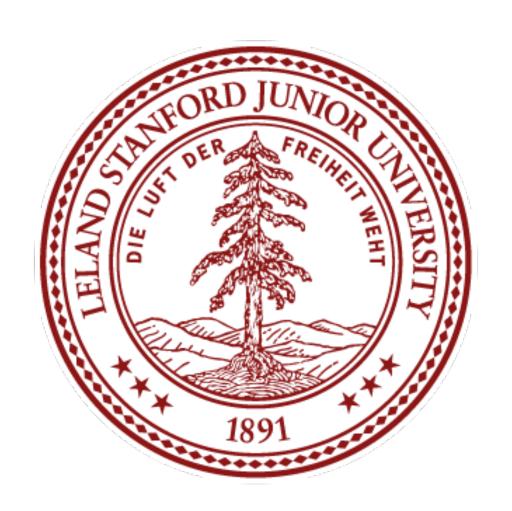
What Statistics and Al Offer Each Other?



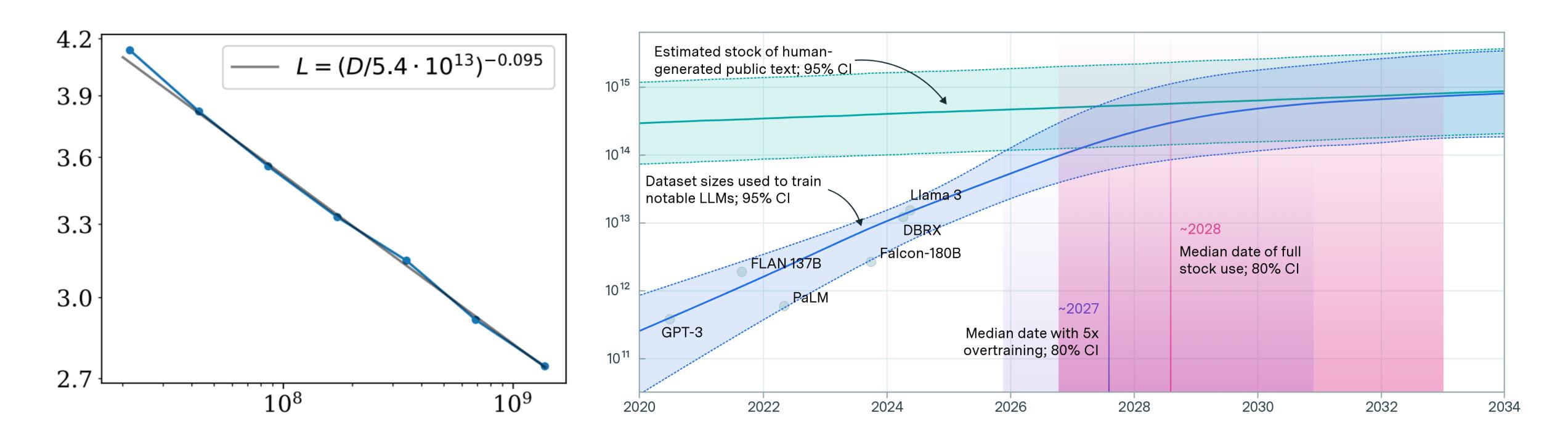
Emmanuel Candès

Asilomar Conference on Signals, Systems, and Computers, Oct 27,

2025

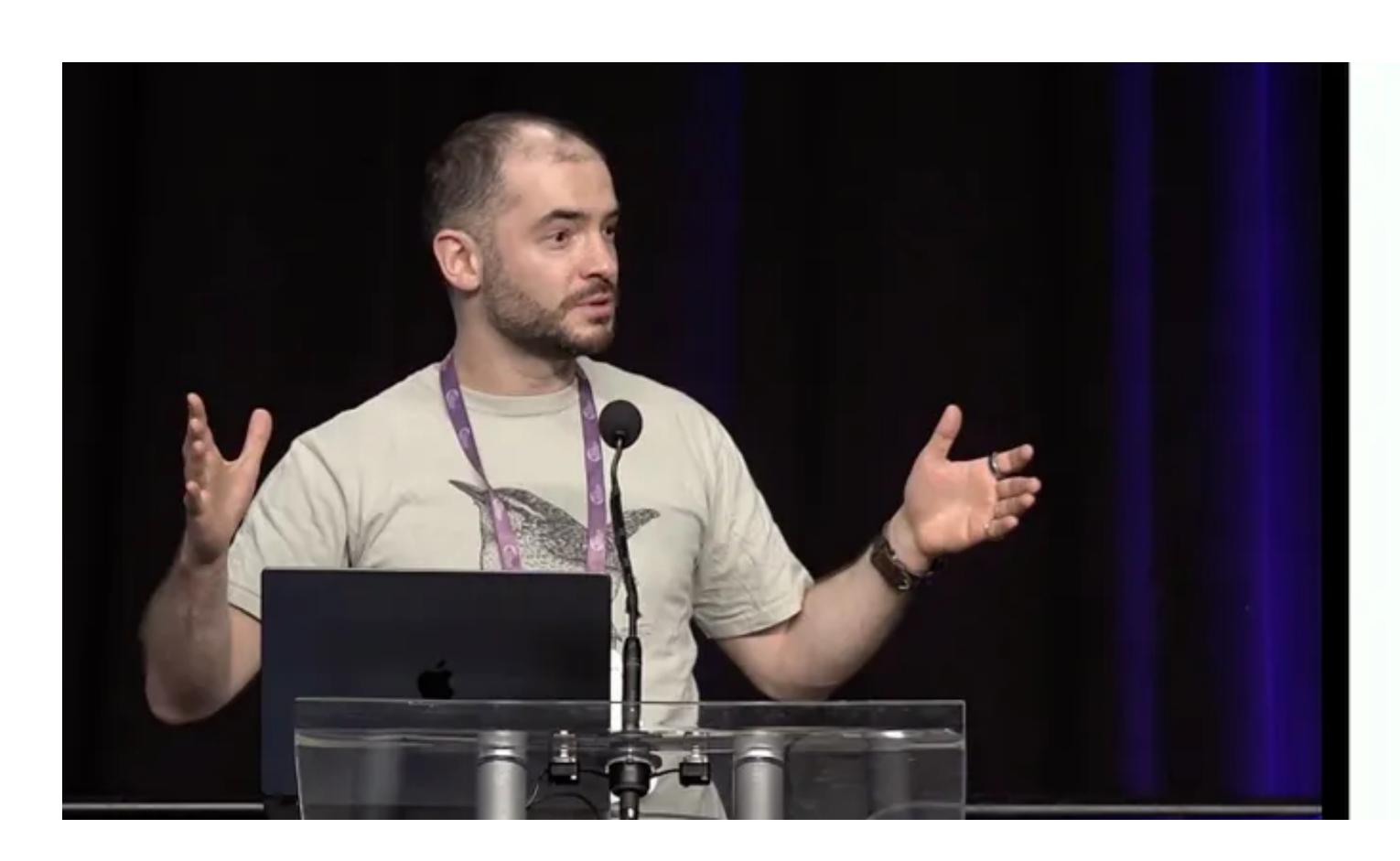
Scaling wall

- ► Scaling law (2020-2025): more data ==> more powerful Al
- Scaling wall (2025+): frontier AI has exhausted internet data



Our take: use existing data more effectively

Ilya Sutskever @NeurlPS 2024 — We have but one internet!



Pre-training as we know it will end

Compute is growing:

- Better hardware
- Better algorithms
- Larger clusters

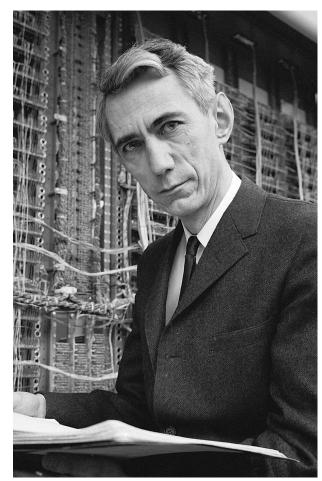
Data is not growing:

- We have but one internet
- The fossil fuel of Al

Language model pretraining

Max likelihood:

$$maximize_{\theta} \sum_{docs} \log p_{\theta} (doc)$$







Ronald Fisher

Extending pretraining beyond the scaling wall via synthetic data

- Synthetic continued pretraining

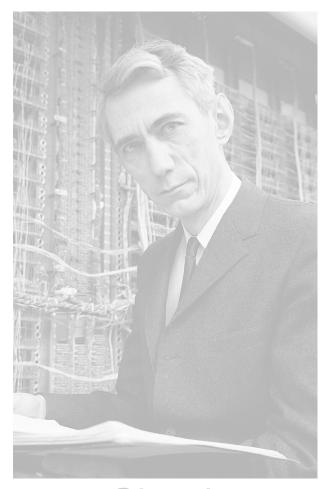
 Zitong Yang, Neil Band, Shuangping Li, Emmanuel Candès, Tatsunori Hashimoto
- Synthetic bootstrapped pretraining

Zitong Yang, Aonan Zhang, Hong Liu, Tatsunori Hashimoto, Emmanuel Candès, Chong Wang, Ruoming Pang

Language model pretraining

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Synthetic continued pretraining

Goal: teach model the knowledge from a niche domain consisting of few "source documents"

Step 1: Generate synthetic text based on the source documents

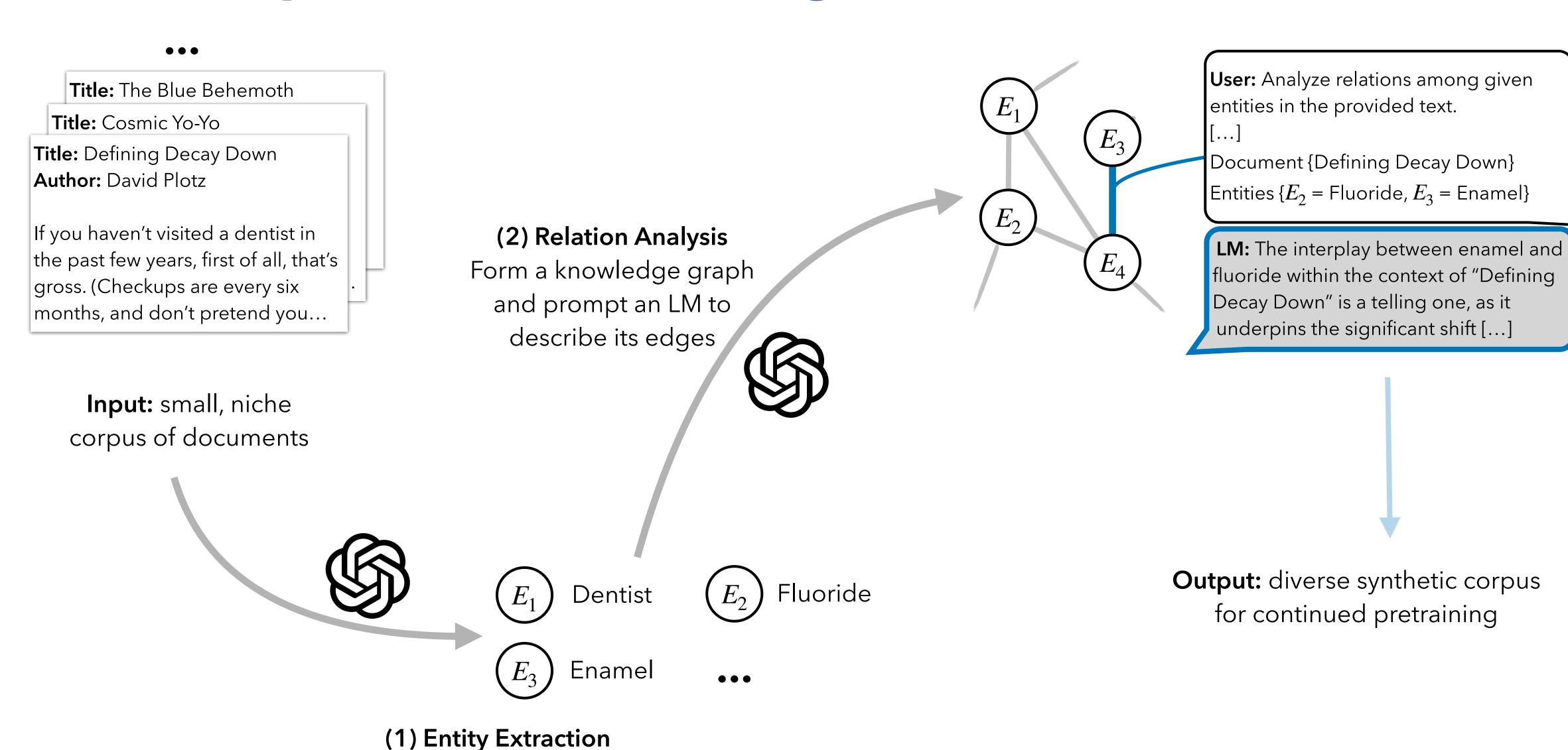
Step 2: Continually pretrain (finetune) the model on generated text

Experiment setup

- Niche source documents (not something model already knows)
- A task that tests a model's knowledge about the source documents

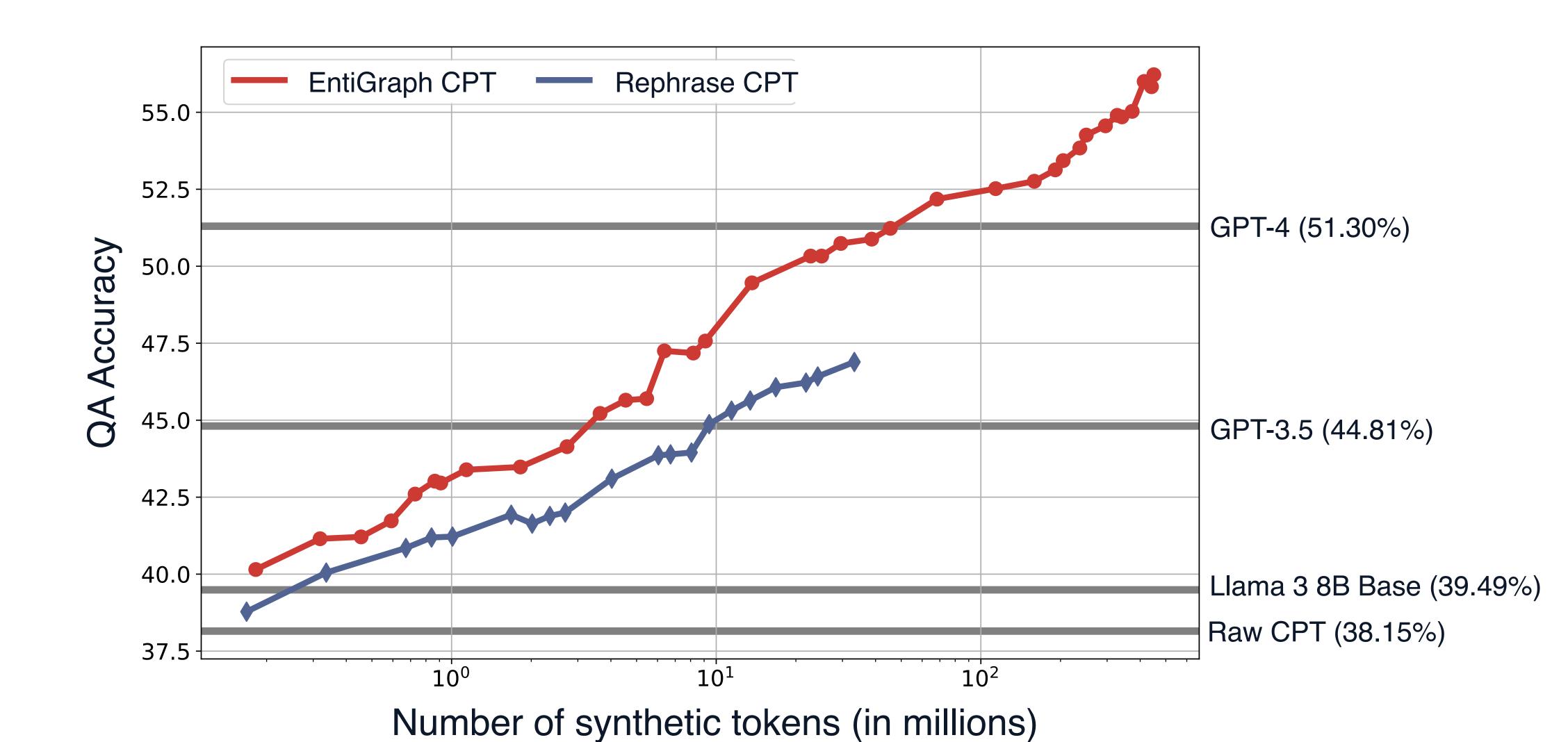
How to generate synthetic data?

EntiGraph: scalable data generator



For each document, extract a list of entities

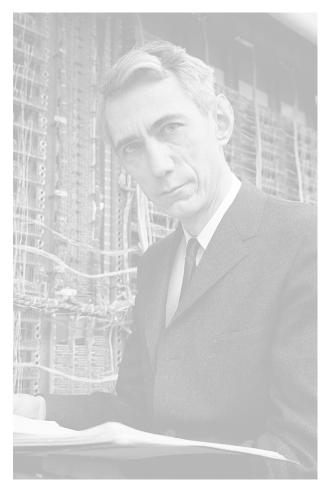
Scaling on synthetic data



Language model pretraining

Max likelihood:

$$maximize_{\theta} \sum_{docs} log p_{\theta} (doc)$$







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Synthetic bootstrapped pretraining

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1. Nearest-neighbor pairing: DCLM subset and Qwen-0.6B-Embedding

a news event All pretraining documents the Harry Potter book HP movie commentary the transformer paper HP book analysis a lesser known novel PyTorch attention implementation a lesser known paper embedding space

Examples of related documents

doc1

The Cultural Sites of Iran

With 196 countries and countless exciting destinations worldwide, there is so much to see in a very limited time. Even the most well-traveled person hardly gets to visit all and has to be selective. So, why should you consider visiting a country like Iran, especially when it comes to all those negative news and stereotypes surrounding it?

Here we're here to give you the reasons and to help you overcome your doubts and even encourage you to consider your next trip to Iran, this mysterious land as soon as you return to your home country!

Beautiful cities, friendly people, fabulous food, glorious architecture, Iran has delighted visitors for centuries with its World Heritage Sites, friendly towns and inspiring desert landscapes.

Things to Do in Iran – Activities & Attractions

Iran is the land of four seasons, history and culture, souvenir and authenticity. This is not a tourism slogan, this is the reality inferred from the experience of visitors who have been impressed by Iran's beauties and amazing attractions.

Antiquity and richness of the Cultural Sites of Iran and civilization, the variety of natural and geographical attractions, four – season climate,

. . .

History of Iran

doc2

Query Text: Home > FAQ Login / Register

Why should we spend our holiday in Iran?

Iran is a country, located in the Middle East, which can meet the various needs of tourists and satisfy their different tastes, due to its rich civilization, historical sites, geographic location, nature of the four seasons and diverse tourist attractions. Therefore, considering the high security and low cost of travel to the country, it is introduced as one of the major tourist destinations to spend holidays in.

Is Iran a safe travel destination?

One of the wrong assumptions about the country of Iran is in terms of its security. Despite its location in Asia and the Middle East, and neighboring countries like Iraq, Afghanistan and Pakistan, Iran is considered as one of the safest countries in the region. According to the international data, security in Iran is much more than a touristic country such as Turkey.

To confirm the statements made above, refer to websites like www.travelriskmap.com.

What does "the rich civilization" mean, as mentioned about Iran?

According to documentation in some of the world history references,

. . .

