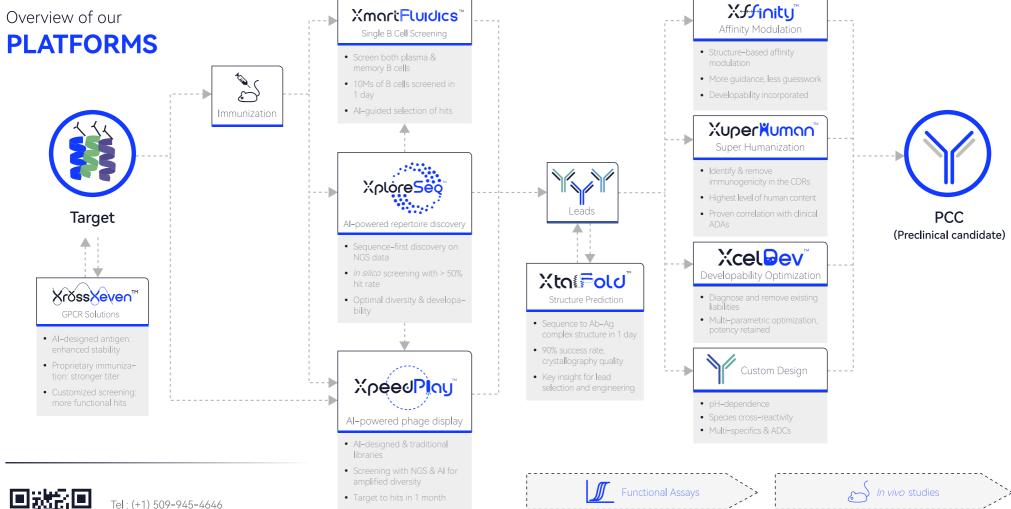


## XupremAb Al-Powered Next-Gen Antibody Discovery Platform





E-mail: ab.bd@xtalpi.com
Global BD Center: 245 Main St, 11th Floor,

Cambridge, MA 02142



Developability Assessment

• Predict 15+ developability metrics in silico

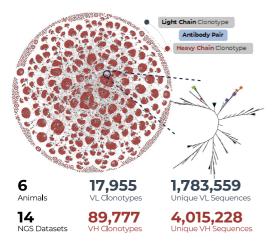
Comprehensive wet-lab characterization

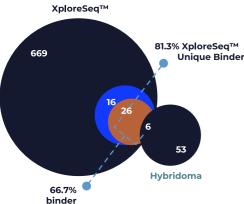
#### Hit-Gen

Tap into greater search space for more quality hits

01

In silico screening of millions of antibody sequences in the immune repertoire





XploreSeg<sup>™</sup> routinely sequences millions of BCRs and identifies hits with high confidence (above). In this project, XploreSeq™ predicted 717 binders, from which 48 were randomly chosen for expression and testing. 66.7% (32/48) were confirmed to be binders, and only 18.6% (6/32) binders overlap with hybridoma hits (below)

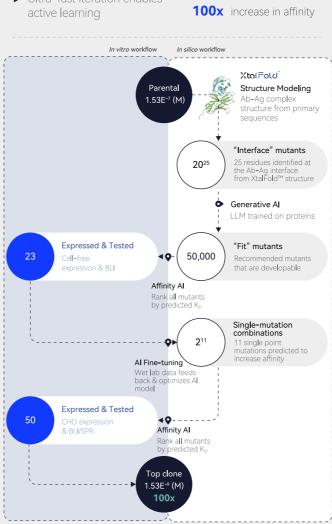
### **Engineering**

Utilize AI to make antibody engineering more efficient, enabling multi-objective optimization

## 02

Affinity 100x ↑ in 3.5 weeks no structure information

- ► Goal: Improve VHH KD from hundred nM to single-digitnM in a short timeframe
- ► Ultra-fast iteration enables active learning
- 2 rounds of optimization
- 73 variants expressed
- 3.5 weeks

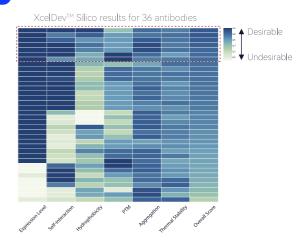


### Developability

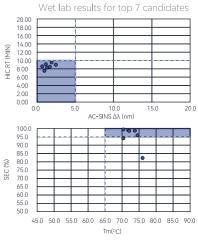
Instill excellent developability into every molecule

# 03

XcelDev™ Silico evaluates and ranks 30+ candidates in terms of developability



36 hits that had passed binding and functional screening were analyzed by XcelDev™ Silico. 6 properties of each antibody was predicted and scored on a scale of 0-1 (displayed in color gradient).



An overall developability score was calculated to rank all 36 hits. To verify the effectiveness of the ranking, the top 7 (in the red box) were expressed and subject to a battery of in vitro developability assays. All 7 performed well in these assays, their Tm, SEC, AC-SINS and HIC results shown above.