# Target ID and drug repurposing

Apply ML to a universe of targets to identify novel targets that fit your selection criteria

### **Description**



Uncover new drug targets or new uses for established targets via exhaustive network analysis leveraging Clarivate's biomedical Knowledge Graph

## **Expert-curated datasets containing**



4.6M+ Molecular interactions



1,500+ pathway maps



746K+ drugs and biologics



2.7M+ gene-disease associations



10M+ patents

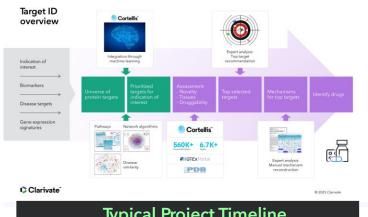


### Methodology

- An extensive set of features is computed proteome-wide describing each protein's
  - Network proximity to disease biomarkers
  - Involvement in pathways and processes implicated in the disease
  - Existing drug development in other molecularly similar indications
- Given known validated targets of the disease, a predictive model is trained to prioritize targets most similar to validated targets across the feature space
- Top targets are further characterized and annotated with information on target biology and competitive landscape
- Expert-led manual biological interpretation is conducted through literature review for top targets to build further scientific rationale supporting the protein as a potential target for the indication

#### Output

- Table of prioritized targets with annotations relevant to the indication of interest
- Supporting evidence reports for top targets providing details of target biology, relevance to the disease and competitive landscape
- Literature review report summarizing available scientific rationale for the target modulation in the disease of interest



# **Typical Project Timeline**

From 8 weeks, subject to complexity

Select case studies available upon request