



Test-Time View Selection for Multi-Modal Decision Making

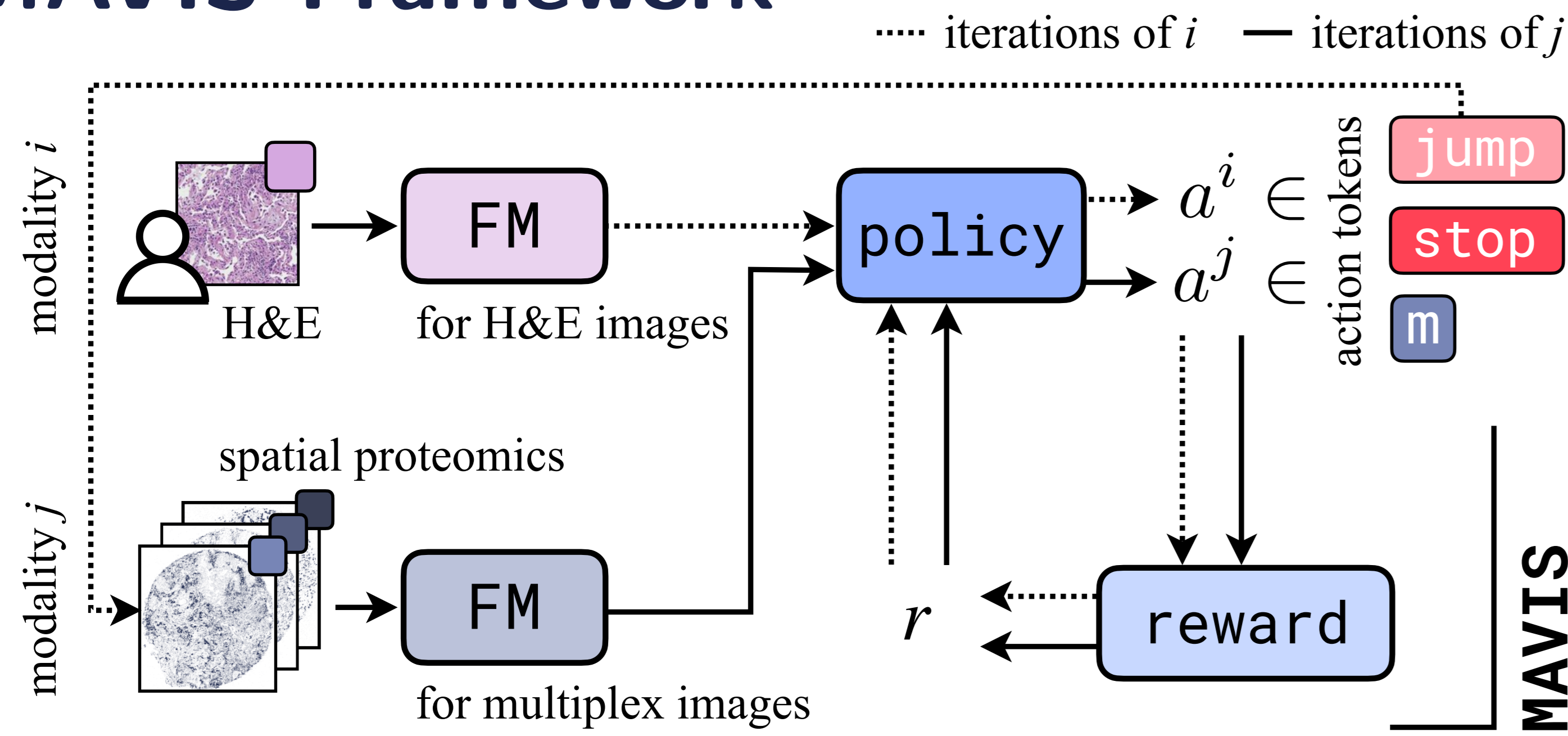
Eeshaan Jain, Johann Wenckstern, Benedikt von Querfurth, Charlotte Bunne



