

# Combining Kernel and Model-Based Reinforcement Learning for HIV Therapy Selection

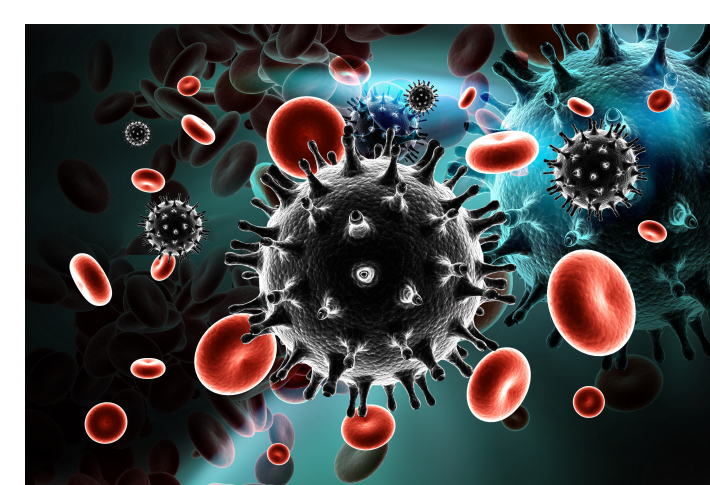


Sonali Parbhoo<sup>1</sup>, Jasmina Bogojeska<sup>2</sup>, Volker Roth<sup>1</sup> and Finale Doshi-Velez<sup>3</sup>

Department of Mathematics and Computer Science, University of Basel, Switzerland<sup>1</sup>  
 IBM Research, Zurich, Switzerland<sup>2</sup>  
 Harvard University, John A. Paulson School of Engineering and Applied Sciences, Cambridge, MA.<sup>3</sup>

## Motivation

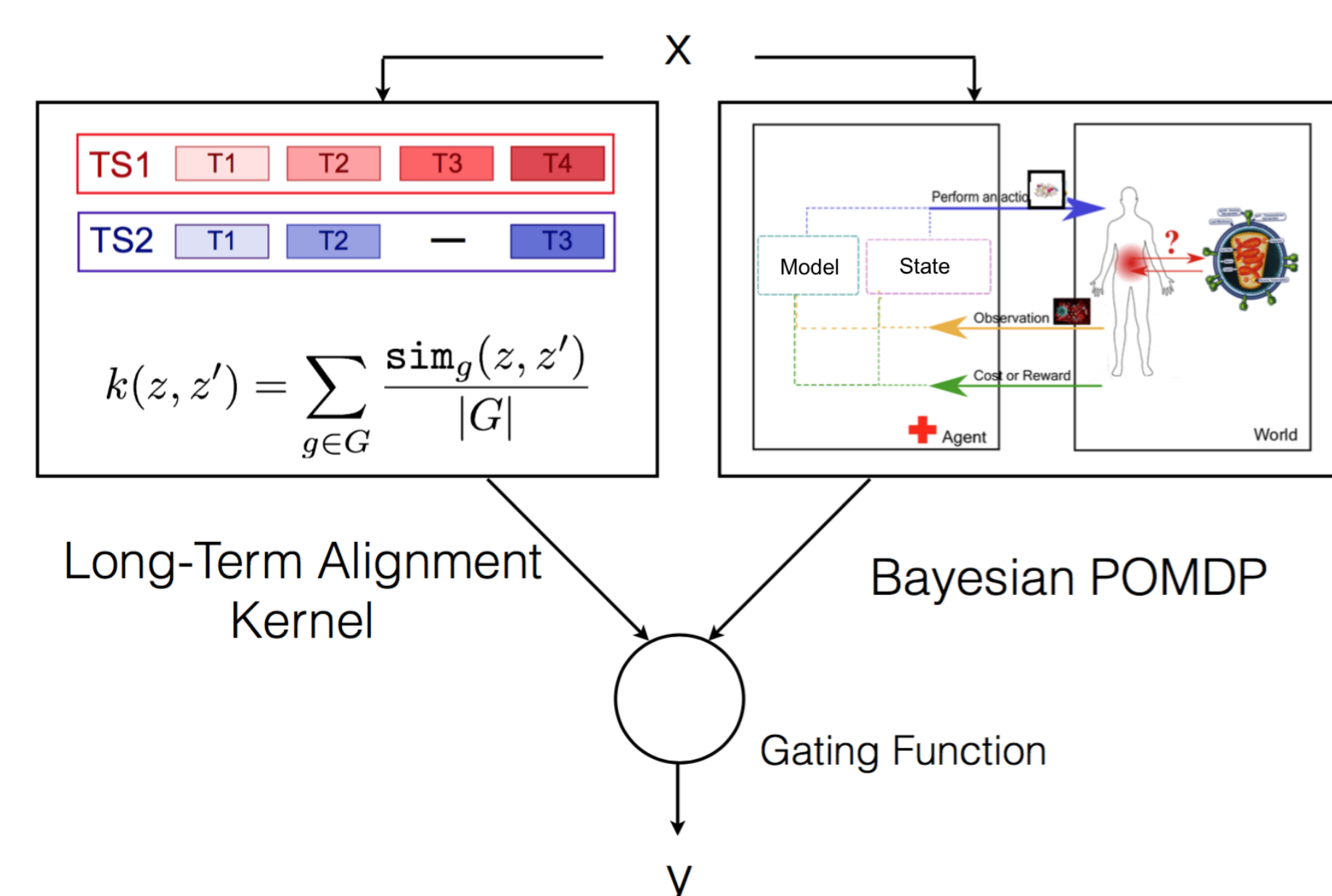
Human Immunodeficiency Virus (HIV) infects immune cells and causes AIDS. The virus is highly mutagenic, so choosing a therapy is difficult.



