

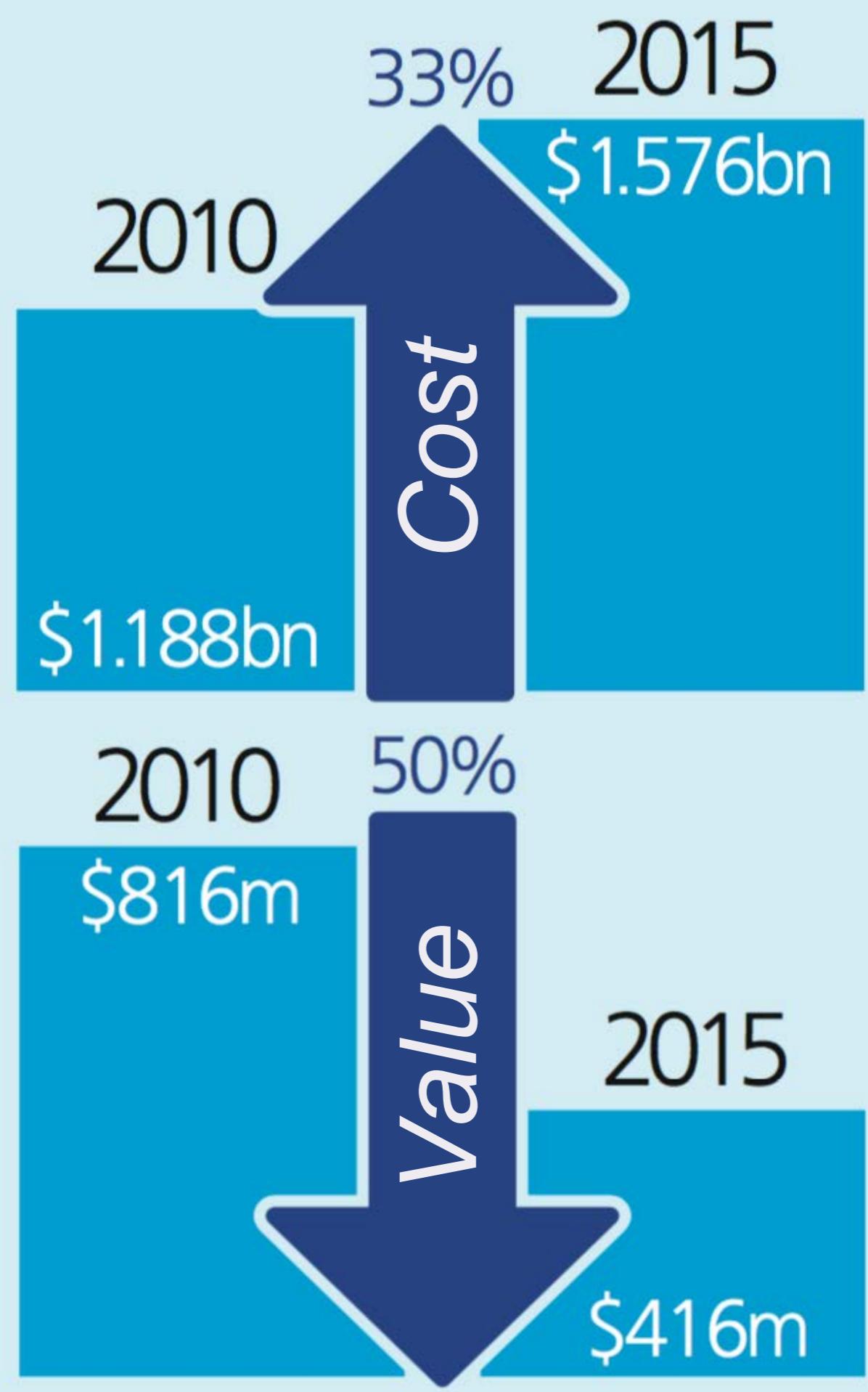
New Targets, New Modalities, New Challenges – The *Inconvenient* Path of Human Genetics in Drug Discovery

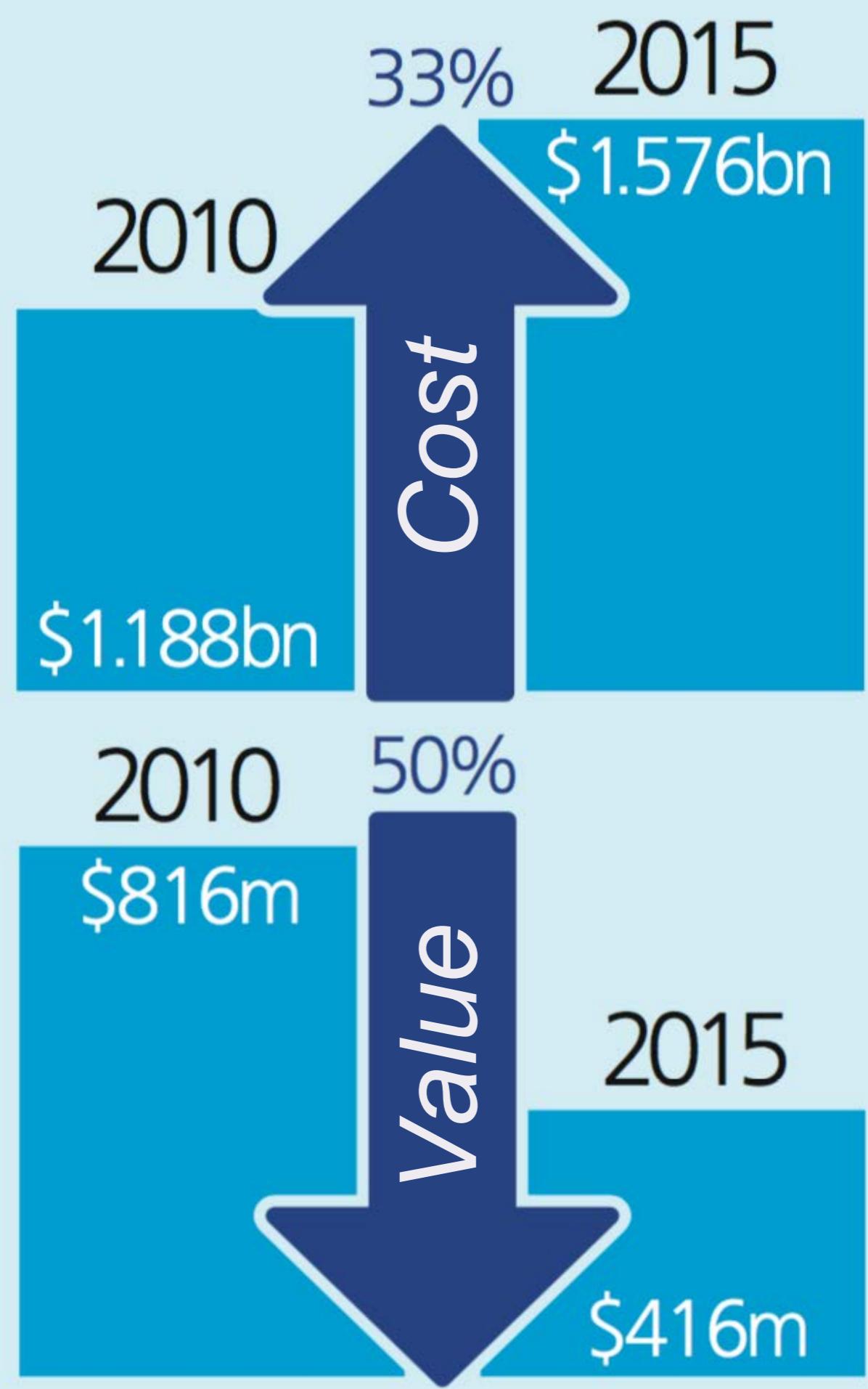
Enabling Precision Medicine: The Role of Genetics in Clinical Drug Development – A Workshop

