

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)
- 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_{inact}/K_l 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

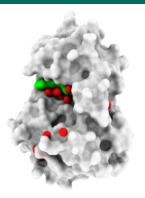
Proteome-Wide Covalent discovery

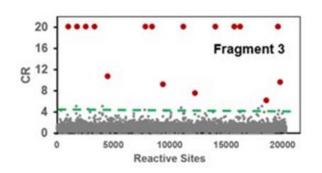
- **51,000** sites/**10,500** proteins, <u>13,000</u> proprietary sites/<u>430</u> proteins & growing; proprietary ligandable site dataset
- Cysteome, and beyond
- · Oncology, CNS, immunology
- In-house proteomics hit-finding library 100 compounds*
- Curated database of sites/targets liganded in literature
- 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

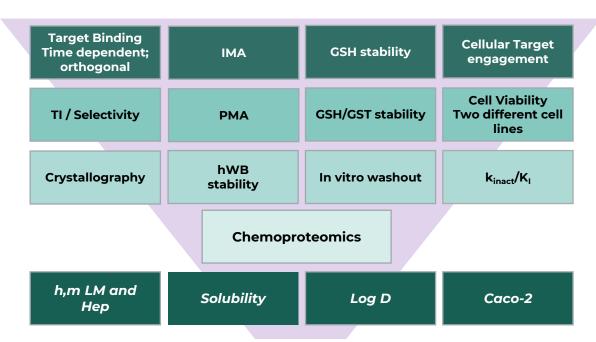




- Proprietary, cutting-edge covalent drug discovery engine powered by X-valent chemoproteomics and X-valent target discovery workflows
- **Proteome-wide** screening (target-directed and target-agnostic)
- Target-oriented high-throughput workflows
- Identification of **cryptic pockets** on desired targets
- 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

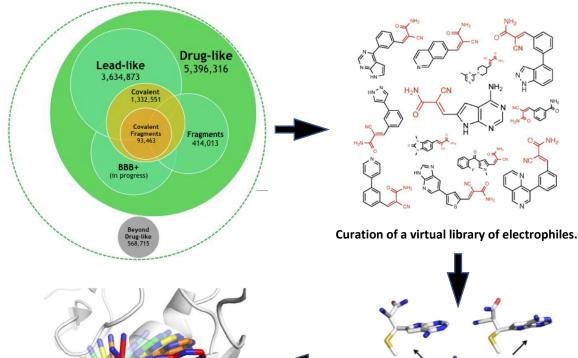
Covalent Virtual Screening

Biophysical/Biochemical



Proprietary chemical space of compounds

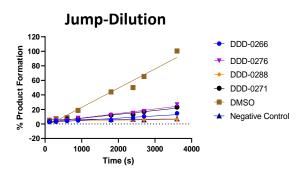
- D3 SPACE: 1.3M molecules with covalent warhead(s)
- CovDock by Schrödinger: Virtual Screening (VS) mode

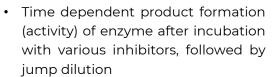


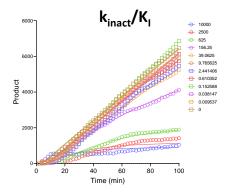
Covalent docking: ligand sampling with covalent bond formation to a target residue

Generation of stereoisomers, ionization states and ligand conformations

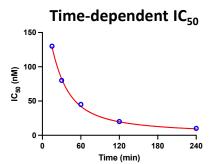
Biochemical: Time-dependent Covalent Engagement Characterization







• Assessment of a covalent modifier using a continuous enzyme assay enabling $\mathbf{k}_{\text{inact}}/\mathbf{K}_{\text{I}}$ 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

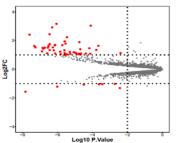
MS and (Chemo)Proteomics Capabilities:

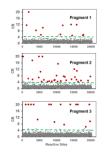


Proteomics Capabilities	Applications	TurnKey Advantage
Chemoproteomics	Target Site Identification, Selectivity Determination	Deep Profiling Of The Reactive Proteome (~20,000 Reactive Sites)
Whole-proteome Profiling	Mechanism Of Action Studies, Selectivity Determination	Deep Proteome Profiling (>5,000 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

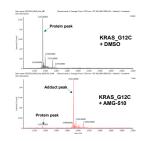
Chemoproteomics Discovery Platform

- **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

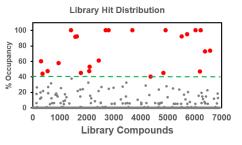




Intact Mass Analysis Platform

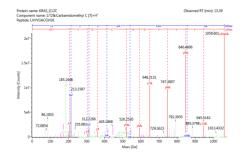


 Determination of %occupancy of AMG-510 on KRas G12C by IMA



• IMA screening of ~7,000 lead-like covalent molecules

Peptide Mapping Analysis



- 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

Drug discovery expertise of world's top firms

Highly skilled scientists
State-of-the-art instrumentation

COST-EFFECTIVE

Highly competitive pricing enabled by our unique **Turn-Key Model** Objective and milestone-oriented

THE

PREFERRED PARTNER FOR

DRIVING FAST-PACED

INNOVATIVE

PROGRAMS

IN SMALL MOLECULE DRUG
DISCOVERY

AGILE AND RAPID EXECUTION

Fastest turn-around times Full customization of R&D Highest partner engagement

COMPREHENSIVE

The most comprehensive support from discovery to candidate nomination

Top partner support





Our Values





Creative



THE MOST COMPREHENSIVE SMALL MOLECULE DRUG DISCOVERY R&D SUPPORT



Evaluate

program science, IP & business objectives



Custom-build

internal R&D teams, program cascades & assays aligned with IP & business strategy



Efficiently execute

R&D across Biology Biochemistry Proteomics Chemistry

ADME CADD 4

Manage

the entire or partial program including transparent outsourcing



Support

further technology development, IP & due diligence processes

The Turn-Key™ Model



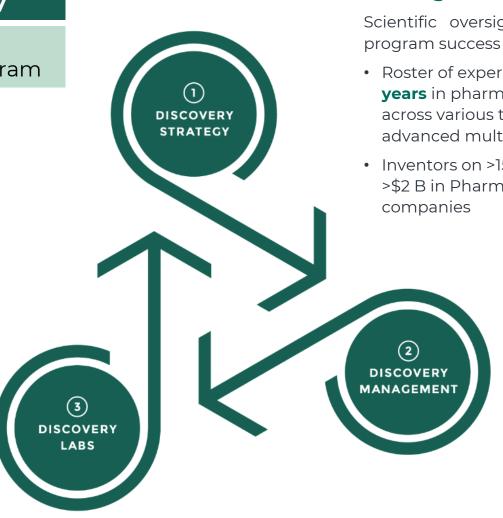
From concept to PDC-ready

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 PDC

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



Medicinal Chemistry





Jeff



Mark ADME / DMPK



Tom

Harpreet



Kevin

Collectively: Over 275 years of drug discovery and specialty experience

Across 100+ different discovery programs & 6+ major therapy areas, including:

Immuno-oncology

Biochemistry / Biophysics

- Inflammation & Immunology
- Cardiovascular & Metabolic Diseases
- Neuroscience
- Oncology
- Anti-infectives

Proteomics

Biology/Pharmacology





lain

Jeff



Diana

Computational Chemistry





Mike

Background:

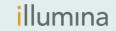




Mohammad



Frosty



Uros













































For the full deck and case studies



Thank You!

For more information please contact:

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