

A stylized, light yellow human figure stands centrally against a light blue background. The figure is surrounded by several circular target symbols, each with a red outer ring and a blue center. The figure's arms and legs are slightly outstretched, and its torso is marked with faint red lines. The overall composition suggests a focus on human health and medical research.

# 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

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March 8, 2017

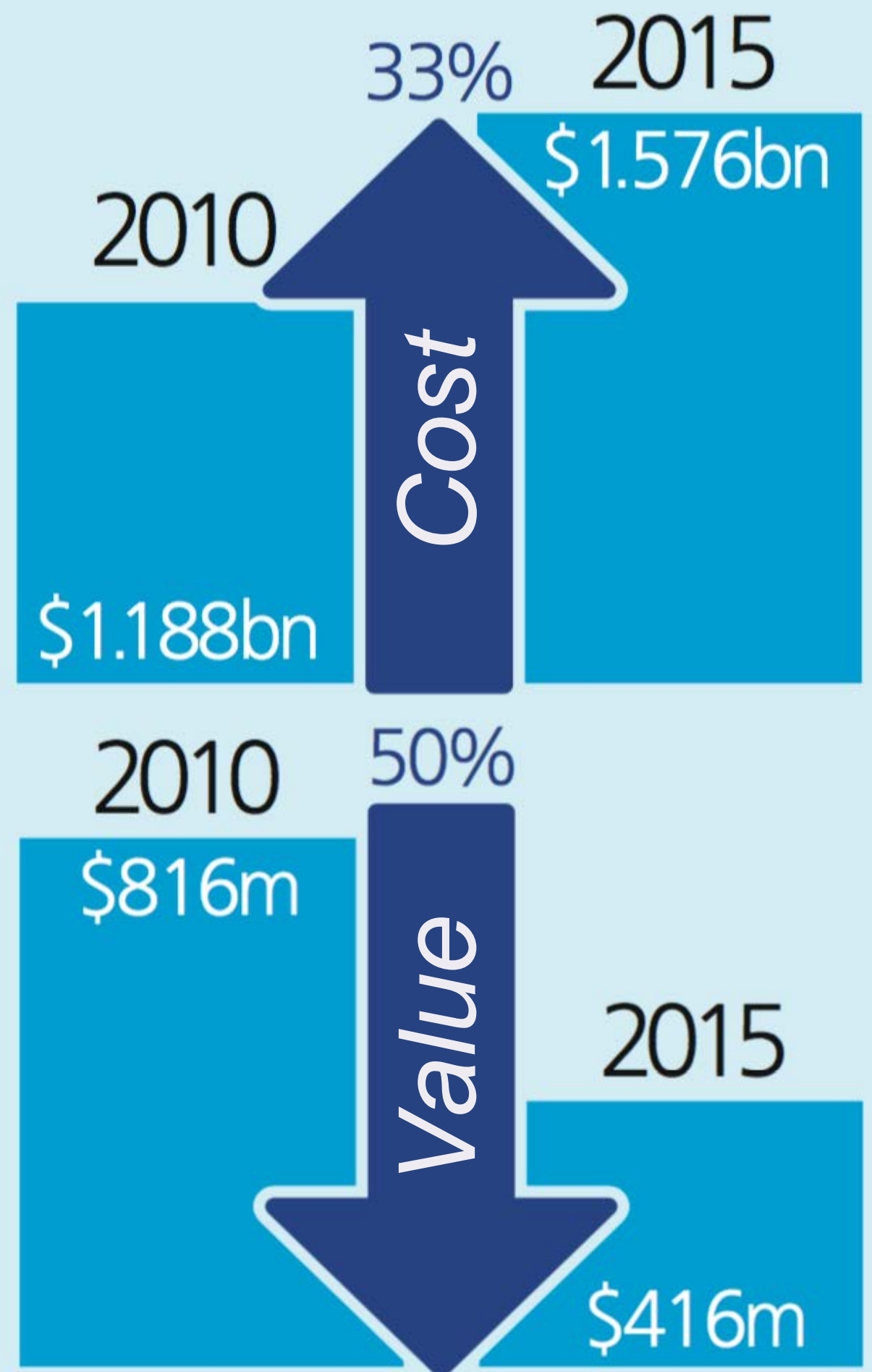


# Overall message

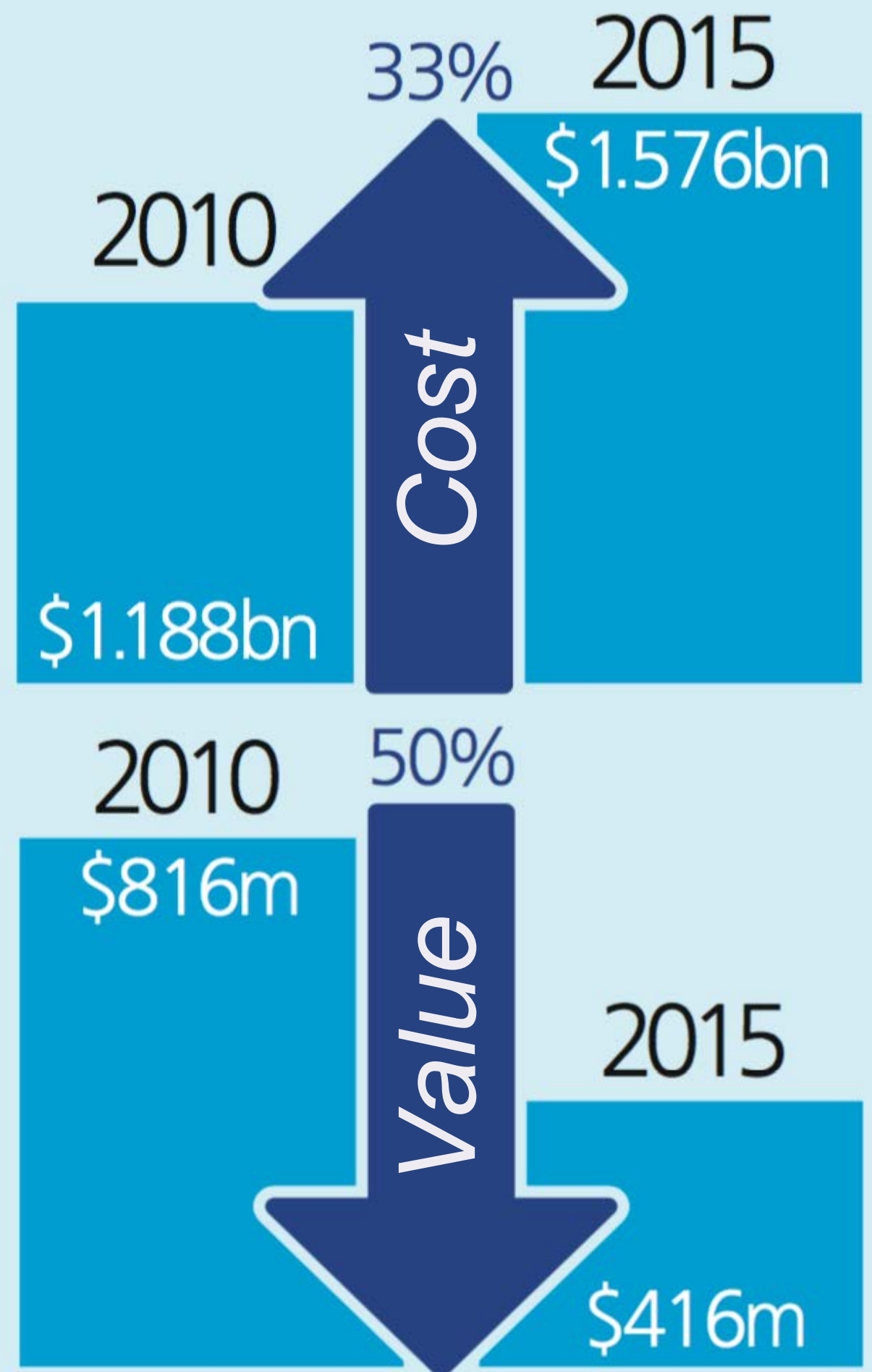
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- ❖ precision medicine: *patient subsets for whom therapeutic intervention works better*
- ❖ few approved drugs will benefit from precision medicine
- ❖ greatest impact will be to guide new drug development, which will be tested and approved in patient 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

