

# Unlocking Covalent Discovery Programs Concept to PDC

**Summary Deck** 

### **World-leading Covalent Capabilities**



#### Deep covalent discovery expertise from target validation, Hit ID, H2L and Lead Op

#### Target-oriented in Vitro Covalent Discovery

- >10,000 cpd/week high-throughput covalent screening (IMA)
- >15 K in-house covalent library (cys, lys +)
- Intact mass analysis (IMA) with detergent samples
- Peptide mapping analysis (PMA) follow-up site detection
- Protein scrubs generation
- 10,000 cpd/week intrinsic reactivity profiling (various nucleophiles)
- \$1.4 M of globally accessible covalent compounds; Curated 100K of lead like diversity screening set; curated 1K of diversity fragment set In-house covalent library;
- GSH/GST; plasma, whole blood, hepatocytes (human, dog, murine)
- k<sub>inact</sub>/K<sub>I</sub> high-throughput determination
- Multiple orthogonal methods for kinetics and binding
- Covalent crystallography
- Multiple clinical and preclinical benchmarks profiled for reference
- Cell profiling; wash-out, protein half-life; target engagement and selectivity in chemoproteomics
- Covalent CADD capabilities

- Expert team who prosecuted >20 covalent discovery programs
- Top industry insight on the covalent DC and TP profiles, Hitfinding strategies, solving selectivity and metabolic stability challenges

#### **Proteome-Wide Covalent discovery**

- **51,000 sites/10,500 proteins**, <u>13,000 proprietary sites/430</u> proteins & growing; proprietary ligandable site dataset
- Cysteome, and beyond
- Oncology, CNS, immunology
- In-house proteomics hit-finding library 100 compounds\*
- Curated database of sites/targets liganded in literature
- Ultra-deep target-hit profiling
- High-throughput screening
- Targeted proteomics (low-abundance sites)
- Lead covalent selectivity profiling (proteome-wide)
- Proprietary high-throughput data processing & pocket mapping algorithms

### X-Valent Discovery Platform to Candidate: Covalent Drug Cascades



Unlocking holy-grail targets to small molecule therapies Identifying novel targets ideally positioned for covalent therapeutics





- **Proprietary**, cutting-edge covalent drug discovery engine powered by *X-valent* chemoproteomics and *X-valent* target discovery workflows
- **Proteome-wide** screening (target-directed and target-agnostic)
- Target-oriented high-throughput workflows
- Identification of cryptic pockets on desired targets
- **In-cell, accelerated**, Hit and Lead generation for **novel targets** tractable to covalent modulation
- Complete **integrated** workflow for covalent drug discovery to Candidate nomination
- World's top expert team in covalent drug discovery

### X-Valent Screening Platform to Candidate: Covalent Drug Cascades

• Our covalent screening cascade has a strong track record of bringing discovery programs to development candidate stage.



- GSH and hWB stability benchmarked versus known, marketed drugs
- GSH/GST stability vs GSH alone provides further information.
- *In vitro* washout experiments help inform duration of action, protein resynthesis and predict in vivo requirements
- LM and Hep stability (Met ID) also help prioritize compounds

## **Industry-leading Covalent Libraries**



> 15,000 ready to screen covalent library collection and growing
 Target-oriented *in vitro* and whole cell Covalent Discovery platforms enabled by world-leading proprietary Covalent Libraries

- Proprietary, cutting-edge covalent libraries incorporating diverse warhead coverage
- Modular libraries (0.4-8.4K individual modules) including commercial collections and rationally curated warhead designed arrays, clinical competitors and reactive probes
- Library Enhancement Program (LEP) rapidly incorporates emerging (pre)clinical, chemical biology probe and covalent drug-like space arrays:
  - Expanded "under exemplified" warheads
  - Warheads in clinic/preclinical but not available in commercial libraries 🔫
  - Increased Fsp3
  - Augmenting with more annotated (Lead-like MWt 300-400) arrays
- **CADD-enabled** covalent Drug Discovery, including covalent VS with curated 100K lead like set
- Dalriada covalent library is powered by *iCLASS* chemoproteomics and *X-valent* target discovery workflows



### **Covalent Virtual Screening**

## **Biophysical/Biochemical**

#### **Proprietary chemical space of compounds**

- D3 SPACE: 1.3M molecules with covalent warhead(s)
- CovDock by Schrödinger: Virtual Screening (VS) mode



Covalent docking: ligand sampling with covalent bond formation to a target residue



Curation of a virtual library of electrophiles.



Generation of stereoisomers, ionization states and ligand conformations

### **Biochemical: Time-dependent Covalent Engagement** Characterization



• Time dependent product formation (activity) of enzyme after incubation with various inhibitors, followed by jump dilution



Assessment of a covalent modifier using a continuous enzyme assay enabling k<sub>inact</sub>/K<sub>I</sub> studies



Time-dependent inhibition of Enzyme by a compound.  $IC_{50}$  values are shown to decrease as pre-incubation times increase.