Contributions

- ... unified framework for modality and feature selection
- ... view selection *at inference time* without ground-truth preferences
- ... learning to guide **test-time selection** through **foundation models**

MAVIS Framework



Aim: learn the best selection rule σ^ℓ for any patient starting from any view ℓ

Challenges: inferring the **optimal** selection rule is $\Omega(2^{|\mathcal{V}|})$
optimal selection rules different for every patient

Metrics: $Acc_k = \mathbb{E}_{p \sim \mathcal{P}} \left[\mathbb{E}_{y \sim \mathcal{H}_k^\ell} [\mathbb{I}\{y = y_p\}] \right]$ foundation model

$$Unc_k = 1 - TCP_k \left((h \circ F) (\{V_{\sigma^\ell(p)(1)}, \dots, V_{\sigma^\ell(p)(k)}\}) \right)$$

σ^ℓ is k -optimal: If \forall selection rules $\rho^\ell: Acc_k(\sigma^\ell) \geq Acc_k(\rho^\ell)$

σ^ℓ is optimal: If σ^ℓ is k -optimal for every $k \leq |\mathcal{V}|$

↳ If exists, is necessarily greedy!

$$\sigma^\ell(p)(k) = \arg \max_{i \in [|\mathcal{V}|] \setminus \cup_{j=1}^{k-1} \{\sigma^\ell(p)(j)\}} (h \circ F) (\{V_{\sigma^\ell(p)(1)}, \dots, V_{\sigma^\ell(p)(k-1)}, V_i\})(y_p)$$

Learning k -optimal selection rules for $k \ll |\mathcal{V}|$:

$$c_t(s_t, i) = \beta \left(\log \left[(h \circ F) (\{V_{\sigma_t^\ell(1)}, \dots, V_{\sigma_t^\ell(t)}, V_i\})(y) \right] \right) \leftarrow \text{instantaneous reward}$$

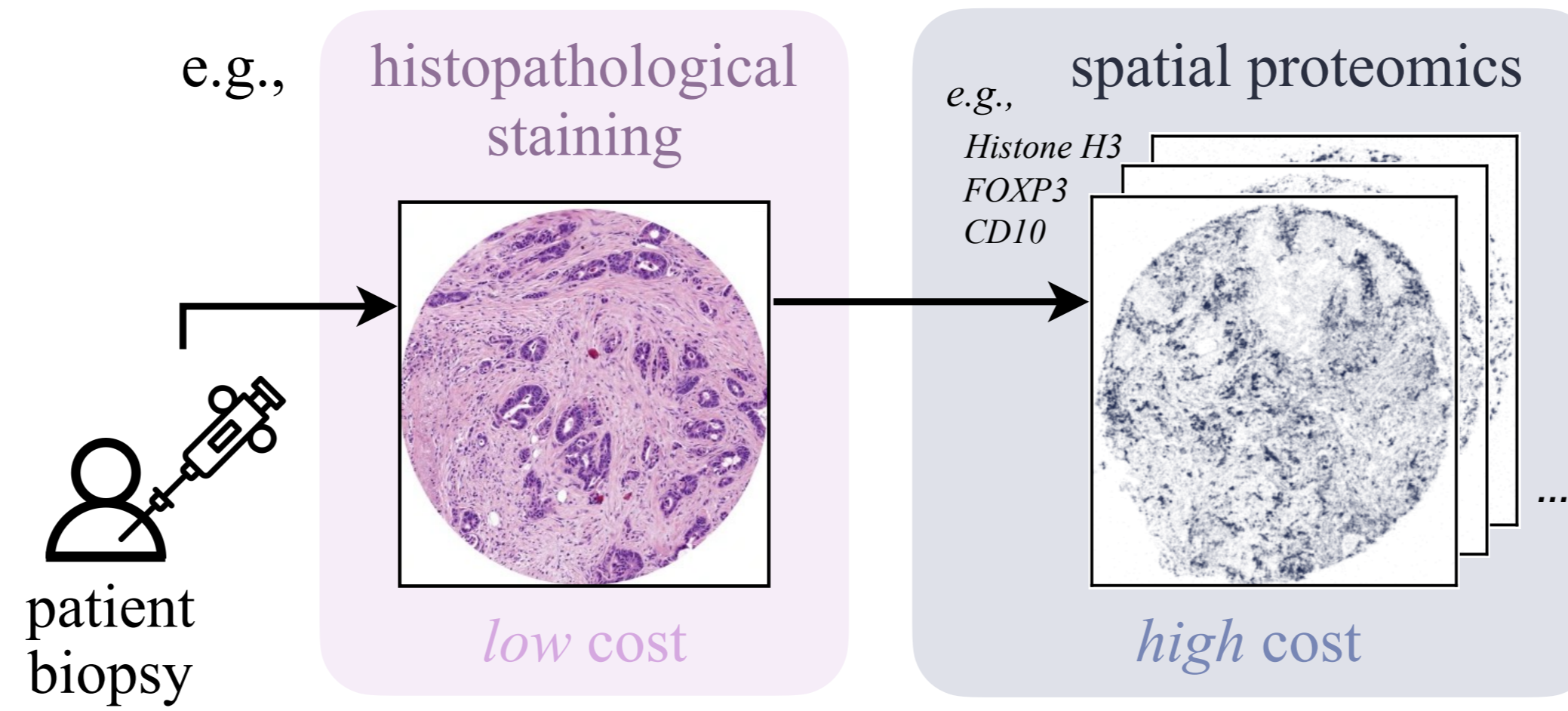
$$- \log \left[(h \circ F) (\{V_{\sigma_t^\ell(1)}, \dots, V_{\sigma_t^\ell(t)}\})(y) \right], \leftarrow \text{episodic reward}$$