Robert M. Plenge

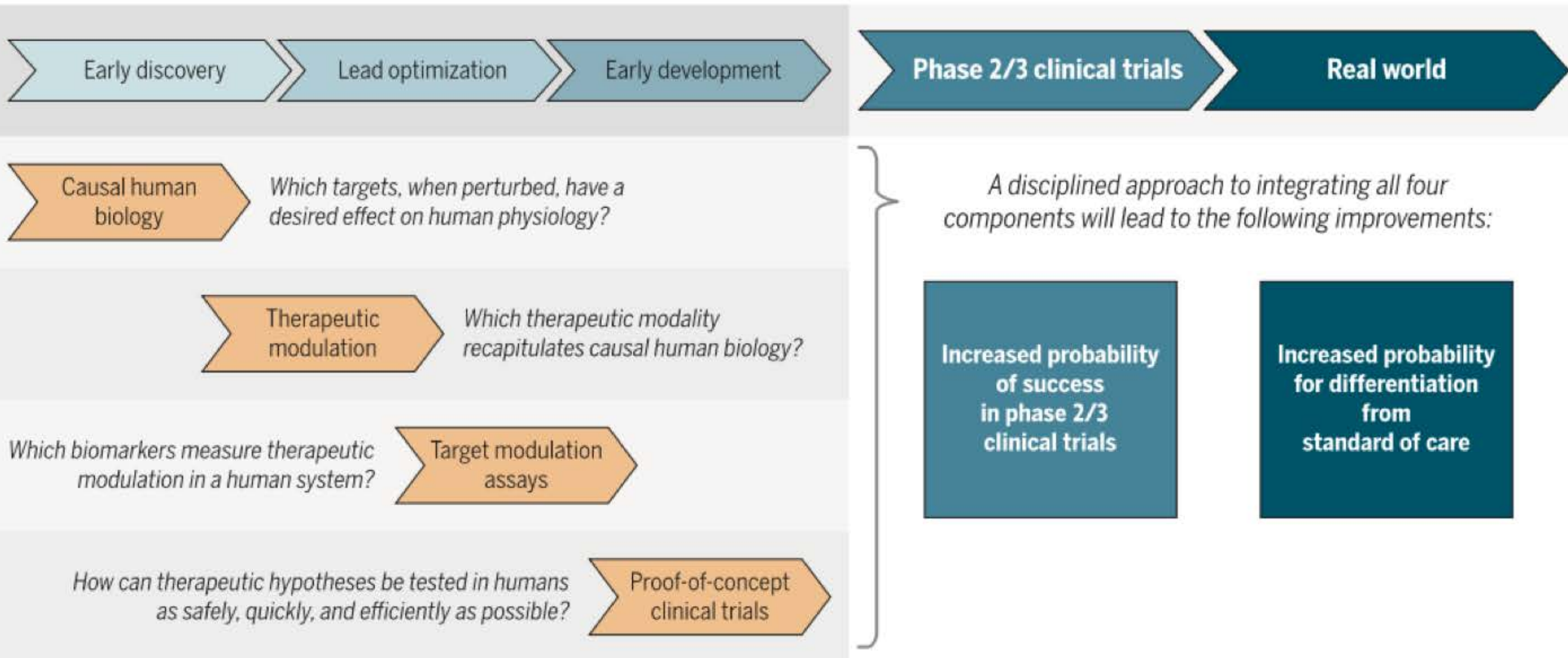




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# Why is cost increasing?



