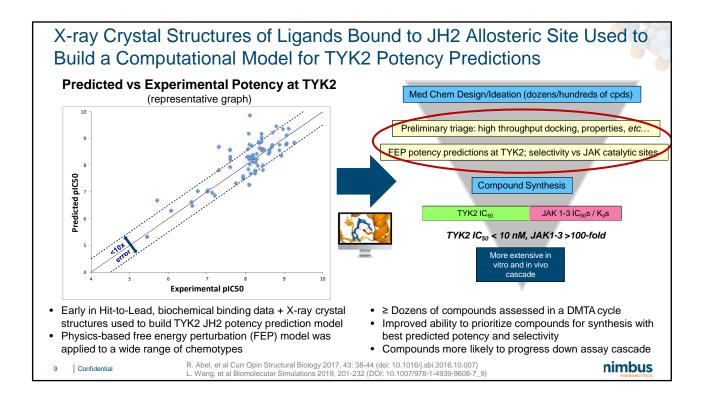
Crystal structures of TYK2, JAK1, JAK2, and JAK3 with tofacitinib in catalytic JH1 site2 Very Similar Binding Pockets

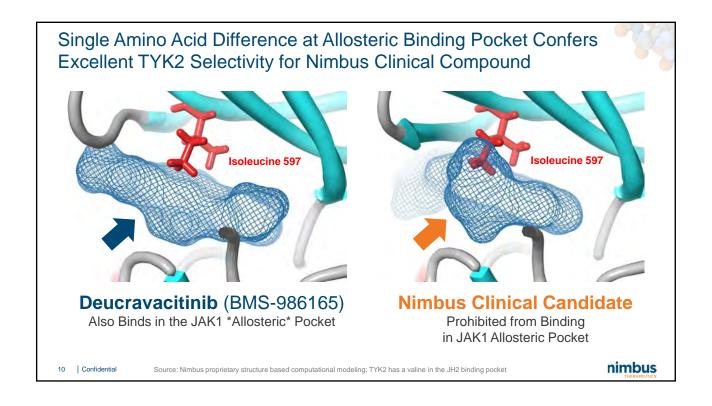
In contrast to the catalytic site inhibitors, allosteric JH2 inhibitors have been described with excellent biochemical selectivity vs JAK catalytic sites (e.g. BMS-986165 = deucravacitinib, >10,000-fold selective vs JAK1/2/3 catalytic domains)2,3

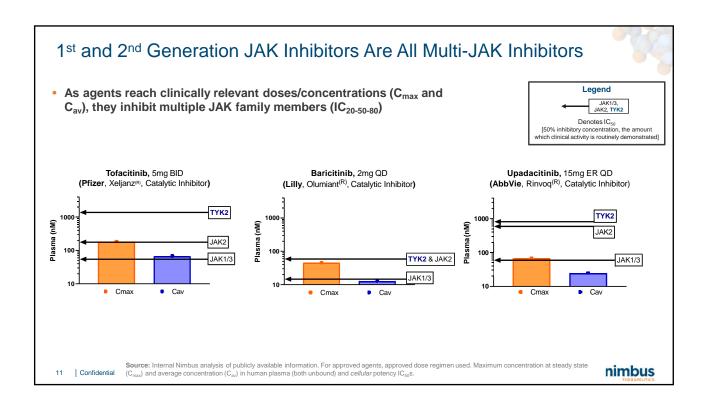
- Deucravacitinib exhibits 17-fold selectivity over JAK1 JH2 site: TYK2 JH2 K_i = 0.02 nM; JAK1 JH2 K_d = 0.33 nM³
 - S.T. Wrobleski, et al. J. Med. Chem. 2019; 62; 8973-8995
 - R. Abel, et al Curr Opin Structural Biology 2017, 43: 38-44.
 J.R. Burke, et al. Sci. Transl. Med. 2019; 11; eaaw1736.