Combination therapy is prescribed to overcome drug resistance. Existing therapy selection approaches based on regression are problematic:

- Do not account for **long-term** effects of therapies.
- Do not account for **patient heterogeneity**.

Trade-off between model generality vs specificity.



We propose a **Mixture-of-Experts (MoE)** model to overcome this.

**Goal:** Automatic optimal therapy selection

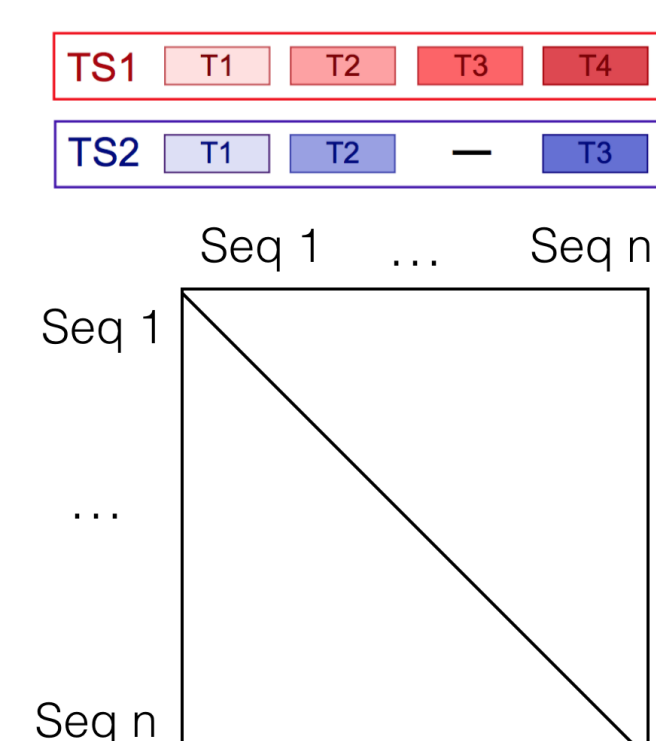
## The History Alignment Kernel

Patients with similar treatment histories respond similarly.

- Alignment of therapies [1]:

$$k(z, z') = \sum_{g \in G} \frac{\text{sim}_g(z, z')}{|G|}; \quad \text{sim}_g(z, z') = \frac{u_{zg}^\top u_{z'g}}{\max(\|u_{zg}\|^2, \|u_{z'g}\|^2)}$$

- Alignment of therapy sequences with Needleman Wunsch.



