

#### HEALNet: Multimodal Fusion for Heterogeneous Biomedical Data

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#### Why fusion?





Late Fusion



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## Open challenges for fusion models in biomedical domains

Information from different biological scales can provide crucial predictive context...



...but often require trading off modality-specific and shared information



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## HEALNet uses shared and modality-specific parameter spaces





#### HEALNet preserves modality-specific structural information

Table 1: Mean and standard deviation of the concordance Index on four survival risk categories. We trained HEALNet and all baselines on four TCGA tasks and report the performance on the test set across five folds. HEALNet outperforms all of its multimodal baselines and three out of four unimodal baselines in absolute c-Index performance.

Model	BLCA	BRCA	KIRP	UCEC
Uni-modal (Omics)	$0.606\pm0.019$	$0.580\pm0.027$	$0.780\pm0.035$	$0.550\pm0.026$
Uni-modal (WSI)	$0.556 \pm 0.039$	$0.550 \pm 0.037$	$0.533 \pm 0.099$	$0.630 \pm 0.028$
Porpoise (Late)	$0.620\pm0.048$	$0.630\pm0.040$	$0.790\pm0.041$	$0.590\pm0.034$
MCAT (Intermediate)	$0.620\pm0.040$	$0.589 \pm 0.073$	$0.789 \pm 0.087$	$0.589\pm0.062$
MOTCAT (Intermediate)	$0.631 \pm 0.051$	$0.607\pm0.069$	$0.810\pm0.062$	$0.587 \pm 0.083$
MultiModN (Sequential Fusion)	$0.551\pm0.060$	$0.582\pm0.084$	$0.753\pm0.152$	$0.610\pm0.121$
Perceiver (Early Fusion)	$0.565\pm0.042$	$0.566\pm0.068$	$0.783\pm0.135$	$0.623\pm0.107$
HEALNet (ours)	$0.668 \pm 0.036$	$\textbf{0.638} \pm \textbf{0.073}$	$\textbf{0.812} \pm \textbf{0.055}$	$\textbf{0.626} \pm \textbf{0.037}$

Learns cross-modal interactions

Effective handling of missing modalities

Easy inspection using modalityspecific attention weights





## Iterative attention setup allows to skip updates for missing data

Table 2: Analysis of the performance of HEALNet, trained on all modalities, in scenarios with missing modalities at inference, compared to unimodal baselines. Each test sample contains only one of either the Omic or WSI modality. The scenarios include test sets consisting of samples with only Omic modality, only WSI modality or a combination of both (at random). HEALNet achieves a higher c-Index across datasets, implying effective encoding of cross-modal information and handling different amounts of data with missing modalities.

Test	100% Omics		100% WSI		50%WSI + 50% Omics		WSI+Omic
	Uni-modal	HEALNet	Uni-modal	HEALNet	Uni-modal	HEALNet	HEALNet
BLCA	0.606	<u>0.618</u>	0.487	0.501	0.547	<u>0.612</u>	0.668
BRCA	0.556	<u>0.571</u>	0.529	0.539	0.543	<u>0.541</u>	0.638
KIRP	0.771	<u>0.773</u>	0.518	0.526	0.644	0.714	0.812
UCEC	0.509	0.529	0.558	0.584	0.533	<u>0.580</u>	0.626

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## Attention-based design allows some explainability



Figure 3: Illustration of model's inspection capabilities using HEALNet on a high-risk patient of the UCEC study. We use the mean modality-specific attention weights across layers to highlight high-risk regions and inspect high-attention omic features. Individual patches can be used for further clinical or computational post-hoc analysis such as nucleus segmentation. We observe that the high-risk regions exhibit a very high concentration and different arrangement of epithelial cells (red) which is commonly associated with the origin of various cancer types [Coradini et al., 2011].

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#### Thank you





Code

Paper



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