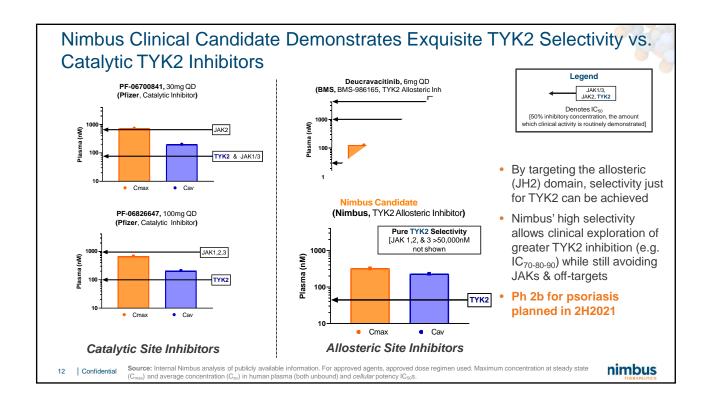
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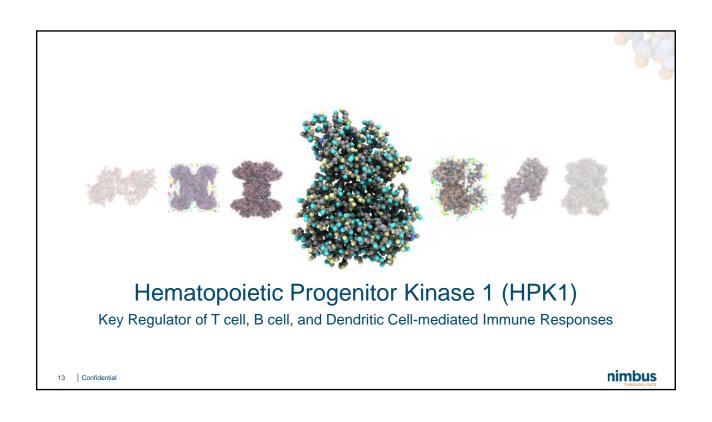
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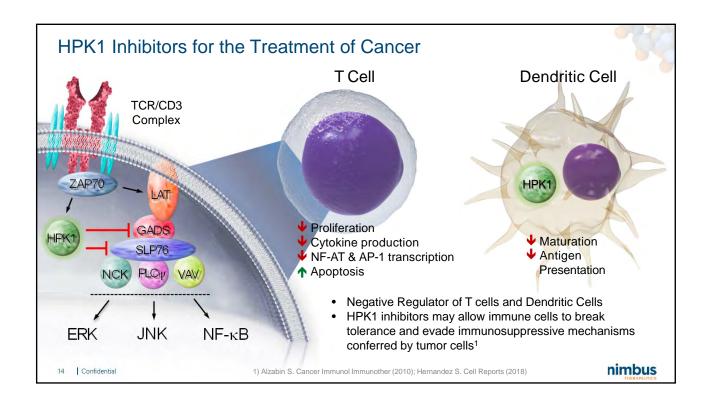








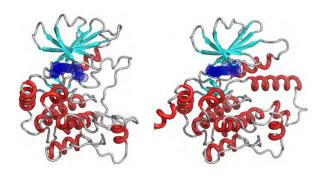




HPK1 Inhibitor Designs Enabled by Advances in Crystallography



GLK Co-Crystal



- Proprietary crystal structures of HPK1 and other MAP4K family members such as GLK
 - Co-crystals of the off-targets applied to design out undesired activities
- Protein/ligand structures guided SBDD and FEP+ to improve HPK1 potency and selectivity
- Further synthesis and optimization yielded novel ligands
- Improved biochemical specificity resulted in robust immune activation responses

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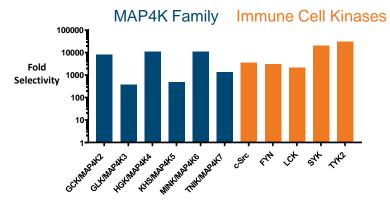
NMBS-2 is a Potent HPK1 Inhibitor with Excellent Selectivity Against MAP4K Family Members and Immune Cell Kinases