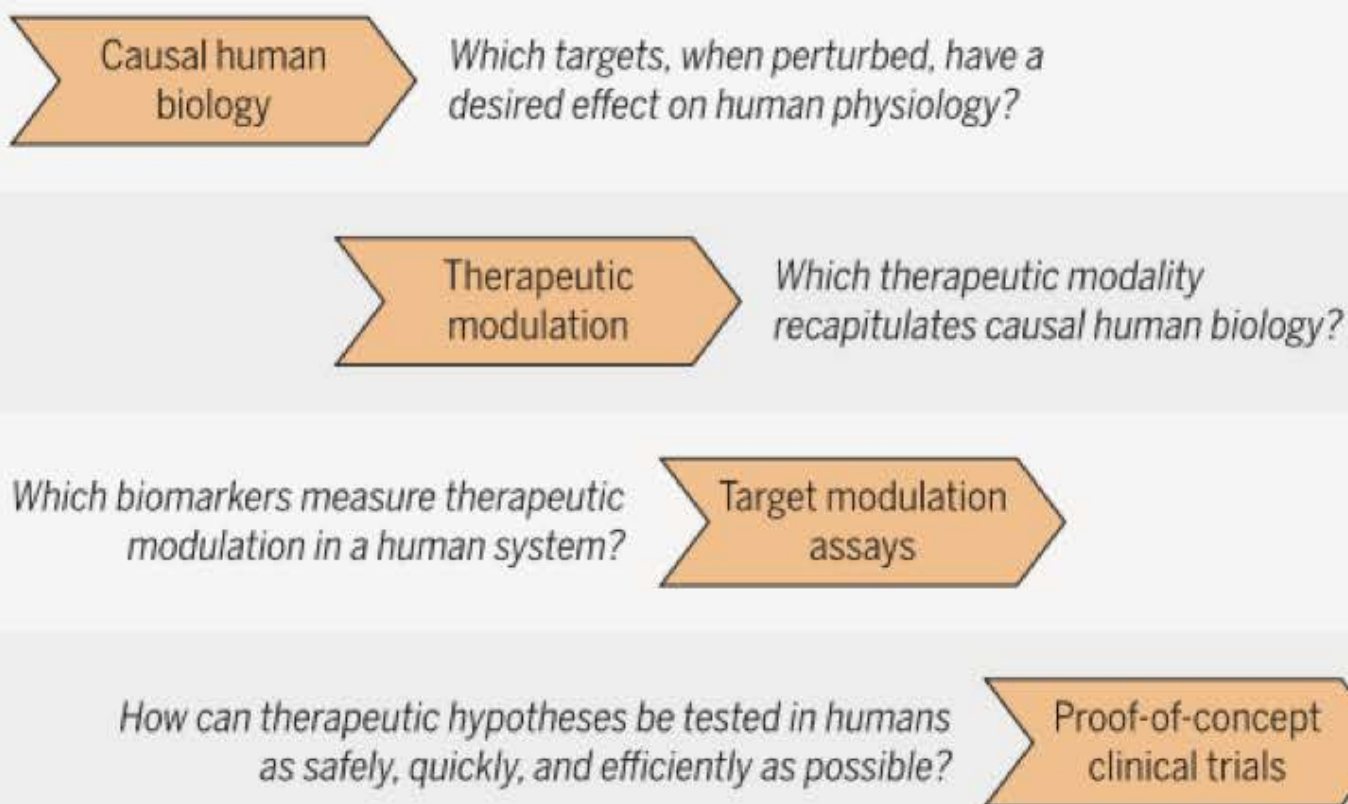
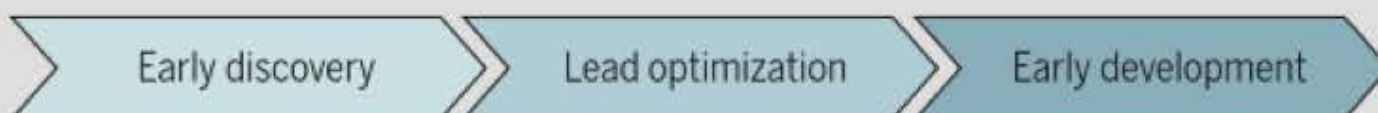
1. Too many Phase 2/3 studies fail
2. Time between discovery and PoC is too long



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# Why don't drugs differentiate?



A disciplined approach to integrating all four components will lead to the following improvements:

Increased probability of success in phase 2/3 clinical trials

Increased probability for differentiation from standard of care

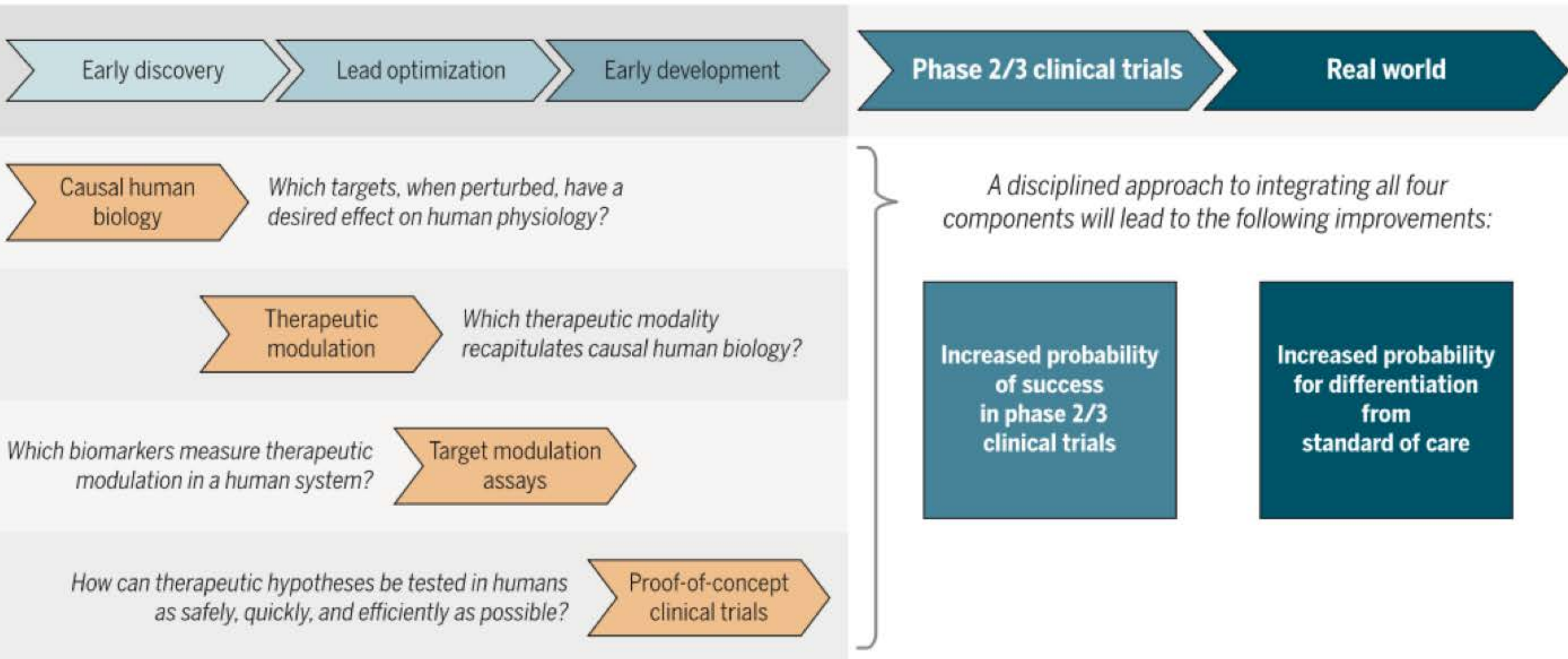
# 1. Not enough sound therapeutic hypotheses!



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# How will *precision medicine* help?