More common drugs/mutations  $\implies$  more similar treatments.

Better alignment  $\implies$  more similar histories.

**Short-Term Therapy Success:**

Viral RNA < 400 copies/mL

## Long-Term Therapy Success

- Immediate reward criterion:

$$r_t = \begin{cases} -0.7 \log V_t + 0.6 \log T_t - 0.2|M|, & \text{if } V_t \text{ is above detection limits} \\ 5 + 0.6 \log T_t - 0.2|M|, & \text{otherwise} \end{cases}$$

Long-term success sums criterion over patient's future history.

## Bayesian POMDP

Models **sequential decision-making** explicitly.

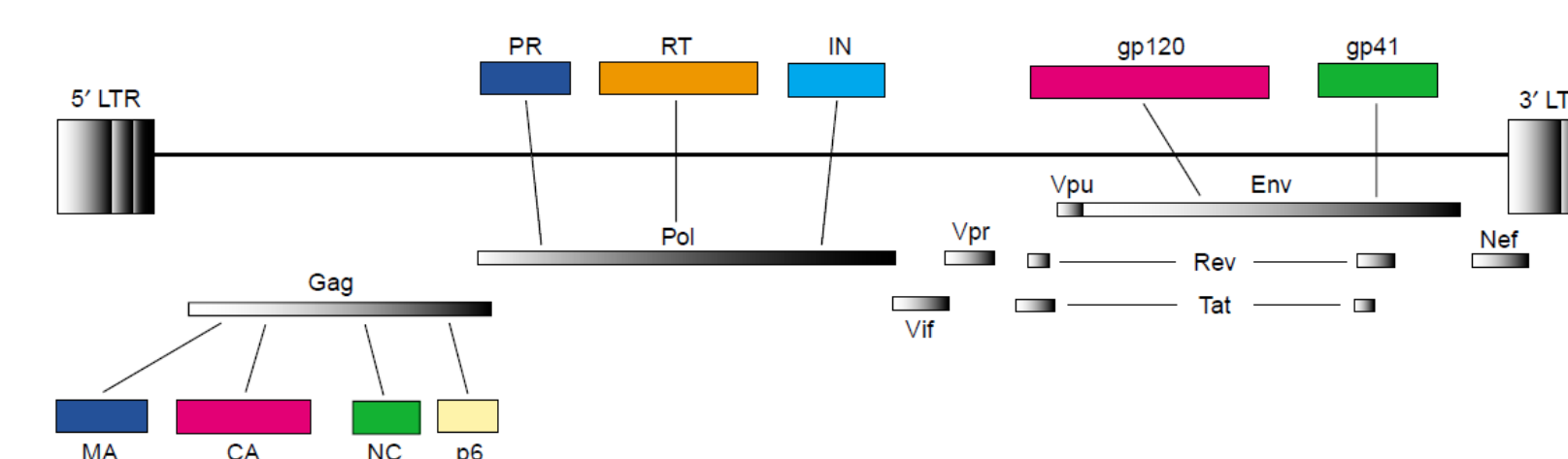
- 6 month time increments
- 7 hidden physiological states; 70 mutations and binned viral RNA values
- Transition and observation parameters drawn from Dirichlet
- Sample  $m$  models and update beliefs accordingly.

How do we know what action to choose?

- Build forward search tree: Search therapies to optimise outcomes over 5 years.
- Evaluate policies with off-policy evaluation.

## Data

- 32 960 patients' Therapy Change Episode (TCE) and clinical data
- 312 most frequently occurring drug combinations
- Genotype resistance data from *Pol* and *Env* regions [2].



The viral genome.