$$c_T = \delta(2\mathbb{I}((h \circ F) (\{V_{\sigma_T(1)}, \dots, V_{\sigma_T(T)}\}) = y) - 1)$$

Acknowledgements: Yexiang Cheng (EPFL), Phil Cheng (HUG)

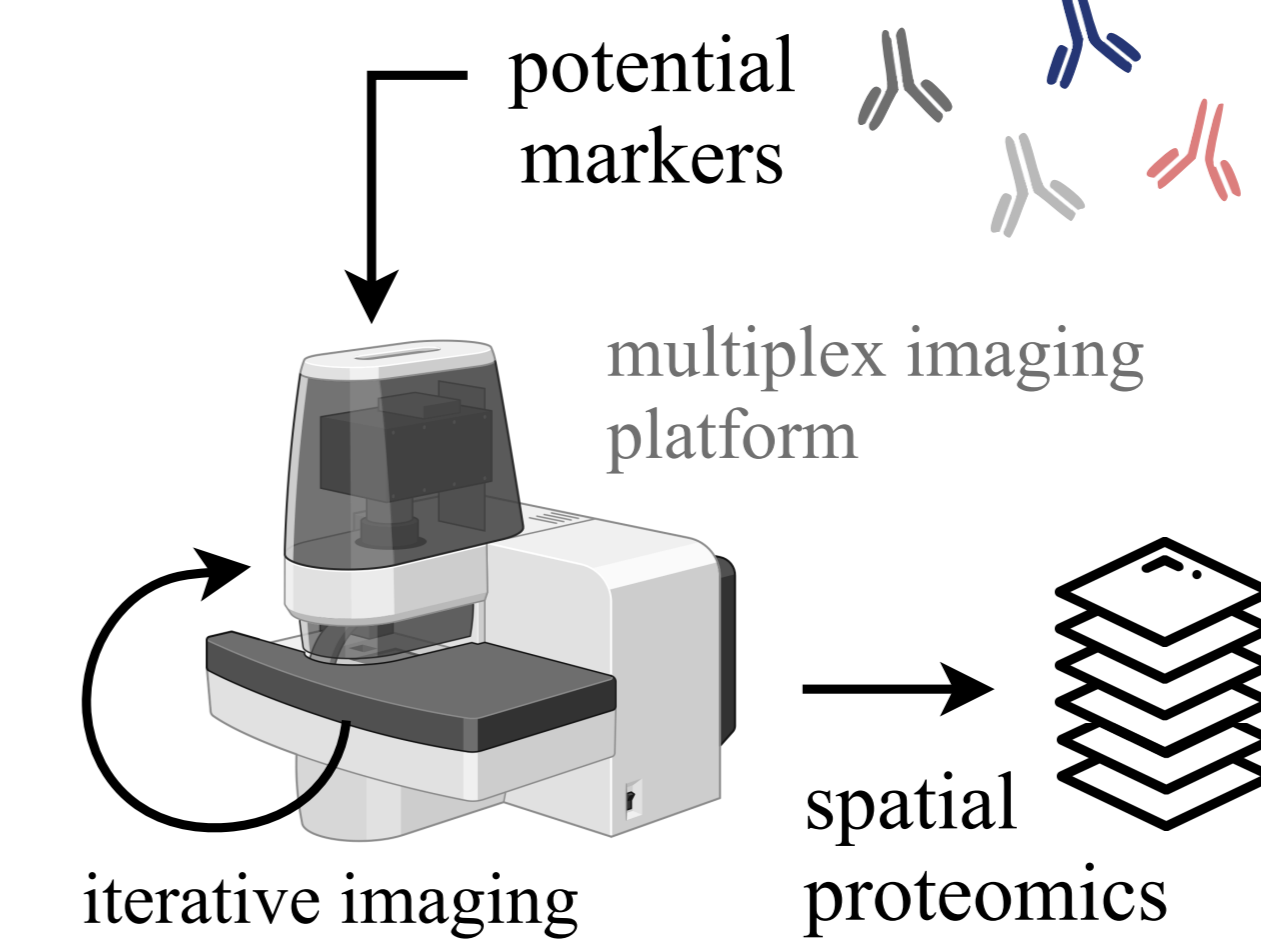
Modality Selection

Which diagnostic test to select?