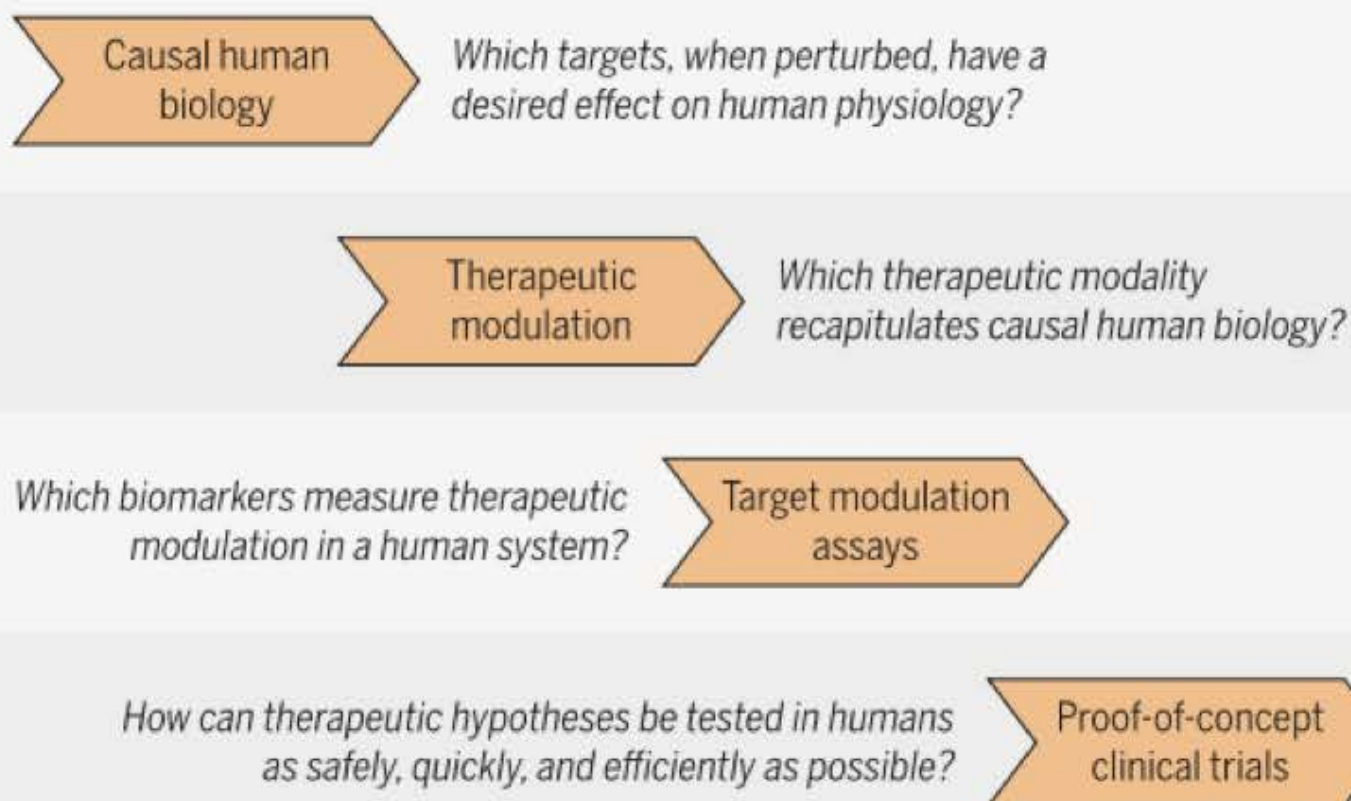
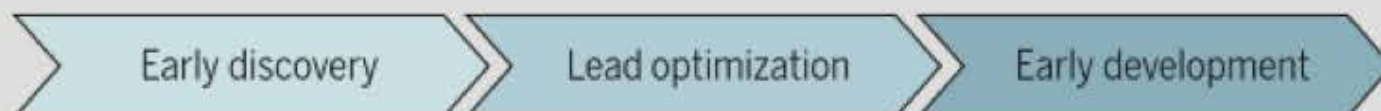


1. Improve cycle time from therapeutic hypothesis to clinical PoC...*efficiently and safely*



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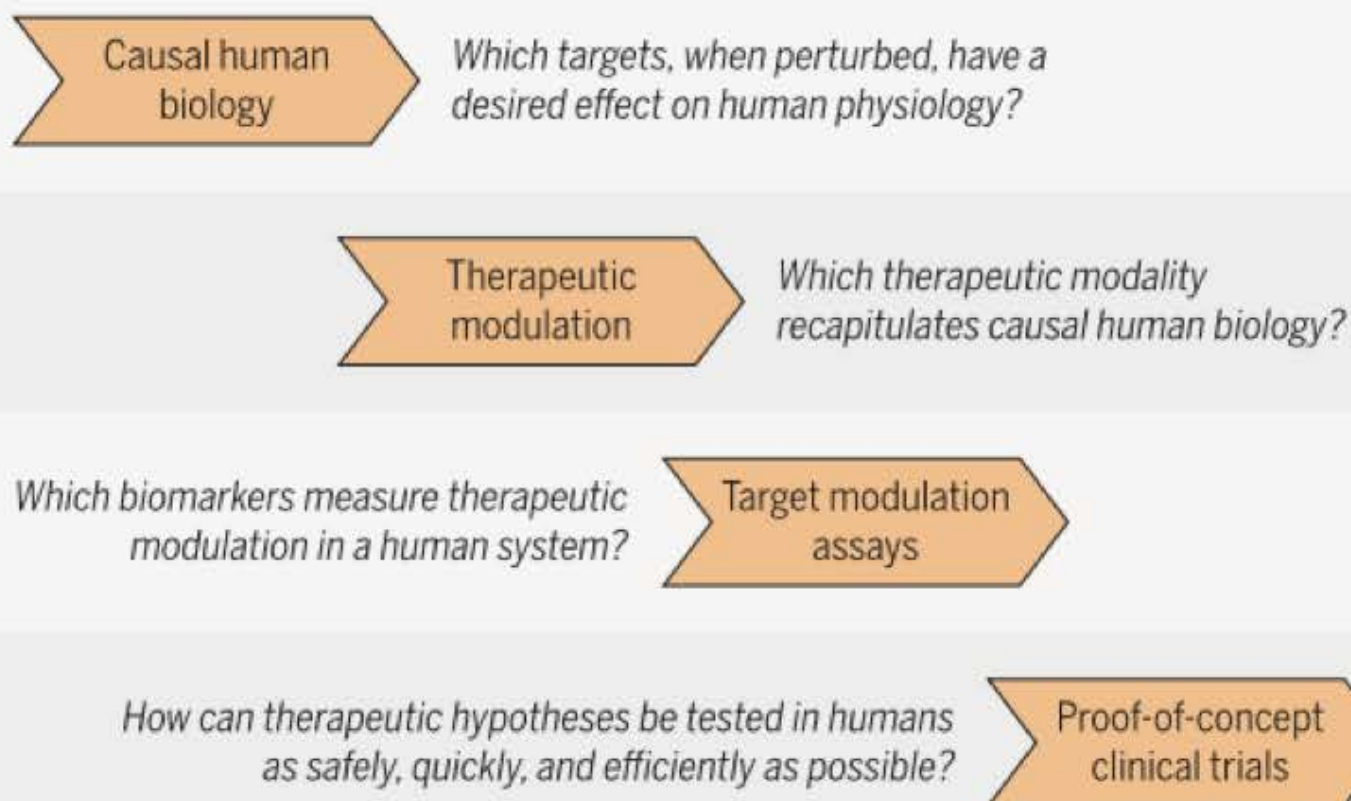
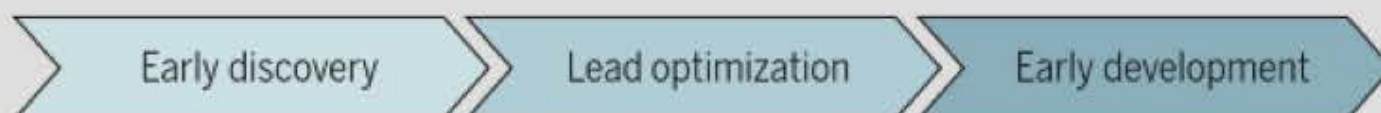
Increased probability  
for differentiation  
from  
standard of care

