

Clinical Trials Are the New Bottleneck AI Drug Discovery Has Created an Evidence Infrastructure Crisis

APR 01, 2026 • PAID



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Abstract

The core argument: AI has dramatically compressed preclinical drug discovery, but clinical development timelines remain stuck. The bottleneck has shifted from molecule identification to evidence generation. The next durable health-tech companies won't discover drugs. They'll prove they work.

Key claims:

- AI-driven structure-based and generative methods have increased preclinical throughput substantially, pushing more candidates into already-strained development pipelines
- Long-run clinical success rates only recently began recovering after decades of decline, per 2025 Nature Communications data – meaning the industry hasn't seen translational efficiency at scale
- FDA's 2025 draft guidance on externally controlled trials effectively outlines a technical stack (phenotype normalization, covariate harmonization, temporal alignment, endpoint ontology mapping) that nobody has fully built yet
- A 2025 Nature Medicine TrialTranslator study showed real-world oncology surgery is often ~6 months worse than RCT results, and ~1 in 5 real-world patients don't qualify for phase 3 trials

- 2025 FedECA paper in Nature Communications introduces federated external control arms for distributed settings – a direct blueprint for privacy-preserving comparator networks
- TrialGPT (Nature Communications, 2024) and successor systems suggest patient trial matching is solvable, but recruitment alone is too narrow a moat without a broader trial-state architecture
- Five infrastructure layers where durable category-defining companies will likely form: comparator infrastructure, phenotype infrastructure, continuous measure infrastructure, model assurance infrastructure, and adaptive protocol infrastructure

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The paradox nobody talks about out loud

Here's a thing that should be more disorienting than it is: the AI-in-biopharma narrative has largely convinced the industry, investors, and the trade press that the hard part of drug development is finding good molecules. Structure-based design, generative chemistry, AlphaFold derivatives, target ID from genomic embedding – of it has gotten genuinely impressive. Not vaporware impressive. Actually impressive. Some of these tools are running live in discovery programs at major sponsors right now.

The problem is that making the front end of discovery faster is a little like widening the on-ramp to a highway that's already gridlocked. You don't actually get more to their destination any faster. You just fill up the backup.

Clinical development is the backup. It takes, on average, somewhere between six to ten years from first-in-human to approval, and the bottleneck during most of that period is not computational. It's not even primarily biological in the narrow sense of evidentiary. The industry is slow at assembling the kind of regulatory-grade, causal, defensible, generalizable evidence packages that the FDA and its international equivalents actually need to say yes with confidence. And as AI accelerates the front end, that evidentiary bottleneck becomes more acute, not less.

This is the paradox that people in clinical development talk about quietly but that rarely makes it into the funding narratives for health-tech companies. Everyone wants to fund the "AI for drug discovery" story because it maps to a familiar venture pattern: scientific insight, clever model, platform play, licensing revenue or acquisition. The evidence generation story is harder to pitch because the product is less romantic. It involves phenotype normalization, external control cohort assessment, federated data governance, digital twin validation frameworks, and adaptive machine learning protocol software. None of that fits on a TED slide.

But that's exactly where the alpha is.

Why this bottleneck exists now

To understand why this is happening now specifically, it helps to separate two different timelines that are running at very different speeds.

The first timeline is discovery throughput. Over the past five or six years, the combination of structure prediction, generative molecular design, and large-scale biological embedding has genuinely changed the rate at which credible drug candidates can be identified. The number of AI-discovered compounds entering clinical trials – while still small relative to the total industry pipeline – is growing

More importantly, the capital and talent flowing into “AI-native” biotech is enormous, which means the pipeline of candidates heading toward IND filings is expanding.

The second timeline is development infrastructure. This one has barely moved. The FDA’s average review timeline hasn’t compressed dramatically. Phase 2 and phase 3 success rates, while having shown some recent improvement after decades of decline, per 2025 Nature Communications analysis, are still deeply sobering – roughly 10 percent of candidates that enter phase 1 ultimately reach approval depending on therapeutic area. Enrollment velocity is still plagued by the same problems it was twenty years ago: sites are overextended, patients are hard to identify, eligibility criteria are often written for clean populations that don’t exist in the wild, and sponsors regularly discover late in development that their trial population doesn’t look much like the patients who will actually use the drug if it’s approved.

The structural cause of this divergence is that discovery innovation has been driven by computational methods that are relatively cheap to deploy and iterate, while development innovation requires touching the actual trial infrastructure: sites, patient populations, regulatory submissions, comparator data sets, endpoint definitions. That stuff is slow, bureaucratic, and deeply institutional. It’s not the kind of thing you can iterate on quickly with gradient descent.