March 8, 2017

Overall message

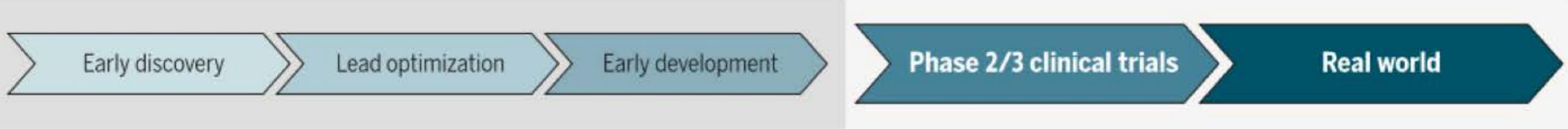
- ✿ precision medicine: *patient subsets for whom therapeutic intervention works better*
- ✿ few approved drugs will benefit from precision medicine
- ✿ greatest impact will be to guide new drug development, which will be tested and approved in patient subsets
- ✿ However, this path is *inconvenient*, and will require biological insight into targets, new therapeutic modalities, and a more creative approaches to clinical development





Disciplined approach to drug discovery and early development

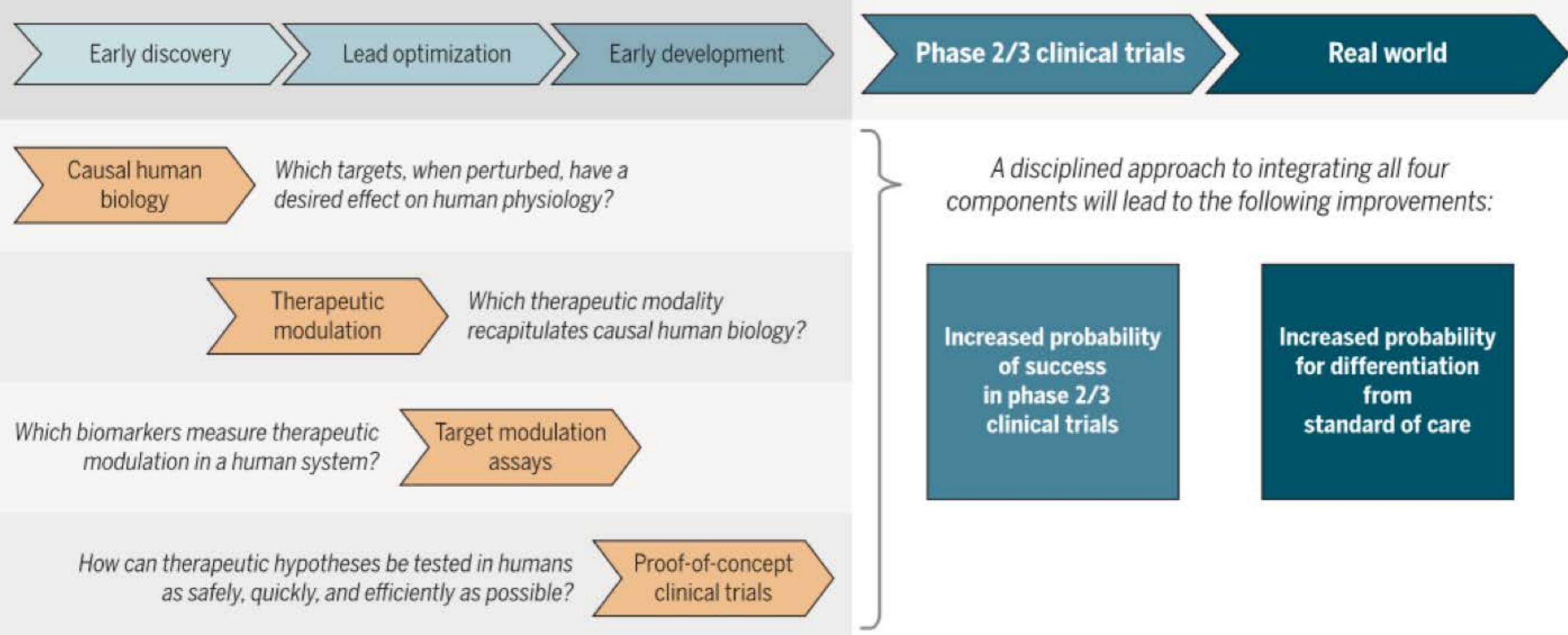
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Why is cost increasing?

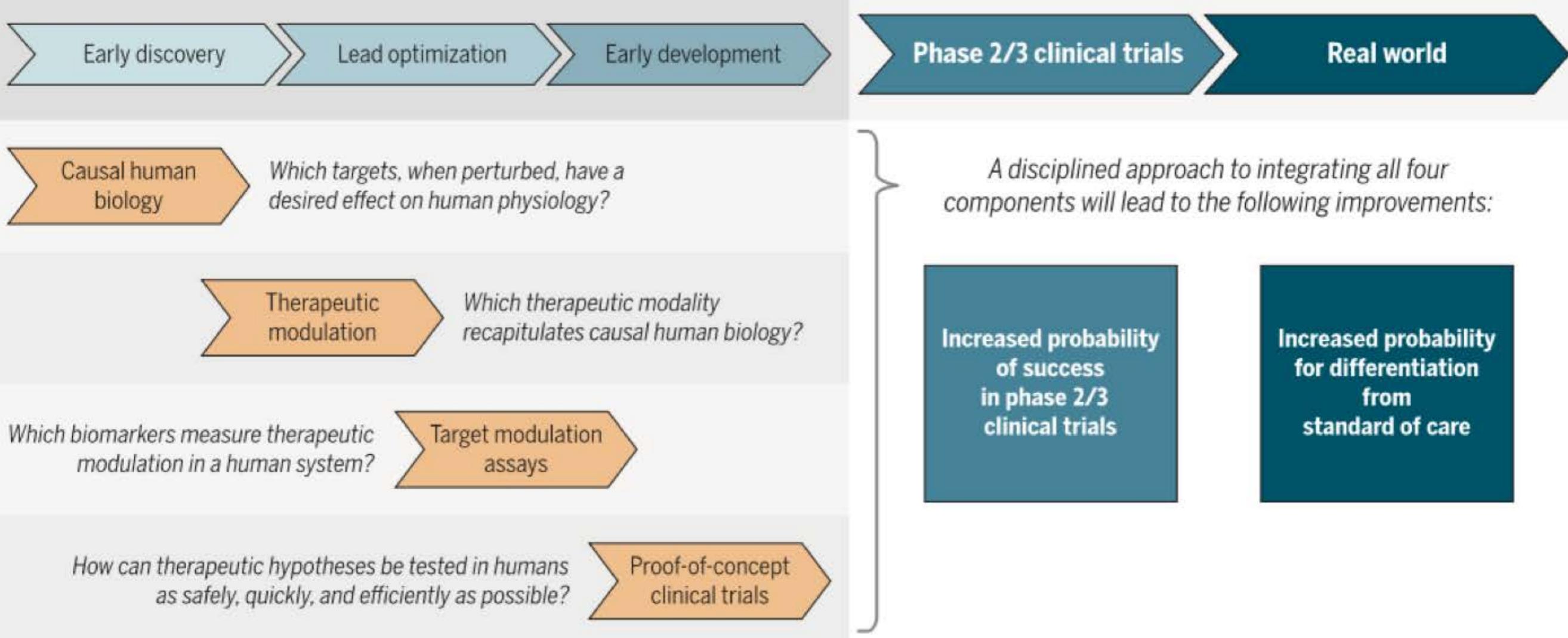


1. Too many Phase 2/3 studies fail
2. Time between discovery and PoC is too long

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Why don't drugs differentiate?

