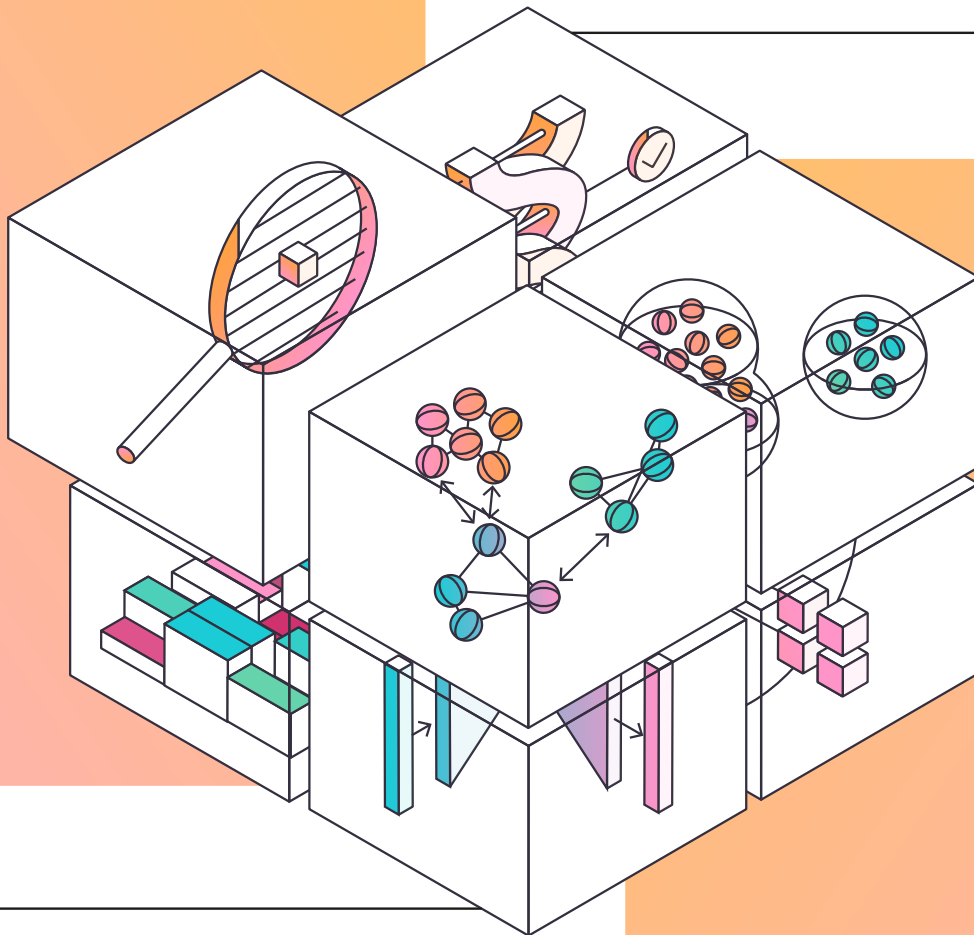




immunaï

A Target Discovery Workflow Designed for Precision Medicine

Immunaï delivers novel targets
with precise therapeutic hypotheses
and detailed data packages to enable
our partners' pipeline decisions



We implement proprietary machine-learning approaches on high-resolution data from patients and model systems to **discover and validate targets** that matter most in disease

Our Approach

Evidence based

A patient-derived single cell atlas powers target discovery, ensuring insights are directly informed by human disease biology

Disease relevant

Dissecting disease mechanisms at high resolution enables testing of preclinical hypotheses in relevant model systems

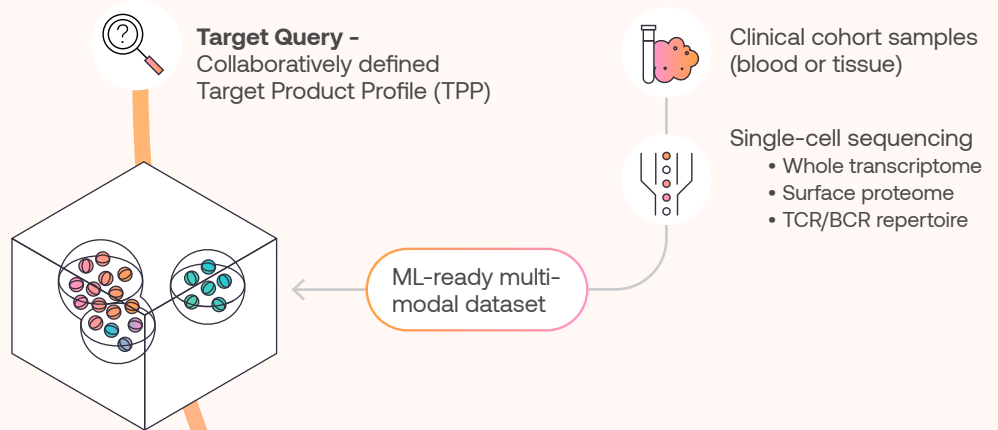
Data-driven

Clinically relevant target validation ensures our data-driven recommendations have a higher likelihood of successful translation to the clinic

01. GENERATE

Deep, Multi-omic Data

Starting from customer clinical samples, we apply single cell sequencing to GENERATE high quality, **high resolution multi-omic data**.

