

10× HIGHER HIT RATES AT 3–10× LOWER COSTS: ACCELERATING LIGAND DISCOVERY WITH ML-GUIDED SYNTHESIS & SCREENING

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Rapid & reliable hit generation with Direct-to-biology-SpaceM1

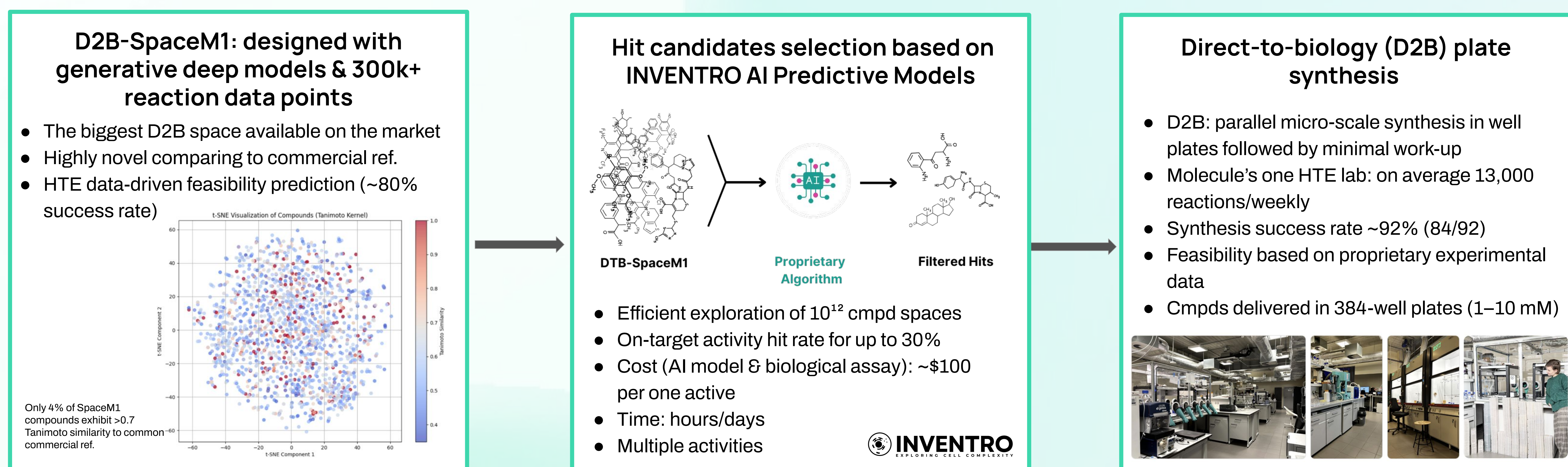
- Unique & highly novel molecular space of ~100M compounds and still growing
- **First chemical space built using deep learning models trained on proprietary HTE lab data of over 300,000 reactions**
- Rapid turnaround (fastest subset from order to measured biological activity in **4 weeks**)
- Compounds starting from <\$10

In comparative virtual screening[#] on 12 protein targets with ~395 randomly selected compounds from commercial reference space and D2B-SpaceM1; D2B-SpaceM1 showed on average higher hit rates:

Target (examples)	D2B-SpaceM1 docking hit rate**	C-Ref* docking hit rate**	Novel to C-Ref (Tanimoto < 0.75)	Highly novel to C-Ref (Tanimoto < 0.5 & Different Scaffold)
PRMT5	10.4%	1.0%	60.0%	10.8%
KRAS(G12C)	6.6%	0.0%	48.6%	18.9%
LRRK2	4.3%	0.3%	54.5%	21.2%
mGluR5-5M	8.1%	0.5%	43.5%	10.1%
BTK	24.3%	3.6%	46.7%	15.6%

[#]Based on ligand efficiency; *C-Ref - commercial reference; **Hit rates calculated using ligand efficiency with threshold of activity was set dynamically as at max(0.4, thr_chembl), where thr_chembl is median ligand efficiency of top 10% most active compounds stored in ChEMBL. Compounds were randomly sampled from each space.