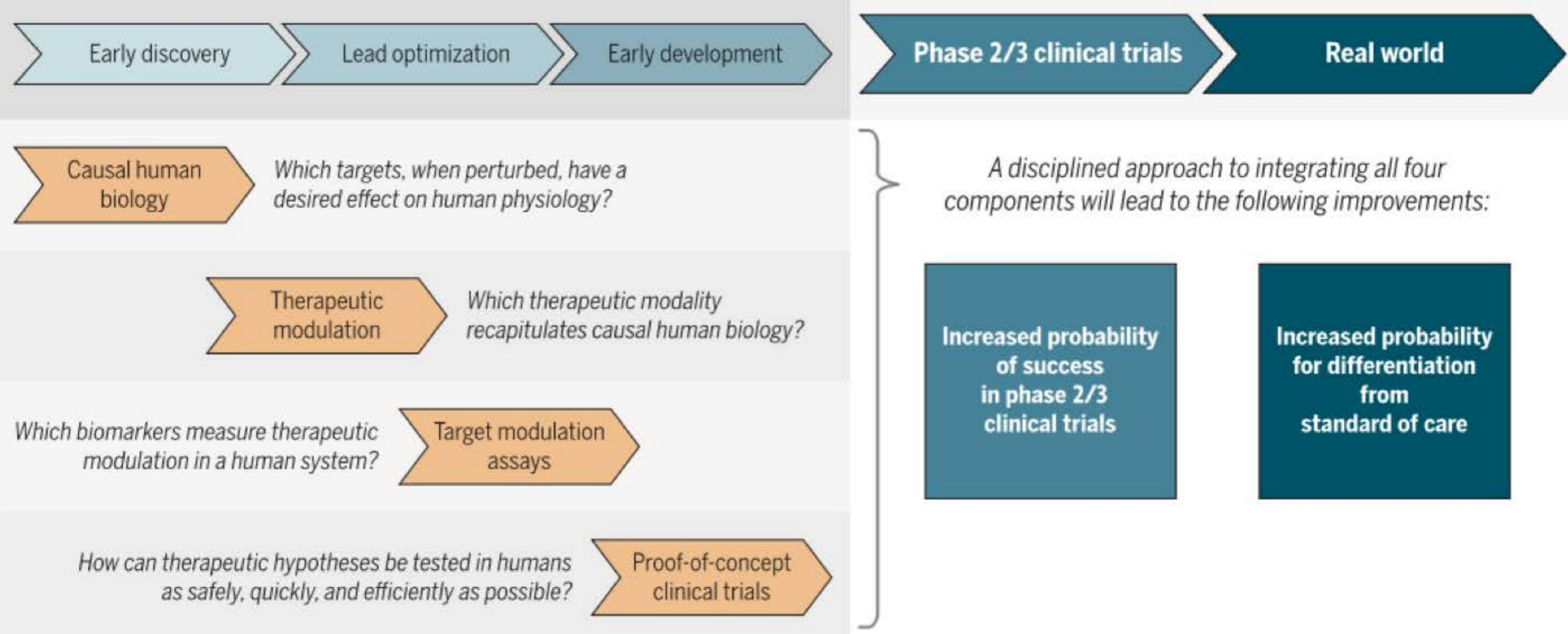


1. Not enough sound therapeutic hypotheses!

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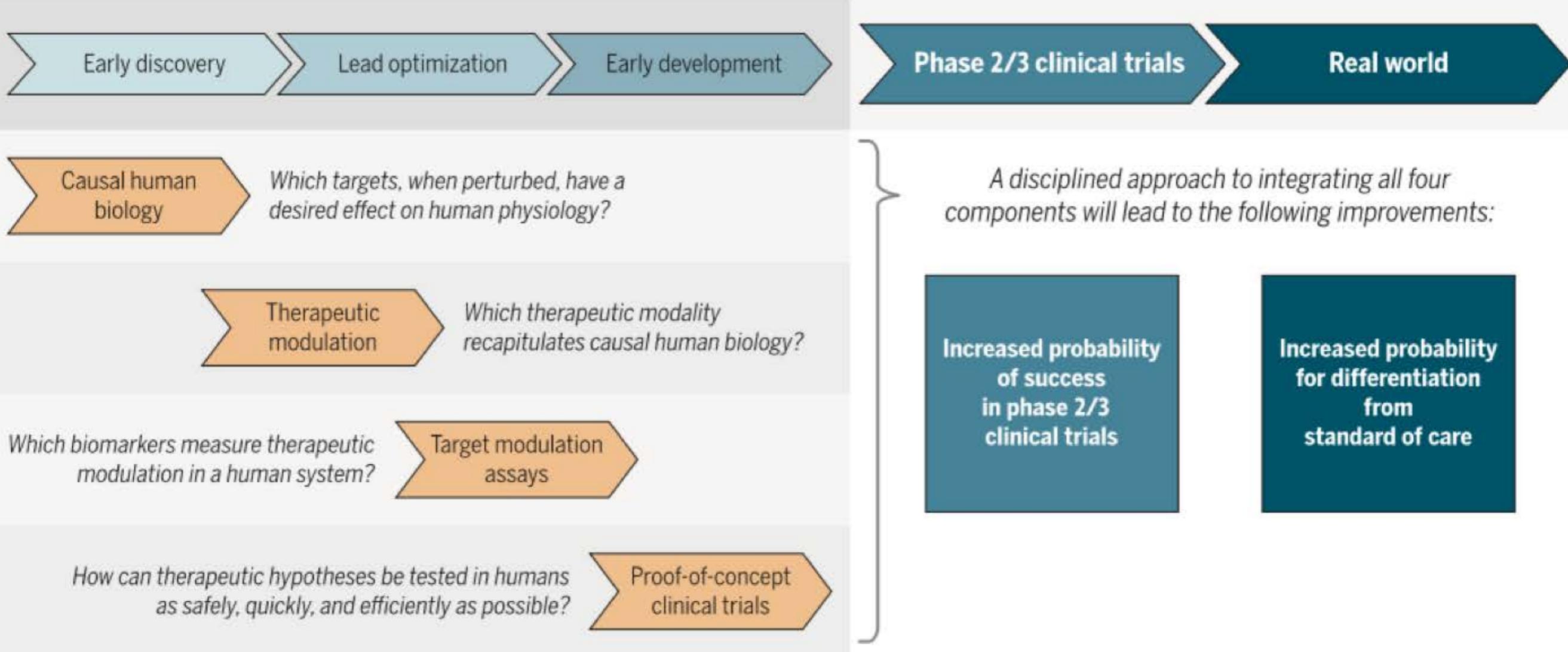
How will *precision medicine* help?



