

Redefining Chemistry: How AI Shrinks Small-Molecule Discovery from Years to Weeks

SAN FRANCISCO, CA, UNITED STATES, September 25,

2025 /EINPresswire.com/ -- Ainnocence today unveiled

a major advance in Al-driven small-molecule R&D, demonstrating how its platform compresses the upfront discovery cycle from years to mere weeks, without compromising rigor on potency, selectivity, and developability.



Chemistry shouldn't take years to show you what's possible"

Dr. Lurong Pan, CEO of Ainnocence

Built on Ainnocence's third-generation AI system and its CarbonAI design engine, the platform unifies generative chemistry, physics-aware scoring, ADME/Tox prediction, and rapid iteration into a closed-loop workflow. In internal and partner programs, this has translated into swift progression from target hypothesis to synthesized, test-ready lead series in a fraction of traditional timelines.

"Chemistry shouldn't take years to show you what's possible," said Dr. Lurong Pan, CEO of Ainnocence. "By combining large-scale generative models with multi-objective optimization and instant manufacturability checks, we turn weeks into the new unit of progress for small-molecule discovery."

What's new

- Weeks, not years: Rapid progression from initial design brief to prioritized lead series via continuous Al-guided cycles.
- Massive search at practical cost: Virtual screening and design exploration across libraries up to 10^10 compounds within hours, focusing compute where it matters most.
- True multi-objective design: Simultaneous optimization for potency, selectivity, ADME/Tox, IP-friendly novelty, synthetic accessibility, and cost-of-goods, all scored in one loop.
- CarbonAl® for small molecules & PROTACs: CarbonAl® is a de novo small-molecule and PROTAC Al design engine capable of screening billions of compounds in mere days, simultaneously optimizing multiple pharmacological properties for lead generation and optimization. It delivers 50 candidates in 3 iterations for wet-lab testing, with milestone success achieved.

How it works

- 1. Design brief in: Define target, desired MoA, liabilities to avoid, and developability constraints, all captured in a structed digital template.
- 2. Generate & score: Physics-informed generative models propose candidates; property predictors and AI/ML scorecards rank them.
- 3. Select & synthesize: Shortlisted compounds move directly into synthesis with vendor-ready routes and procurement lists.
- 4. Close the loop: Wet-lab results feed back to the models for rapid, data-driven improvement ensuring each cycle is faster and smarter than the last.

Early access & partnering

Ainnocence is opening CarbonAl® Accelerator slots for biopharma, specialty chemicals, and advanced materials teams seeking to compress timelines or rescue stalled programs. Partners gain hands-on access to the platform, optional on-prem deployment, and dedicated Al-chemistry support.

About Ainnocence

Ainnocence is a next-generation biotech and deep-tech company building AI for molecular discovery across

therapeutics, synthetic biology, and advanced materials. Its third-generation AI platform delivers lightning-fast virtual screening and multi-objective profile optimization, with computational capacity reaching billions of sequences or compounds within hours, enabling high wet-lab hit rates and dramatically reduced discovery time and cost. Learn more at www.ainnocence.com.

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Carbon Al® Small Molecule Design Engine **Lead Generation** · Scaffold Hopping de novo Generation
• Lead Optimization Analog **Application Scenarios** PROTAC de novo Generation **Lead Optimization** ADME (Absorption, Distribution, Metabolism, Excretion) Prediction

Toxicity Prediction

DMPK Prediction **Target Binding** Why Choose Ainnocence? Optimization · Protein Binding Specificity No structure required Optimization Target Binding Pocket
 Detection
 Molecular Docking Wet-lab validated Ultra-high throughput Rapid Turnaround: from a few hours to two weeks Off-Target Prediction > Compound Off-target Toxicity Ainnocence's CarbonAl® Platform for AI-Powered Small Molecule Drug Design — Enabling De Novo

Generation, Lead Optimization, and
Off-Target Prediction Without
Structural Data

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