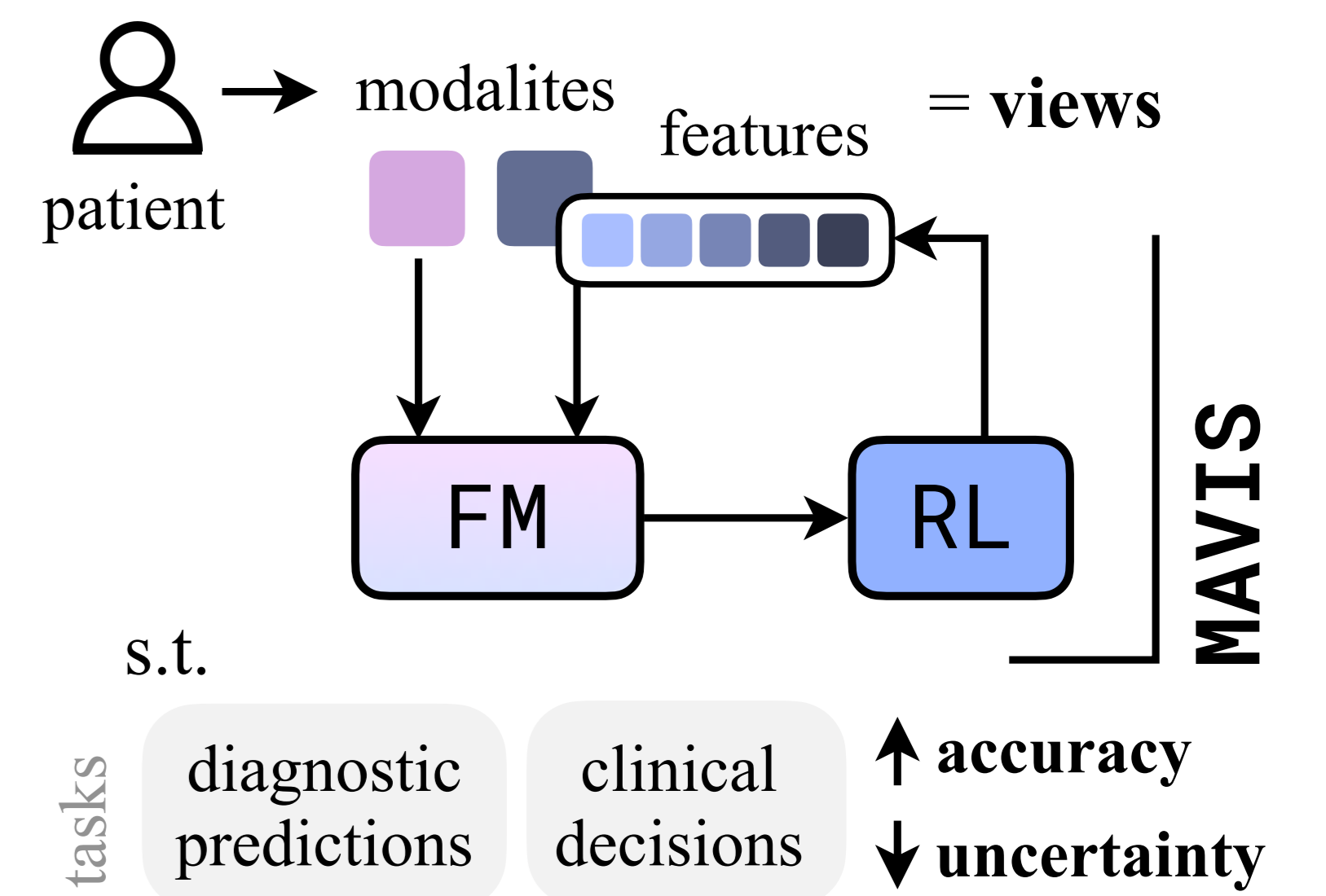


Experimental Design

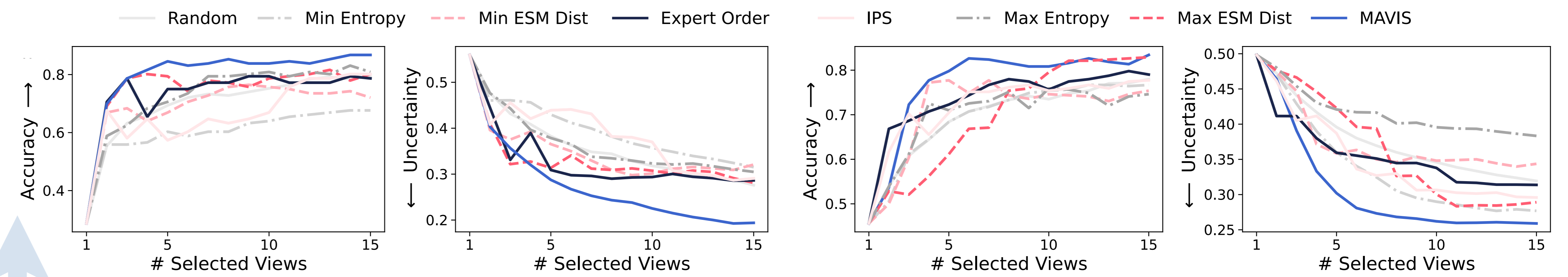
Which biomarkers to measure?