1. Improve cycle time from therapeutic hypothesis to clinical PoC... *efficiently and safely*

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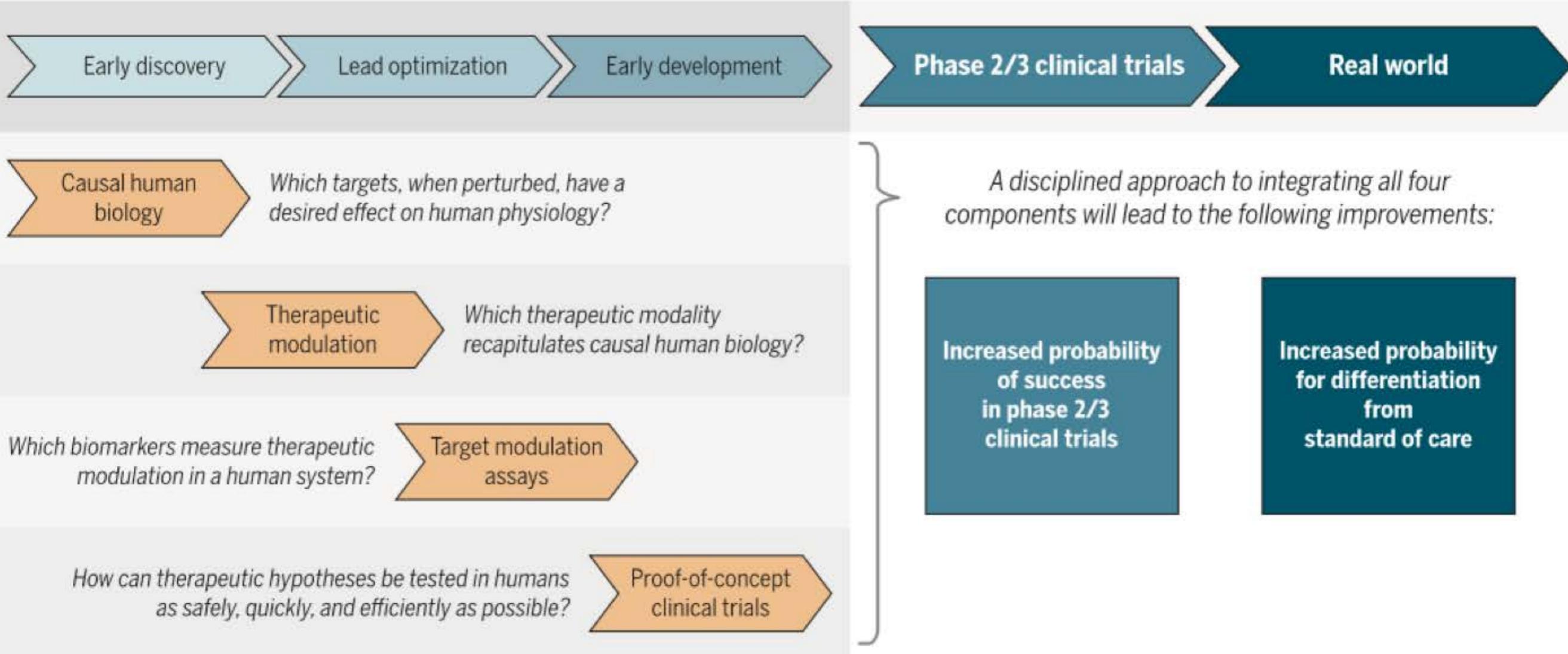
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Today: 5-7 years

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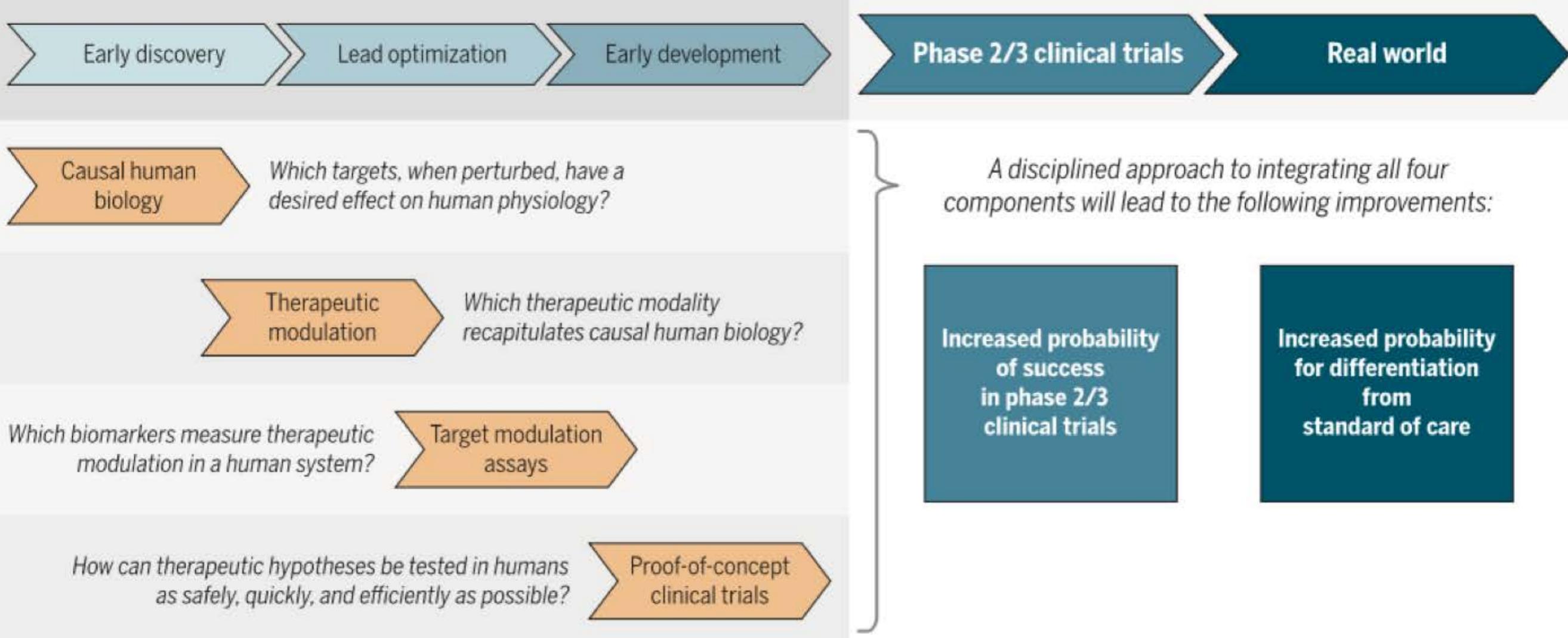