Travel guide to Iran

ahaalmaint

- 1. Nearest-neighbor pairing: DCLM subset and Qwen-0.6B-Embedding
- 2. Synthesizer tuning: SFT-like objective $p_{\theta}(d_2 \mid d_1)$ initialized at pretrained

the Harry Potter book

HP movie commentary

the transformer paper

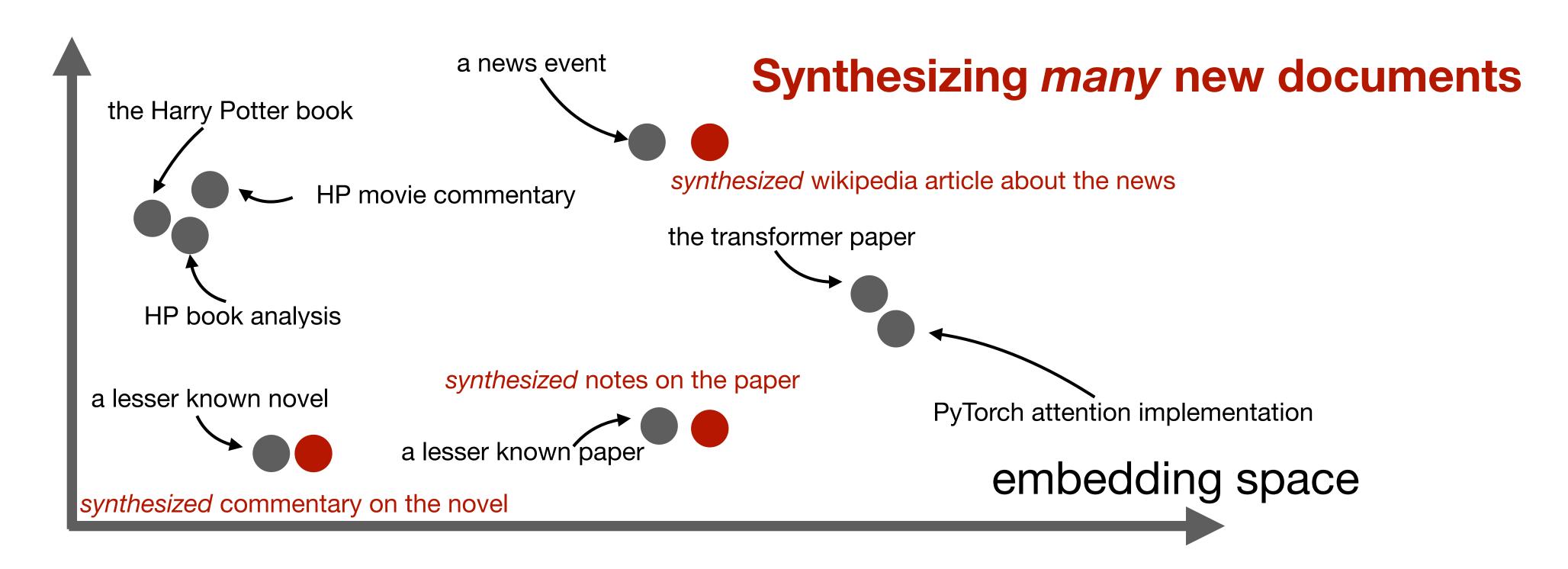
HP book analysis

a lesser known novel

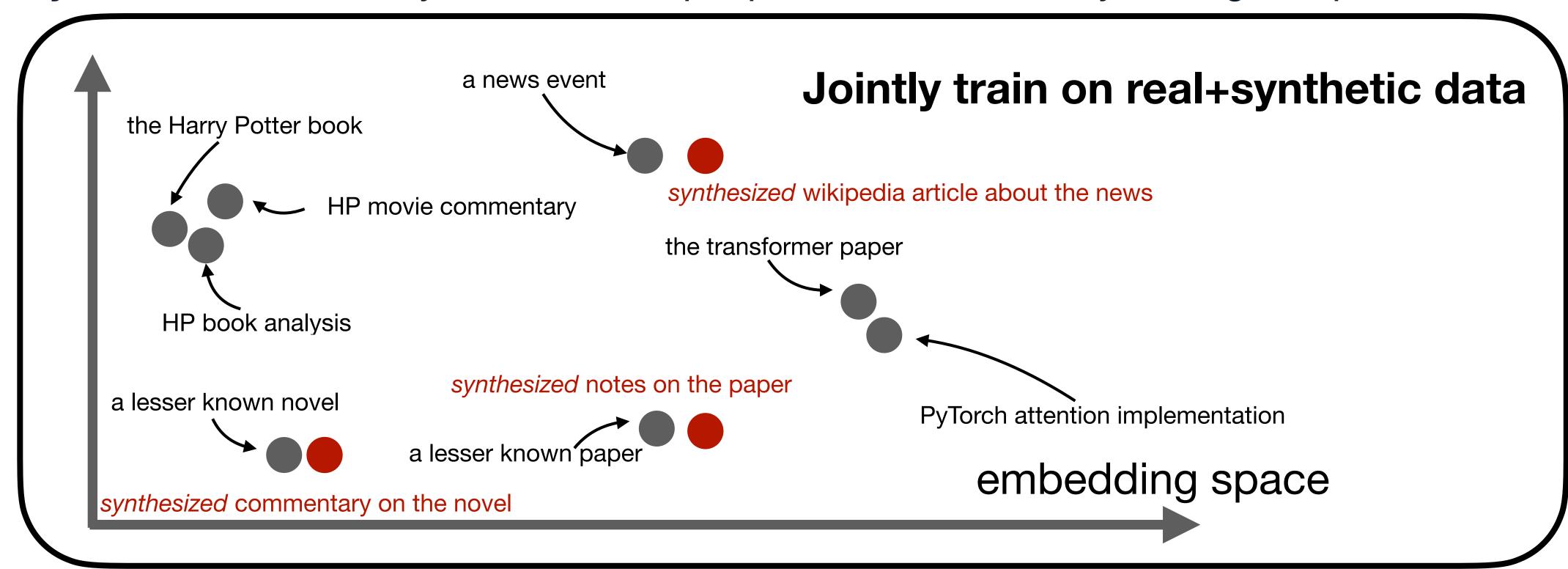
PyTorch attention implementation

embedding space

- 1. Nearest-neighbor pairing: DCLM subset and Qwen-0.6B-Embedding
- 2. Synthesizer tuning: SFT-like objective $p_{\theta}(d_2 \mid d_1)$ initialized at pretrained checkpoint
- 3. Synthesis at scale: Synthesize multiple paired documents by setting Temp=1



- 1. Nearest-neighbor pairing: DCLM subset and Qwen-0.6B-Embedding
- 2. Synthesizer tuning: SFT-like objective $p_{\theta}(d_2 \mid d_1)$ initialized at pretrained checkpoint
- 3. Synthesis at scale: Synthesize multiple paired documents by setting Temp=1



Examples of synthesis

Real document

HOME > Chowhound > California >

Coffeehouses in SD

I am always looking for good coffeehouses in San Diego. I am looking for coffeehouses with (obviously) excellent coffeebeans, baristas who know their business (good microfoam, good crema, they should know the different ratios for steamed milk, foamed milk, espresso for the cappuccino, latte etc., etc), good snacks (paninis, cakes etc), good (cold) flavored coffeedrinks and good ambience. So far I haven't found many who are good but would like to hear more suggestion.

Caffe Calabria: By far the best coffeehouse in SD. Excellent coffee and baristas (best cappuccino in SD,), great paninis. Only downside is that they close very early.

Café Lestat: Good example what happens with good coffee and average baristas. Lestat is also using coffeebeans from Caffé Calabria but it is sad to see that they don't know how to make excellent espresso, cappuccino.

E Street Café: Average espresso but strange flavored ..

Synthesis I

Coffee = Love

On a recent trip to San Diego, one thing I wanted to find was coffee. I was worried it might be difficult with the huge decrease in Asian flavors, but I was pleasantly surprised. Many of the Mexican, Italian, and Cuban flavors that are pretty common in California are now in the mix. While I did find a few highly recommended spots, the majority of the locations I checked out were nothing to write home about. At the least, they were not being sold out, which I consider a plus.

One major factor that really needs to be taken into account is the type of machine being used. There are two types of espresso machines: the cheap ones that spit out lots of pressure and don't have any control over the shot, and the expensive ones that create amazing shots and utilize the types of superfine grinds you usually find in Italy. The latter types are much harder to find and cost significantly more.

Another factor to consider is the type of beans being used. If they are done correctly, great espresso can be made from any type of bean, but it's a rare experience to find a place that does ...

Synthesis II

Discuss as:

Coffeehouses: San Diego. The best?

Café Lestat, by far, is a stand-out. (photo by Thomas Moriarty)

If the coffee scene in San Diego sounds like the one in New York, there's a reason. The upper echelons of NYC's drink scene are the same ones who have taken over the majority of the coffee houses in San Diego. This is probably partly due to the fact that both cities have large student populations, both are home to large coffee drinker bases, both have a large number of espresso shops, and both offer top-notch, hand-crafted cappuccinos. But if there is one city in America that deserves to have its name above that of New York, it's San Diego.

There are just under 100 coffee shops in San Diego, with almost half of them located on University Ave. alone. So finding the perfect coffee shop is crucial. We spent a whole day just roaming around the area, hunting for the best.

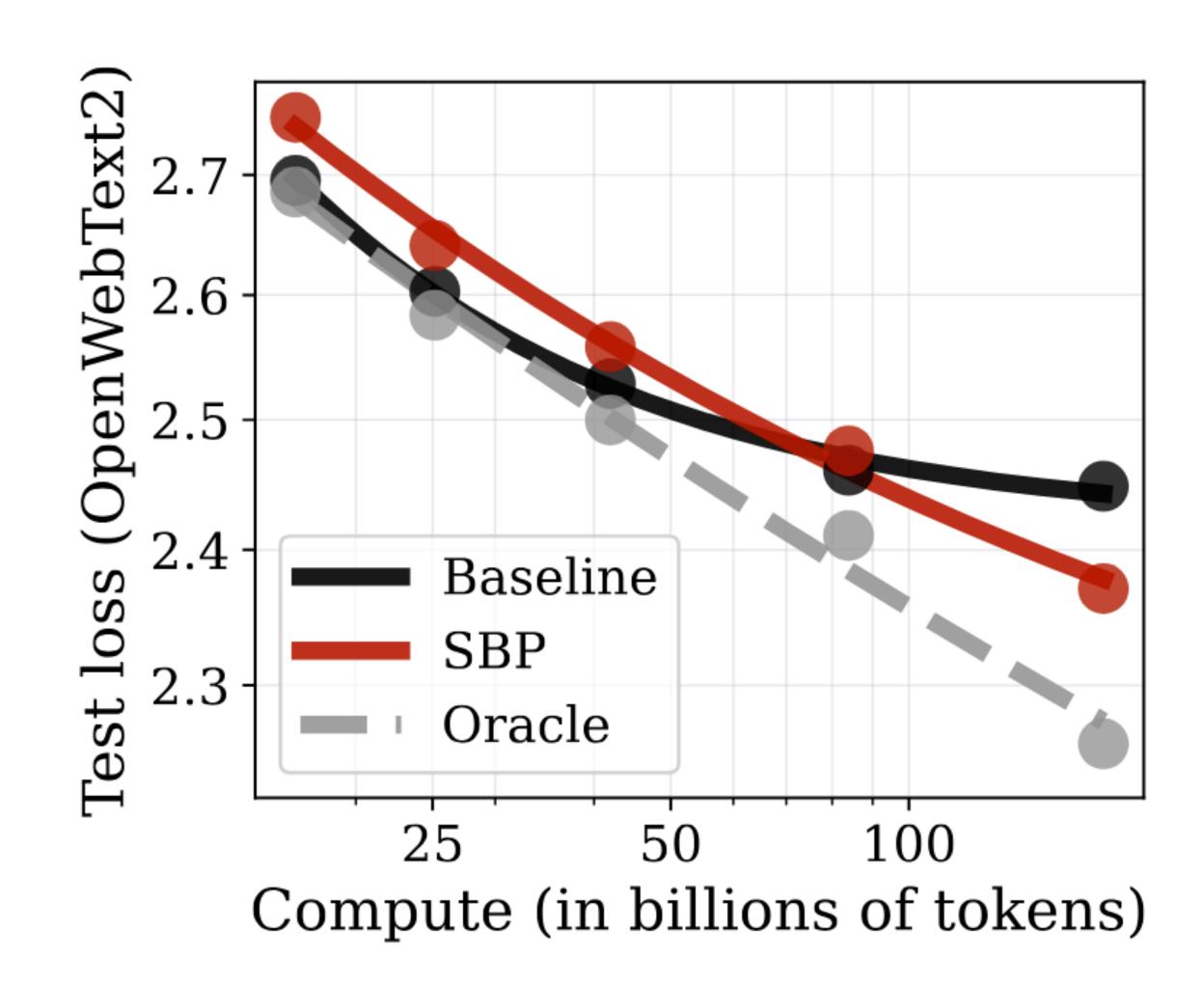
In terms of the coffee itself, it's hard to beat Café Lestat. The baristas are amazing and their methods are pristine ...

40% of improvement attained by oracle with 20x data

	200B-scale		1T-scale			
Benchmark	Baseline	SBP	Oracle	Baseline	SBP	Oracle
	Perplexity on held-out data↓					
OpenWebText2	5.74	-0.53	-1.02	4.51	-0.02	-0.12
LAMBADA	6.87	-0.85	-1.86	4.33	-0.03	-0.22
Five-shot MMLU	3.83	-0.36	-0.51	3.17	-0.06	-0.05
	QA accuracy ↑					
ARC-Challenge (0-shot)	35.32	+1.28	+2.82	42.66	+1.62	+3.84
ARC-Easy (0-shot)	68.94	+2.65	+4.29	75.63	+0.42	+2.11
SciQ (0-shot)	90.50	+1.00	+2.40	93.20	+0.80	+0.50
Winogrande (0-shot)	60.14	+1.90	+5.53	65.19	+1.42	+2.92
TriviaQA (1-shot)	22.51	+3.36	+7.37	36.07	+0.25	+0.59
WebQS (1-shot)	8.56	+3.74	+10.83	19.34	+0.54	+0.44
Average QA accuracy	47.66	+2.32	+5.54	55.35	+0.84	+1.73

Training dynamics

- Initially, baseline and oracle perform similarly. SBP is worse because it uses synthetic data
- <u>Later on</u>, baseline and oracle diverge;
 SBP follows a linear trend
- Near the end, baseline plateaus while SBP continues to improve

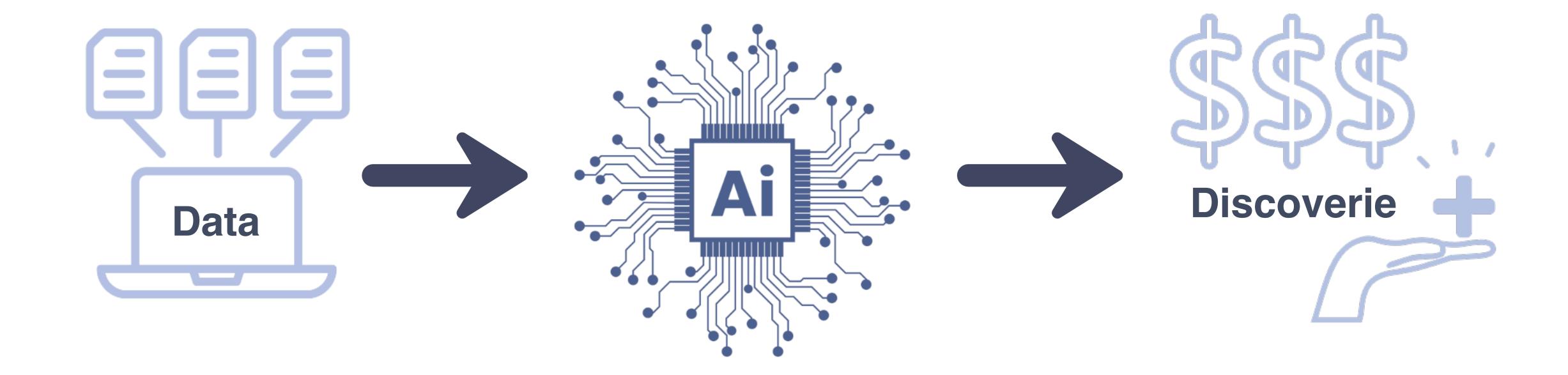


Synthetic continued/bootstrapped pretraining: summary

Biggest reservoir of machine learnable knowledge resides in unsupervised learning (Y. leCun) — witness LLM from GPT-3 on

As we run out of internet data, we propose a form of self-supervision weaker than next-token prediction, exploiting existing knowledge/correlations on the internet

Modern discovery pipeline

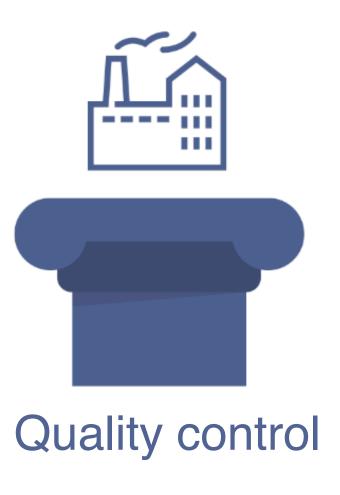


Thesis: Thinking carefully about Al inputs and outputs yields more powerful, safer Al

Agenda: vignettes on three pillars

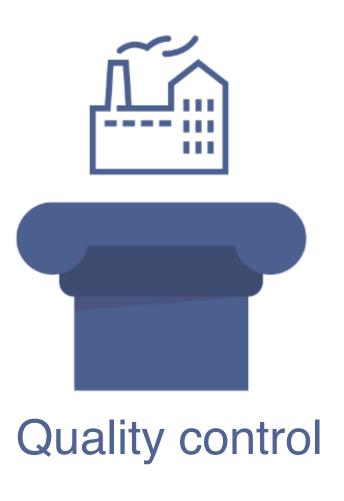










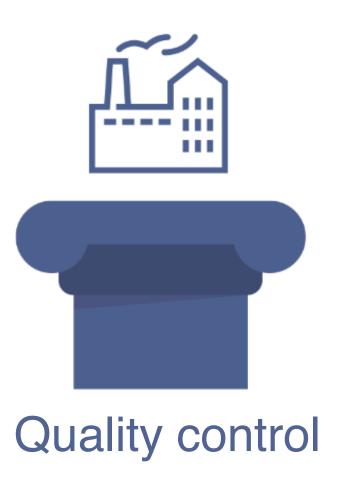




synthetic pretraining datamodels s1









synthetic pretraining datamodels s1



Al-powered inference

Increasing life expectancy

NEW YORK, WEDNESDAY, APRIL 13, 1955.

Timer Square New York 26, N. Y. Telephone Lackswamps 4-1000

FIVE CENTS

dual Approaches Urged Integration of Schoolsegro Lawyers Opposed

y LUTHER A. HUSTON

Special to The New York Times. ASHINGTON, April 12tesmen for South Carolina Virginia told the Supreme t today that their people id not obey a decree ordering mmediate end to racial segtion in the public schools. hen Chief Justice Earl Warasked S. E. Rogers, repreing Clarendon County, S. C. s were willing to say that 'honest attempt" would be to conform to whatever ee the court might issue, Mr. ers said;

et's get that word 'honest of there. It would depend the kind of decree. The

THEOURT HEARS SALK POLIO VACCINE PROVES SUCCESS; OUTH WILL DEFY MILLIONS WILL BE IMMUNIZED SOON; CITY SCHOOLS BEGIN SHOTS APRIL 25



TRIAL DATA GIVEN

Efficacy of 80 to 90% Shown—Salk Sees Further Advance

Abstract of report, summary of data on tests, Page 22.

By WILLIAM L. LAURENCE Special to The New York Times. ANN ARBOR, Mich., April 12 -The world learned today that its hopes for finding an effective weapon against paralytic polio had been realized.

No perfect mechanistic understandings of vaccines \rightarrow we must learn from data

How sure are we of what we learn from data?