- **DSF experiments** to determine protein stability
- **DSLS experiments** to determine stability of proteins
- ITD experiments to determine Denaturation Rate and Half Life
- ITC to determine thermodynamics of binding
- **CD** to determine 2° structures of proteins
- **SPR** experiments to determine thermodynamics of binding

### MS and (Chemo)Proteomics Capabilities:



Proteomics Capabilities	Applications	TurnKey Advantage
Chemoproteomics	Target Site Identification, Selectivity Determination	Deep Profiling Of The Reactive Proteome (~20,000 Reactive Sites)
Whole-proteome Profiling	Mechanism Of Action Studies, Selectivity Determination	Deep Proteome Profiling (>5,000 Proteins) High Quantitative precision (CV < 4%)
Pulsed SILAC	Determination of protein homeostasis and turnover	High-throughput, high precision (Integration of SILAC with TMT technology)
Intact Mass Protein Analysis	Protein-compound Binding, Covalency Determination	Ultra-high Throughput Screening Capacity (>7000 Compounds/Week)
Peptide Mapping Analysis	Protein Sequencing, PTM Profiling, Binding Site Determination	High Sequence Coverage
Bioinformatics	Biostatistical Analysis, Network Enrichment, Hierarchical clustering	Comprehensive Data Analysis, Cloud-based/Interactive Data Visualization And Data Sharing

#### **Chemoproteomics Discovery Platform**

- 51,000 reactive protein sites mapped
- >10,500 proteins with reactive sites mapped
- >13,000 new reactive sites identified
- >440 new proteins with reactive sites identified





• Determination of %occupancy of AMG-510 on KRas G12C by IMA

#### **Peptide Mapping Analysis**







- IMA screening of ~7,000 lead-like covalent molecules
- Determining covalent binding site and characterization of recombinant protein sequence, isoforms, PTM, etc.
- A high-throughput covalent PMA platform enabled by plate-based sample processing, micro-flow chromatography, and the in-house PMA data analysis and QC tool.

### Who We Are - Headquartered in Canada Supporting Programs Globally



EXPERT & TOP • QUALITY	THE	
Drug discovery expertise of world's top firms Highly skilled scientists State-of-the-art instrumentation	PREFERRED PARTNER FOR	Fastest turn-around times Full customization of R&D Highest partner engagement
	DRIVING FAST-PACED	
	INNOVATIVE	
	PROGRAMS	
<b>COST-EFFECTIVE</b> Highly competitive pricing enabled by our unique <b>Turn-Key Model</b> Objective and milestone-oriented	IN SMALL MOLECULE DRUG DISCOVERY	• <b>COMPREHENSIVE</b> The most comprehensive support from discovery to candidate nomination
		Top partner support



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### THE MOST COMPREHENSIVE SMALL MOLECULE DRUG DISCOVERY R&D SUPPORT





### From concept to IND-ready

Unparalleled value-build and efficiency for any discovery program

### Leading Bench-Side Expertise, Capabilities And Turn Around Times

A track-record for executing top-quality R&D for even the most complex programs at the bench-side

- Canadian team of > 70% PhDs across chemistry, CADD, *in vitro* biology, DMPK, and Proteomics
- 500+ publications; >300 years combined R&D experience
- Interdisciplinary teams working side
  by side for fully integrated programs
- Fastest turn around cycles
- State-of-the-art instrumentation and facilities



### Strategic Guidance From Concept To IND

Scientific oversight from industry experts to maximize program success

- Roster of experienced veterans and niche KOLS >300 years in pharma, biotech completed 100+ programs across various therapeutic areas and targets classes, and advanced multiple programs to DC, clinic and market
- Inventors on >150 patents; raised over \$100 M, executed >\$2 B in Pharma partnerships and founded multiple companies

#### Seamless Execution And Complete Integration

Dedicated, experienced program leaders and managers to deliver ultimate collaborative experience and rapid program progression

- Delivery of consistent two-week design-make-test-analyze cycles
- Selection and management of subcontractors for niche needs and costeffective solutions from an **expansive network** for complete integration.

### **Dalriada's IDD Leadership Experience and Philosophy**



**Dalriada's Discovery Strategy leadership team** that will be assigned to your project have 140+ years integrated drug discovery (IDD) experience (in Pharma, CRO & Biotech) successfully leading small molecule projects through hit validation, H2L and LeadOp phases, identifying 41 Development Candidates.

- Project Led on 180+ IDD programs
- Delivering on challenging target classes including PPIs, Covalency, PROTACs and CNS
- Track record of delivering development-quality Candidates with clinical progress-ability; Over 20 molecules delivered to the clinic

Discovery lab teams are fully enabled to efficiently deliver IDD projects

### Dalriada's shared IDD philosophy incorporates the following pillars to drive project outcomes

#### **Rapid DMTA cycles**

Co-localization, effective processes & monitoring – continuous improvement – maximizes learning iterations for the end-to-end R&D

### Hypothesis driven design

Utilizing SBDD, CADD, pharmacophore, conformation and mechanism info – every compound counts and should address a question Physiologically relevant endpoints & translation Monitor formation of ternary complex & protein homeostasis, considering translation to patient group

#### Project back-planning

Visibility to # iterations and triggers for data collection, ensures **delivery focus on our shared goals** and value inflection points

### **Drug Discovery and R&D Leadership**







### For the full deck and case studies



## **Thank You!**

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