<1 yr

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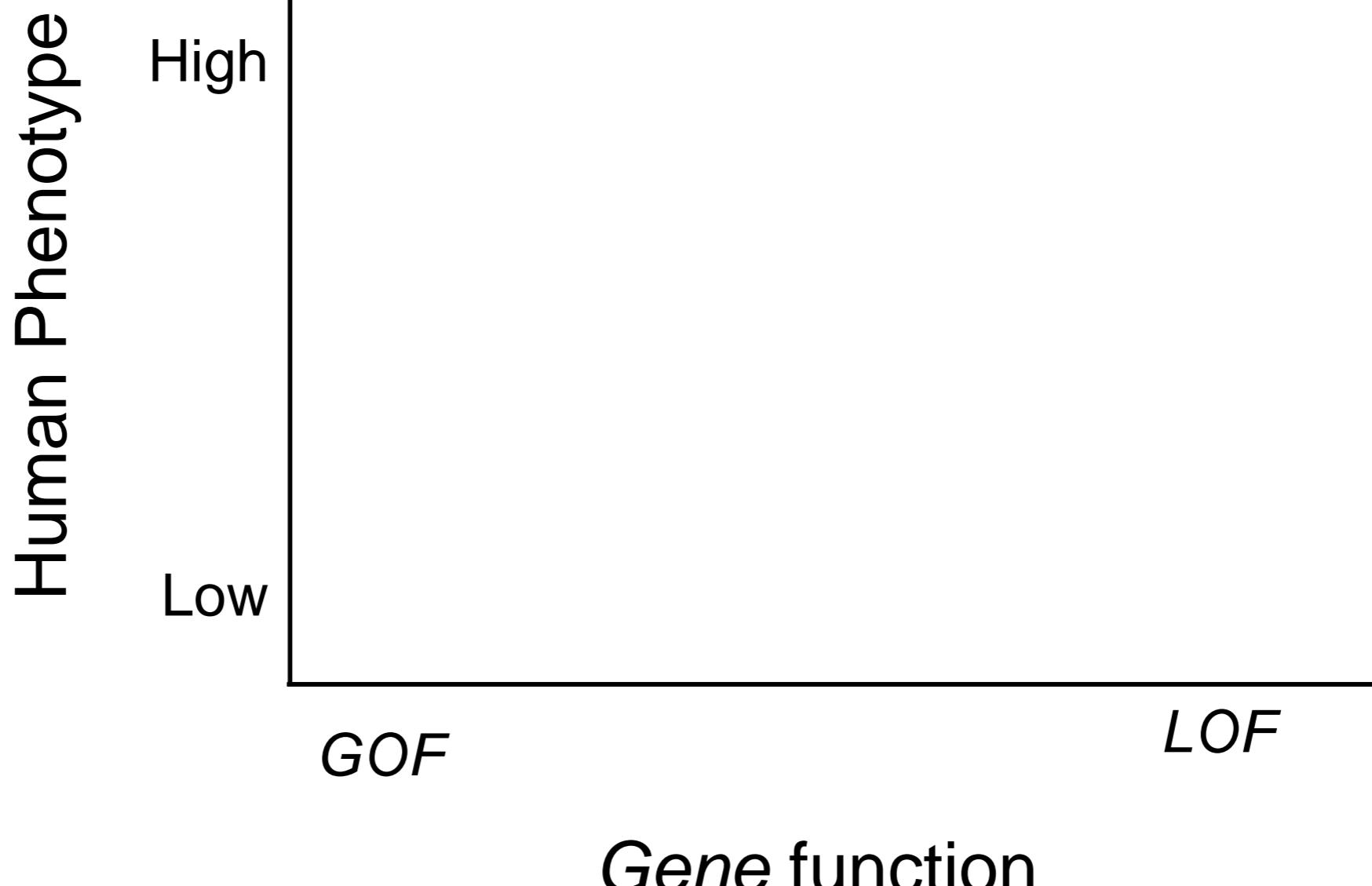
What are potential solutions?