Hit identification & confirmation from virtual screen in 5 weeks

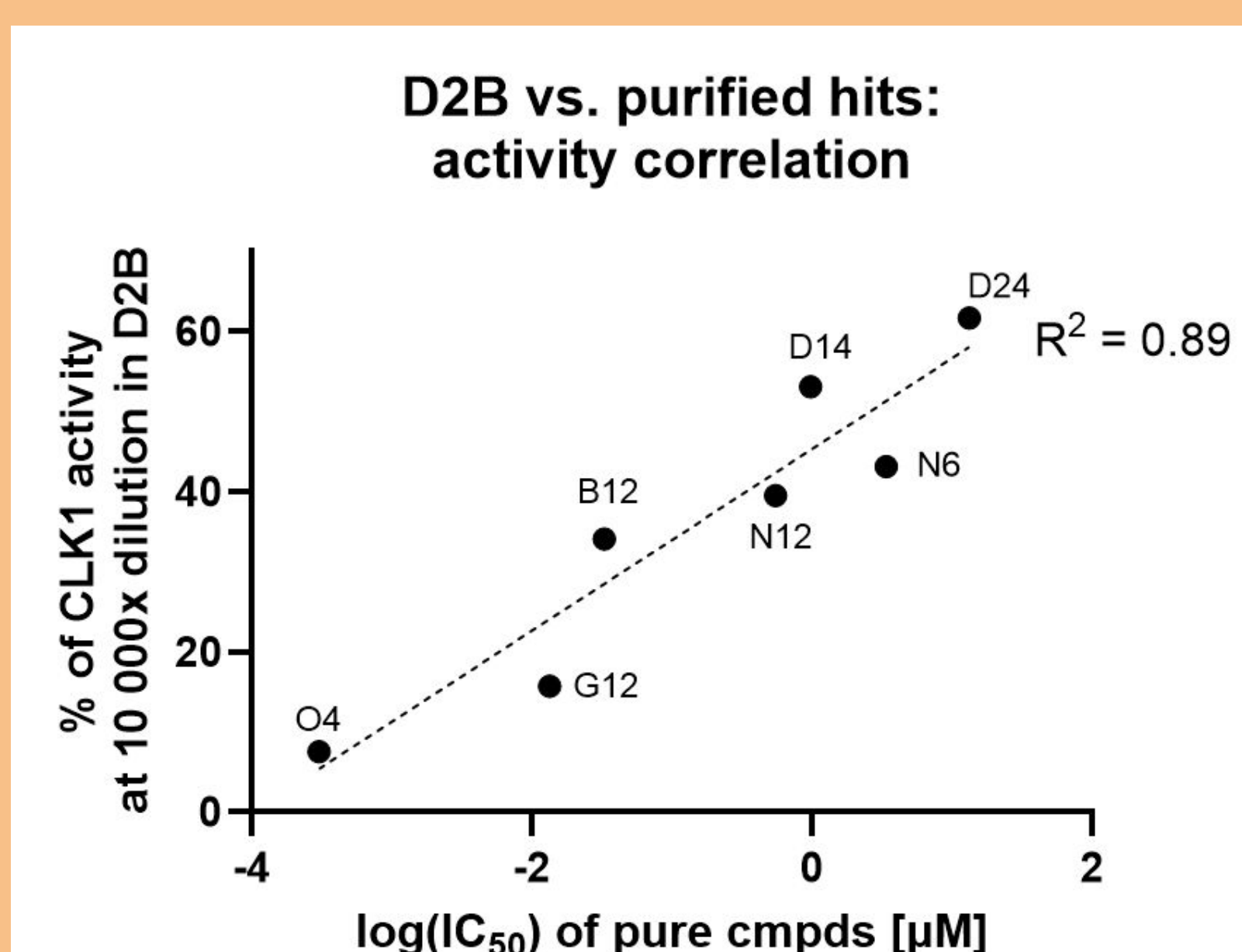


Summary:

- **Largest & most novel D2B space**, built on extensive HTE reaction data
- **Strong (0.89 R²) correlation between D2B crude activity and purified IC₅₀ ranking**
- Hit activity up to **sub-nM level**, hit rate 18%
- End-to-end workflow is **very rapid**, enabling experimental hitID and confirmation within **weeks, not months**
- Workflow applicable also to hit expansion and hit2lead