What makes this moment distinct is that multiple academic and regulatory threads are now converging on a shared diagnosis, and regulators themselves are beginning to acknowledge that the evidence generation stack needs to be rebuilt rather than patched. That combination – a growing pipeline of AI-discovered candidates, a strained development infrastructure, and a regulatory environment signaling conditional openness to new methodologies – is the setup for a very large business opportunity.

The technical stack academia is quietly building

The most interesting thing happening in the academic literature right now isn’t in discovery papers. It’s in the clinical and translational work on external controls,

federated learning, digital twins, and trial generalizability. These are the papers tend to get cited in regulatory guidance but not in Andreessen Horowitz market which is itself a signal worth paying attention to.

Start with external controls, which are quickly moving from statistical afterthought to computable infrastructure. The FDA's 2025 draft guidance on externally controlled trials is notable not because it blesses the approach uncritically, but because of what it demands. External control and trial populations must be closely matched on baseline characteristics, disease severity, standard of care, prognostic factors, outcomes definitions, and follow-up procedures. That sounds like standard epidemiology, but if you read it more carefully and what it describes is a substantial technical stack: phenotype normalization across source systems, covariate harmonization across sites and time periods, temporal alignment to account for different eras of standard of care, endpoint ontology mapping, and robust causal adjustment for measured confounders. Building this reliably and in a way that satisfies an FDA statistician under adverse review conditions is genuinely hard.

The more interesting development on the external control front is federated. A 2024 Nature Communications paper introducing FedECA presents a federated learning approach for constructing external control arms with time-to-event outcomes in distributed data settings. The motivation is important: most of the data you'd want to use as external comparators lives in health systems that can't or won't share raw records. Privacy law, data use agreements, institutional politics – all of it creates friction that centralized data pooling approaches hit immediately. Federated methods that can train and validate comparator models without moving raw data aren't just a technical curiosity; they're a necessary condition for scaling this infrastructure across the real-world data landscape.

The entrepreneurial implication here is underappreciated. The most valuable evidence company of the next cycle might not be a CRO or a data broker. It might be a privacy-preserving cross-institution evidence network – something like a distributed proof system for regulatory comparators – that can assemble external control cohorts without ever centralizing the underlying records. That's a different business model than what most people are building.

On generalizability, a 2025 Nature Medicine study called TrialTranslator is one of the cleaner demonstrations of the problem and the solution space. The authors show that real-world survival outcomes for oncology patients are often substantially worse than RCT results, citing median overall survival reductions of roughly six months in some settings from prior work. They also find that about one in five real-world oncology patients wouldn't qualify for a phase 3 trial, and they argue that eligibility criteria alone don't explain the gap – selective recruitment of better-prognosis patients into trials is also a contributor. Their methodological response is to use ML-based prognostic phenotyping combined with trial emulation to test how well landmark RCTs generalize across different real-world risk strata.

The investment framing here is worth sitting with. If a trial result isn't a scalar – but a distribution over subpopulations and care settings, some of which see something close to the trial effect and some of which see much less – then the ability to systematically quantify where a drug's efficacy attenuates in the real world is genuinely valuable. It changes launch forecasting. It changes payer strategy. It changes medical affairs planning. It potentially changes M&A valuation in situations where the sponsor is trying to figure out how much of a phase 3 result will survive contact with real-world prescribing patterns. Trial informatics reframed as asset diligence infrastructure is a very different market than trial informatics reframed as work automation.

Digital twins are the most overhyped piece of this stack, and also the one that's most consequential if the validation problem gets solved. The marketing use of “digital twin” in health and life sciences covers a huge range of things, most of which are not twins in any meaningful modeling sense. But the academic literature is getting more specific. A 2025 npj Systems Biology and Applications paper argues that properly constructed twins can improve trial efficiency, enable early adverse event detection and contribute to adaptive trial design – while also flagging serious ethical and validation concerns that remain unresolved. A translational example cited in the npj Drug Discovery review involved systems-based digital twins in a Phase I mosunetuzumab study in NHL to characterize dose-response and identify predictive biomarkers.

biomarkers, which then informed adaptive dosing protocols. That's a concrete clinical application, not a marketing slide.

The catch, which any serious investor should dwell on, is that twin technology is investable at scale if verification, validation, and uncertainty quantification (VVUQ) matures. A 2025 npj Digital Medicine survey on this topic makes the point clear: VVUQ is not optional infrastructure for a twin that's going to support safety- and efficacy-relevant decisions in a regulated environment. It's the central requirement. The moat in digital twins isn't the simulation demo. It's the validation framework, uncertainty propagation methodology, and the decision thresholding protocol that makes the model usable under regulatory scrutiny. That's a much more defensible, much harder thing to build, and it's not what most current twin vendors are selling.