1. Pick targets based on human biology
2. Programmable therapeutics to test PoC

human genetics to pick targets

Pick a human phenotype for drug efficacy



Pick a human phenotype for drug efficacy

Human Phenotype

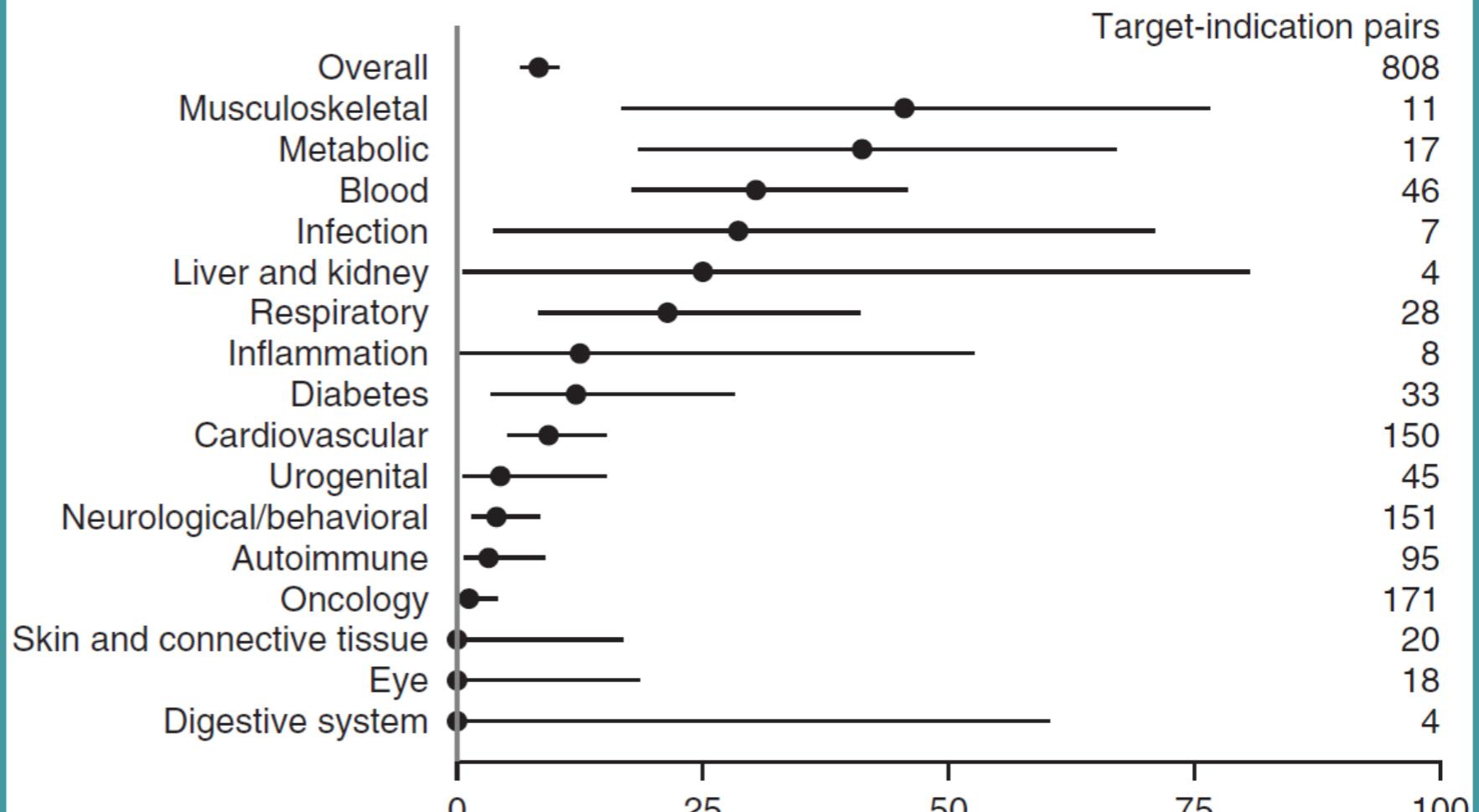
High
Low

GOF

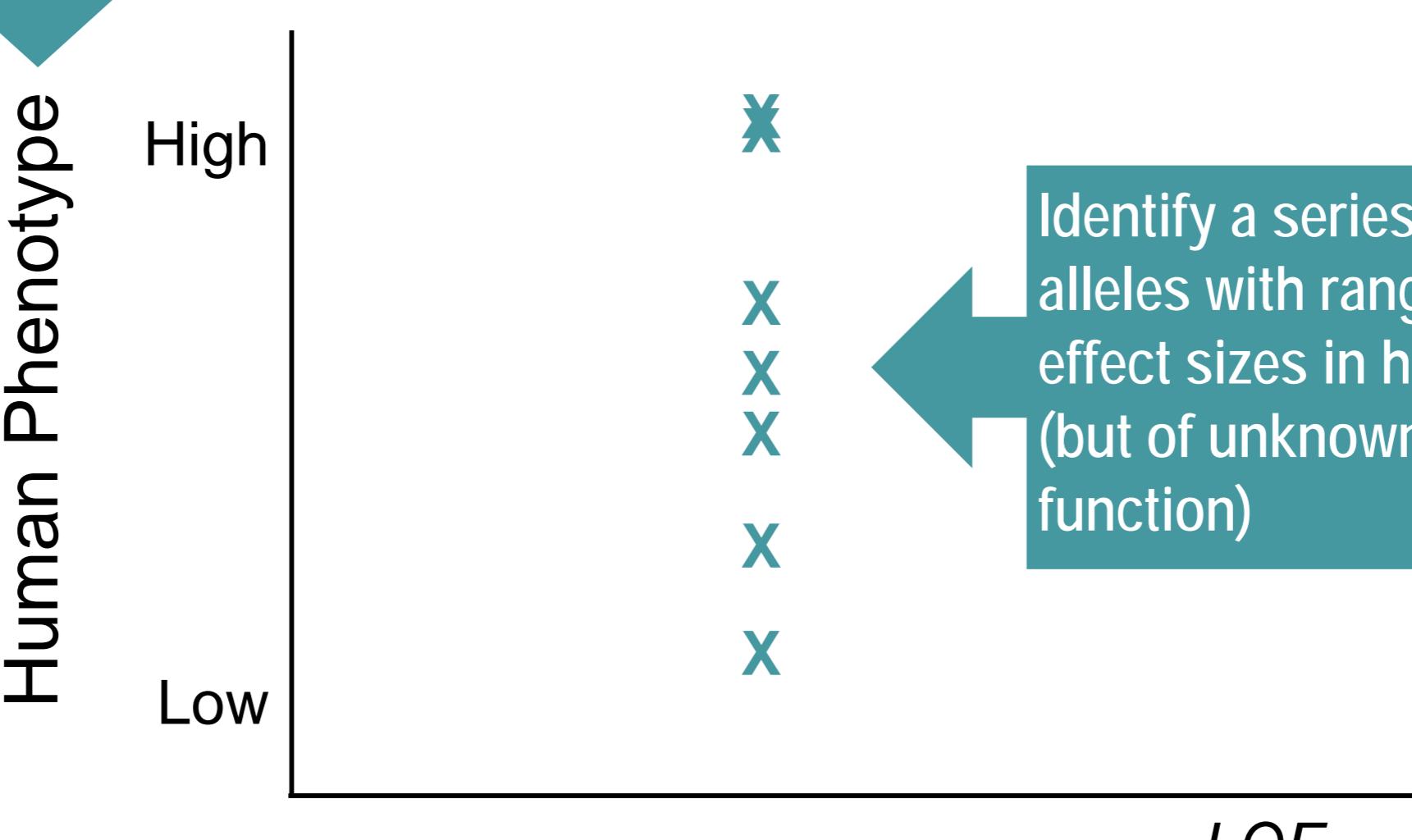
LOF

Gene function

Nelson et al *NG* 2015



Pick a human phenotype for drug efficacy



Identify a series of alleles with range of effect sizes in humans (but of unknown function)

Gene function

Association of Rare and Common Variation in the Lipoprotein Lipase Gene With Coronary Artery Disease

Amit V. Khera, MD; Hong-Hee Won, PhD; Gina M. Peloso, PhD; Colm O'Dushlaine, PhD; Dajiang Liu, PhD; Nathan O. Stitziel, MD, PhD; Pradeep Natarajan, MD; Akihiro Nomura, MD; Connor A. Emdin, DPhil; Namrata Gupta, PhD; Ingrid B. Borecki, PhD; Rosanna Asselta, PhD; Stefano Duga, PhD; Piera Angelica Merlini, MD; Adolfo Correa, MD; Thorsten Kessler, MD; James G. Wilson, MD; Matthew J. Bown, MD; Alistair S. Hall, MD; Peter S. Braund, PhD; David J. Carey, PhD; Michael F. Murray, MD; H. Lester Kirchner, PhD; Joseph B. Leader, BA; Daniel R. Lavage, BS; J. Neil Manus, BS; Dustin N. Hartzel, BS; Nilesh J. Samani, MD; Heribert Schunkert, MD; Jaume Marrugat, MD, PhD; Roberto Elosua, MD, PhD; Ruth McPherson, MD; Martin Farrall, FRCPath; Hugh Watkins, MD, PhD; Eric S. Lander, PhD; Daniel J. Rader, MD; John Danesh, FMedSci; Diego Ardissono, MD; Stacey Gabriel, PhD; Cristen Willer, PhD; Gonçalo R. Abecasis, PhD; Danish Saleheen, MD; Frederick E. Dewey, MD; Sekar Kathiresan, MD; for the Myocardial Infarction Genetics Consortium, DiscovEHR Study Group, CARDioGRAM Exome Consortium, and Global Lipids Genetics Consortium

