

# TIPIT

Towards an Integrative approach for Precision ImmunoTherapy

## LITO lab meeting

Nicolas Captier

21/12/2023

Fondation  
pour la recherche  
sur le cancer



PR[AI]RIE  
PaRis Artificial Intelligence Research Institute

institut  
Curie



Inserm  
La science pour la santé  
From science to health



institutCurie

# Outlines

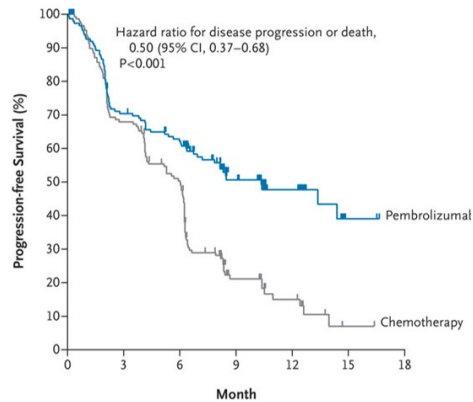
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- 1. Introduction:** build new biomarkers for immunotherapy outcome in Non-Small Cell Lung Cancer
- 2. Materials and Methods:** A retrospective multimodal cohort to develop multimodal predictors
- 3. Results:** New proofs of the benefits of multimodal machine learning to build accurate prognostic models
- 4. Discussion:** What can we do next ?

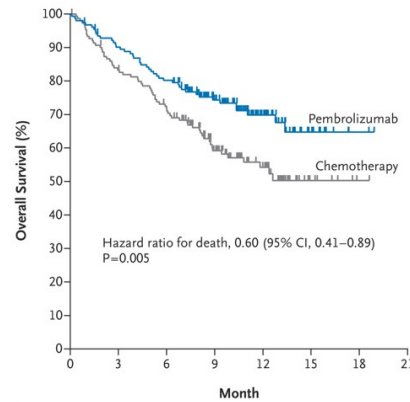
# Multimodality to predict immunotherapy outcome in lung cancer

- Immunotherapy is the standard-of-care for metastatic Non-Small Cell Lung Cancer (NSCLC)
- Highly variable responses + only 40% of patients are alive at 2 years
- Established univariate biomarkers are very few with limited power.

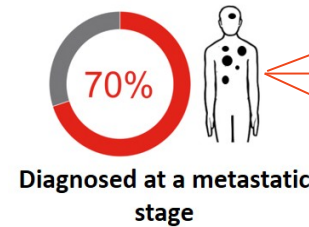
## Standard-of-care for advanced NSCLC in Europe



|               |     |     |    |    |    |   |   |
|---------------|-----|-----|----|----|----|---|---|
| No. at Risk   |     |     |    |    |    |   |   |
| Pembrolizumab | 154 | 104 | 89 | 44 | 22 | 3 | 1 |
| Chemotherapy  | 151 | 99  | 70 | 18 | 9  | 1 | 0 |

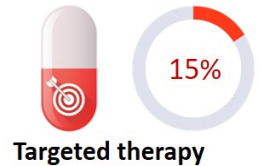


|               |     |     |     |    |    |    |   |
|---------------|-----|-----|-----|----|----|----|---|
| No. at Risk   |     |     |     |    |    |    |   |
| Pembrolizumab | 154 | 136 | 121 | 82 | 39 | 11 | 2 |
| Chemotherapy  | 151 | 123 | 106 | 64 | 34 | 7  | 1 |



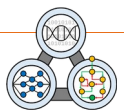
Expression of PD-L1 in  $\geq$  50% cancer cells

Expression of PD-L1 in < 50% cancer cells



Anti PD1\PD-L1

Anti PD1\PD-L1 + chemo



# Multimodality to predict immunotherapy outcome in lung cancer

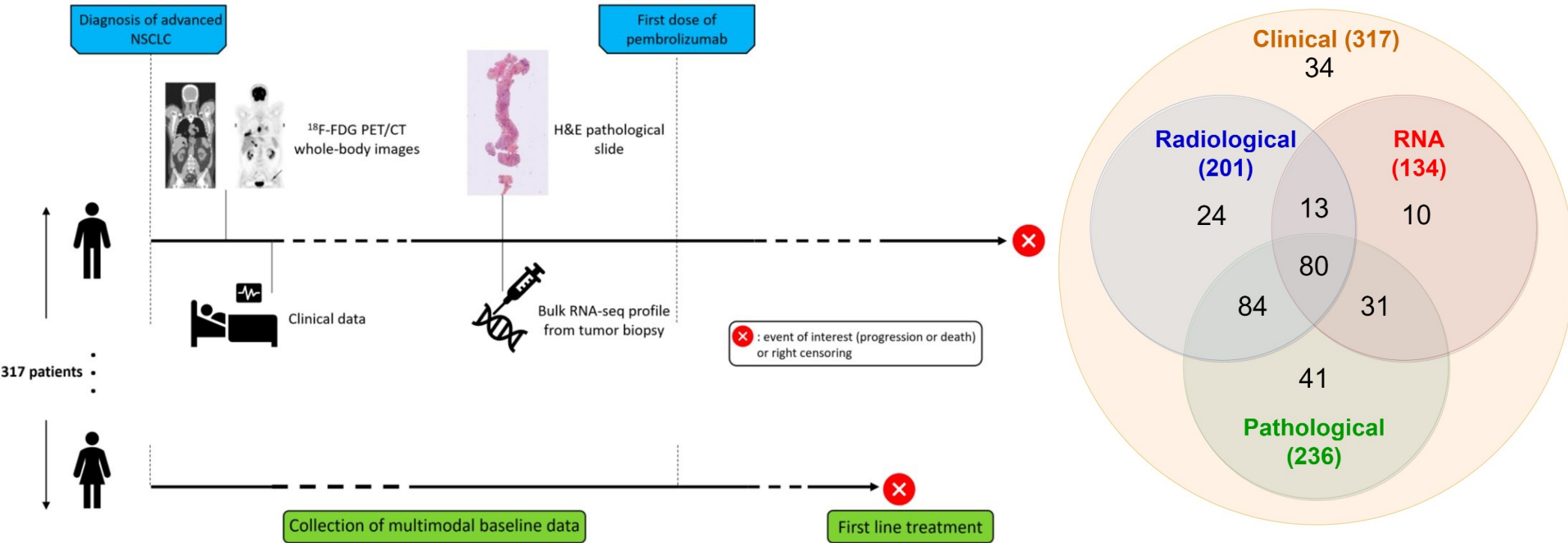
1. **Methodological question:** Can multimodal machine learning (& modelling) approaches build more accurate prognostic (potentially predictive) signatures ?
2. **Biological question:** Can we identify mechanisms associated with immunotherapy response with multimodal analyses ?