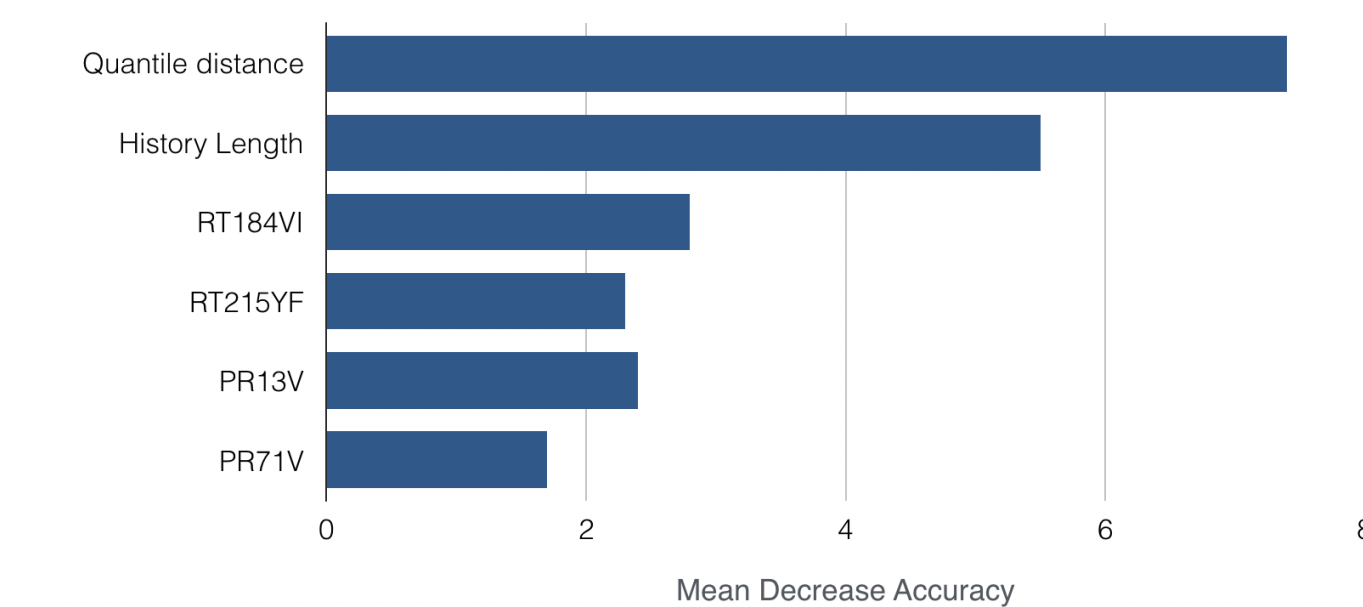
## Experiments

	Doubly Robust	IS	Weighted IS
Short-term Alignment	2.17 ± 1.47	2.14 ± 1.22	2.15 ± 1.16
Long-term Alignment	9.48 ± 1.90	5.42 ± 1.93	6.74 ± 1.89
POMDP	6.34 ± 2.15	4.36 ± 2.38	6.76 ± 2.24
<b>MoE</b>	<b>11.47 ± 1.38</b>	<b>12.25 ± 1.41</b>	<b>11.23 ± 1.40</b>
Truncated MoE	4.61 ± 2.35	4.73 ± 2.49	4.71 ± 2.18