Assay NMBS-2

HPK1 Caliper IC₅₀ <1 nM

@ 1mM ATP <1 nM

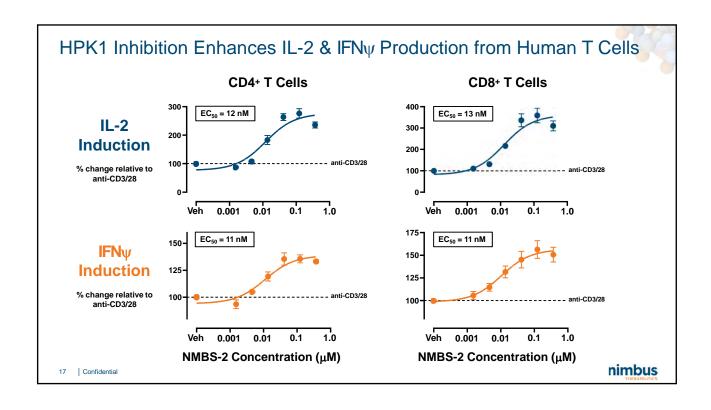
pSLP-76 Cell IC₅₀ 42 nM

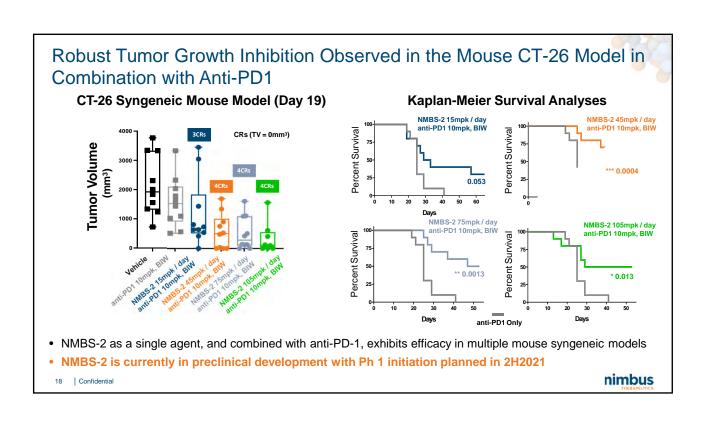


- Assessed against broad panel of >300 kinases; highly selective for HPK1 only
- · High selectivity required for robust immune cell activation

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Werner Syndrome Helicase (WRN)



Casitas B lymphoma-B E3 Ligase (Cbl-b)



CTP Synthase 1 (CTPS1)

Selectively Targeting a Synthetic Lethal Dependency of Microsatellite Instable Tumors

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Multiple Promising Targets For Nimbus' SBDD Approach



Werner Syndrome Helicase (WRN)



Casitas B lymphoma-B E3 ligase (Cbl-b)

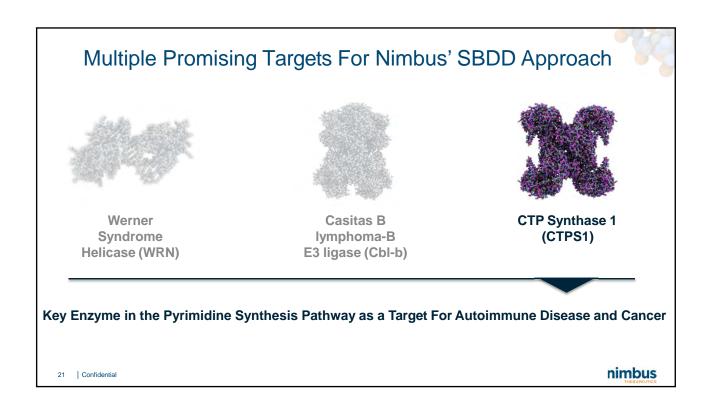


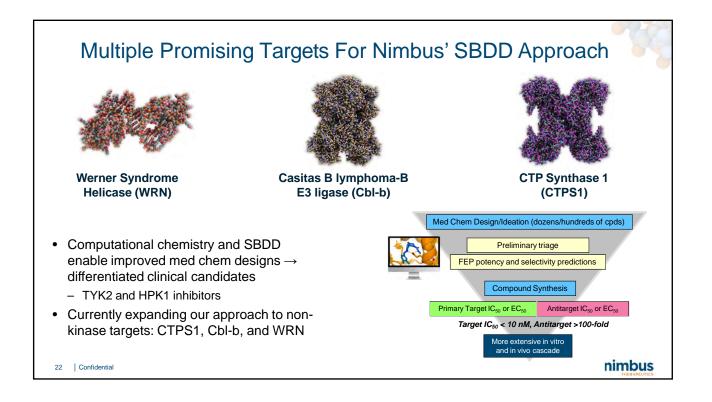
CTP Synthase 1 (CTPS1)