Achieve the gold standard, i.e. produce confidence bounds L, U from data such that

Prob(L < object of inference < U) = 95%



level of deforestation in the Amazon effectiveness of a vaccine side effect of a drug

Warning! before clinical trials Rasmussen (2008)

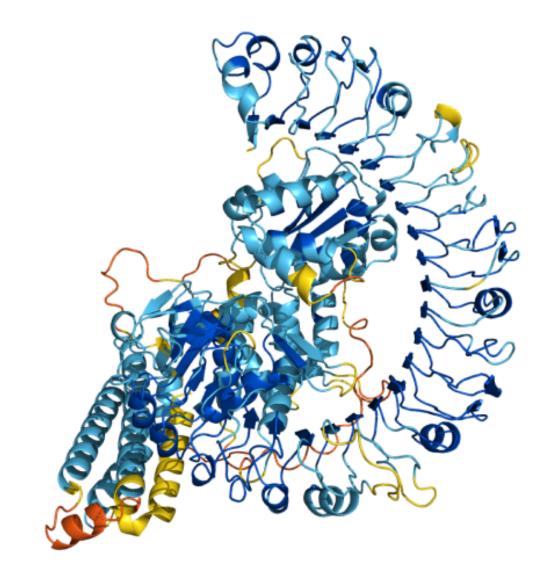
"Large quantities of amphetamines were dispensed in the 1960s by weight loss clinics."

One estimate = **2 billion tablets annually**; millions suffered cardiovascular damage. \implies reliable results matter!

Can we leverage imperfect Al predictions to get gold-standard results?



How much deforestation is in the Amazon?



Which fraction of proteins have a certain property?



How much has political sentiment in the media changed?

Setup

features labels

X_1	?
X_2	?
X_{N-1}	?
X_N	?

N unlabeled data points

satellite images deforestation % X_1 X_2 X_{N-1}

Goal: learn scientific target $T = \text{mean}(Y_1, ..., Y_N)$

T = deforestation rate in the Amazon

Setup

features labels

X_1	Y_1
•	•
X_n	Y_n
	•
X_{N-1}	?
X_N	?

We can collect at most n expert labels Y_i (much fewer than N)

N-n unlabeled data points

deforestation % satellite images X_1 15% X_2 7% X_{N-1} X_N

Goal: learn scientific target $T = \text{mean}(Y_1, ..., Y_N)$

T = deforestation rate in the Amazon

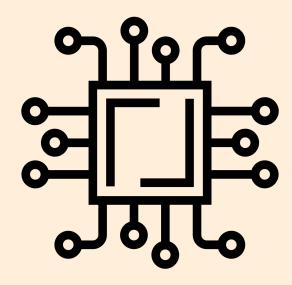
Setup

features labels

X_1	$Y_1 \hat{Y}_1$
•	
X_n	$Y_n \hat{Y}_n$
X_{N-1}	\hat{Y}_{N-1}
X_N	\hat{Y}_N

We can collect at most n expert labels Y_i (much fewer than N)

N predicted labels



Al model

produces informative but imperfect predictions $\hat{Y}_1, \ldots, \hat{Y}_N$

model can be fine tuned on same labels

Goal: learn scientific target $T = \text{mean}(Y_1, ..., Y_N)$

Gold-standard results with Al

Step 1: Collect n randomly chosen expert labels Y_i

Step 2: Given $(X_1, Y_1), \ldots, (X_n, Y_n), X_{n+1}, \ldots, X_N$, estimate the scientific target T

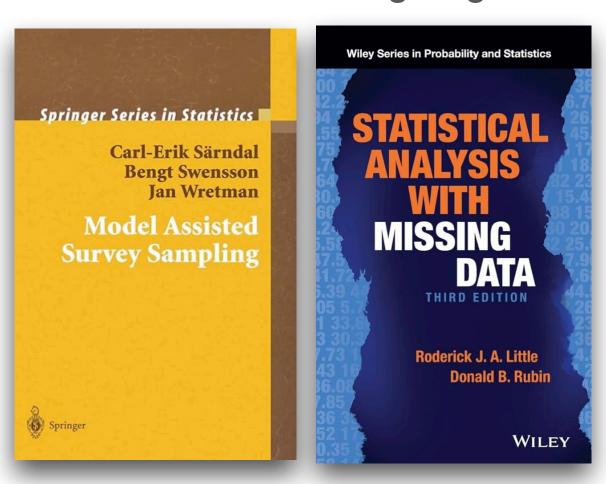
$$T^{\text{PP}} = \operatorname{mean}(\hat{Y}_1, \dots, \hat{Y}_N) - \operatorname{mean}(\hat{Y}_1 - Y_1, \dots, \hat{Y}_n - Y_n)$$
 if we pretended AI bias of AI predictions were correct predictions

No matter the Al bias, can make sure we achieve the gold standard!

Angelopoulos et al. (2023), Zrnic, Candès (2024a)



*This trick is based on classical statistical thinking, e.g.,



Gold-standard results with Al

Classical inference

$$\hat{\theta}^{\text{CL}} = \frac{1}{n} \sum_{i=1}^{n} Y_i$$

Prediction-powered inference (PPI)

$$\hat{\theta}^{\text{PP}} = \frac{1}{N} \sum_{i=1}^{N} f\left(X_{i}^{\text{unlabeled}}\right) + \frac{1}{n} \sum_{i=1}^{n} \left(Y_{i} - f\left(X_{i}^{\text{labeled}}\right)\right) \quad \hat{Y} = f(X)$$
 blackbox prediction

Unbiased 🚫

Unbiased 🕢

Variance: $\operatorname{Var}\left(\hat{\theta}^{\operatorname{CL}}\right) = \frac{1}{n}\operatorname{Var}(Y)$ $\operatorname{Var}\left(\hat{\theta}^{\operatorname{PP}}\right) \approx \frac{1}{n}\operatorname{Var}(Y - f(X))$

$$\operatorname{Var}\left(\hat{\theta}^{\operatorname{PP}}\right) \approx \frac{1}{n} \operatorname{Var}(Y - f(X))$$

 \rightarrow When predictions are good and $N \gg n$, $\hat{\theta}^{PP}$ has lower variance!

Proteomics with AlphaFold

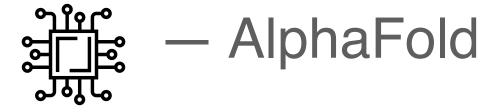
PLOS BIOLOGY

 X_i — protein sequences, Y_i — indicator of disorder (IDR)

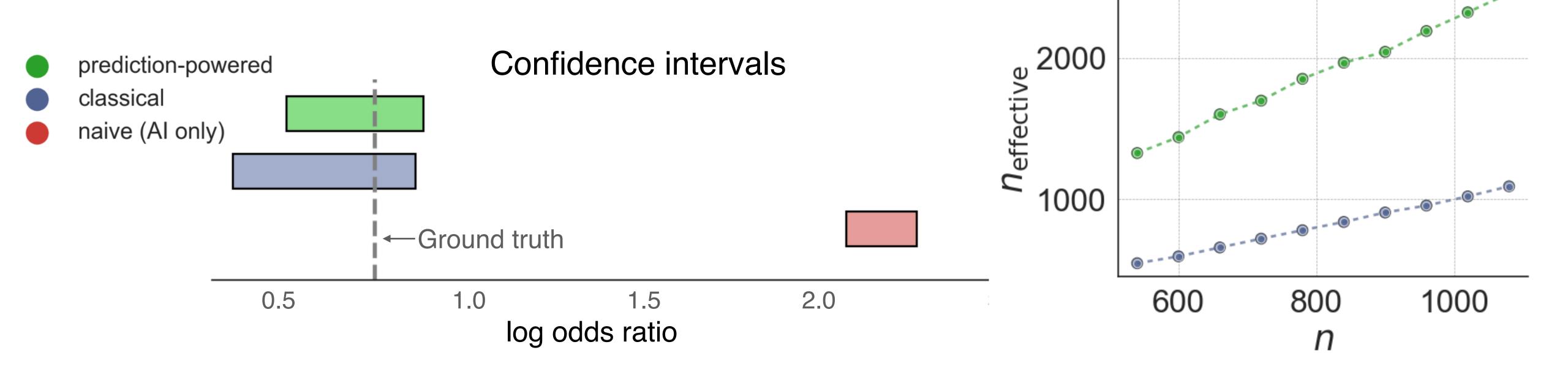
T — relationship between disorder and phosphorylation

The structural context of posttranslational modifications at a proteome-wide scale

Isabell Bludau¹, Sander Willems¹, Wen-Feng Zeng¹, Maximilian T. Strauss², Fynn M. Hansen¹, Maria C. Tanzer¹, Ozge Karayel¹, Brenda A. Schulman³, Matthias Mann^{1,2}*



Protein disorder vs phosphorylation



Which labels are most informative?

X_1	?
X_2	?
X_{N-1}	?
X_N	?

--- should prioritize collecting expert labels where AI makes a mistake

→ learn more given the same budget

want p_i = Prob(collect label Y_i) small for "easy-to-predict" data points and large for "hard" ones

$$T^{\text{adaptive}} = \text{mean}(\hat{Y}_1, ..., \hat{Y}_N) - \text{mean}\left(\frac{1}{p_1}(\hat{Y}_1 - Y_1), ..., \frac{1}{p_n}(\hat{Y}_n - Y_n)\right)$$

bias of Al predictions when expert labels are collected adaptively

No matter the Al bias, can achieve the gold standard!

Data-adaptive sampling with active inference

$$\hat{\theta}^{\text{Active}} = \frac{1}{N} \sum_{i=1}^{N} f\left(X_i^{\text{unlabeled}}\right) + \frac{1}{n} \sum_{i=1}^{n} \frac{\xi_i}{\pi(X_i)} \left(Y_i - f\left(X_i^{\text{labeled}}\right)\right)$$

$$\hat{Y} = f(X) \qquad \text{blackbox prediction}$$

$$\xi_i \sim \text{Bern}(\pi(X_i)) \text{ indicates whether label } Y_i \text{ is collected}$$

Labeling policy: $\pi(X_i) \propto \text{uncertainty } u(X_i)$

$$u(X_i) \propto \begin{cases} \sqrt{E(Y_i - f(X_i))^2 \mid X_i} & \text{(regression)} \\ 2\min(f(X_i), 1 - f(X_i)) & f(x_i) = \hat{P}(Y_i = 1 \mid X_i) & \text{(classification)} \end{cases}$$

This can be done sequentially: fine-tune f after collecting some labels

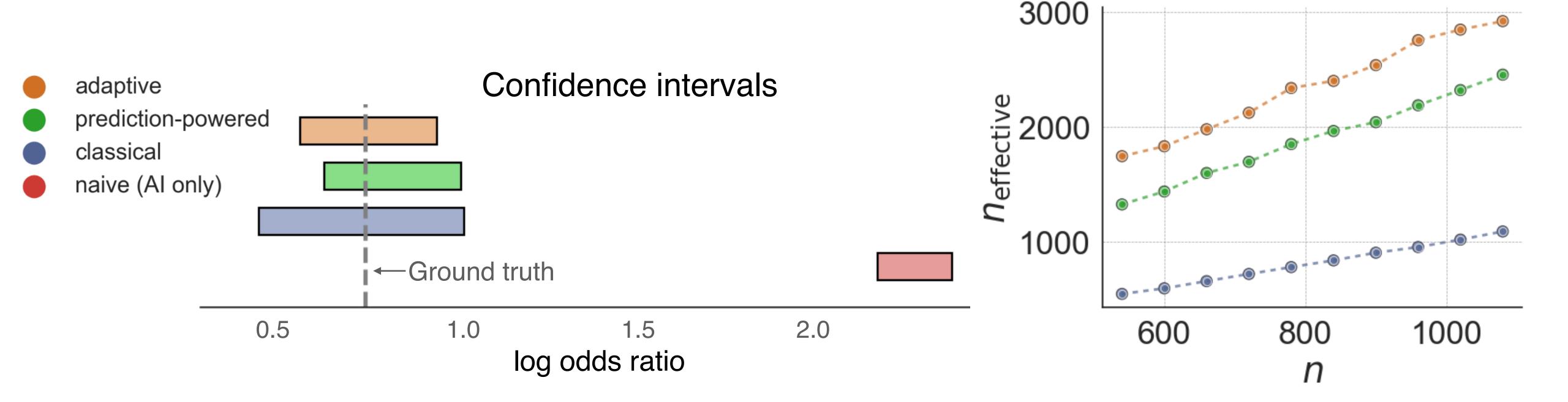
Proteomics with AlphaFold

 X_i — protein sequences, Y_i — indicator of disorder (IDR)

T — relationship between disorder and phosphorylation



Protein disorder vs phosphorylation



All-purpose efficient dataset labeling

X_1 X_2	$egin{array}{c} ilde{Y}_1 \ ilde{Y}_2 \end{array}$
X_{N-1}	\tilde{Y}_{N-1}
X_N	$ ilde{ ilde{Y}_N}$

N unlabeled data points

Want to collect labels \tilde{Y}_i that are both accurate and cheap

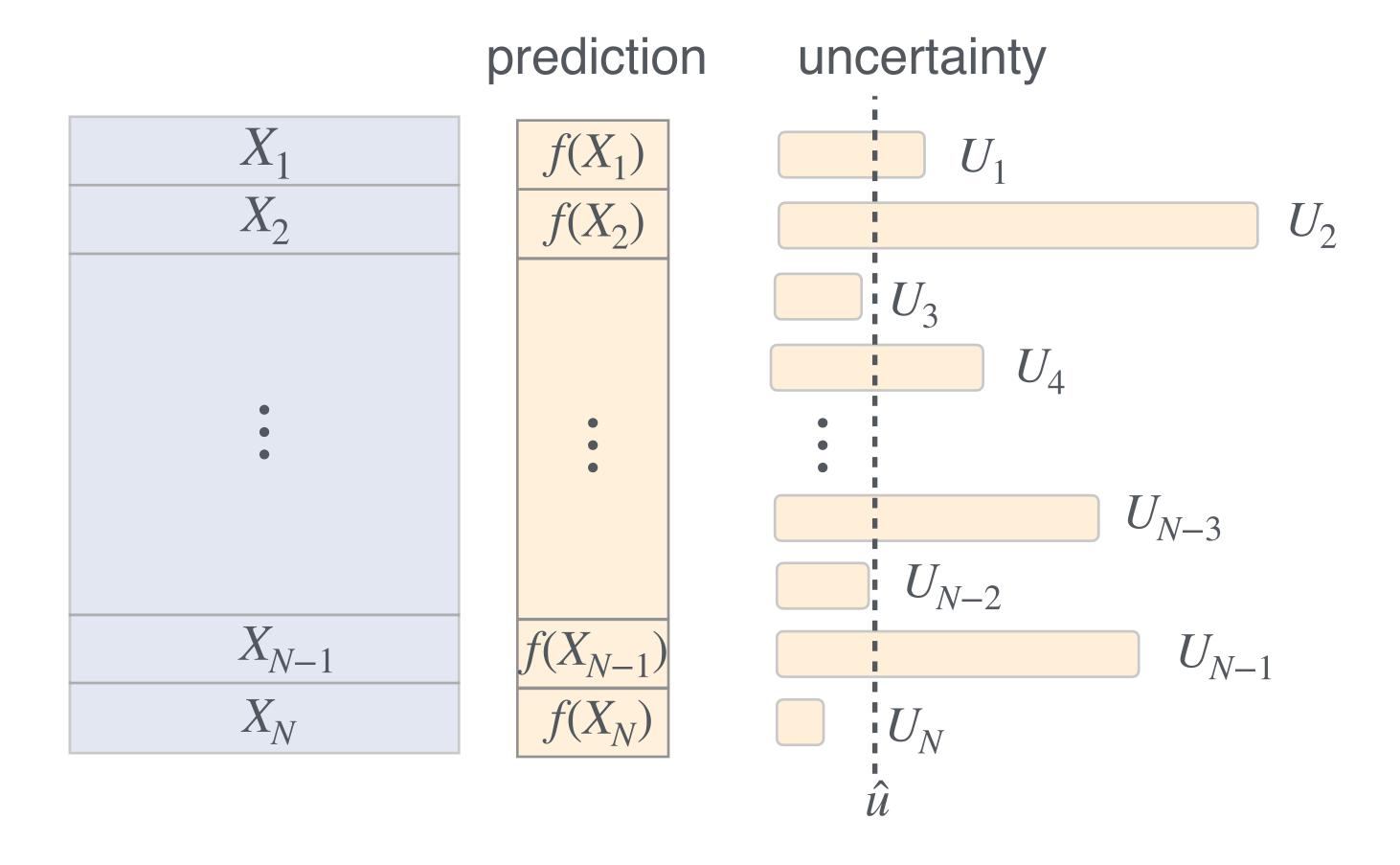
 \tilde{Y}_i can be either expert label Y_i (expensive) or Al prediction \hat{Y}_i (cheap)

$$\tilde{Y}_i^u = Y_i \cdot \mathbf{1} \{ U_i \ge u \} + \hat{Y}_i \cdot \mathbf{1} \{ U_i < u \}$$

Goal: return labeled dataset $(X_1, \tilde{Y}_1), \ldots, (X_N, \tilde{Y}_N)$, while collecting as few expert Y_i as possible, such that

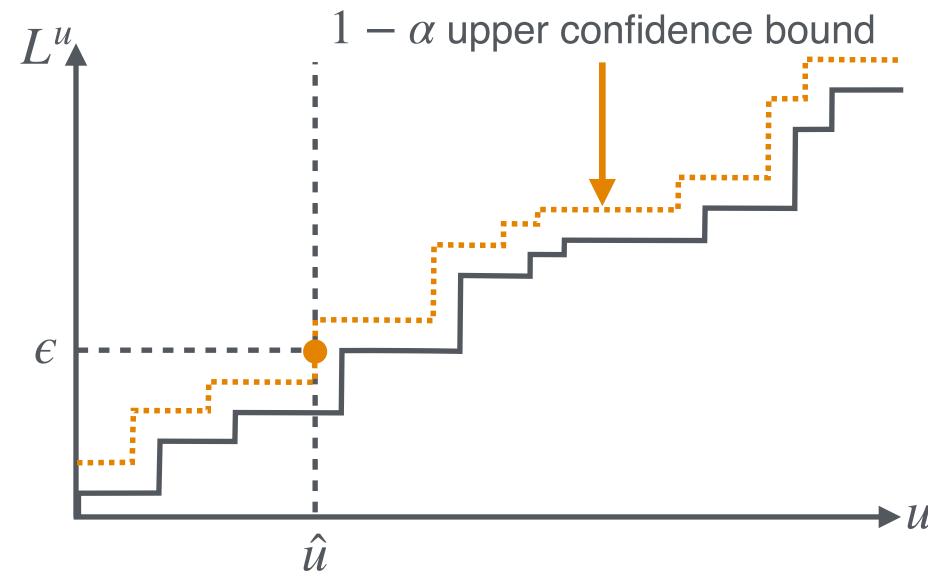
 $\tilde{Y}_i = Y_i$ for 95% of the dataset, with 95% probability

PAC labeling method



$$\tilde{Y}_i^u = Y_i \cdot \mathbf{1} \{ U_i \ge u \} + f(X_i) \cdot \mathbf{1} \{ U_i < u \}$$

$$L^u = \frac{1}{N} \sum_{i=1}^N \mathcal{L}(Y_i, \tilde{Y}_i^u)$$



Theorem $ilde{Y}_1^{\hat{u}},..., ilde{Y}_N^{\hat{u}}$ are PAC labels

All-purpose social science PAC labelling with GPT-40

Dataset	Metric	Method	
		Our approach 🧼	ChatGPT only
Misinformation [1]	Budget save (%) Error	(18.12 ±4.93)% 3.80%	18.56%
Stance on global warming [2]	Budget save (%) Error	(28.09 ±3.28)% 4.57%	24.79%
Media bias [3]	Budget save (%) Error	(13.79 ±3.38)% 4.10%	37.72%