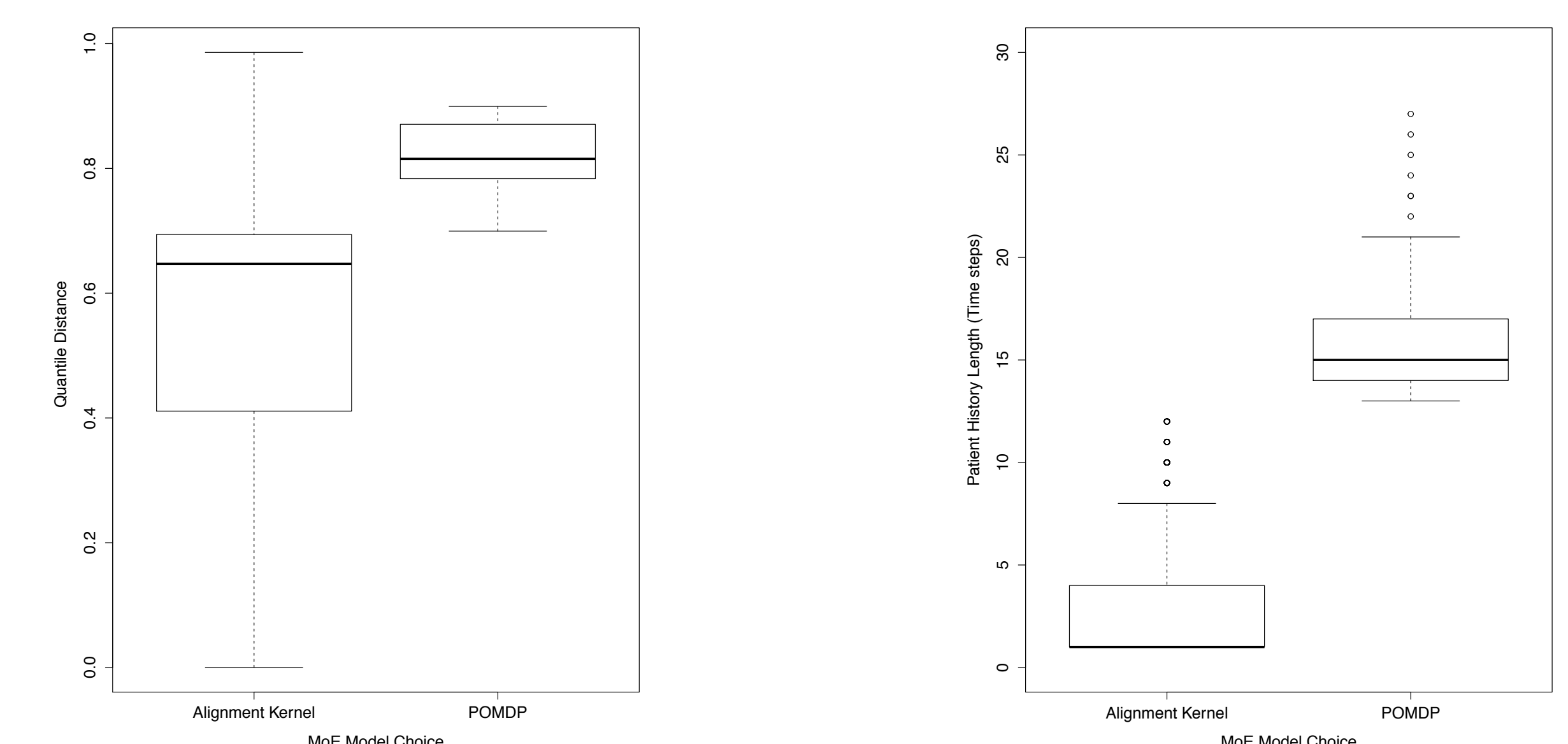
Off-policy evaluation for therapy selection models.

- Mixture-of-Experts selects therapy policy over 5 year horizon.
- Mixture-of-Experts chooses the POMDP 26% vs. the Long-Term Alignment 74%.
- Off-policy evaluation corrects distributional mismatch between data policy and learned policy.

When does the Mixture-of-Experts pick each model?



Feature importance for Mixture-of-Experts.



Mixture-of-Experts model choice over a) distances to closest neighbour b) history lengths.

## Conclusions

- Mixture-of-experts addresses trade-off between model **generality** and **specificity**.
- Different policies** produced when optimising over long-term vs. short-term.
- Mixture-of-Experts yields **best** policies.
- Expert chosen based on **clustering characteristics**.

## References

- Jasmina Bogojeska, Daniel Stöckel, Maurizio Zazzi, Rolf Kaiser, Francesca Incardona, Michal Rosen-Zvi, and Thomas Lengauer. History-alignment models for bias-aware prediction of virological response to hiv combination therapy. In *AISTATS*, pages 118–126, 2012.
- Annemarie M Wensing, Vincent Calvez, Huldrych F Günthard, Victoria A Johnson, Roger Paredes, Deenan Pillay, Robert W Shafer, and Douglas D Richman. Special contribution 2014 update of the drug resistance mutations in hiv-1. *Topics in antiviral medicine*, page 642, 2014.

**Acknowledgement:** We thank the Swiss National Science Foundation for financial support and EuResist Integrated Database for providing the data used in this research.