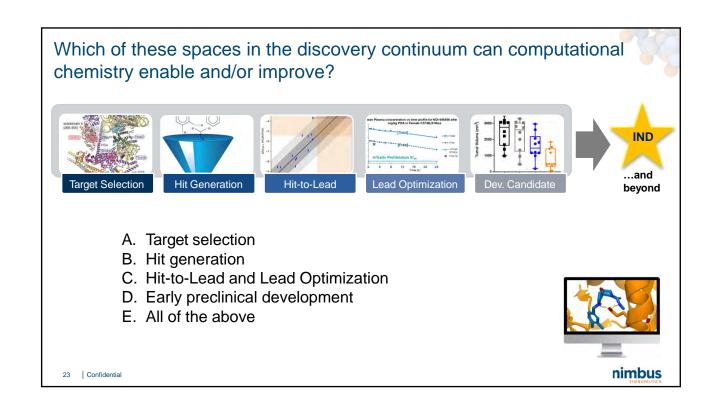
A Negative Regulator of Anti-tumor Immune Responses as a Target for Immuno-oncology

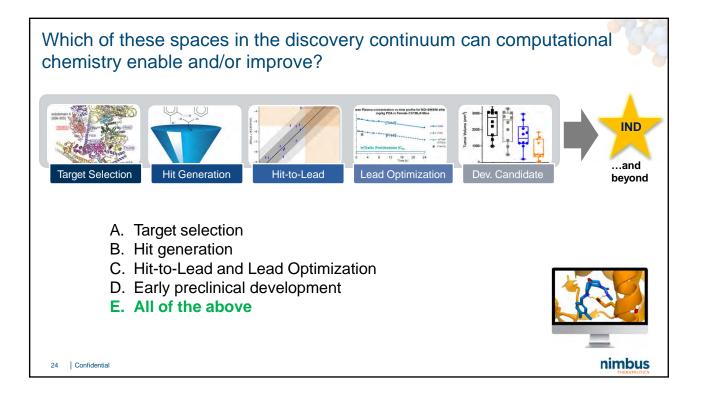
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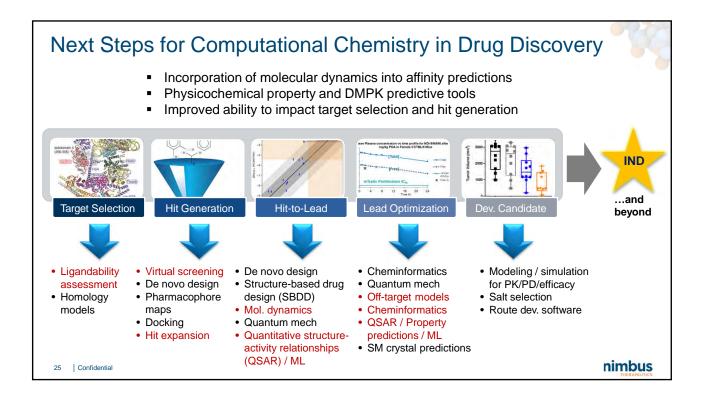
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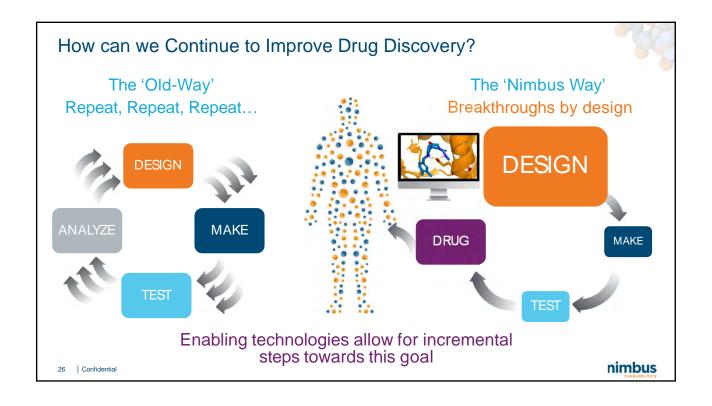






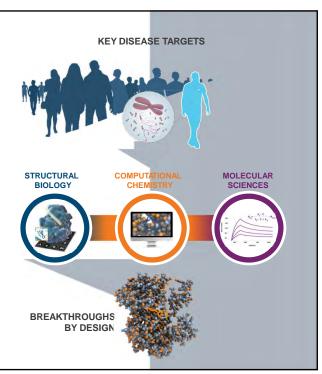






We Design Breakthrough Medicines

- Medicines are our mission
 - Targets with highly validated disease roles
 - Couple structure-based expertise with cutting-edge computational tools
 - Progress key programs into clinical development
- Track record of success propelling us forward
 - Multiple programs to the clinic:
 - · ACC inhibitor for NASH
 - TYK2 inhibitor for psoriasis
 - Discovery engine has produced a pipeline of desired, difficult to drug targets, progressing to the clinic, including HPK1
 - Next generation of targets already on the horizon