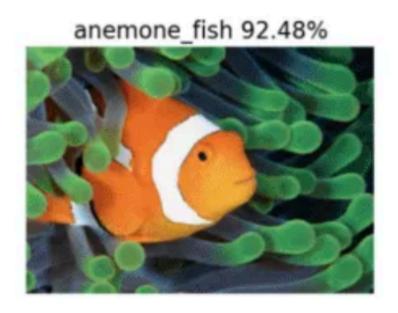
^[1] Misinfo Reaction Frames: Reasoning about Readers' Reactions to News Headlines. Gabriel et al. (2022)

^[2] Detecting Stance in Media on Global Warming. Luo et al. (2020)

^[3] We Can Detect Your Bias: Predicting the Political Ideology of News Articles. Baly et al. (2020)

Image classification with ResNet

ballplayer 69.22%







 $Y_i \in 1000 \text{ ImageNet labels}$











ImageNetV2

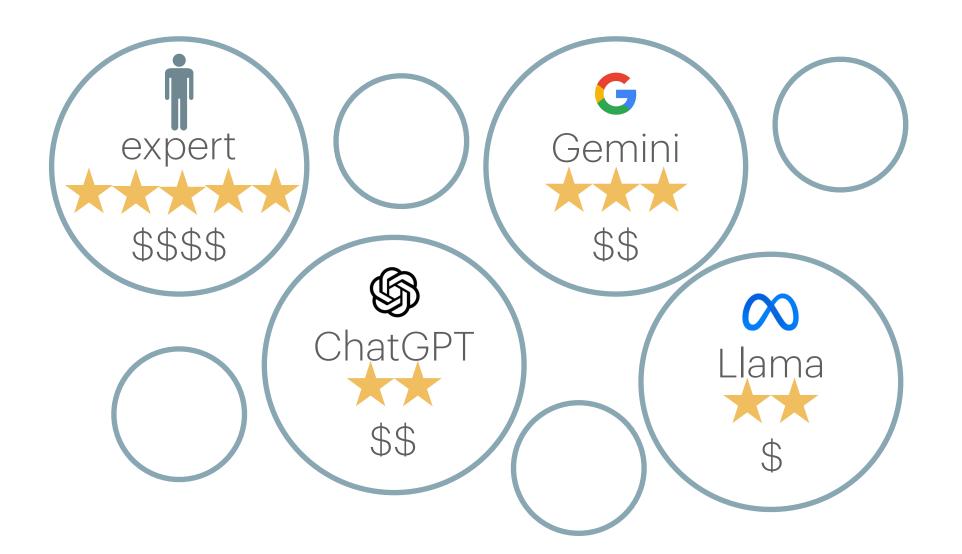
Ben Recht Ludwig Schmidt Rebecca Roelofs Vaishaal Shankar

PAC labelling image datasets with ResNet-152

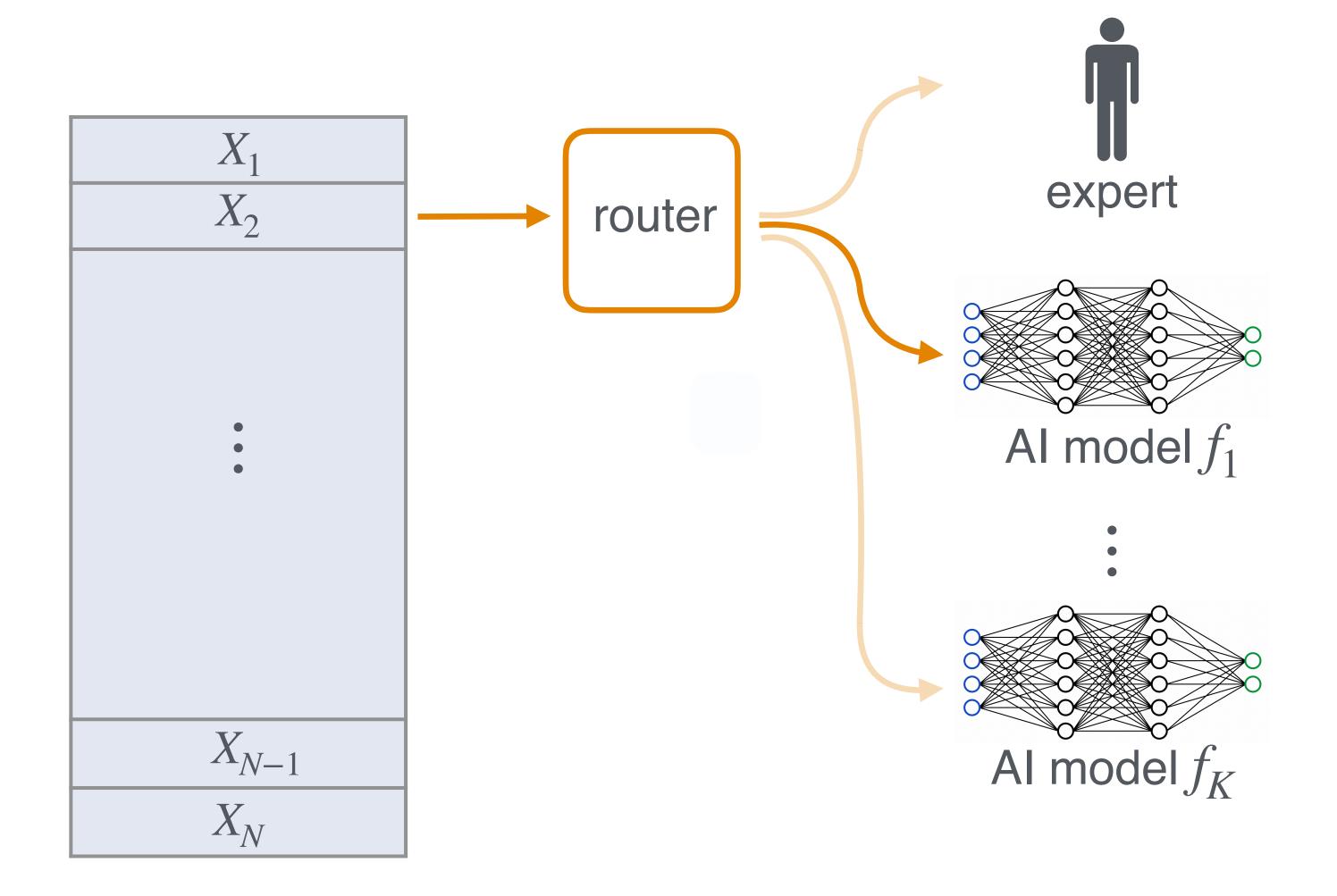
Dataset	Metric	Method	
		Our approach	ResNet only
ImageNet [1]	Budget save (%) Error	(59.64 ±1.49)% 4.73%	21.69%
ImageNet v2 [2]	Budget save (%) Error	(39.07 ±2.67)% 4.74%	35.33%

Labeling with multiple models

How do we reliably trade off between data sources of varying qualities and costs?



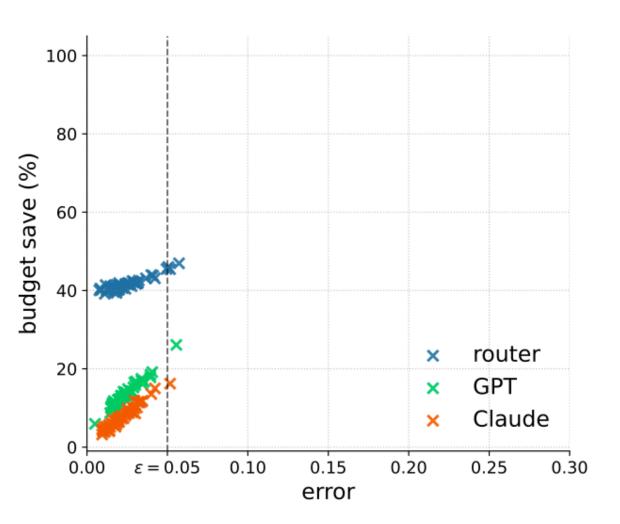
PAC labeling with multiple models

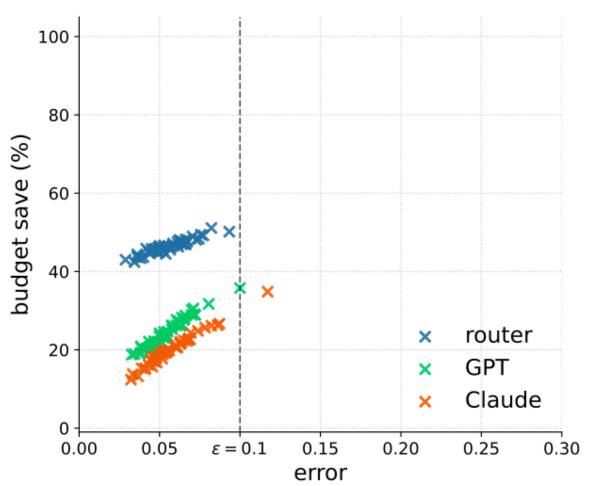


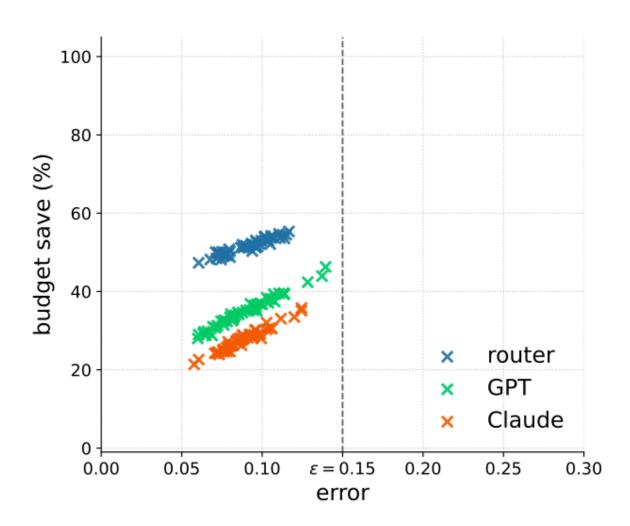
Router optimizes final labeling cost

PAC labeling with multiple models

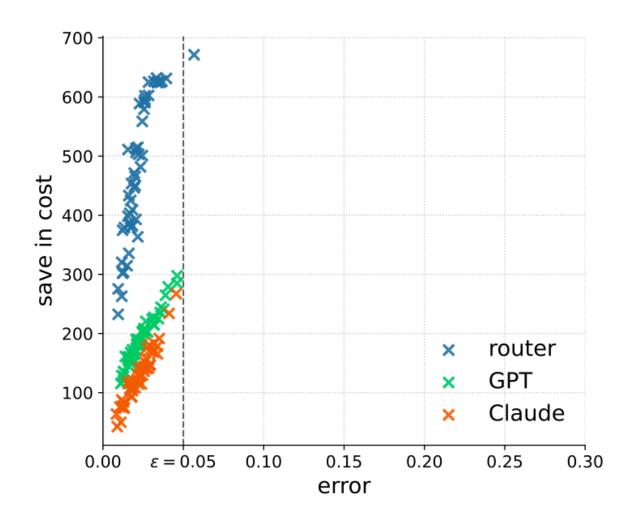
Media bias (with router)

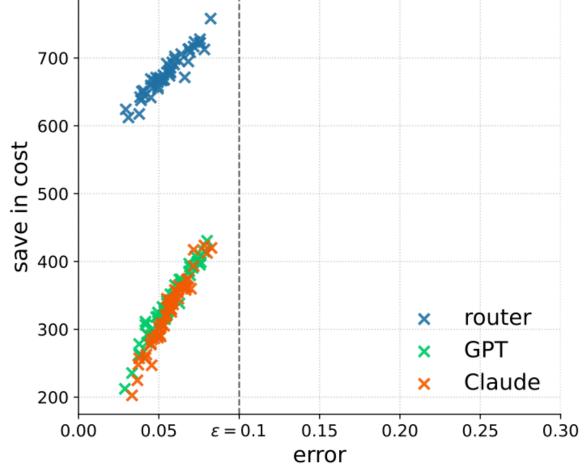


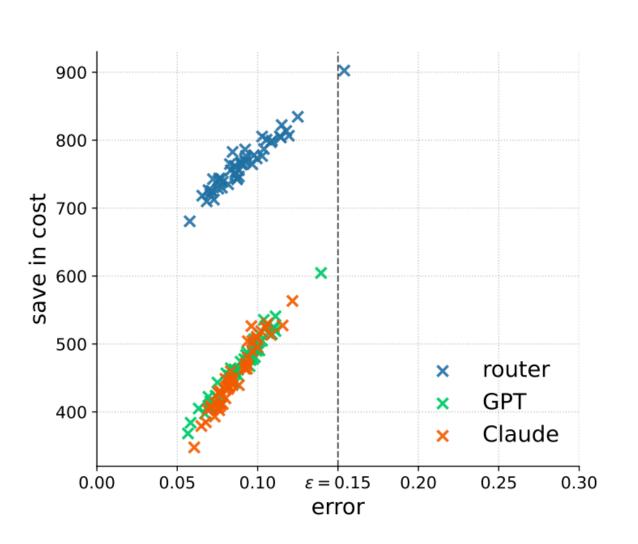




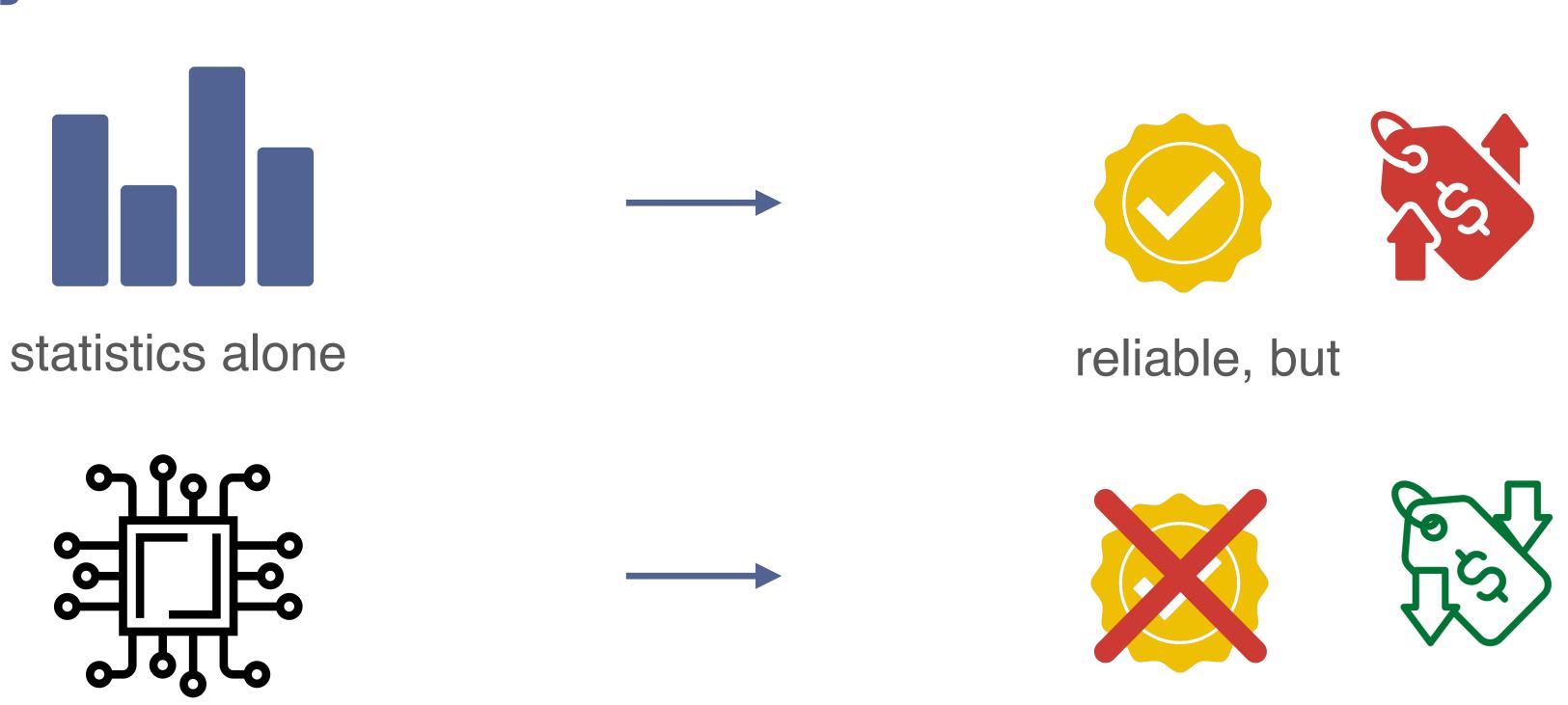
Cost-sensitive (\$) routing:

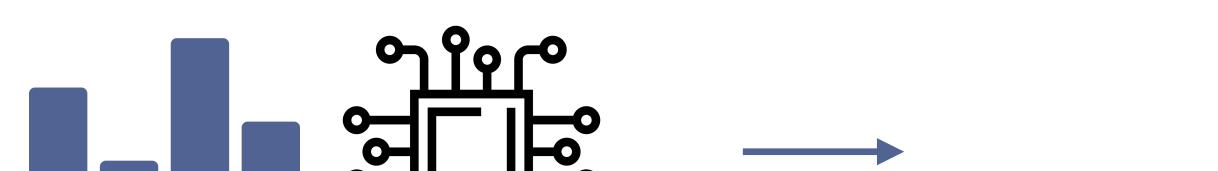






Takeaway





Al alone

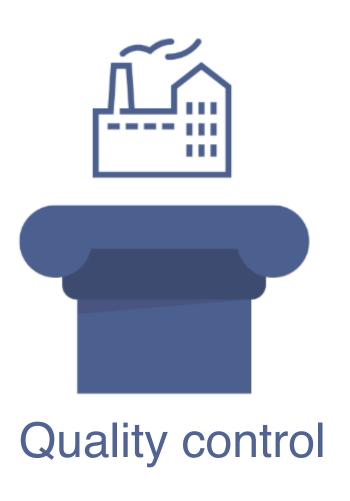
statistics + Al



not always reliable, but powerful and







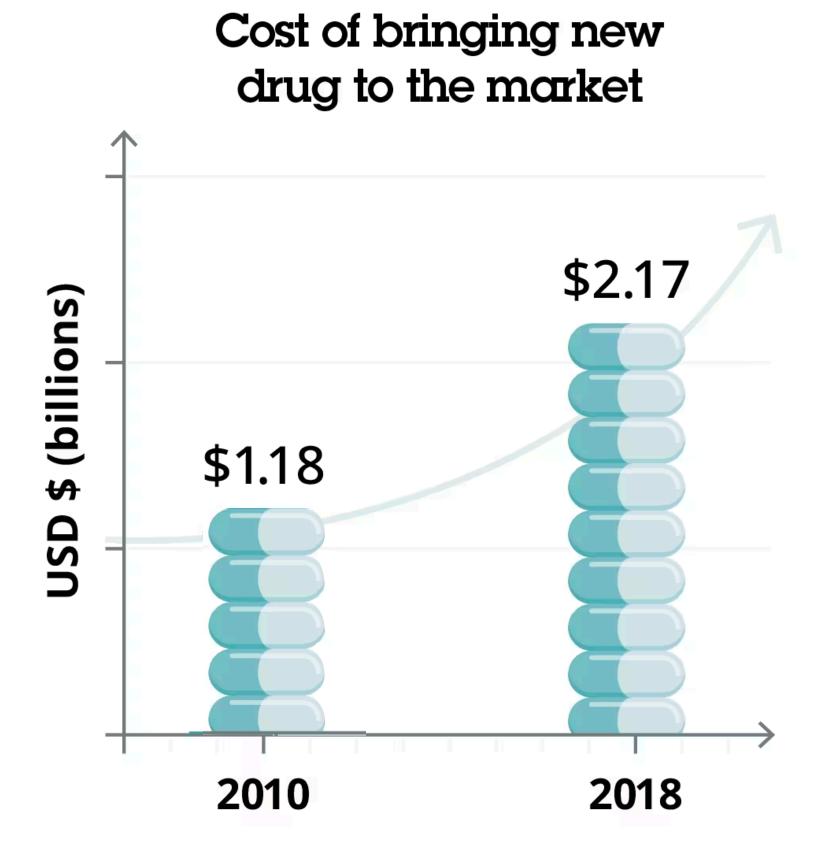


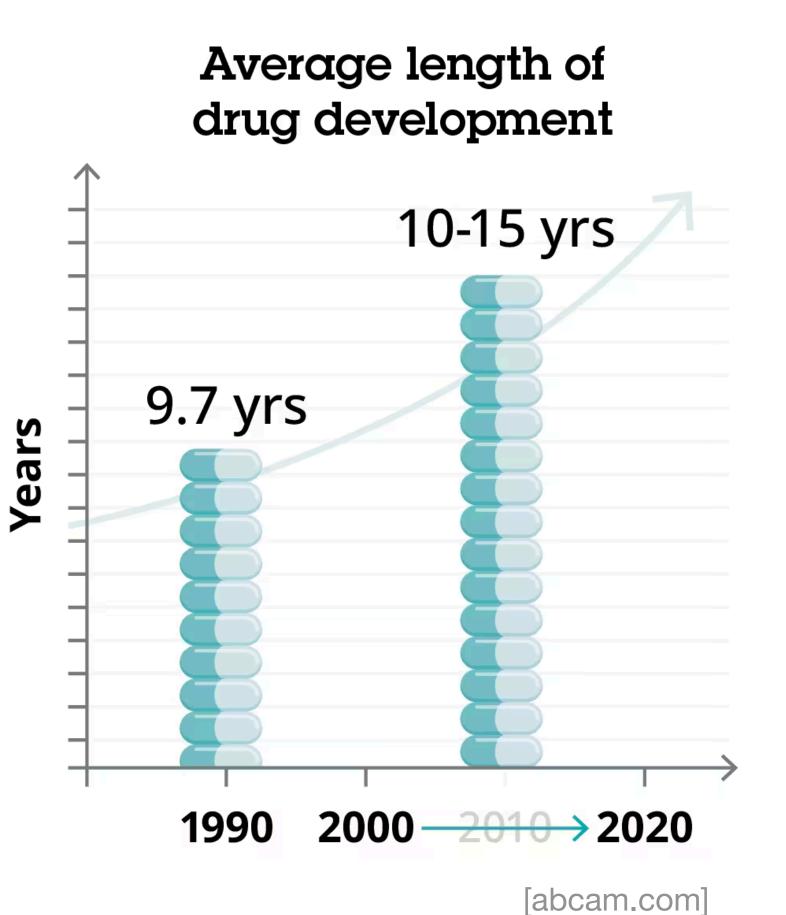


synthetic pretraining Al-powered inference datamodels **s1**

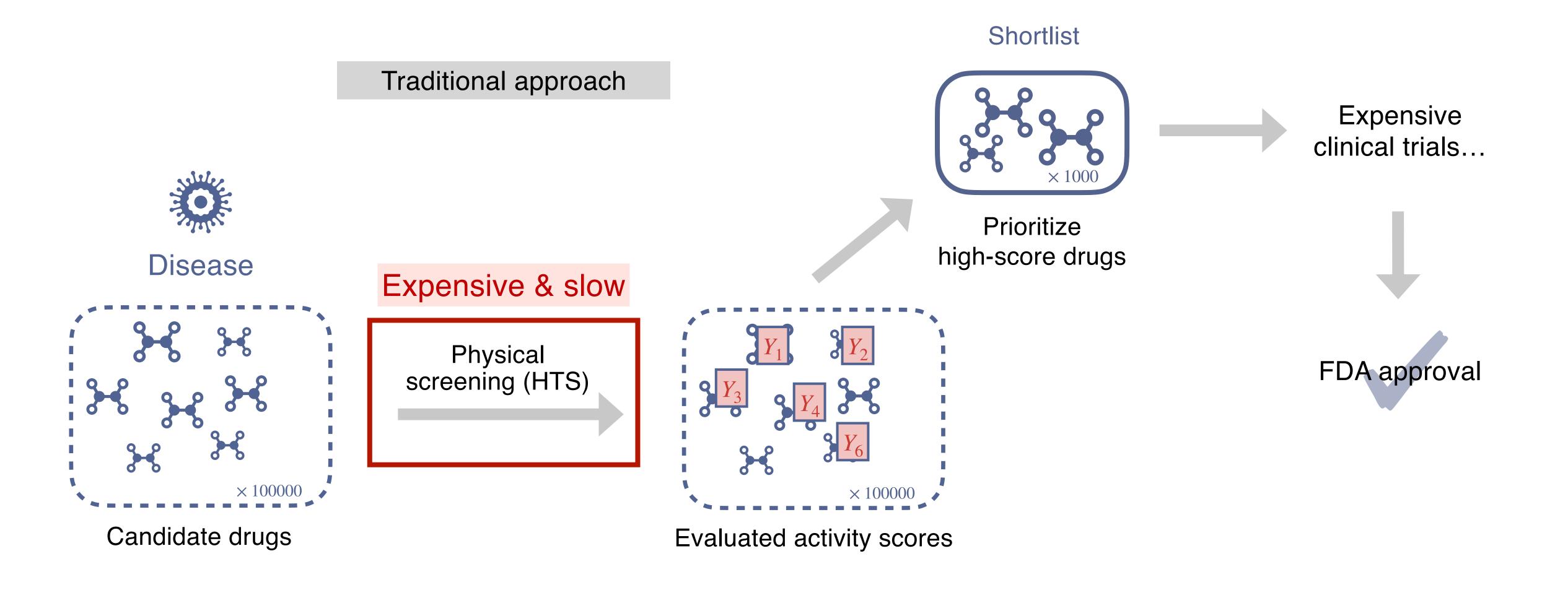
Al-powered drug discovery

Accelerating drug discovery?

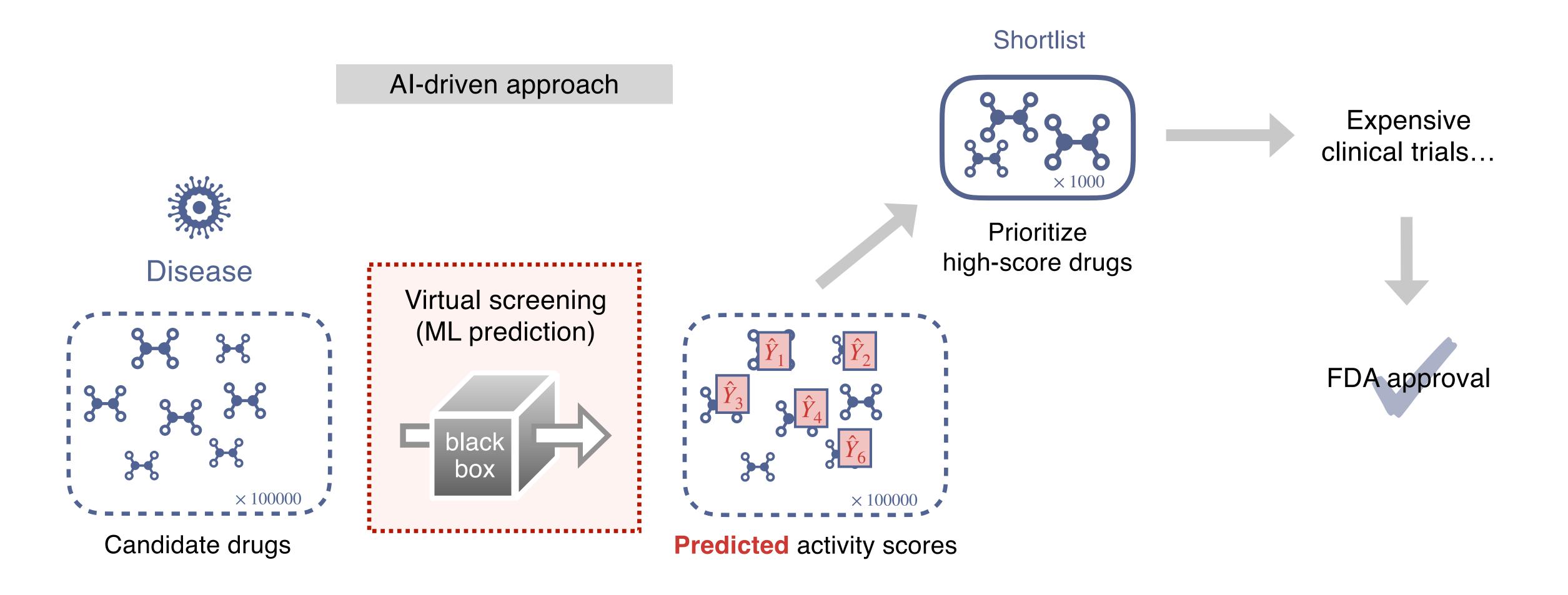




Drug discovery pipeline



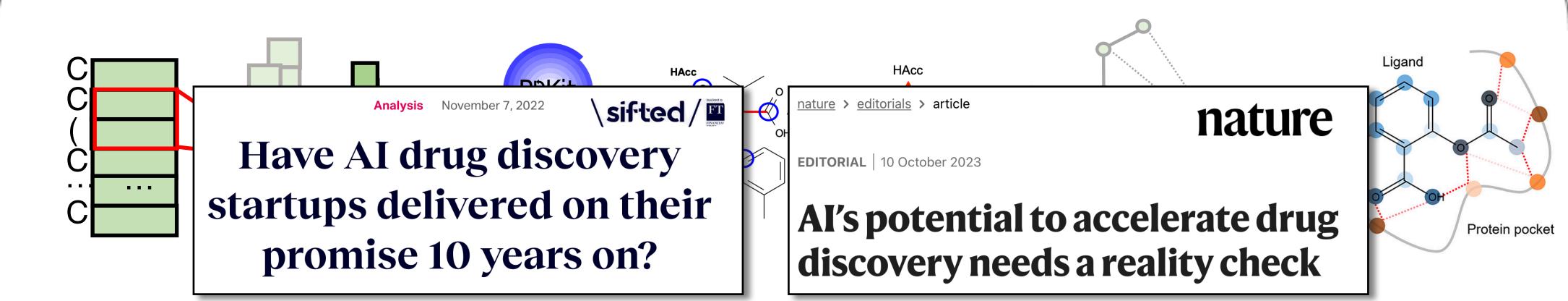
Drug discovery pipeline



Al as imperfect scoring?

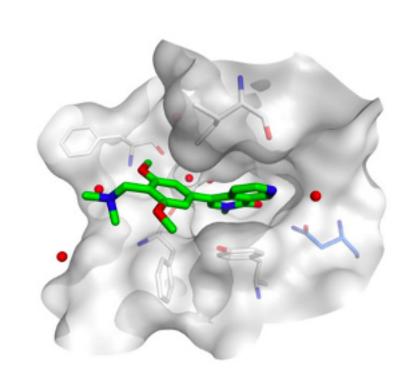
Quality control?



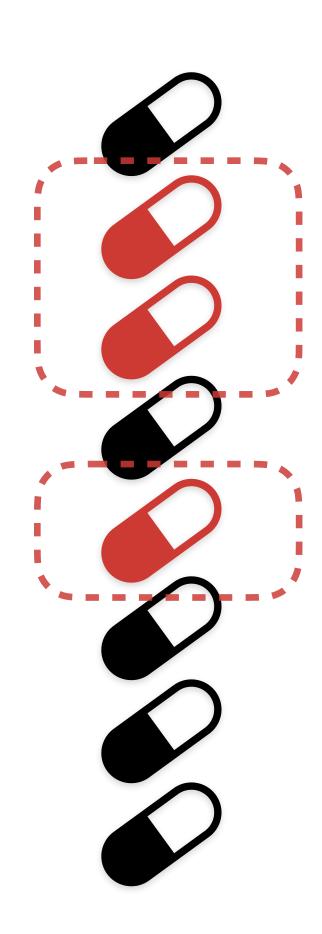


Can we make discoveries with few mistakes?

Goal: finding "actionable" instances



Want drugs with high binding affinities to a disease target

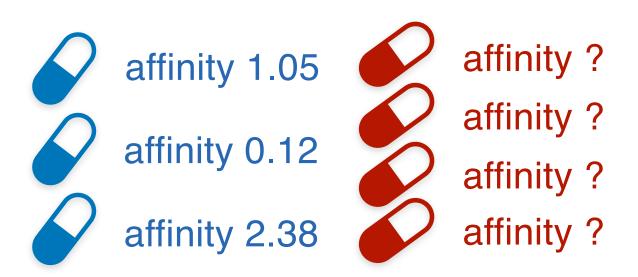




Which drugs are sufficiently active?

Problem setup

- Any pre-trained prediction model $\hat{\mu}\colon \mathcal{X} \to \mathcal{Y}$ (independent of training and test data)
 - lacktriangleright X physical/chemical feature/amino acids of the drug
 - Y binding affinity
 - $\sim Y \in \{0,1\}$: whether the drug binds to the target
 - $\prec Y \in \mathbb{R}$: how well the drug binds to the target

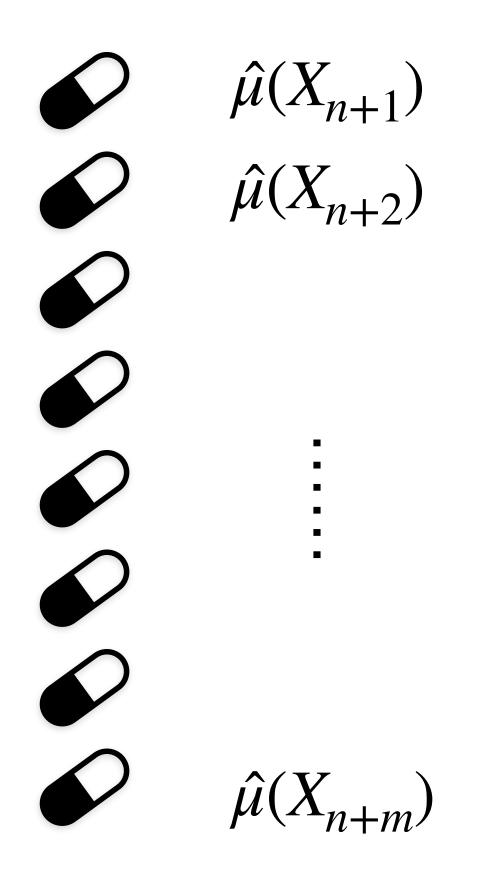


- ► Training data $\{(X_i, Y_i)\}_{i=1}^n$ (screened drugs)
- Test samples $\{(X_{n+j}, Y_{n+j})\}_{j=1}^m$ with unknown $\{Y_{n+j}\}_{j=1}^m$ (new drugs)

Goal: find large outcomes $Y_{n+j} > c_{n+j}$ without too many errors

 \sim user-specified thresholds c_{n+i} to become 'interesting'

Challenges



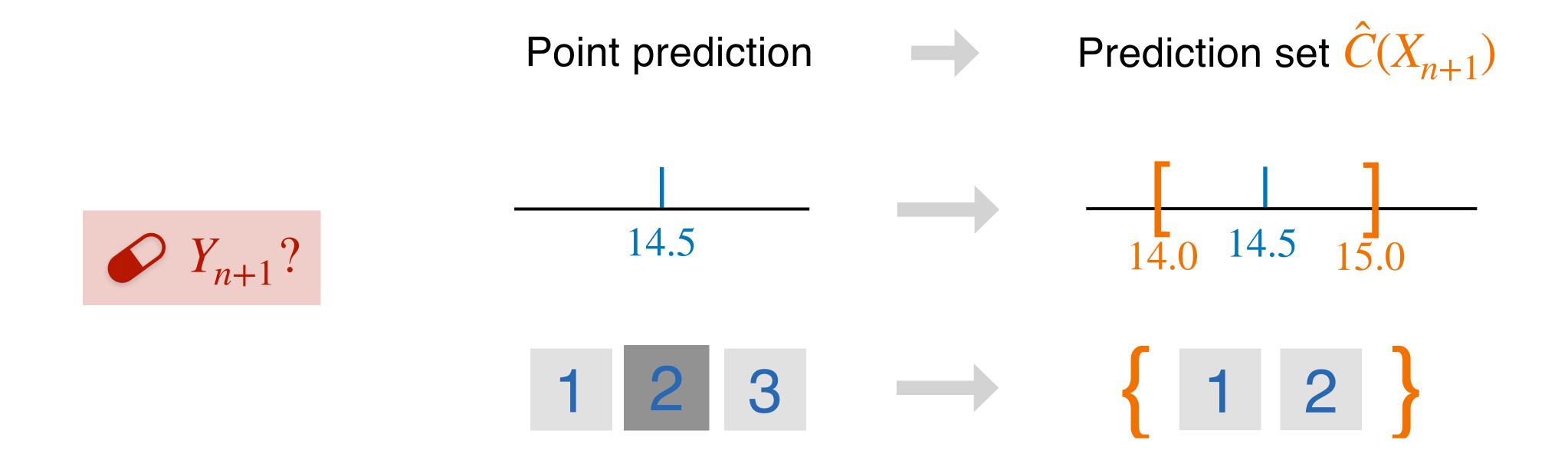
Quantifying uncertainty in point predictions