On recruitment and patient matching, the TrialGPT paper published in Nature Communications in 2024 introduced an end-to-end LLM framework for zero-shot patient-to-trial matching. Subsequent 2025 and 2026 work has pushed into multimodal inputs, ingesting structured EHR data alongside clinical notes and chart artifacts. Progress is real. But the subtler argument is that recruitment alone is probably too narrow a moat for a durable infrastructure business unless it expands into a broader trial-state architecture: protocol criteria management, eligibility evidence tracking, missing data handling, site capability assessment, patient longitudinal state representation, and expected screen-failure risk prediction. The companies that capture enrollment as an entry point and then build upward into a broader state graph are the ones worth watching.

A 2025 npj Digital Medicine scoping review that examined 142 AI-in-trials studies from 2013 through 2024 found growing AI use across safety monitoring (55 studies), efficacy assessment (46), and operational risk management (45). But the same review makes clear that most of this work is retrospective, many datasets are nonrepresentative, and prospective reproducible evidence for these tools is still limited. That's both a problem and an opportunity. The winning businesses here will be the ones with the strongest benchmark on a retrospective dataset. They'll be the ones that have figured out prospective deployment, label scarcity, site workflow integration, and auditability under regulatory review. Those are organizational and

technical challenges simultaneously, which is exactly the kind of problem that's to replicate quickly.

Finally, on platform and adaptive trial architecture, a 2024 Signal Transduction Targeted Therapy review makes the case that basket, umbrella, and platform trial designs are increasingly aligned with the biological logic of precision medicine as multi-omics and sequencing make molecular stratification more granular. Platform trials can dynamically add and drop arms, incorporate biomarker-guided stratification, and use interim evidence to adapt allocation rules. Once the trial architecture becomes this dynamic and biomarker-dependent, software isn't back office tooling anymore. It becomes part of the protocol logic itself. That's a different market than generic eClinical infrastructure, and it's more likely to support businesses with real defensibility.

Regulators are opening the door but raising the bar

The FDA's posture over the last two years is worth reading carefully because it's misread in two opposite directions simultaneously. One camp thinks the agency is becoming more conservative and therefore all this novel methodology work is speculative. The other camp thinks the agency is embracing new tools enthusiastically and that regulatory friction is going away. Both camps are wrong.

What the agency is actually doing is more nuanced: it's conditionally opening the door to external controls, real-world data, digital health technologies, and model-assisted endpoints while simultaneously raising the evidentiary bar for how those tools must be validated and deployed. The 2025 externally controlled trials draft guidance is the clearest example. The agency isn't saying don't use external comparators. It's saying here is the technical specification for what a regulatory-grade external comparator actually requires, and that specification is demanding. The same pattern shows up in FDA guidance on digital health technologies for drug development and in the framework for AI and machine learning in drug development – conditional open

combined with explicit expectations around transparency, validation, and human oversight.

That combination is exactly what creates a wedge for infrastructure companies. If a regulator says “we will accept this approach if you can demonstrate X, Y, and Z” and X, Y, and Z are technically hard and require significant organizational capability to demonstrate, then whoever builds the infrastructure to demonstrate X, Y, and Z reliably becomes a necessary part of every program that wants to use these approaches. That’s a strong structural position.

The European Medicines Agency has been running parallel threads on adaptive and real-world evidence that broadly reinforce this direction, and the International Council for Harmonisation’s E6(R3) update to GCP guidance signals a similar move toward risk-based monitoring and more flexible trial architectures. The regulatory environment isn’t creating a free pass. It’s creating a technical specification for a new generation of development infrastructure, and that specification is detailed enough that building to it is a real engineering problem.

What this means for founders, CROs, and venture underwriting

The practical implications of this analysis split across three groups: founders deciding what to build, CROs thinking about where their business model goes, and VCs figuring out where the durable value accumulates.

For founders, the clearest takeaway is to think carefully about which part of the evidence stack you’re building and whether it’s a wedge or a destination. Recruitment is a real problem and a reasonable entry point, but it’s also the most crowded and most likely to get commoditized by EHR vendors and large CROs with existing relationships. The more durable positions are likely in the layers that sit beneath recruitment: comparator construction, phenotype infrastructure, continuous measurement, model validation, and adaptive protocol software. These are harder to build, slower to sell, and less intuitive to demo, but they also sit closer to the act

evidence package that a sponsor needs to file, which means they're harder to cut the budget when things get tight.