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TYK2 and HPK1 Acknowledgements



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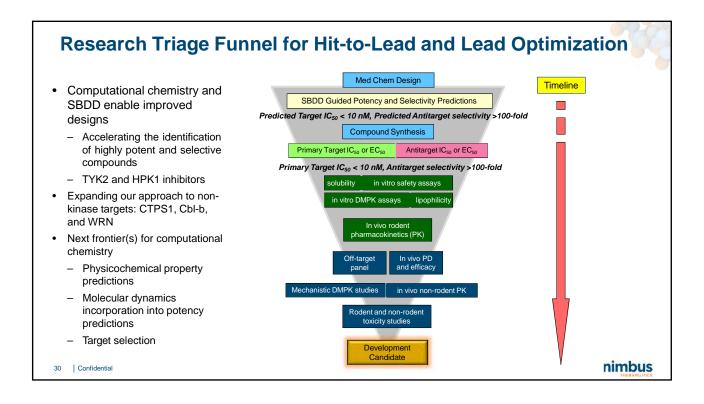






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How Computational Chemistry is Accelerating Drug Discovery





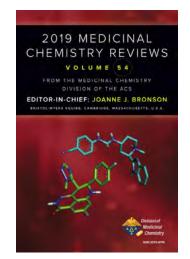
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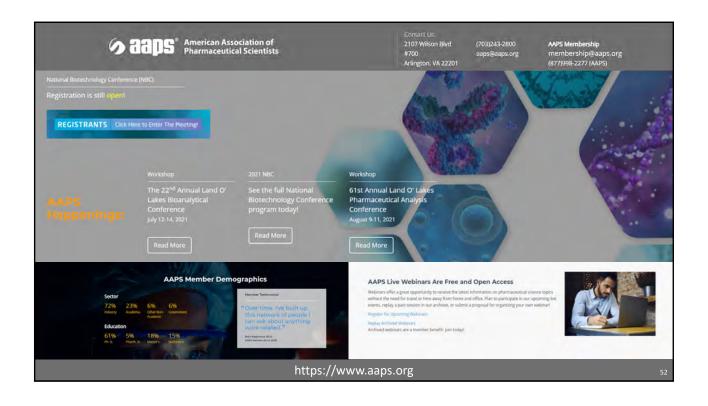




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Chemistry for Life®

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ACS President H.N. Cheng Presents: FRIBAYS

Date: Friday, June 25, 2021 @ 2-3:30pm ET Speaker: Sir Fraser Stoddart, 2016 Nobel Laureate in Chemistry, Board of Trustees Professor of Chemistry, Northwestern University and H.N. Cheng, ACS

Moderator: Young-Shin Jun, Washington University in St. Louis

Register for Free!

What You Will Learn:

- How mechanically interlocked molecules (MIMs) are easily made and how they can be used inn the construction of artificial molecular machines
- How AMMs operate under kinetic control using energy ratchets in a manner similar to that employed by our many biomotors and are at odds with how machines operate in the macroscopic world: the difference could not be

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Chemistry on Capitol Hill

2021 Emerging Policies

Date: Wednesday, June 30, 2021 @ 2-3pm ET Speakers: Caroline Trupp Gil, American Chemical Society / Karen Garcia, American Chemical Society / Carl Maxwell, American Chemical Society

What You Will Learn:

- . How the Biden Administration and 117th Congress are shaping up in terms
- of its STEM priorities

 Which specific pieces of legislation or federal policies will be likely to impact
- How members can become involved

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Designing

Bio-Sourced Polymers that Enable Recycling

Date: Thursday, July 1, 2021 @ 2-3pm ET Speaker: Stefan Mecking, University of Konstanz Moderator: Mark Jones, Dow Chemical (retired)

What You Will Learn:

What is solvolysis and how it can enable plastics recycling

- . How renewable polycarbonates and polyesters with a low density of in-chain functional groups as break points in a polyethylene chain can be recycled
- How long-chain building blocks for polycondensation can be created from common plant oil feedstocks or microalgae oils

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