Today: 5-7 years



# Disciplined approach to drug discovery and early development

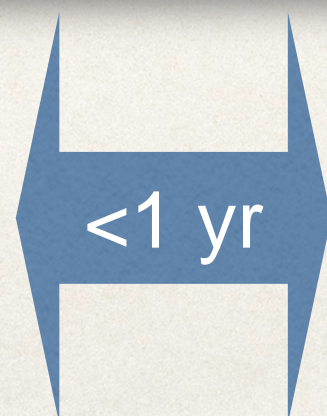
Robert M. Plenge



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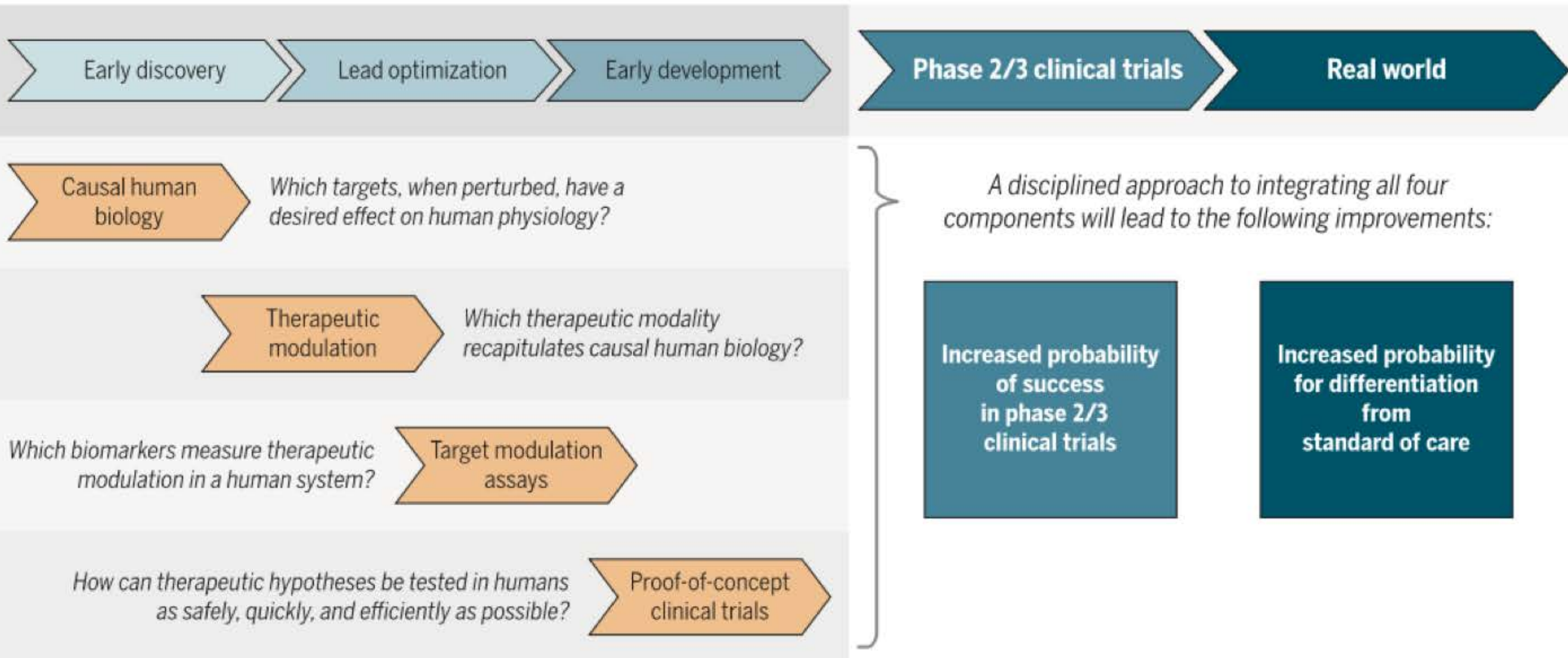




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# What are potential solutions?



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



# human genetics to pick targets

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Pick a human  
phenotype for drug  
efficacy