The federated architecture point deserves particular emphasis for founders. The that would make the best external comparators – rich longitudinal records from health systems covering diverse patient populations – is almost never going to be centralizable in a single commercial database. The legal, political, and competitive dynamics are too complicated. The companies that figure out how to generate regulatory-grade evidence from distributed data without pooling raw records are building something that addresses a structural constraint rather than a workflow inefficiency, and structural constraints tend to create better businesses.

For CROs, this is both an opportunity and an existential threat depending on how they respond. The large CROs have site relationships, operational infrastructure, regulatory track records that are genuinely hard to replicate. But their core competency – running trials as currently defined – is increasingly being challenged by the question of whether trials as currently defined are the right design in the first place. If adaptive master protocols with dynamic arm allocation and biomarker-driven enrollment become more common, the operational model of the large CROs needs to change significantly. The CROs that are building or acquiring the computational infrastructure for external controls, digital endpoints, and adaptive design are positioning well. The ones that are treating software as a commodity layer they can outsource are making a mistake.

For VCs, the most important framing shift is to stop thinking about this market as “for trials” and start thinking about it as evidence infrastructure, which is more analogous to financial market data infrastructure than to SaaS workflow tools. The businesses with the highest long-run value in financial data didn't build fancy analytics. They built the underlying data network – the feeds, the normalization, the audit trail, the compliance layer – that every analytics product depended on. The analogous businesses in clinical development will be those that own the provenance chain from raw clinical and real-world data to regulatory-grade evidence submissions. That chain involves a lot of unglamorous infrastructure: source data verification

protocol versioning, endpoint adjudication documentation, uncertainty reporting, human override tracking. None of it is conference-talk material. All of it is necessary.

The five infrastructure layers that matter

To make this concrete, here's where the category-defining businesses of the next development cycle are most likely to form.

The first layer is comparator infrastructure: external controls, target trial emulation, and federated evidence networks. This is arguably the most technically demanding and the most immediately actionable given the FDA's draft guidance providing an explicit requirements specification. The players that build federated comparator networks with real governance, real privacy architecture, and real regulatory trail records will own a capability that every sponsor running a single-arm trial or a randomized study needs access to.

The second layer is phenotype infrastructure: multimodal risk stratification, biomarker enrichment, and generalizability scoring. The TrialTranslator work points to this directly. If trial results are distributions over subpopulations rather than scalars, the ability to characterize those distributions systematically – using ML phenotyping across structured and unstructured data – creates value at multiple stages of development and commercialization. This layer is also closely connected to the precision medicine thesis, since the biomarker enrichment work that makes platform trials more efficient is the same capability that makes generalizability assessment more rigorous.

The third layer is continuous measurement infrastructure: digital health technologies, remote endpoint capture, and sensor-grade clinical measures. The FDA has been investing significant guidance bandwidth in digital health technologies for drug development, and the clinical community has been developing validation frameworks for patient-reported outcomes and wearable-derived endpoints. The infrastructure play here is less about the device itself and more about the data architecture, validation documentation, and endpoint certification pathway that turns a device-generated signal into a regulatory-grade clinical endpoint.

The fourth layer is model assurance infrastructure: verification, validation, uncertainty quantification, and auditability for twins and predictive models. This is currently the most underbuilt part of the stack. Lots of companies have models, but few have the frameworks to demonstrate that their models behave reliably under distribution shift, can propagate uncertainty in ways that support decision thresholding, and maintain an audit trail that satisfies regulatory review. That gap is the bottleneck for digital twins specifically but also for any predictive model that wants to be used in a regulated decision context.

The fifth layer is adaptive protocol infrastructure: software for master protocols, dynamic allocation, biomarker-driven arm management, and interim evidence integration. As the trial architecture itself becomes more computationally native software that manages the protocol logic – not just the eClinical workflow but the actual decision rules for arm allocation, stopping, and enrichment – becomes critical path. This is still early, but the academic momentum behind basket, umbrella, and platform designs combined with the FDA's growing comfort with adaptive design suggests the market is building.

The unified thesis across all five layers is the same: AI is industrializing hypothesis generation faster than the industry can industrialize evidence generation. The molecule problem, while not fully solved, is no longer the primary constraint. The evidence problem – who can assemble trustworthy, regulatory-grade, causally defensible proofs that a drug does what it's supposed to do in patients who actually exist – is the constraint now. That's where the next generation of durable health-infrastructure companies will be built, and the window to form them is open.



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