

Discovery and Optimization of CBL-B Inhibitors

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Nurix's DELigase Protein Modulation Discovery Platform

DEL molecule Allosteric Inhibitor E3 Ligase

DEL Discovery

> 5 billion drug-like compounds that can be easily screened against hundreds of proteins to identify starting points therapeutic discovery

Rational and Empirical Chemistry



Structure Based Drug Design combined with chemistry automation enables broad exploration of lead-like chemical space for each program

Direct-to-Cell Biology Capabilities



High throughput cellular assays monitor protein levels and biological phenotypes to assess impact on biology

Scaled Screening for in vivo exposure



Capacity to screen for ideal in vivo drug exposure profile and assess impact on disease biology

Nurix Is Advancing Four Wholly Owned Clinical Programs with a Deep Pipeline of Proprietary and Partnered Novel Targets

MOA	Drug Program	Target/ Delivery	Therapeutic Area	Pre-Clinical	Phase 1	Phase 2	Phase 3
TPD	NX-2127 Degrader	BTK-IKZF Oral	B-Cell Malignancies				
	NX-5948 Degrader	BTK Oral	B-Cell Malignancies				
TPE	NX-1607 Inhibitor	CBL-B Oral	Immuno-Oncology				
	DeTIL-0255 Cell Therapy	Adoptive Cell Therapy Ex vivo CBL-B Inhibition	Gynecologic Malignancies				
ТРМ	Wholly owned	5 targets	Multiple				
TPD	Gilead Sciences	5 targets	Multiple				
TPD	Sanofi	5 targets	Multiple				

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CBL-B is a Modulator of Immune Cell Activation

- CBL-B is an E3 ubiquitin ligase highly expressed in cells of the immune system
- CBL-B regulates T, B, and NK cell activation
- Blocking CBL-B removes a brake on the immune system
- *cbl-b* deficient mice demonstrate robust T cell and NK cell-mediated antitumor immunity



CBL-B is a Modulator of Immune Cell Activation

Inactivation or deletion of CBL-B results in hyperactive T cells and inhibition of tumor growth.



IL-2 secretion in KO and ligase inactive T cells ex vivo

Ligase-dead or KO exhibit enhanced and equivalent response to either single- or double stimulation

Ligase-inactive *cbl-b* knock-in mice inhibit tumor growth (TC-1 syngeneic model).



Inactive CBL-B is Autoinhibited



- When Y363 of CBL-B is not phosphorylated, the helix of the LHR domain packs against the TKB domain
- Incapable of binding Ub-E2
- Phosphorylation of Y363 requires dissociation of LHR-RING from TKB

Active CBL-B Binds Ub-loaded E2 Ligases



Multiple Lead-Finding Approaches Afforded CBL-B Binders





CBL-B HTS Triage Revealed a Singleton Hit



CBL-B Phosphorylation FRET assay

- E2~Ub Binding FRET assay
- Src Counter Screen FRET assay

HTS Reveals a Singleton Hit





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Compound Binding to CBL-B by SPR



SPR confirms **NRX-1** binding affinity and stoichiometry to CBL-B SPR binding affinity and biochemical potency in close agreement

NRX-3 is a Specific Inhibitor of CBL-B



NRX-3 is an Intramolecular Glue



Crystal Structure Confirms Binding Mode as Intramolecular Glue





NRX-3 binds to closed-state CBL-B and prevents phosphorylation



Early SAR: Focus on Affinity and Properties











	NRX-3	NRX-4	NRX-5	NRX-6
E2-Ub: IC ₅₀ (μM)	12	0.23	0.092	0.088
Ligand Efficiency	0.29	0.33	0.36	0.37
Cellular Substrate Ub IC ₅₀ (μ M)		7	3	1.7
Microsomes h/m Cl _{int} (mL/min/kg)		20/360	-/500	30/73
Plasma stability m/r $T_{1/2}$ (min)		-	140/-	280/-
Papp MDCK (MDR1) A→B/B→A ratio		26/1	33/1	9/6
Ksol (μM)		250	300	270
LogD _{7.4}		2.6	2.3	1.9

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Early SAR: Focus on Affinity and Properties



	NRX-6	NRX-7	NRX-8
E2-Ub: IC ₅₀ (μM)	0.088	0.038	0.021
Ligand Efficiency	0.37	0.37	0.36
Cellular Substrate Ub IC ₅₀ (μ M)	1.7	0.78	0.79
Microsomes h/m Cl _{int} (mL/min/kg)	30/73	-/67	7/26
Plasma stability m/r T _{1/2} (min)	280/-	>1000/163	>1000/>1000
Papp MDCK (MDR1) $A \rightarrow B/B \rightarrow A$ ratio	9/6	7/7	2/14
Ksol (μM)	270	260	300
LogD _{7.4}	1.9	2.4	1.7



Complex SAR for Rat Plasma Stability





The SAR for rat plasma stability was not predictable by chemists

First observed with low recovery in PPB assays



Machine Learning Model for Rat Plasma Stability

To assist with lead optimization, models were built based on the 104 experimental plasma stability data points available at the time.

Despite the low volume of data, both regression and classification models demonstrated high predictive power and provided key insights driving series progression



Classification SVM Model





NRX-8 Is a Specific Inhibitor of CBL-B



NRX-8 displays clean 1:1 binding stoichiometry with CBL-B and is clean in off-target screening.



Me

RU

NRX-8 Maintains Original Hit Binding Mode







NRX-8 Inhibits Substrate Ub and Stimulates IL-2 Induction





Human T cell assay – IL-2 production



	NRX-8
IL-2 (2.5X over baseline response)	80 nM
Cellular Ubiquitylation of substrate (BT20 – MSD assav)	850 nM



Pharmacologic Inhibition of CBL-B Recapitulates Anti-Tumor Effects of Genetic Model of Ligase Inhibition

Ligase-inactive *cbl-b* knock-in mice inhibit tumor growth in TC1 Syngeneic Model





CT26 Syngeneic Model

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Over 10,000-fold Enzymatic Potency Improvement Achieved While Improving Molecular Properties



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Single-Agent NX-1607 Induces Antitumor Response in Multiple Models



Shaded area indicates dosing period

NX-1607 and Anti-PD-1 Synergize to Enhance Anti-tumor Effects and Survival of Mice in Multiple Tumor Models

Colorectal (CT26)

Colorectal (MC38)



Vehicle
NX-1607
Anti-PD-1
NX-1607+anti-PD-1

Shaded area indicates dosing period: NX-1607 (30 mg/kg, PO daily) and anti-PD-1 twice a week at 10 mg/kg dosing period **Triple-Negative Breast (4T1)**



- CBL-B regulates T, B, and NK cell activation
- Multiple screening approaches afforded validated binders to CBL-B
- Plasma instability may be an under-appreciated liability for amide-containing compounds
- Pharmacological inhibition of CBL-B recapitulates the anti-tumor effects of the genetic model of ligase inhibition
- NRX-8 specifically binds to CBL-B and 'glues' the protein in a closed state, preventing phosphorylation and E2-Ub binding
- Dosing of NRX-8 (45 mg/kg BID) inhibits tumor growth in mice
- Further optimization resulted in NX-1607 with sub-nM affinity and optimal in vivo anti-tumor activity
- Phase 1 clinical trial of NX-1607 in relapsed or refractory tumors is currently ongoing

Thank you

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