Human Phenotype

High

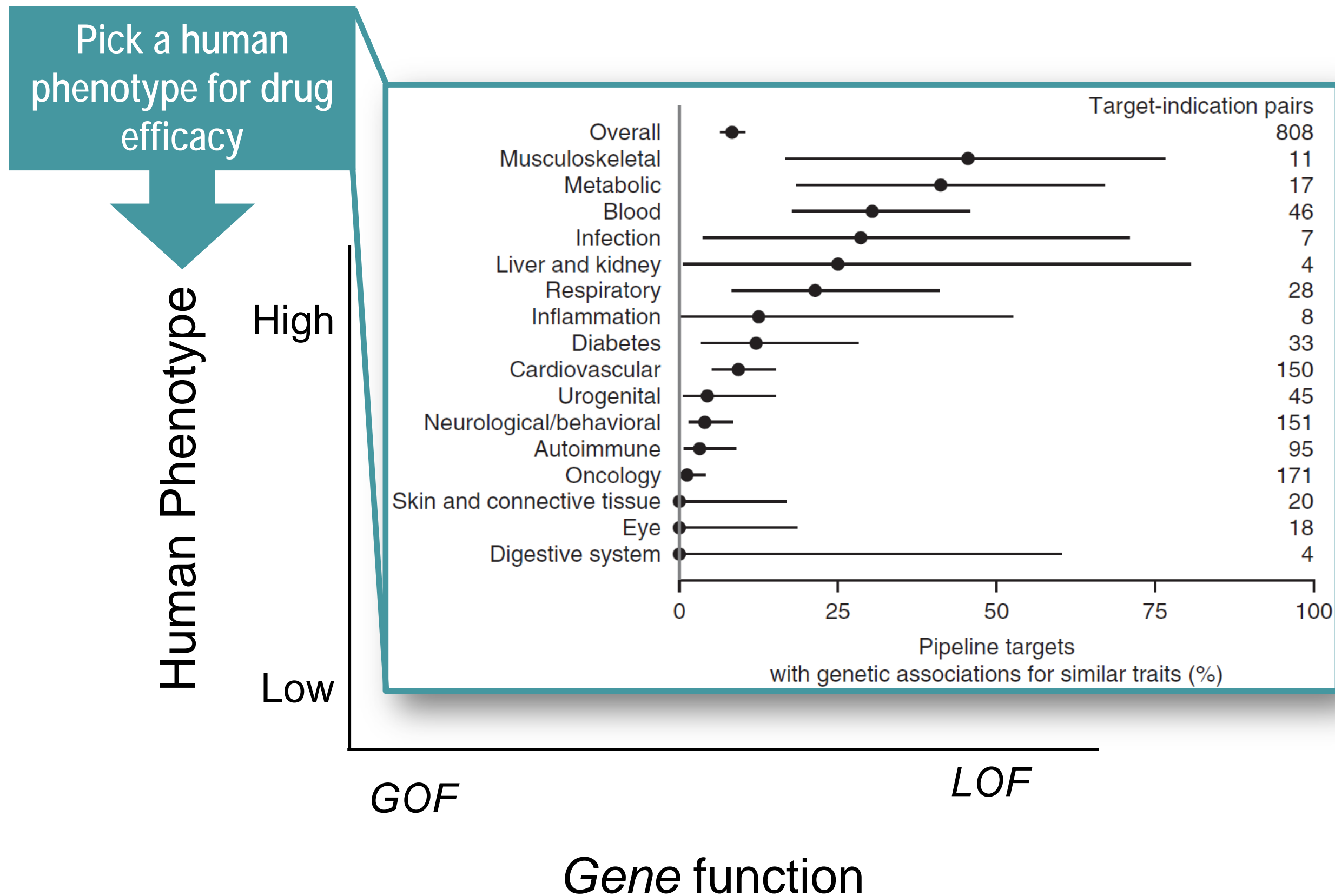
Low

*GOF*

*LOF*

*Gene function*







Pick a human phenotype for drug efficacy



Human Phenotype

High

Low

*GOF*

*LOF*

Gene function

X

X

X

X

X

X

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





Pick a human  
phenotype for drug  
efficacy



Human Phenotype

High

Low

GOF

LOF

JAMA | Original Investigation

## 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. Stitzel, MD, PhD; Pradeep Natarajan, MD; Akihiro Nomura, MD; Connor A. Ermdin, 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 Ardissino, 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