Pick a human phenotype for drug efficacy

Human Phenotype

High

Low

>100 rare variants
6 common variants

GOF

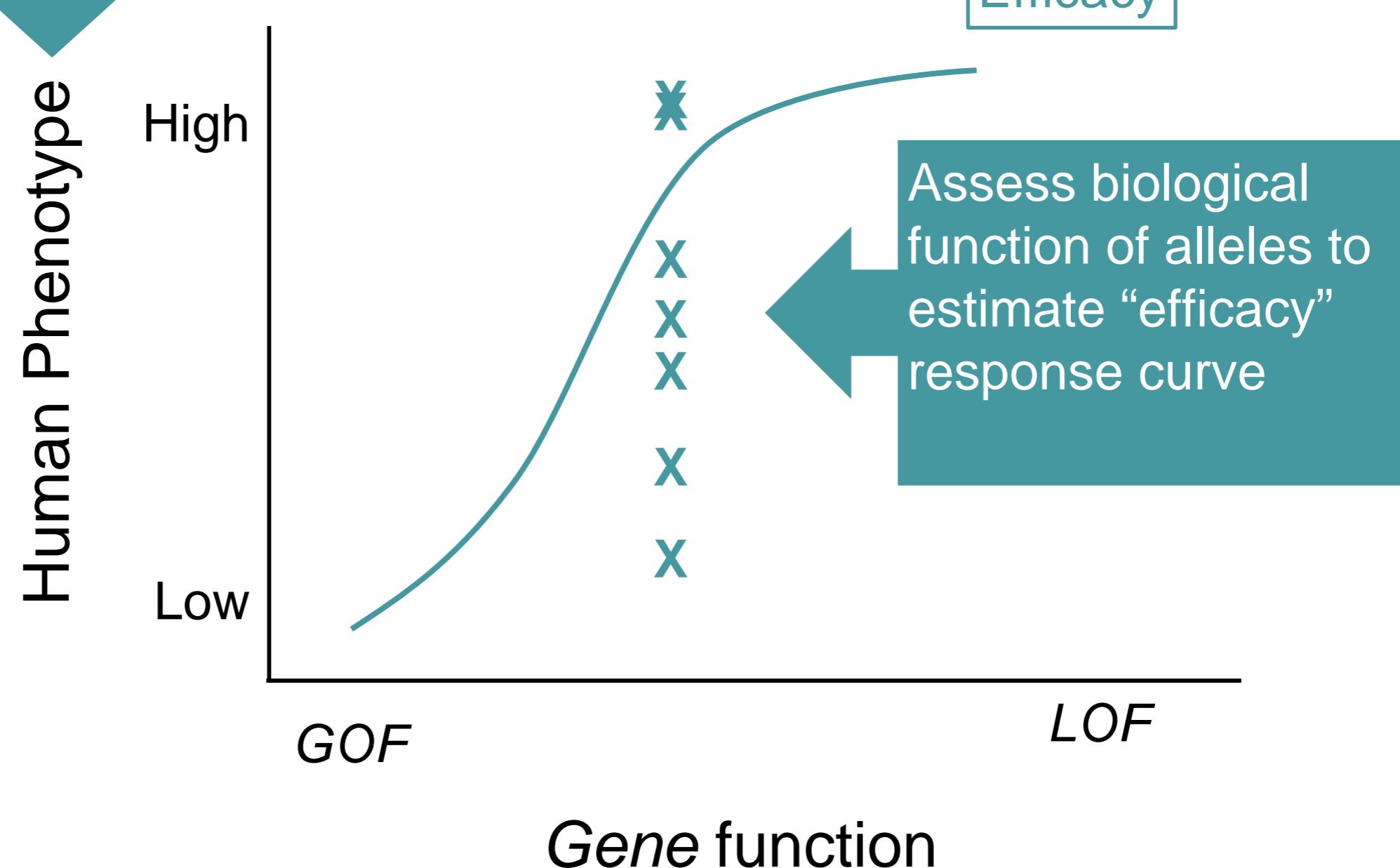
LOF

Gene function

Khera et al JAMA 2017

RESULTS Among 46 891 individuals with *LPL* gene sequencing data available, the mean (SD) age was 50 (12.6) years and 51% were female. A total of 188 participants (0.40%; 95% CI, 0.35%-0.46%) carried a damaging mutation in *LPL*, including 105 of 32 646 control participants (0.32%) and 83 of 14 245 participants with early-onset CAD (0.58%). Compared with 46 703 noncarriers, the 188 heterozygous carriers of an *LPL* damaging mutation displayed higher plasma triglyceride levels (19.6 mg/dL; 95% CI, 4.6-34.6 mg/dL) and higher odds of CAD (odds ratio = 1.84; 95% CI, 1.35-2.51; $P < .001$). An analysis of 6 common *LPL* variants resulted in an odds ratio for CAD of 1.51 (95% CI, 1.39-1.64; $P = 1.1 \times 10^{-22}$) per 1-SD increase in triglycerides.

Pick a human phenotype for drug efficacy



New target for drug screen!

Pick a human phenotype for drug efficacy

Human Phenotype

High
Low

GOF

LOF

Gene function

Efficacy

Toxicity

Assess biological function of target

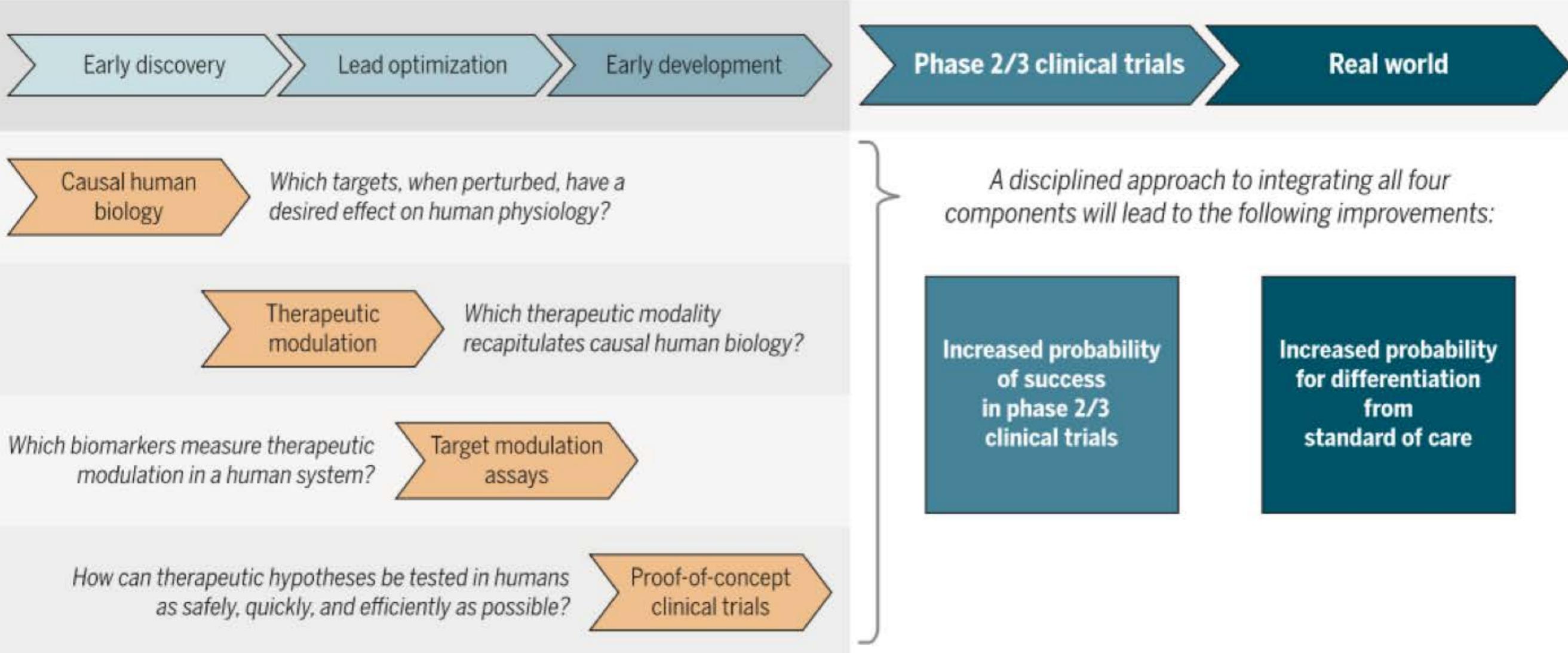
Assess pleiotropy as proxy for ADEs

This provides evidence for the therapeutic window at the time of target ID & validation.