Hit resynthesis & confirmatory assay

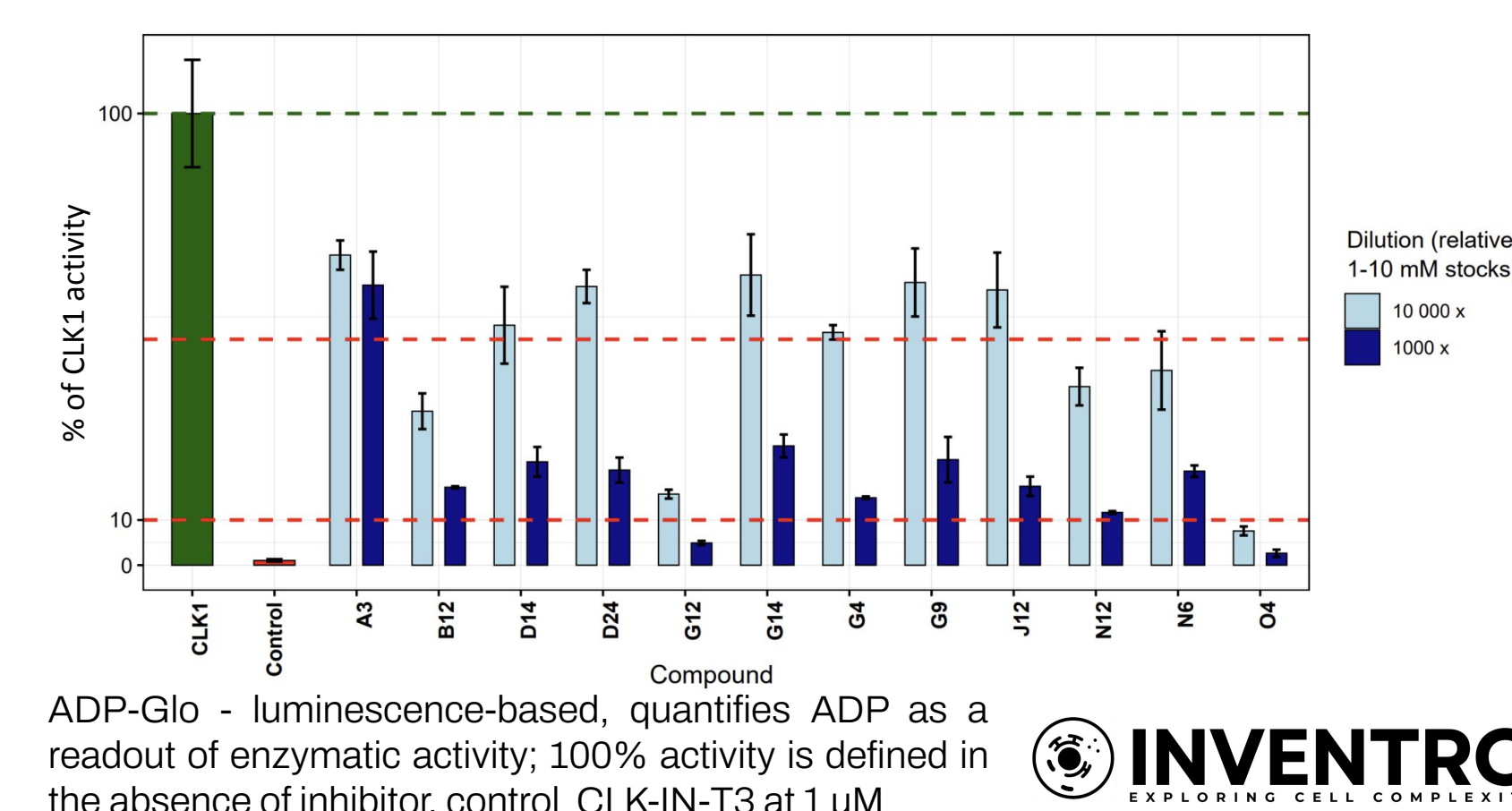
- Fast turnaround: hits resynthesized, purified and delivered within 1 week of D2B assay results
- All purified hits confirmed active at Inventro
- Potent hits* down to sub-nM level (IC₅₀ distribution: 2 at μM, 2 at sub-μM, 2 at nM, 1 at sub-nM)



*No prior literature reports for these structures; for O4, SciFinder similarity 85 to the closest known

Rapid biological screening

- Initial 100× dilution screen in ADP-Glo → 12 compounds with CLK1 activity <10%
- 2nd run at 1,000× and 10,000× dilutions
- Outcome: 11 compounds inhibited CLK1 activity >50%; hit rate: 18% (AI-prioritized), 12% (full grid)



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