**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.

>100 rare variants  
6 common variants

Gene function

Khera et al JAMA 2017



Pick a human phenotype for drug efficacy



Human Phenotype

High

Low

*GOF*

*LOF*

Gene function

Efficacy

Assess biological function of alleles to estimate “efficacy” response curve

x

x

x

x

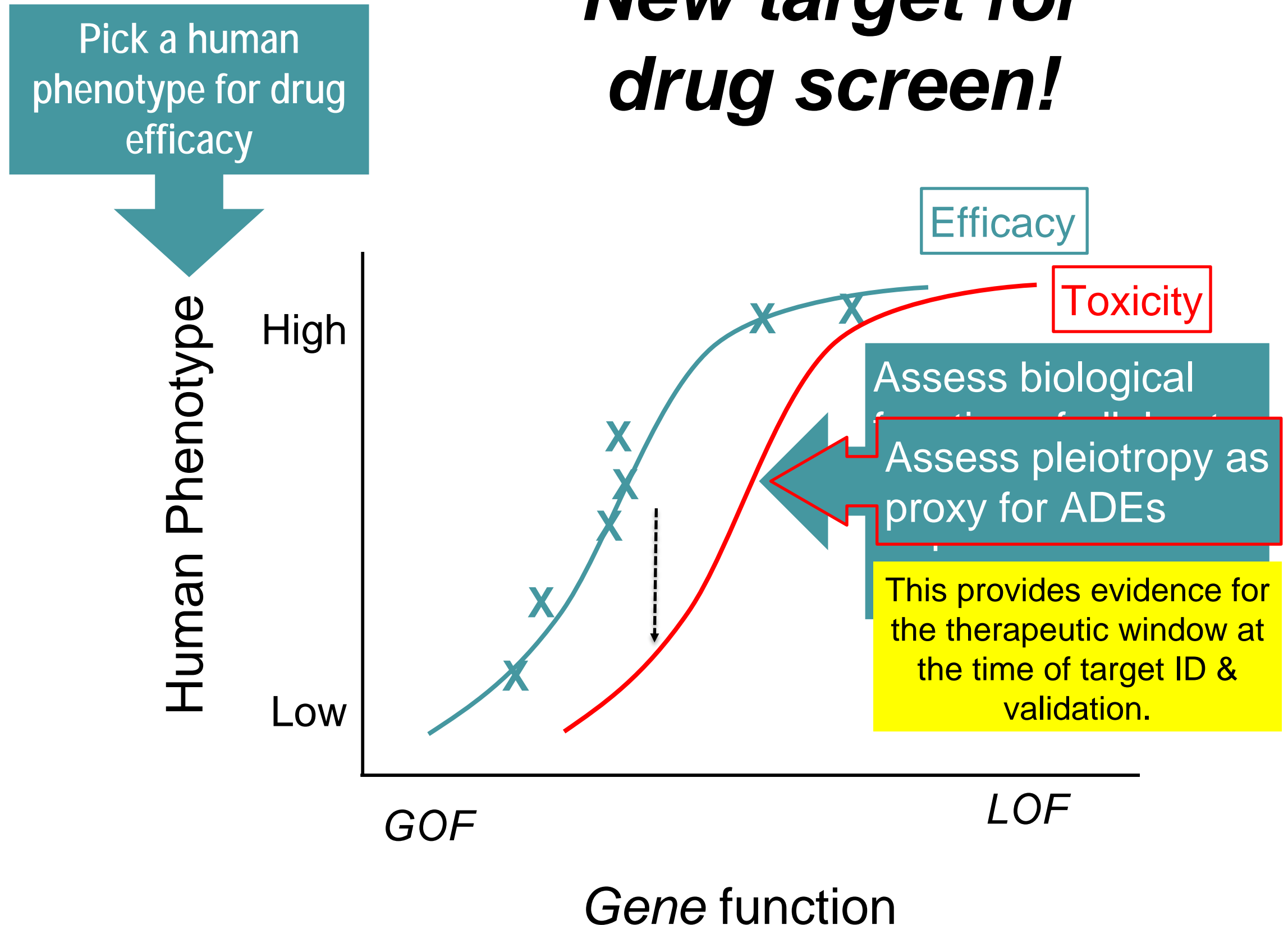
x

x





# ***New target for drug screen!***





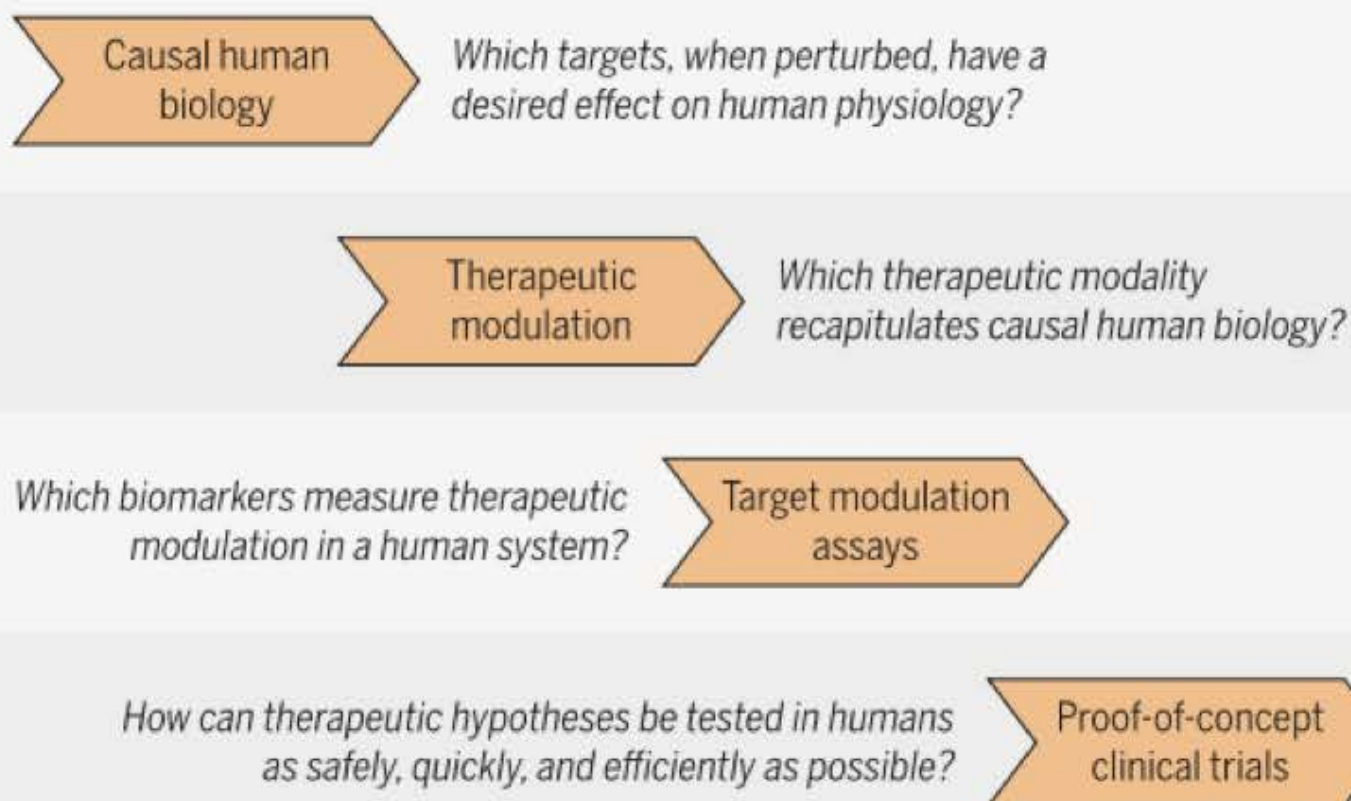
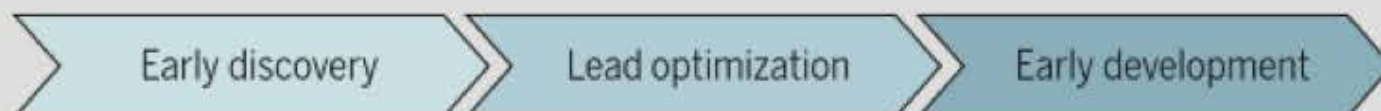
programmable therapeutics to test  
therapeutic hypotheses quickly