programmable therapeutics to test
therapeutic hypotheses quickly

Disciplined approach to drug discovery and early development

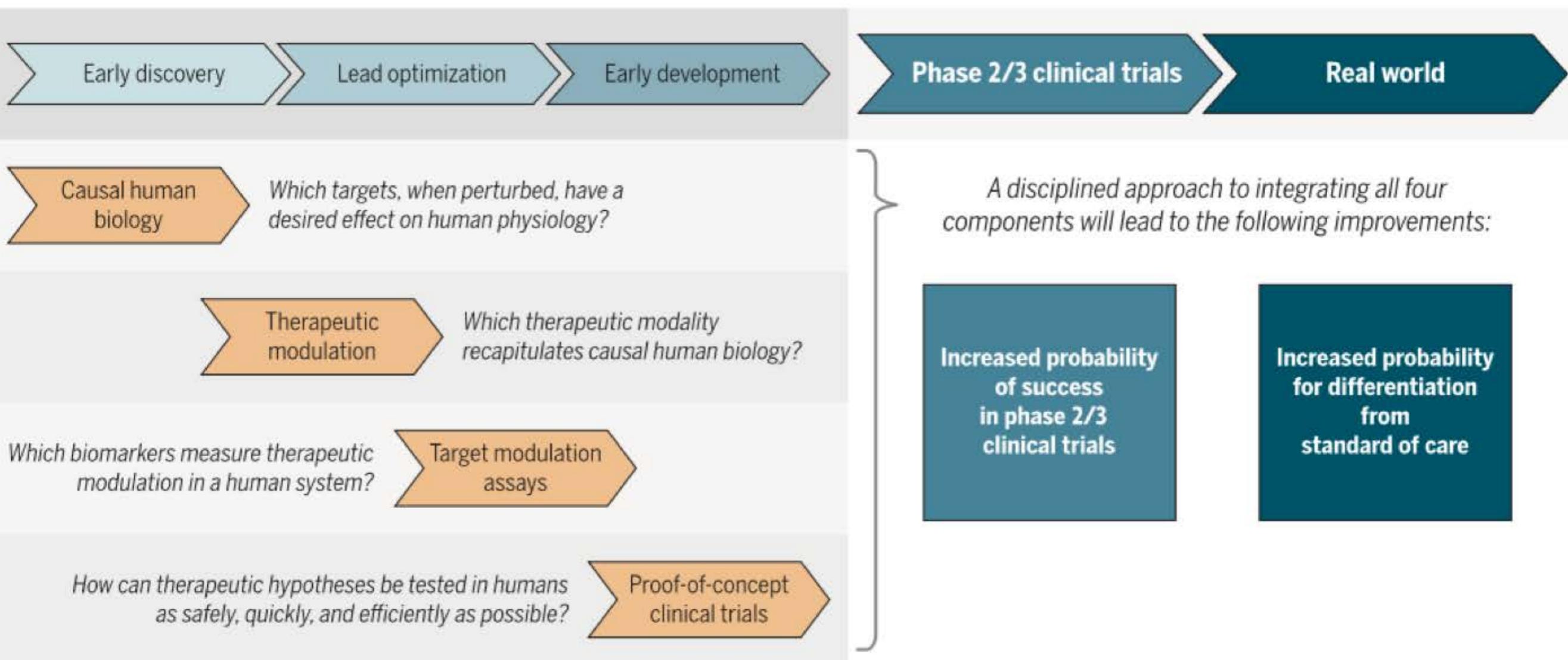
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Today: 5-7 years

Disciplined approach to drug discovery and early development

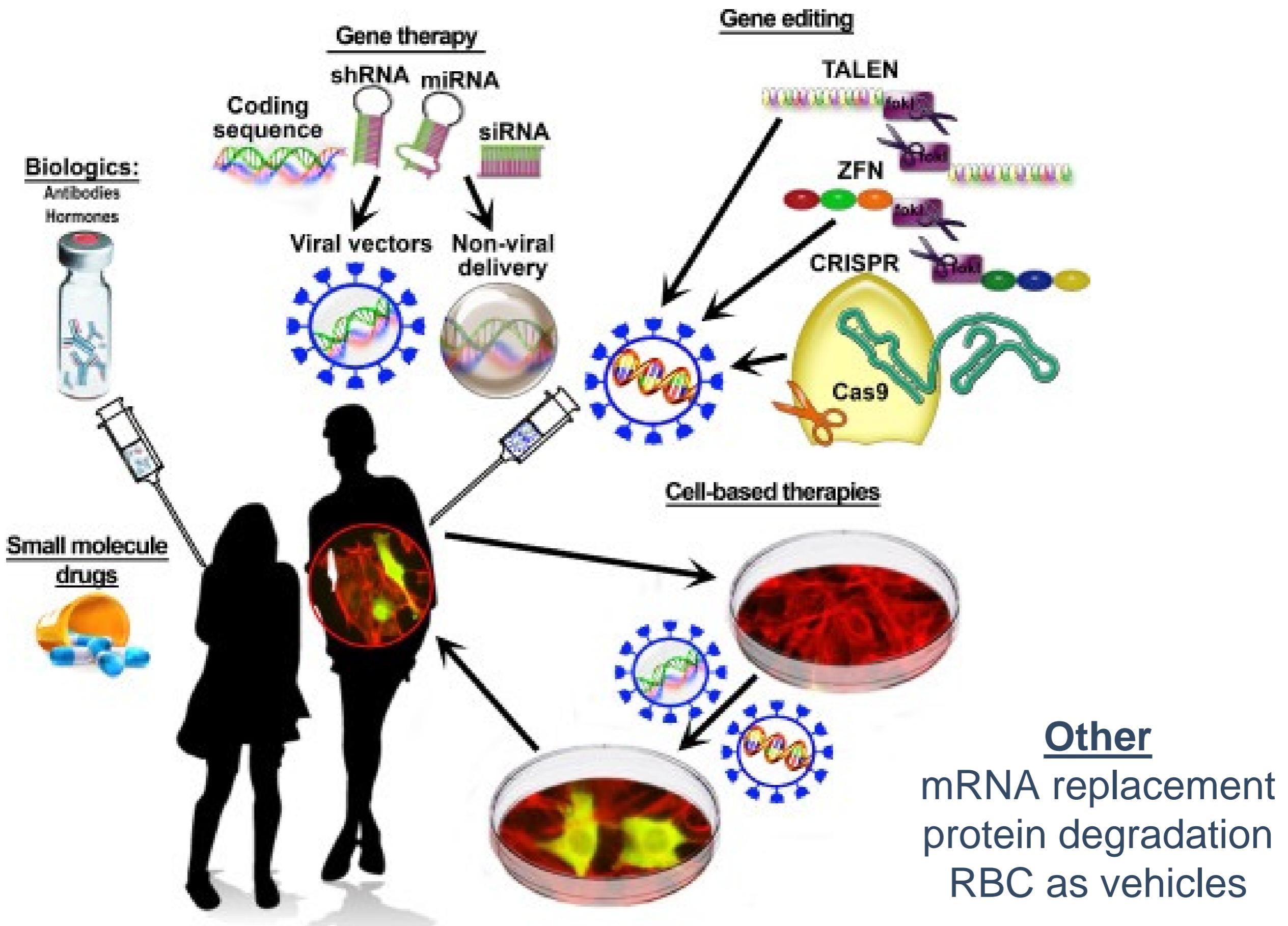
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<1 yr

Limitations that we face today

- ✿ Biological function of associated variants, genes & pathways incompletely understood
- ✿ Conventional modalities (e.g., small molecules, monoclonal antibodies) modulate <20% of targets
- ✿ ***New modalities are desperately needed***, but today are limited by delivery and pharmacological properties



Burning platform for precision medicine

- ✿ understand molecular mechanism of disease-associated variants, genes and pathways
- ✿ recapitulate mechanism with a credible therapeutic molecule
- ✿ shorten cycle time to test therapeutic hypotheses in small PoC trials defined by specific molecular features

Conclusion

Overall message - inconvenient path to precision medicine

- ✿ precision medicine: *patient subsets for whom therapeutic intervention works better*
- ✿ few approved drugs will benefit from precision medicine
- ✿ greatest impact will be to guide new drug development, which will be tested and approved in patient subsets
- ✿ However, this path is *inconvenient*, and will require biological insight into targets, new therapeutic modalities, and a more creative approaches to clinical development