Multi-Modal Active View Selection

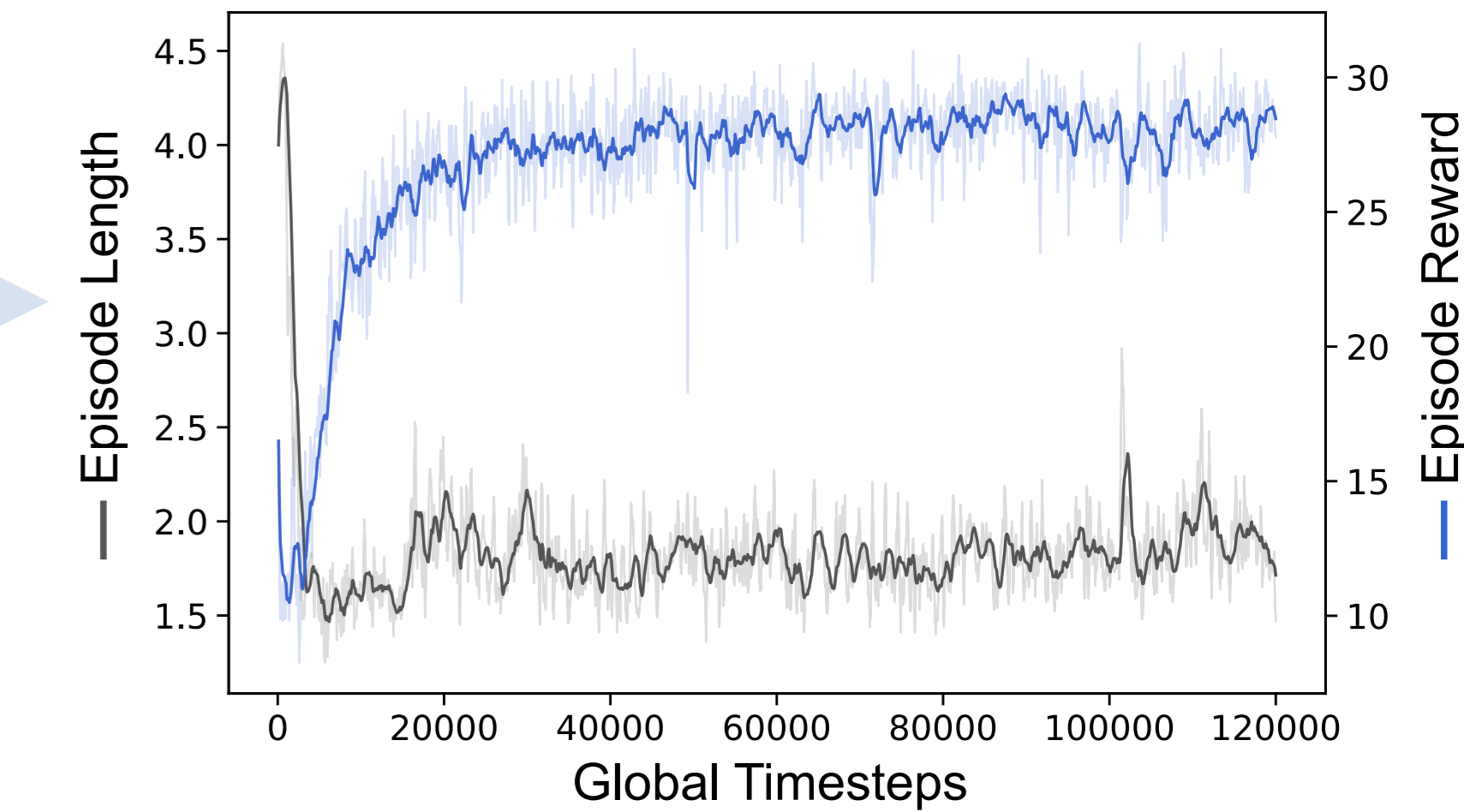


Results: MAVIS learns to select better selection rules



Over 15 marker acquisitions, **MAVIS** achieves faster uncertainty reduction and higher accuracy gains, outperforming global and expert orders.

MAVIS progressively learns better selection rules while learning when to switch to multiplex imaging over H&E.