Model-free

Work for any prediction model No modeling assumptions

Distribution shift

Conformal prediction: model-free uncertainty quantification

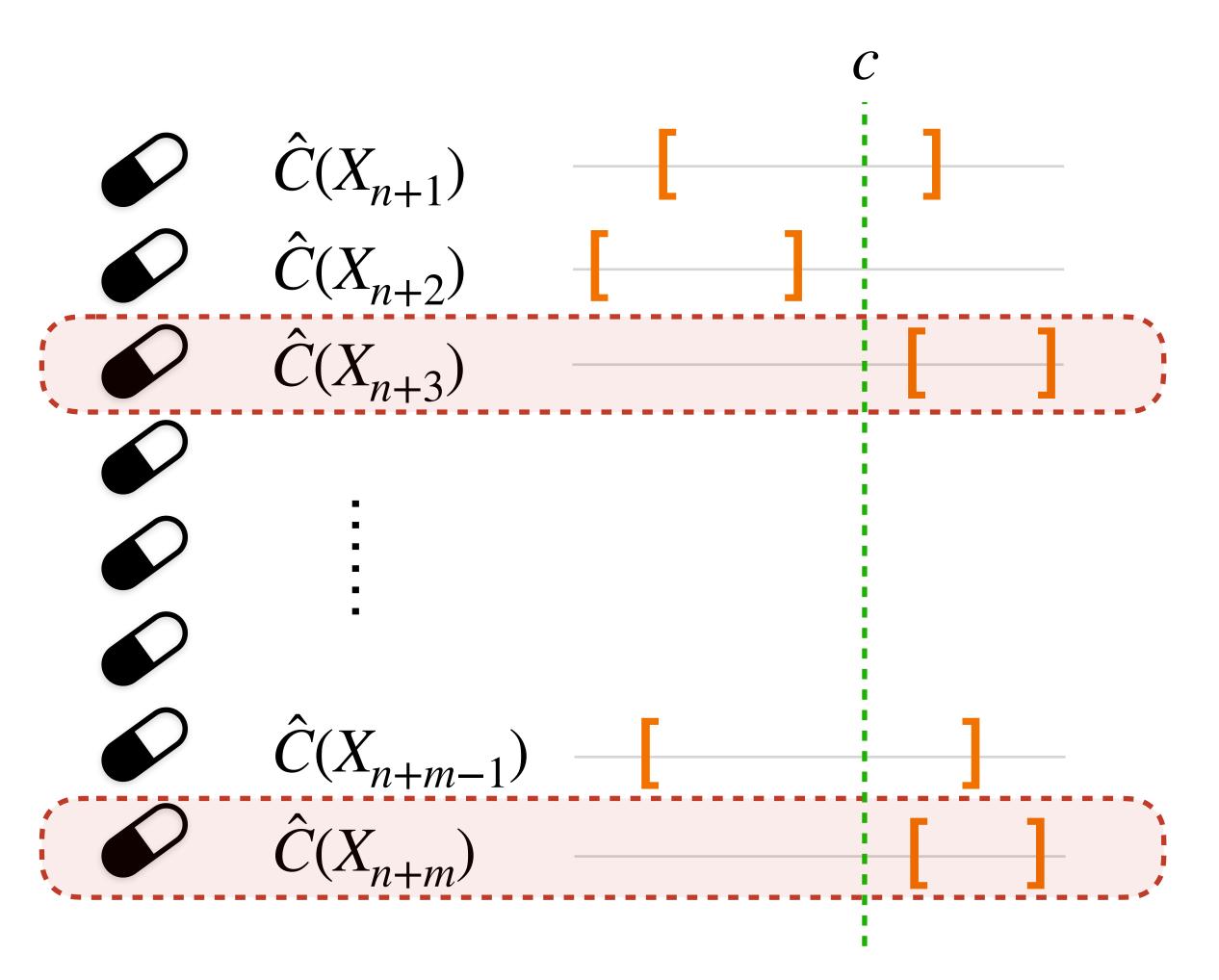


Validity of conformal prediction intervals (PIs) [Vovk et al., 1999]

$$\mathbb{P}\left(Y_{n+1} \in \hat{C}(X_{n+1})\right) \ge 95\%$$

~ Covers 95% of outcomes <u>no matter prediction model</u>

Challenges

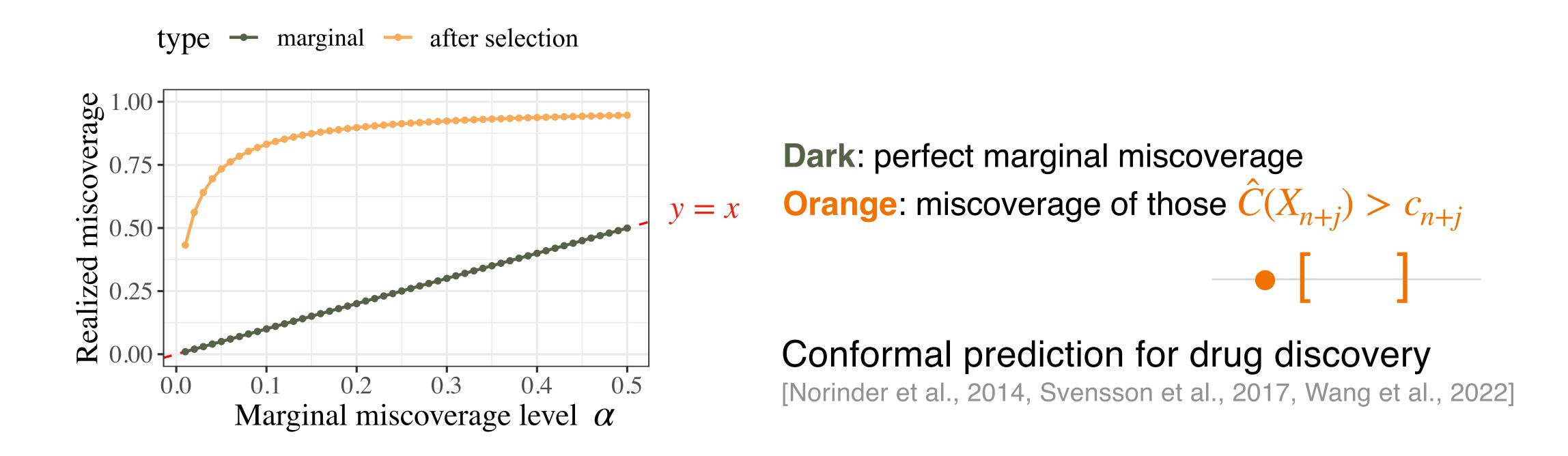


- Uncertainty quantification
- ► Model-free ✓

Can we use them to find interesting instances (drugs)?



Miscoverage of naive selection



1% nominal error, yet >30% error after selection!

This is the winner's curse [Soric, 1989]

Inspired a whole field of research: Selective Inference

[Benjamini and Yekutieli, 2005, Berk et al., 2013, Taylor et al., 2014, Fithian et al., 2014; Storey et al, 2003]

Our proposal: select with guarantees

Find "actionable instances" while controlling fraction of false positive (FDR)

FDR =
$$\mathbb{E}[\text{FDP}]$$
, FDP = $\frac{\#\{\text{false discoveries}\}\}}{\#\{\text{selected instances}\}\}}$

- Control of FDR implies
 - Most Al-powered decisions are correct
 - Resource allocation is efficient

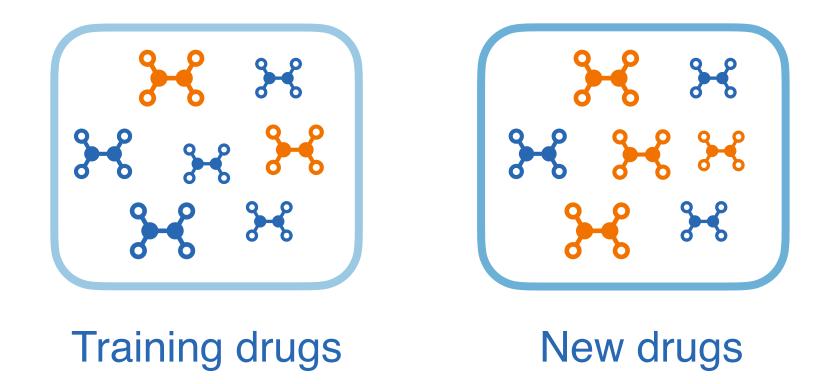


Drugs → 90% active Customers → 90% responding Patients ~ 90% benefiting LLM outputs ~ 90% trustworthy

Controlling the false discovery rate: a practical and powerful approach to multiple testing

Distribution shift

- Are my evaluated drugs comparable to the unknown drugs?
 - No if you preferred drugs with some specific structures, etc



- In reality: <u>distribution shift</u> when generating/exploring new drugs
 - → Similar issues in job hiring, health monitoring, counterfactual inference...

Distribution shift

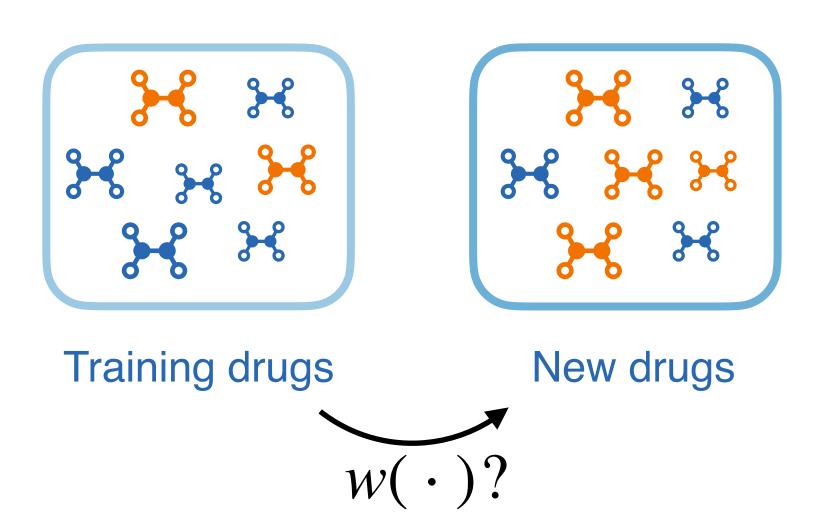
- ► Test data $\{(X_{n+j}, Y_{n+j})\}$ ~ \mathbb{Q} (unknown)
- ▶ Covariate shift: training data $\{(X_i, Y_i)\}$ ~ \mathbb{P} obeying

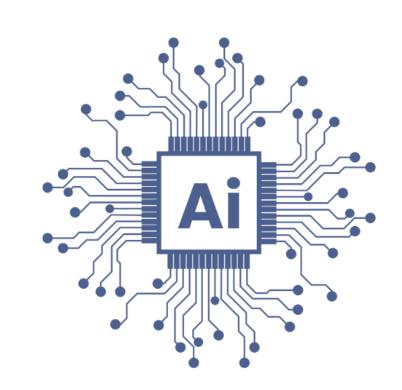
$$\frac{dQ}{dP}(x,y) = w(x)$$

for some (known or estimable) weight function $w\colon \mathscr{X} o \mathbb{R}^+$ [Sugiyama et al., 2007, Tibshirani et al., 2019

lacktriangle Why? Training data collected by looking at X (drugs, job applicants...)

Entropy balancing for distribution shift adjustment





 $\phi(\,\cdot\,)$: hidden embeddings from AI models

Finding "simplest" weights that balance key representations across batches [Hainmueller, 2012]

$$\begin{aligned} & \text{maximize}_{\mathbf{w}} \quad \sum_{i=1}^{n} -w_{i} \log w_{i} \\ & \text{subject to} \quad \left| \frac{1}{n} \sum_{i=1}^{n} w_{i} \phi(X_{i}) - \frac{1}{m} \sum_{j=1}^{m} \phi(X_{n+j}) \right| \leq \delta \\ & w_{i} \geq 0 \quad \frac{1}{n} \sum_{i=1}^{n} w_{i} = 1 \end{aligned}$$

Obtaining valid confidence measures

Weighted conformal p-values:

pprox weighted rank of \hat{V}_{n+j} among training scores $\{V_i\}_{i=1}^n$

$$p_{j} = \frac{\sum_{i=1}^{n} w(X_{i}) \mathbf{1} \{ V_{i} < \hat{V}_{n+j} \} + U_{j} \cdot w(X_{n+j})}{\sum_{i=1}^{n} w(X_{i}) + w(X_{n+j})}, \quad U_{j} \sim \mathsf{Unif}[0,1]$$

- Calibration and test scores: $V_i = Y_i \hat{\mu}(X_i)$ and $\hat{V}_{n+j} = c_{n+j} \hat{\mu}(X_{n+j})$
- ▶ Valid p-value in sense that $\mathbb{P}(p_i \leq \alpha, Y_{n+j} \leq c_{n+j}) \leq \alpha$ if w(X) is known

Takeaway: small $p_j \Longrightarrow$ small $\hat{V}_{n+j} \Longrightarrow$ small $c_{n+j} - \hat{\mu}(X_{n+j}) \Longrightarrow Y_{n+j}$ is likely large and above threshold



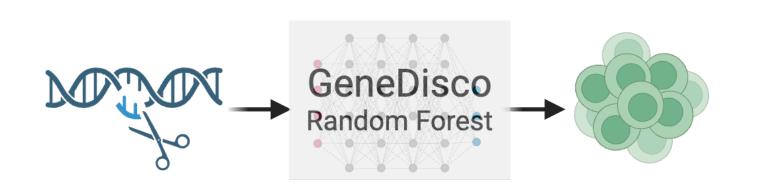
Selection Task (Distribution Shift)

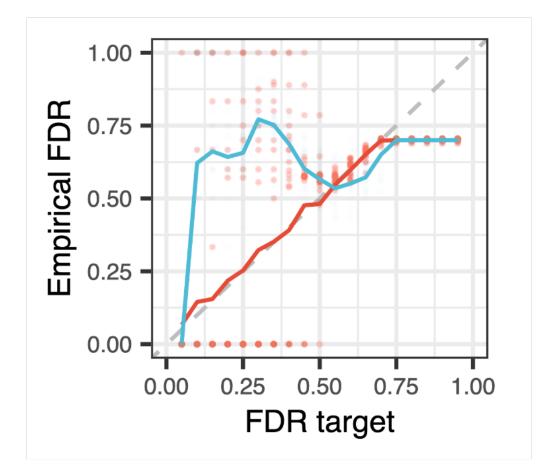
Selection FDR Control

- Conformal-Select
- Baseline

Covariates: learned representation in the hidden layer of neural nets

Select gene perturbations with high T-cell proliferation (Uniform)

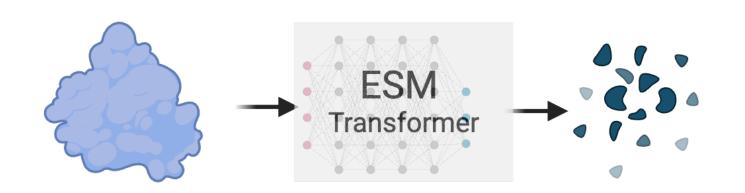


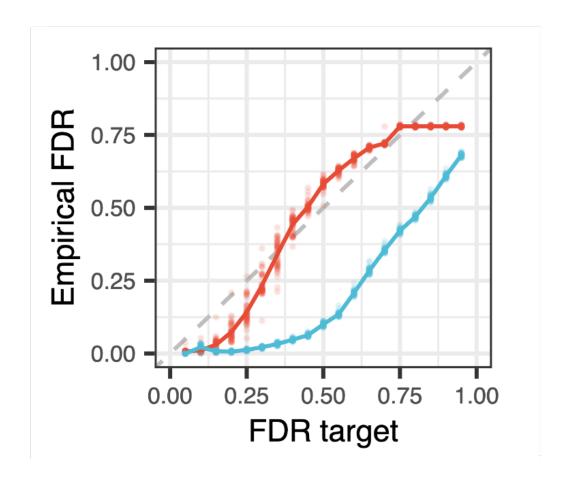


1: Gene perturbation selection

Setup with no shift

Select proteins with high stability (Mutant shift)





2: Protein stability selection

Significant shift from proteins in four rounds of experiments to single-mutation proteins



Selection Task (Distribution Shift)

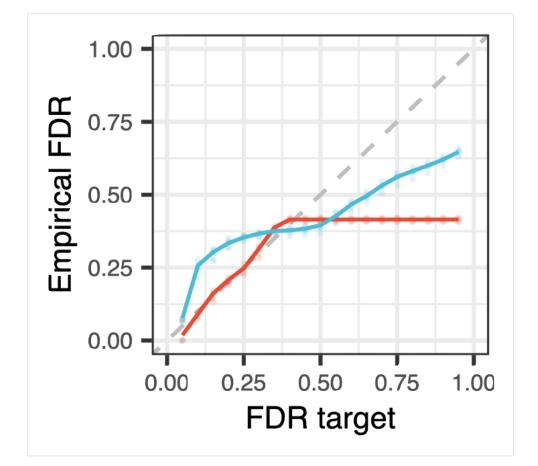
Selection FDR Control

- Conformal-Select
- Baseline

Covariates: learned representation in the hidden layer of neural nets

Select compounds with low CYP2C9 inhibition rate (Scaffold shift)



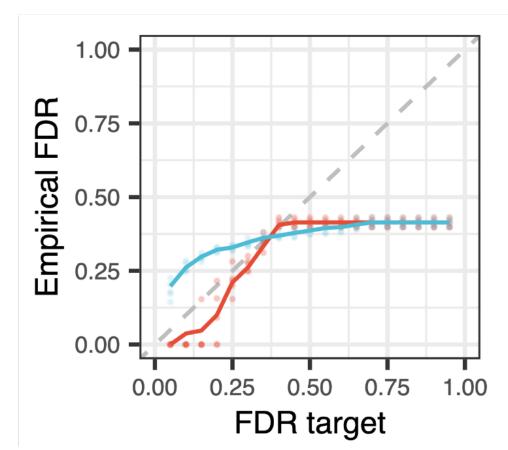


3: Drug property selection

Shift in drug structure (scaffold)

Select clinical trials that meet primary outcome (Temporal shift)

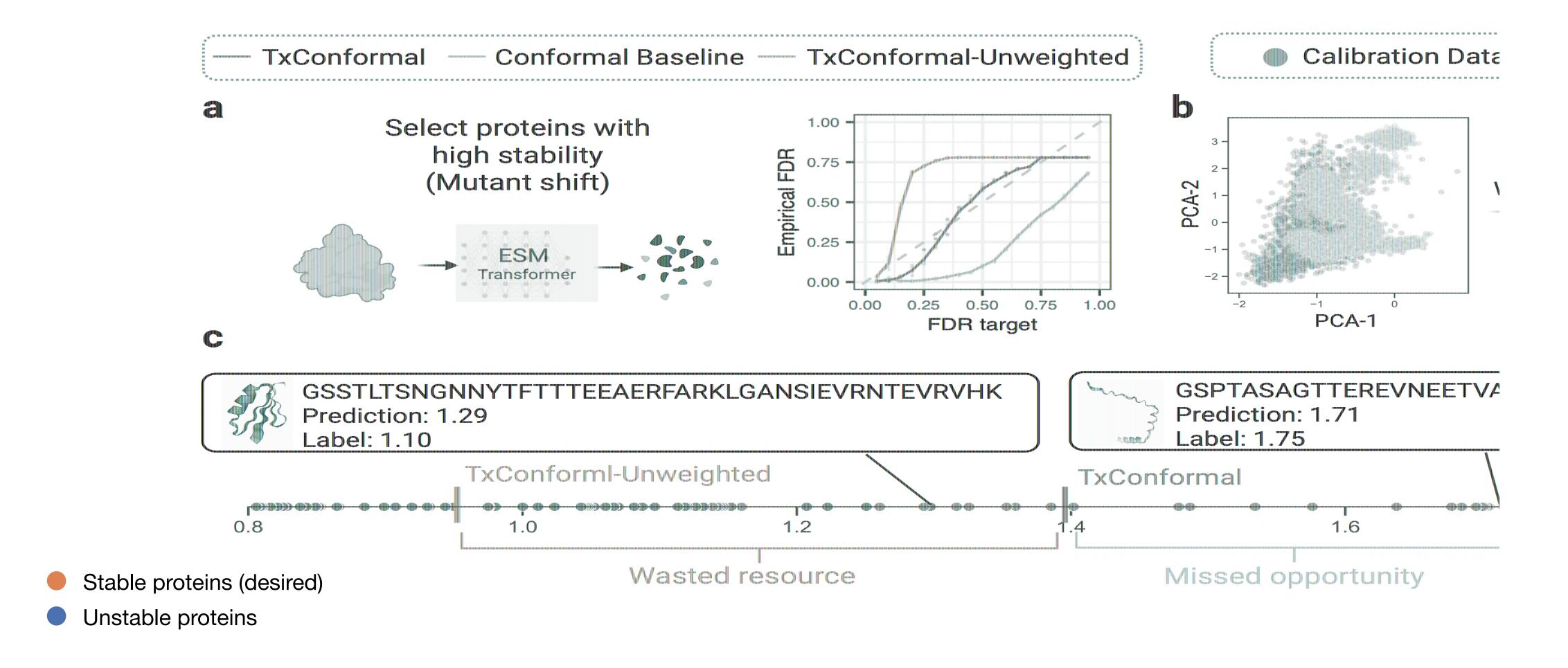




4: Trial outcome prediction

Shift from earlier to future trials





Adjusting for distribution shift yields accurate FDR control



Controlling other metrics than FDR in selecting promising drug candidates

Select compounds with high CYP2D6 inhibition rate (Scaffold shift)



Question

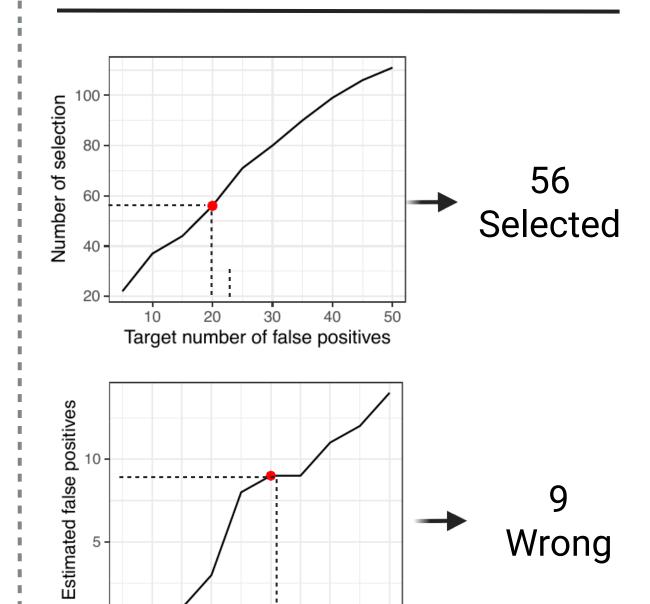
c Scenario 3

How many candidates can I test until I make 20 mistakes?

d Scenario 4

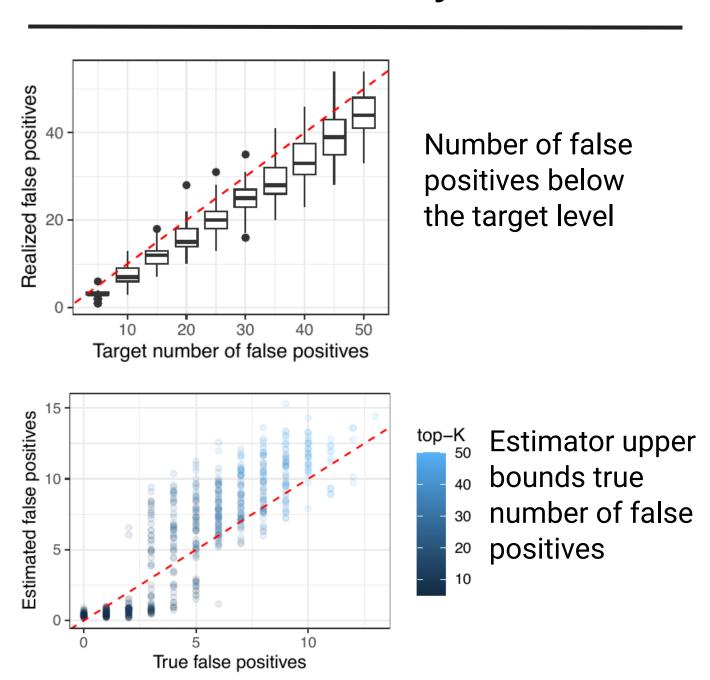
I want to screen my top 30 candidates.
How many of them are expected to be wrong?