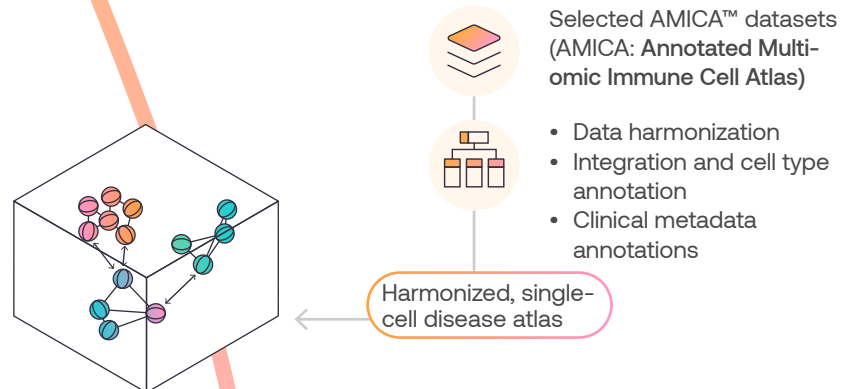


02. AUGMENT

AMICA™-Powered Data Foundation

We AUGMENT customer datasets with studies contained in **AMICA™**, integrating millions of cells into a **highly differentiated data foundation**.

Importantly, we **harmonize clinical metadata** to enable a unified analysis of many disparate studies.



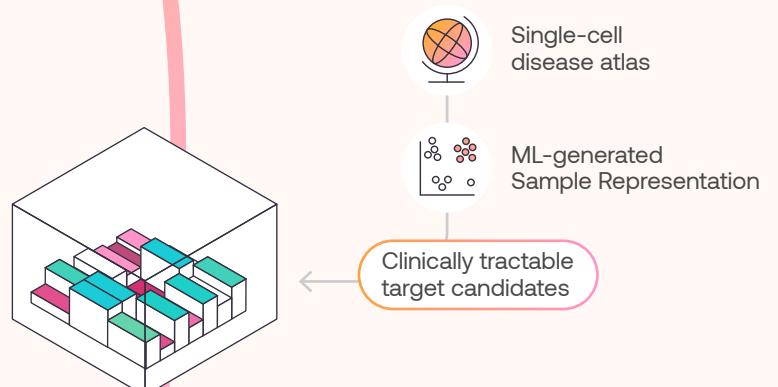
03. COMPUTE

ML-Driven Target Discovery

With our harmonized disease atlas, we apply machine learning to identify candidate target genes.

We COMPUTE a Sample Representation from high dimensional single cell data, in order to **link cell type specific targets to disease and treatment outcomes**.

Our models **rank targets** based on biological and therapeutic relevance, with our experts further refining selection to fit the desired TPP.



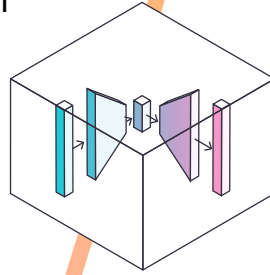
04. VALIDATE

Disease Relevant Target Validation

Next, we **VALIDATE** candidate targets to understand their **impact on the disease process**.

A disease-relevant model system is optimized for each target. **In vitro knockout effects** are characterized through an **integrative analysis** measuring key disease-related processes.

These insights are mapped back to human samples to confirm clinical relevance.



Optimized disease relevant model systems



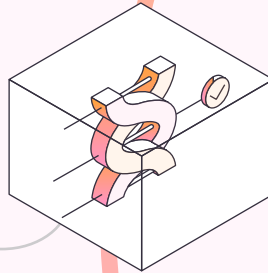
in vitro/ in vivo functional validation

Validated target candidates

05. RECOMMEND

Translating Data into Therapeutic Strategies

Recommendation towards drug development decision making



The right patient population

Example recommendation: A target that is induced in patients following Standard of Care treatment could be strategically modulated after discontinuation to maximize therapeutic benefit.



The right cell type

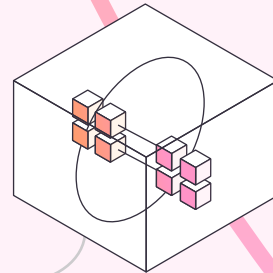
Example recommendation: A target with a pathogenic function in fibroblasts and a protective function in macrophages should be selectively modified in fibroblasts.

06. EXPLAIN

Target Data Package

Comprehensive **data package** for each target **explains** clinical data supporting the therapeutic hypothesis.

Target data package



Expert immunological insight



Key *in vitro/ in vivo* functional validation data



Clinical significance of experimental data to patients

Our Impact

Novel Targets

Our platform identifies and validates novel, actionable, molecular targets

Informed Decision-Making

We empower our partners to make precise, data-driven decisions that de-risk their drug development pipeline

Maximized Success

By prioritizing target candidates with a higher likelihood of success, we help drive more effective therapies

Selected publications

Van de Sande, Bram et al.

Applications of single-cell RNA sequencing in drug discovery and development

Nature reviews. Drug discovery vol. 22,6 (2023): 496-520.

doi:10.1038/s41573-023-00688-4

G De Baets et al.

Inflammatory Bowel Disease single cell atlas construction to enable cell-type-specific target identification

ECCO 2023 DOP40

Abstract citation ID: jjad212.0080

G De Baets et al.

An integrated single-cell atlas enables the discovery and validation of a novel candidate therapeutic target for Inflammatory Bowel Disease

ECCO 2025 DOP132

Abstract citation ID: jjae190.0171