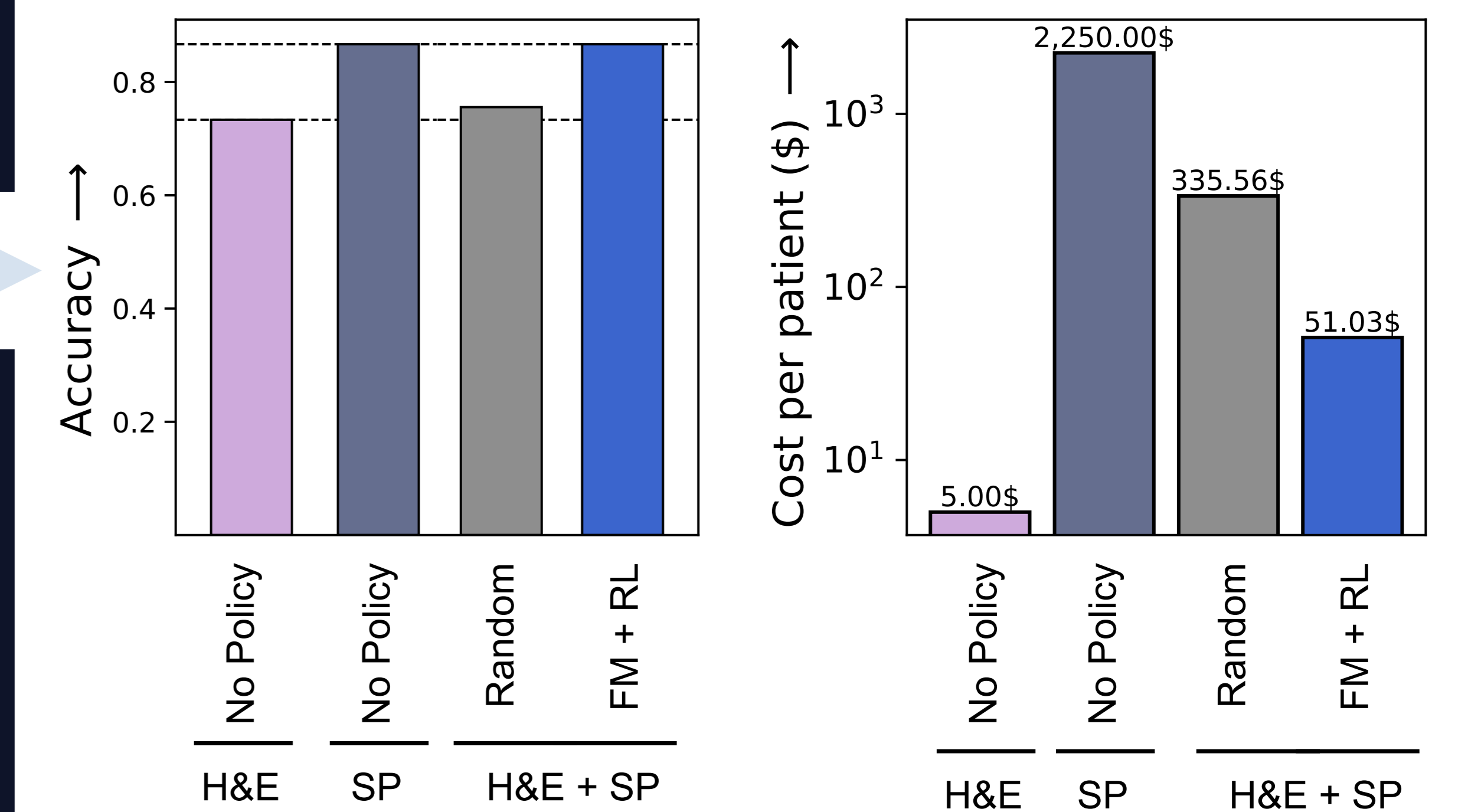


MAVIS achieves accuracy on par with SP, transitioning only those patients from H&E who genuinely benefit from additional multiplex imaging.

MAVIS reduces experimental costs by over 95% and learns to adaptively limit biomarker selection.



switch reward $\rightarrow c = \zeta \delta(2\mathbb{I}(\arg \max(h \circ F) (\{V_{\sigma_k^\ell(1)}, \dots, V_{\sigma_k^\ell(k)}\}) = y) - 1)$



To be presented at **MLGenX (Spotlight) & GemBio, ICLR 2025**

... we introduce **MAVIS** a unified framework for active multi-modal view selection in clinical diagnostics enabling dynamic adaptation to individual patient needs while leveraging FMs to guide test selection.