Result



Number of top-ranking units

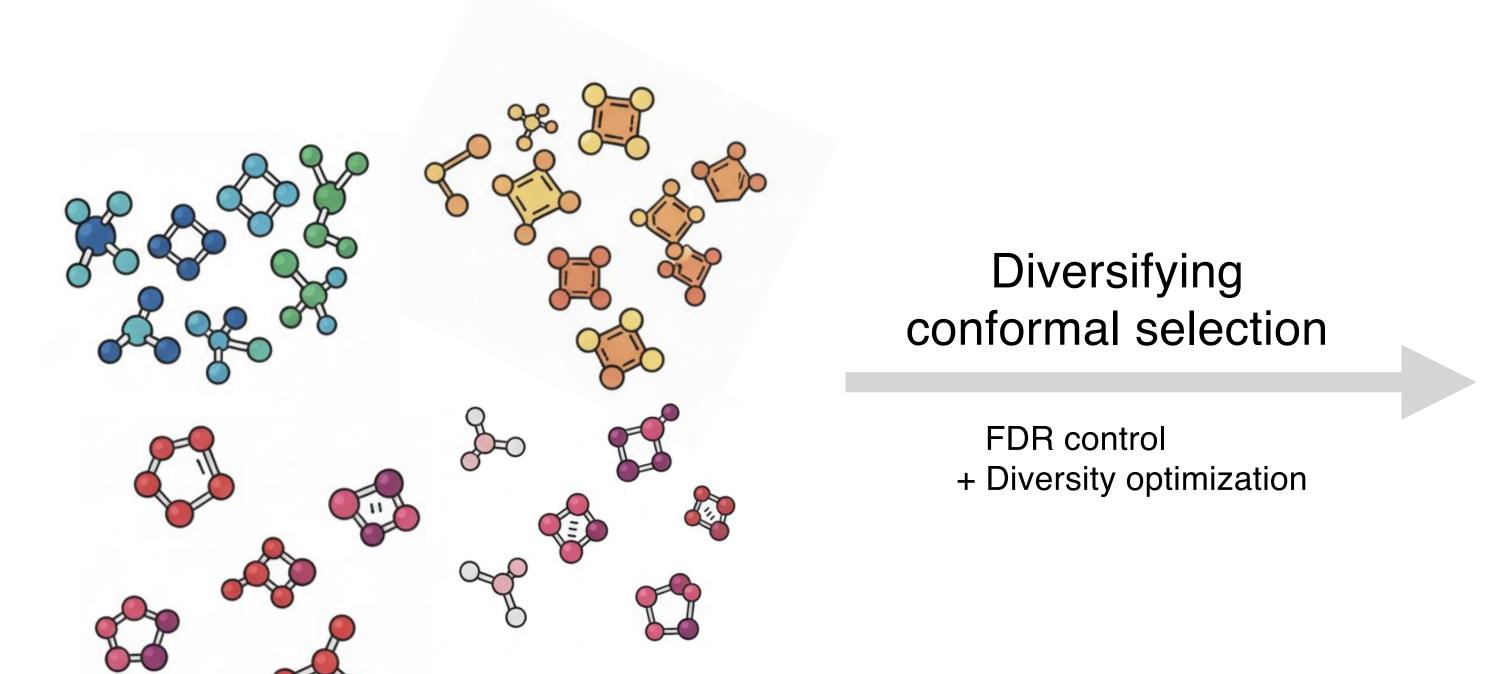
Validity



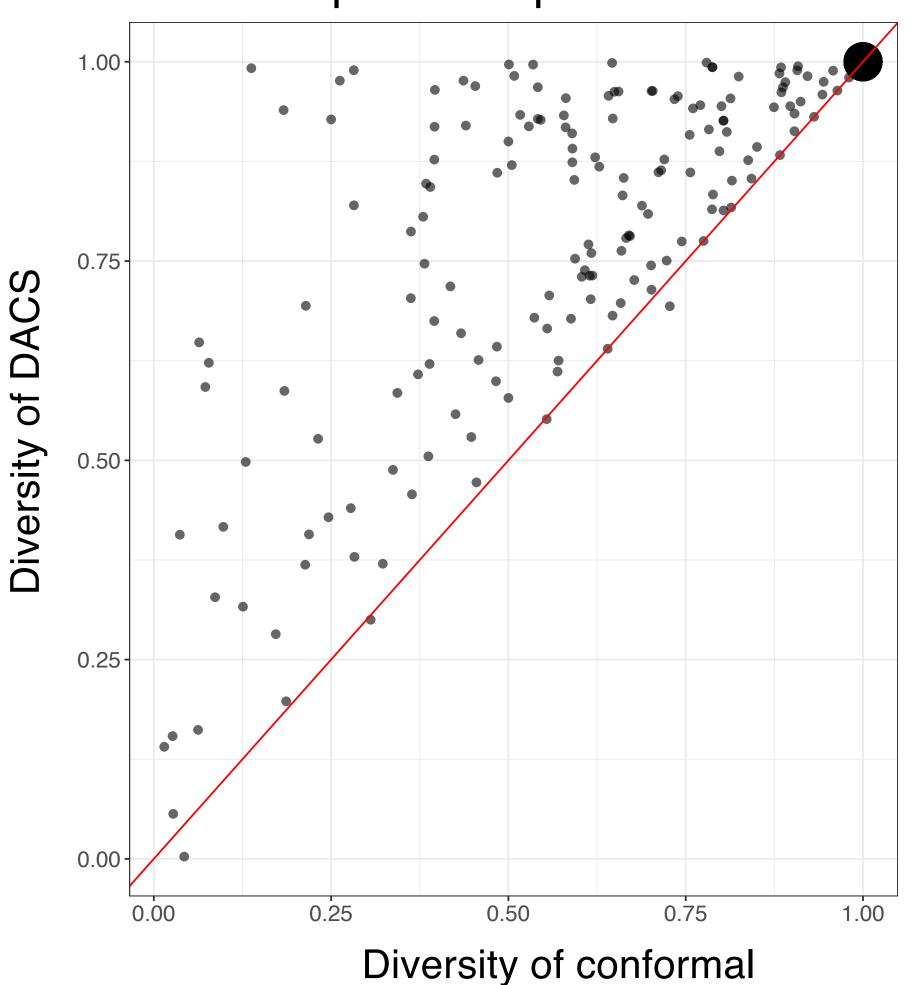
Diversifying conformal selections

How to identify structurally diverse drug candidates with FDR control?

Diversity-aware conformal selection (DACS) Nair et al. (2025)

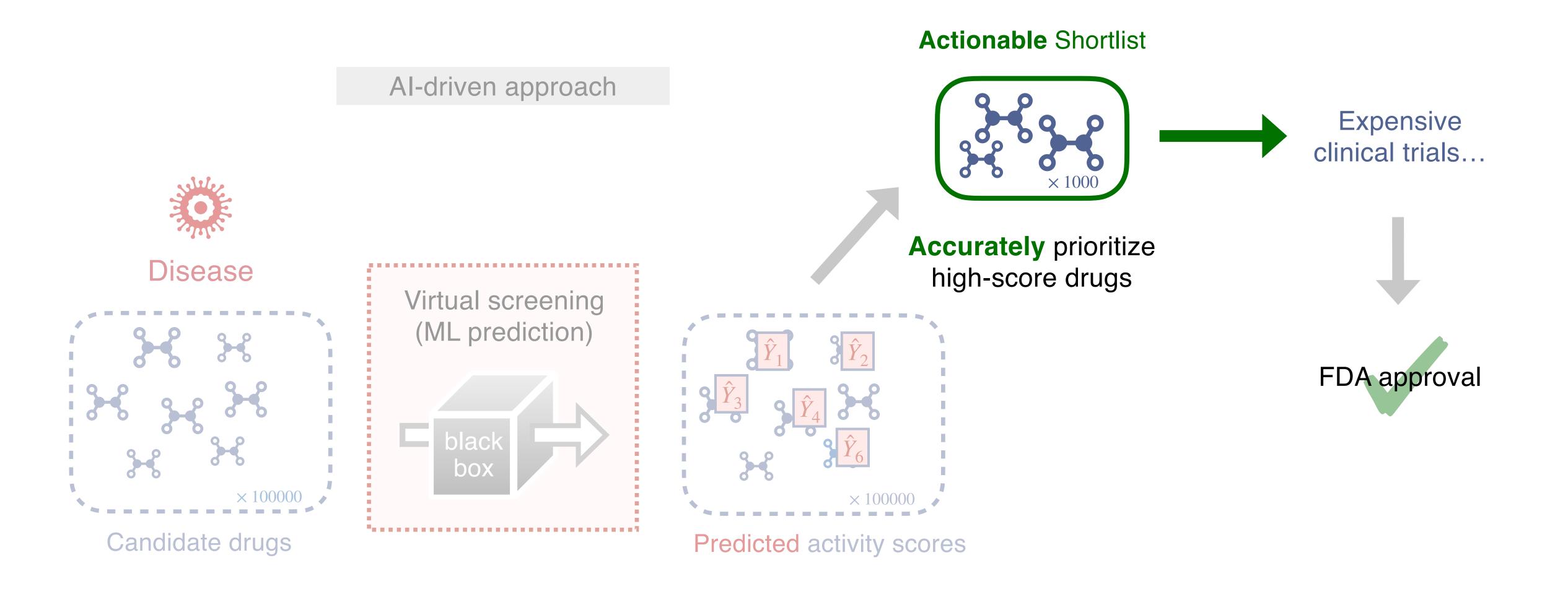


Diversity of selections on G-protein coupled receptor dataset



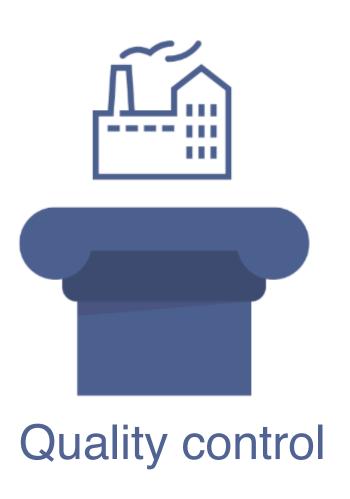
selection

Quality control for drug discovery

















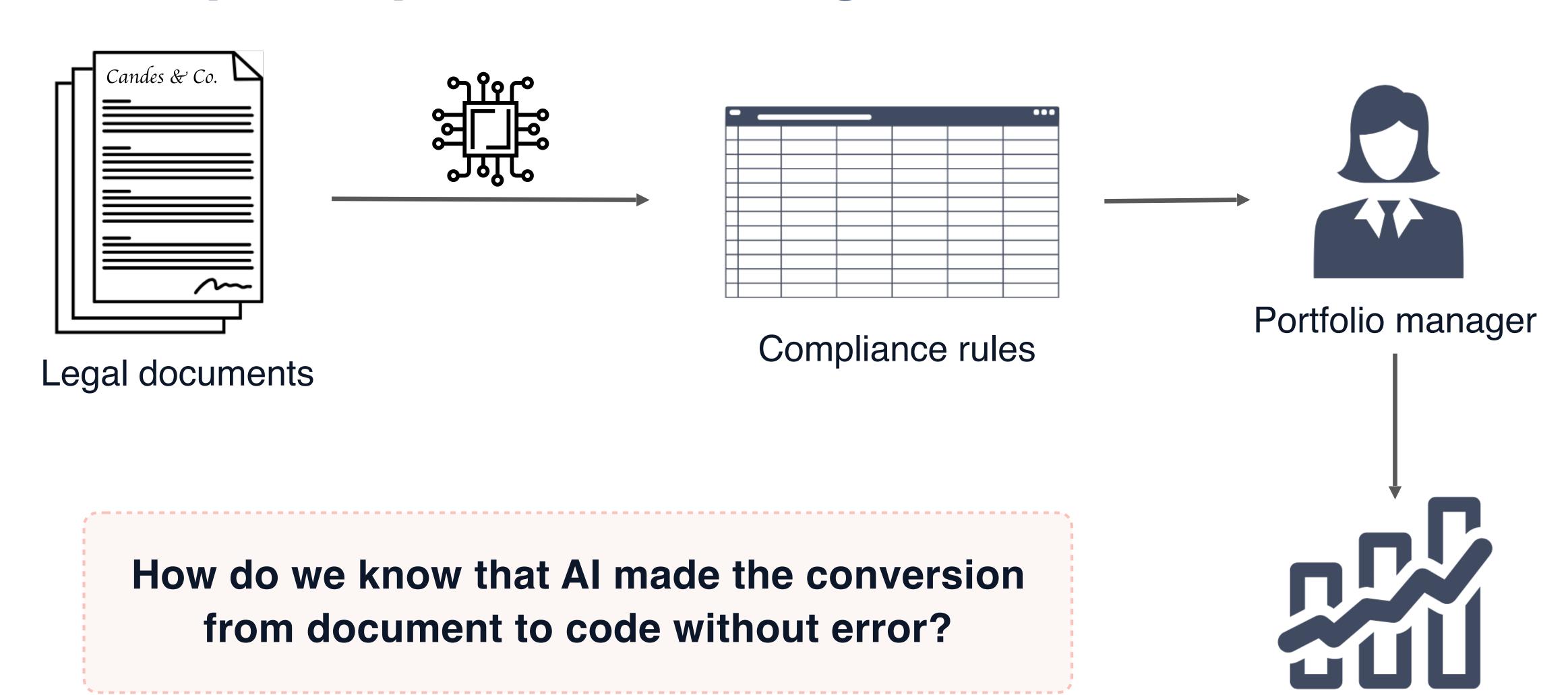
Al-powered inference

Al-powered drug discovery

Factual accuracy calibration for LLMs

synthetic pretraining datamodels s1

Example of production diagram in 2025



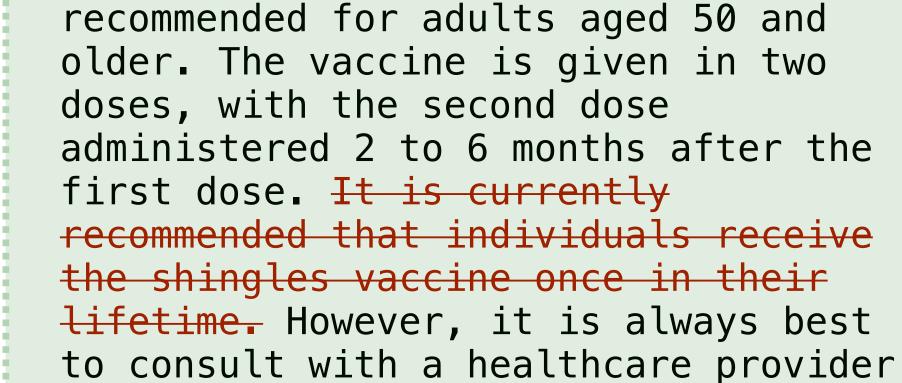
Active investment strategies

Goal: hallucination removal with guarantees

How often is a shingles vaccine required?

The shingles vaccine is typically recommended for adults aged 50 and older. The vaccine is given in two doses, with the second dose administered 2 to 6 months after the first dose. It is currently recommended that individuals receive the shingles vaccine once in their lifetime. However, it is always best to consult with a healthcare provider for personalized recommendations.





for personalized recommendations.

The shingles vaccine is typically

Unreliable

Guaranteed factuality

(with high probability)

Split conformal factuality

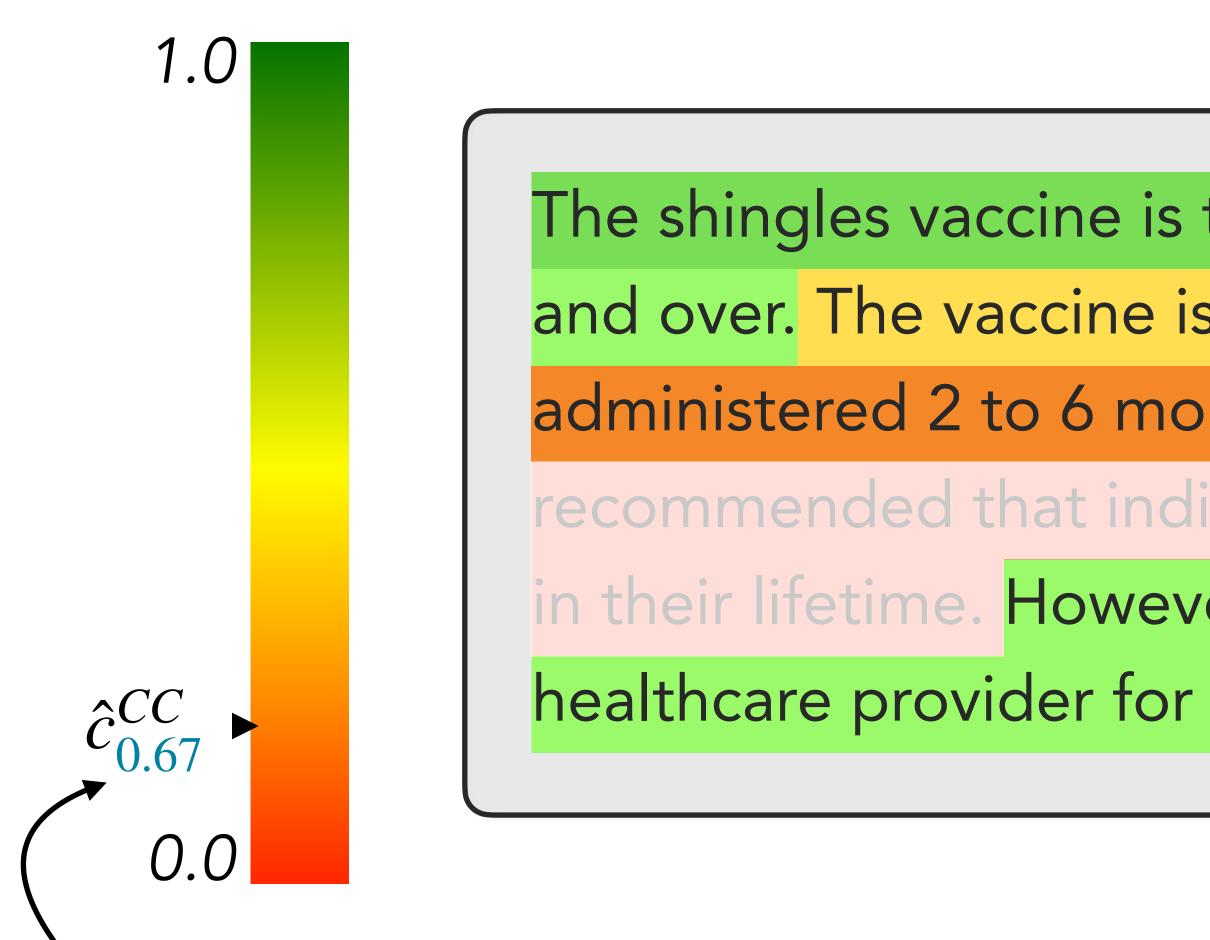
(Mohri & Hashimoto, 2024)

1.0

The shingles vaccine is typically recommended for adults aged 50 and over. The vaccine is given in two doses, with the second dose administered 2 to 6 months after the first dose. It is currently recommended that individuals receive the shingles vaccine once in their lifetime. However, it is always best to consult with a healthcare provider for personalized recommendations.