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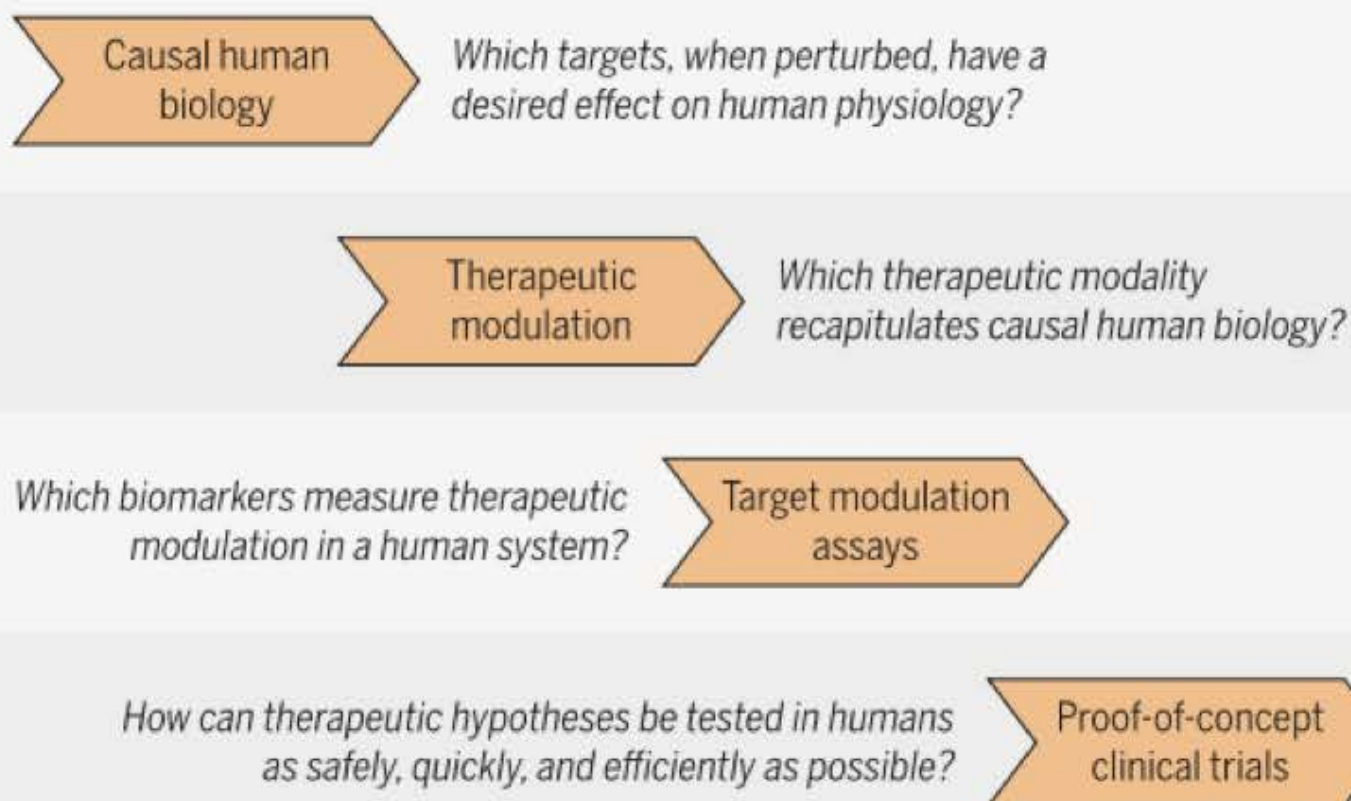
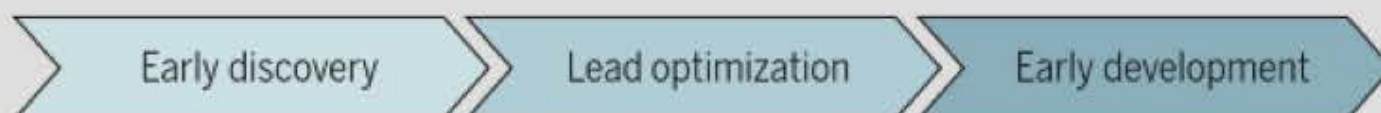
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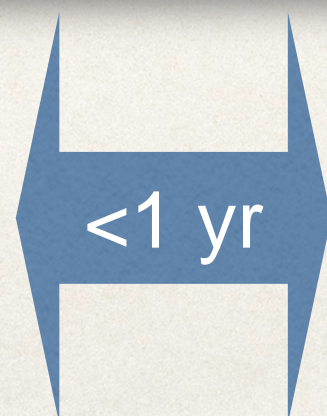
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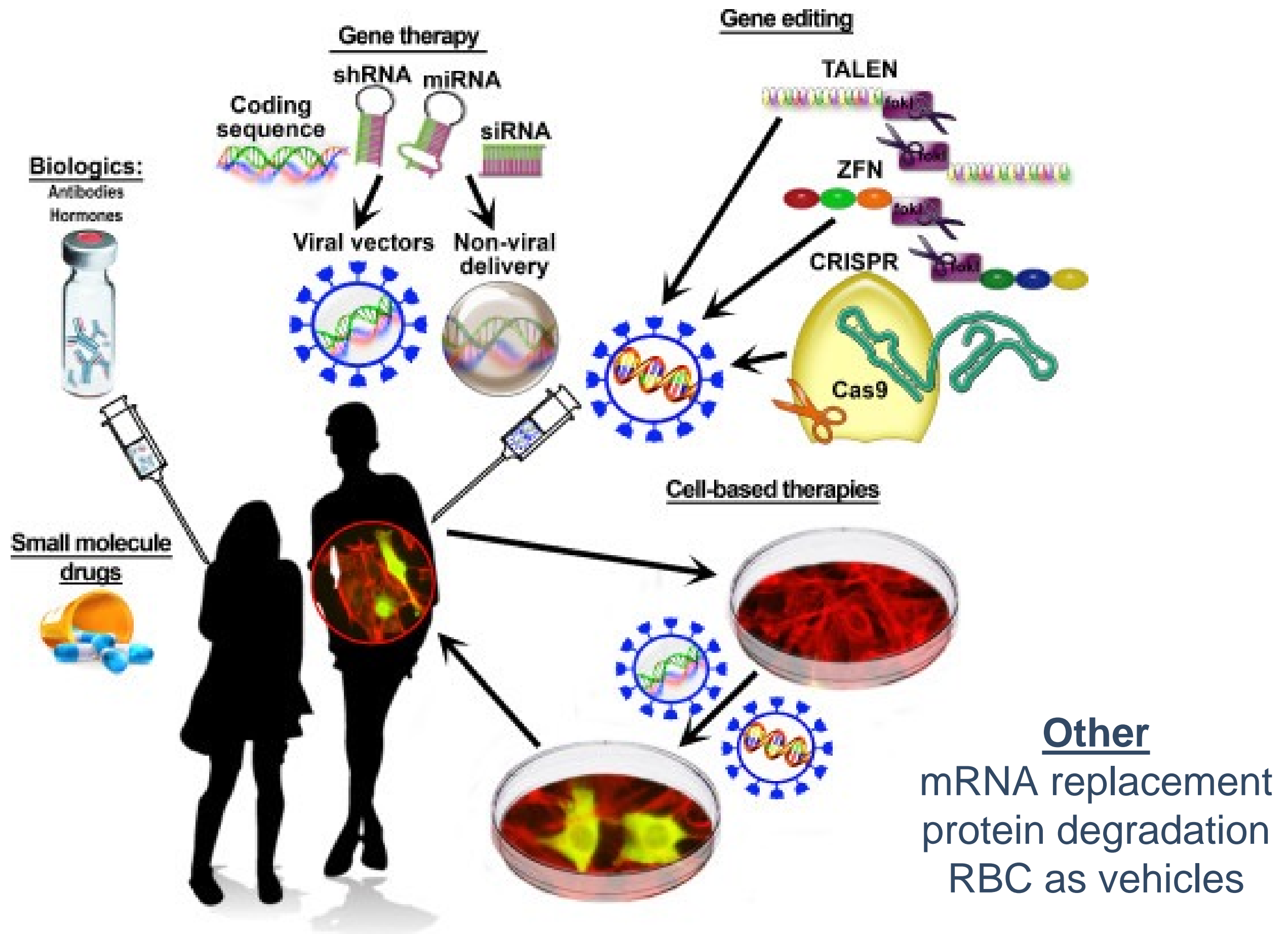


# Limitations that we face today

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- ❖ 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

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- ❖ 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

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# Overall message - inconvenient path to precision medicine

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- ❖ 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