Our approach

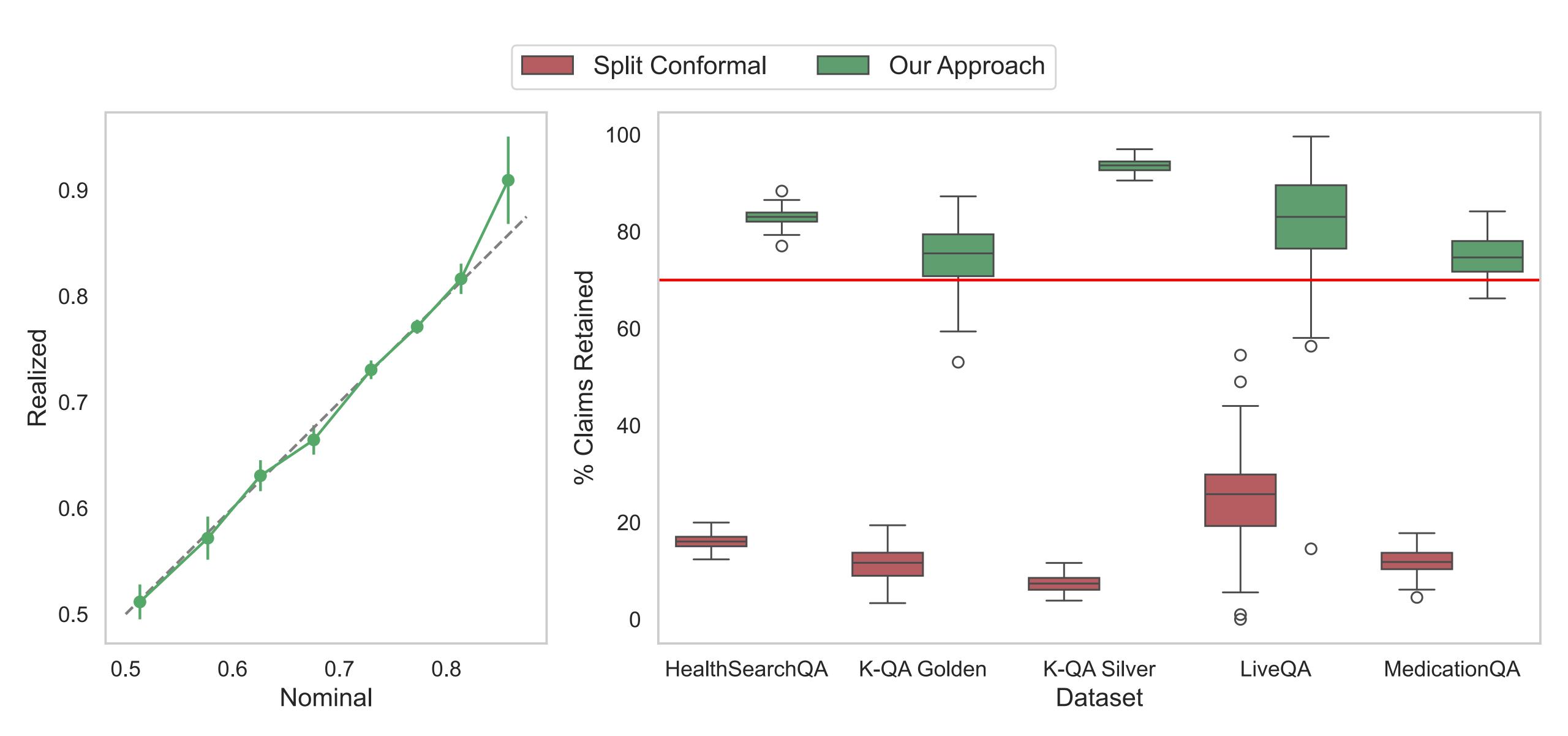
(Gibbs, Cherian, Candes, 2025)



The shingles vaccine is typically recommended for adults aged 50 and over. The vaccine is given in two doses, with the second dose administered 2 to 6 months after the first dose. It is currently recommended that individuals receive the shingles vaccine once in their lifetime. However, it is always best to consult with a healthcare provider for personalized recommendations.

Prompt-dependent guarantee is calibrated and \mathcal{F} -conditionally valid

Calibrated validity

















synthetic pretraining datamodels s1

Al-powered inference

Al-powered drug discovery

Factual accuracy calibration for LLMs

Al and statistics offer a lot to each other















Open-source LLMs

Mathematical reasoning

Efficient training

Synthetizing data

Deforestation
Media bias
Global warming
Protein structures

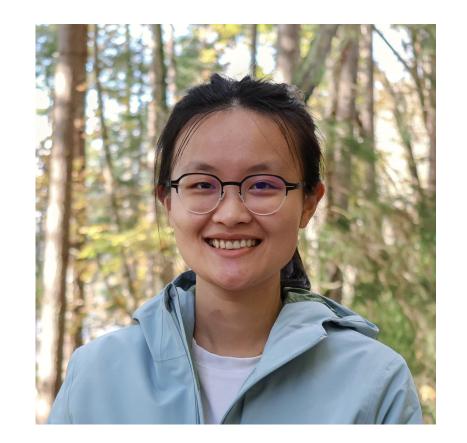
Drug discovery

Clinical trials

Filtering hallucinations
Factual calibration of LLMs
Automating legal compliance
Managing misinformation

Thinking statistically about AI inputs and outputs yields more powerful, safer AI

Close Collaborators



Ying Jin



Isaac Gibbs



John Cherian



Zitong Yang



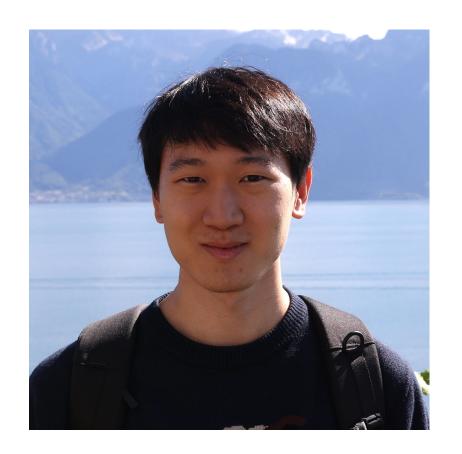
Tijana Zrnic



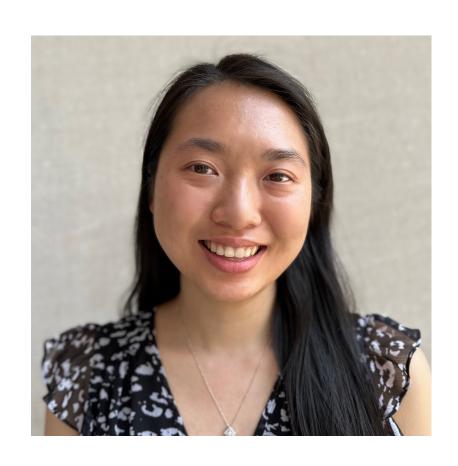
Asher Spector



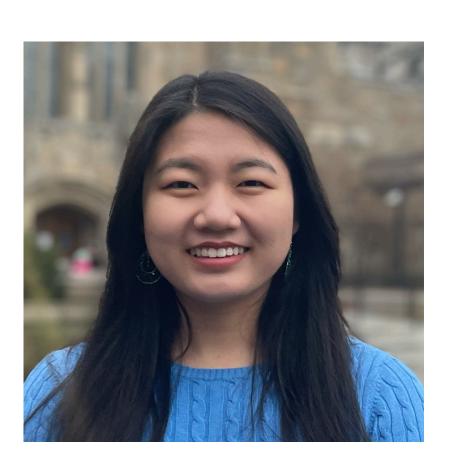
Yash Nair



Joon Lee



Ginnie Ma



Sarah Zhao

Other Collaborators



Anastasios Angelopoulos



Stephen Bates



Clara Fannjiang



Michael Jordan



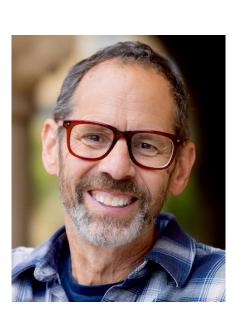
John Duchi



Kristina Gligorić



Cinoo Lee



Dan Jurafsky



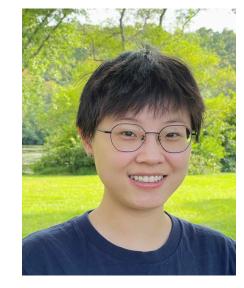
Neil Band



Tatsunori Hashimoto



Percy Liang



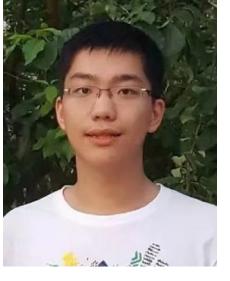
Shuangping Li



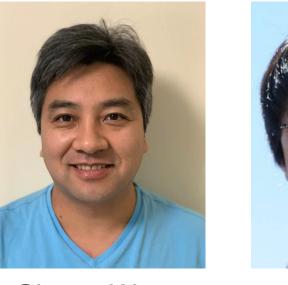
Christopher Mohri



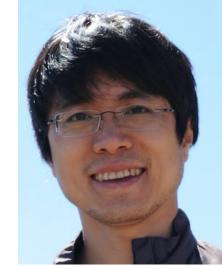
Aonan Zhang



Hong Liu



Chong Wang



Ruoming Pang



Niklas

Muennighoff

Andrew Ilyas



Sung Min (Sam) Park



Logan Engstrom



Kristian Georgiev



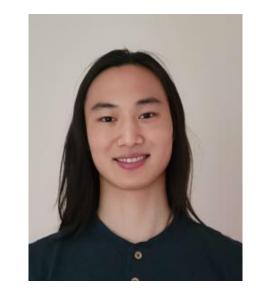
Guillaume Leclerc



Aleksander Madry



Axel Feldmann

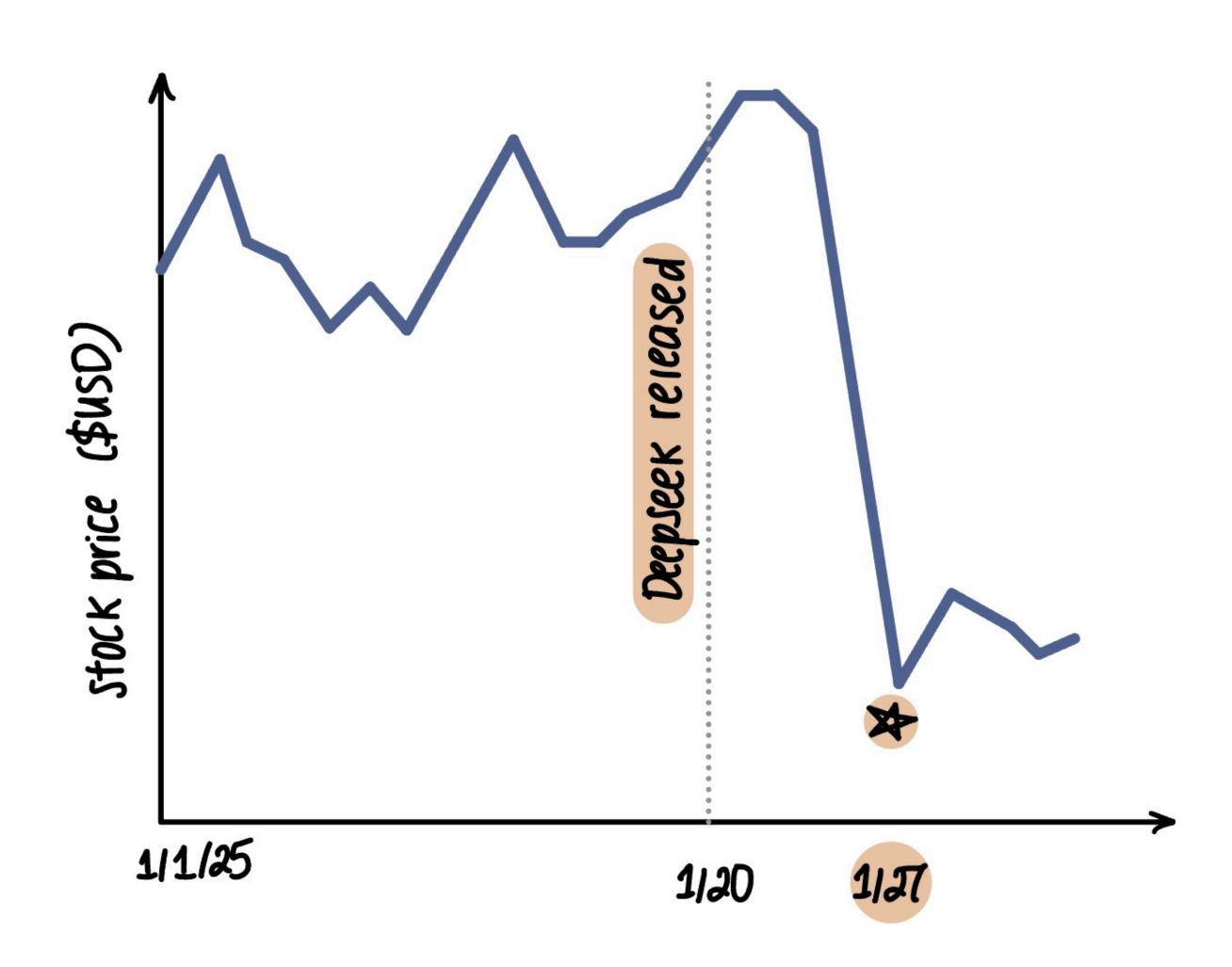


Benjamin Chen

Last week of January



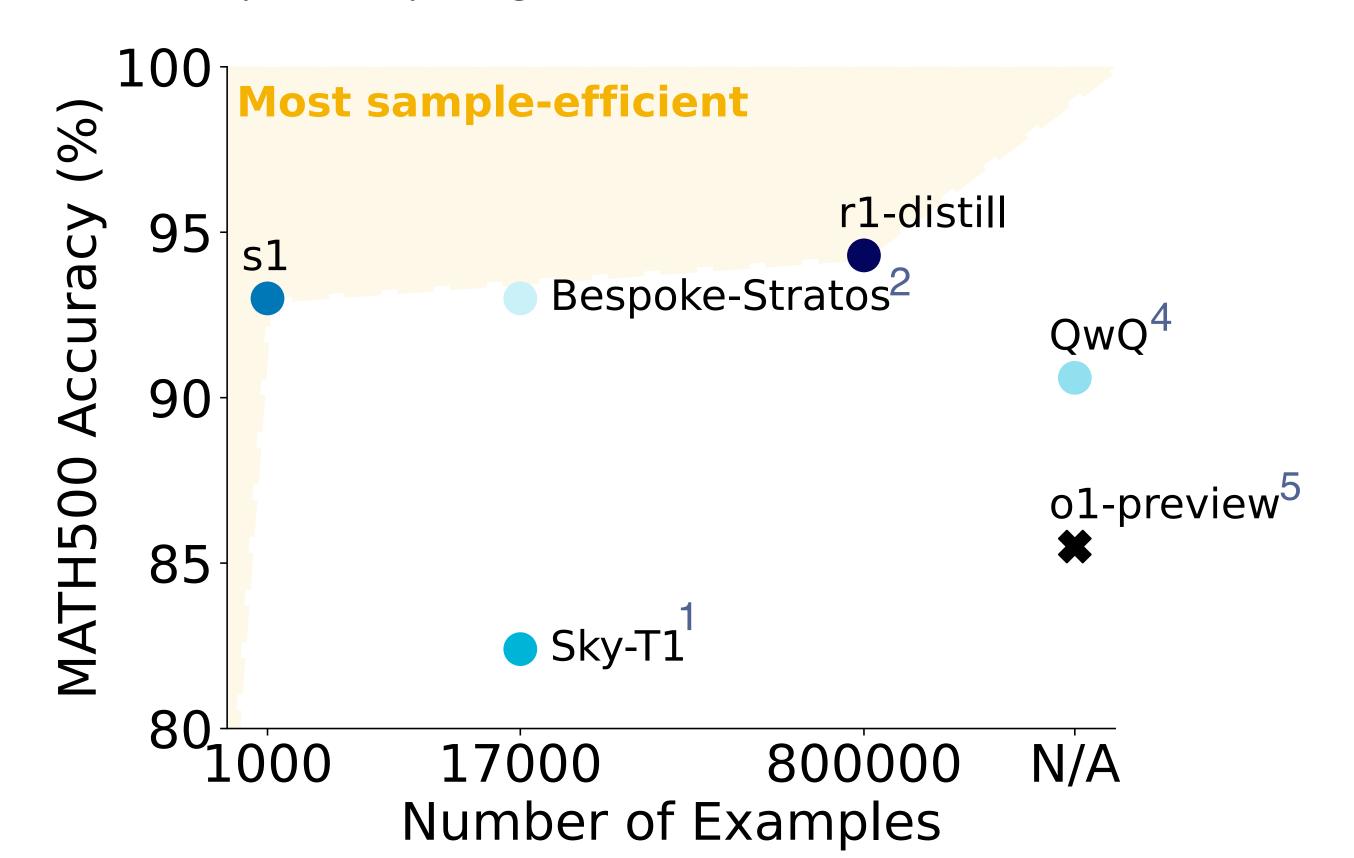




Stanford s1 (first week of February)

s1: Simple test-time scaling

Niklas Muennighoff * 1 3 4 Zitong Yang * 1 Weijia Shi * 2 Xiang Lisa Li * 1 Li Fei-Fei ¹ Hannaneh Hajishirzi ^{2 3} Luke Zettlemoyer ² Percy Liang ¹ Emmanuel Candès ¹ Tatsunori Hashimoto ¹



- 1. Team (2025)
- 2. Labs (2025)
- 3. DeepSeek-Al et al. (2025)
- 4. Qwen et al. (2024)
- 5. OpenAI (2024)

'Science' of LLMs

- Demonstrate test-time scaling
- High performance on small training data sets (S1K)
- Open source/weights/data/ideas/everything