# Outlines

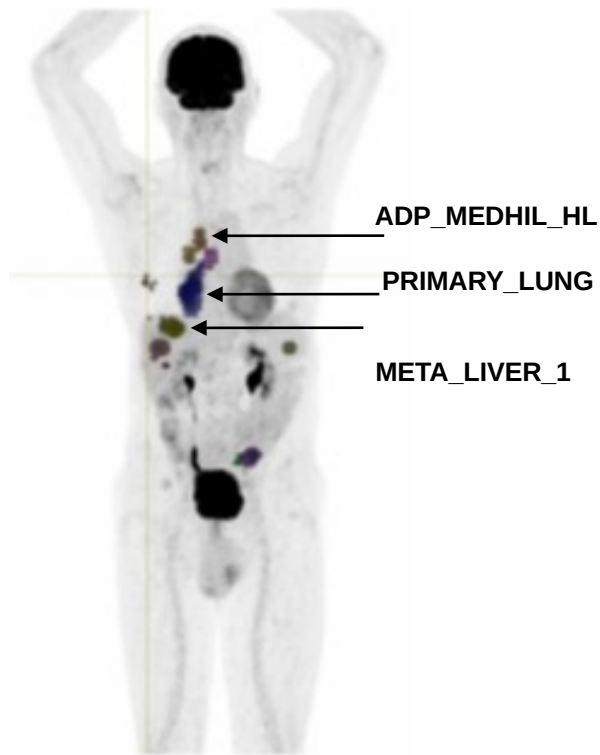
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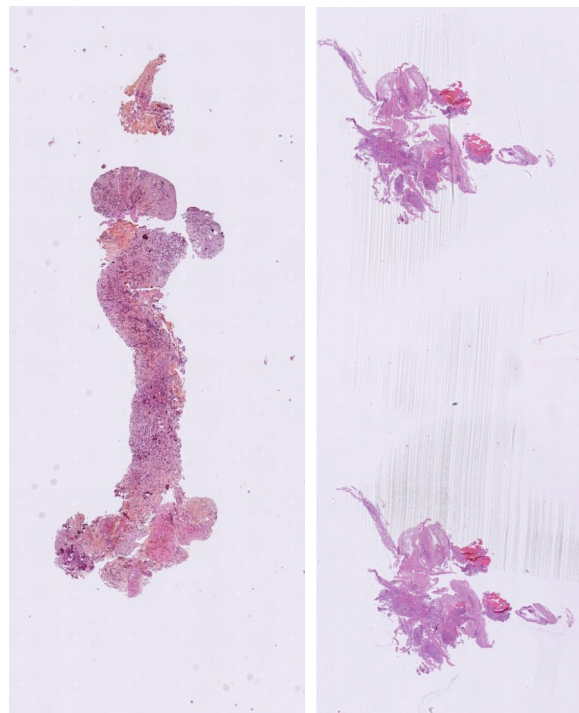
# A retrospective multimodal cohort - I



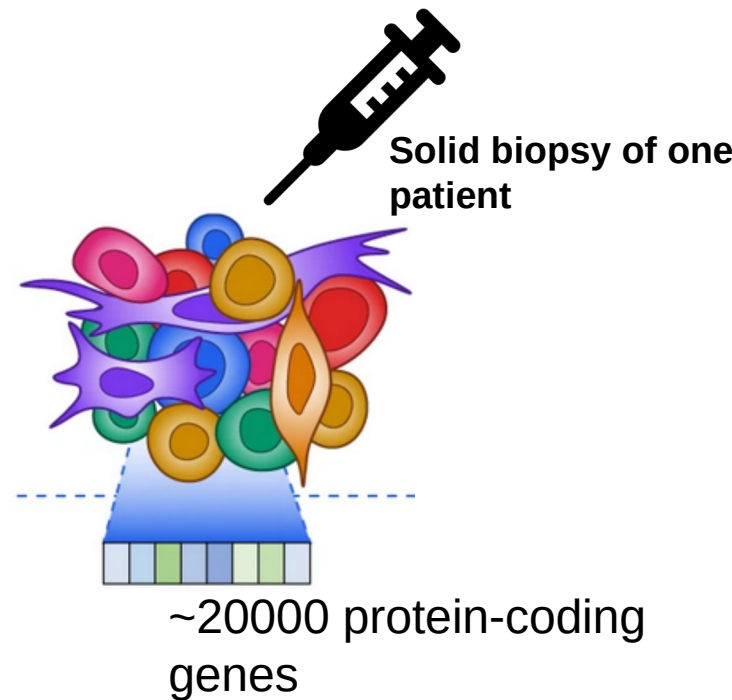
# A retrospective multimodal cohort - II



201 annotated/segmented  
18F-FDG PET/CT

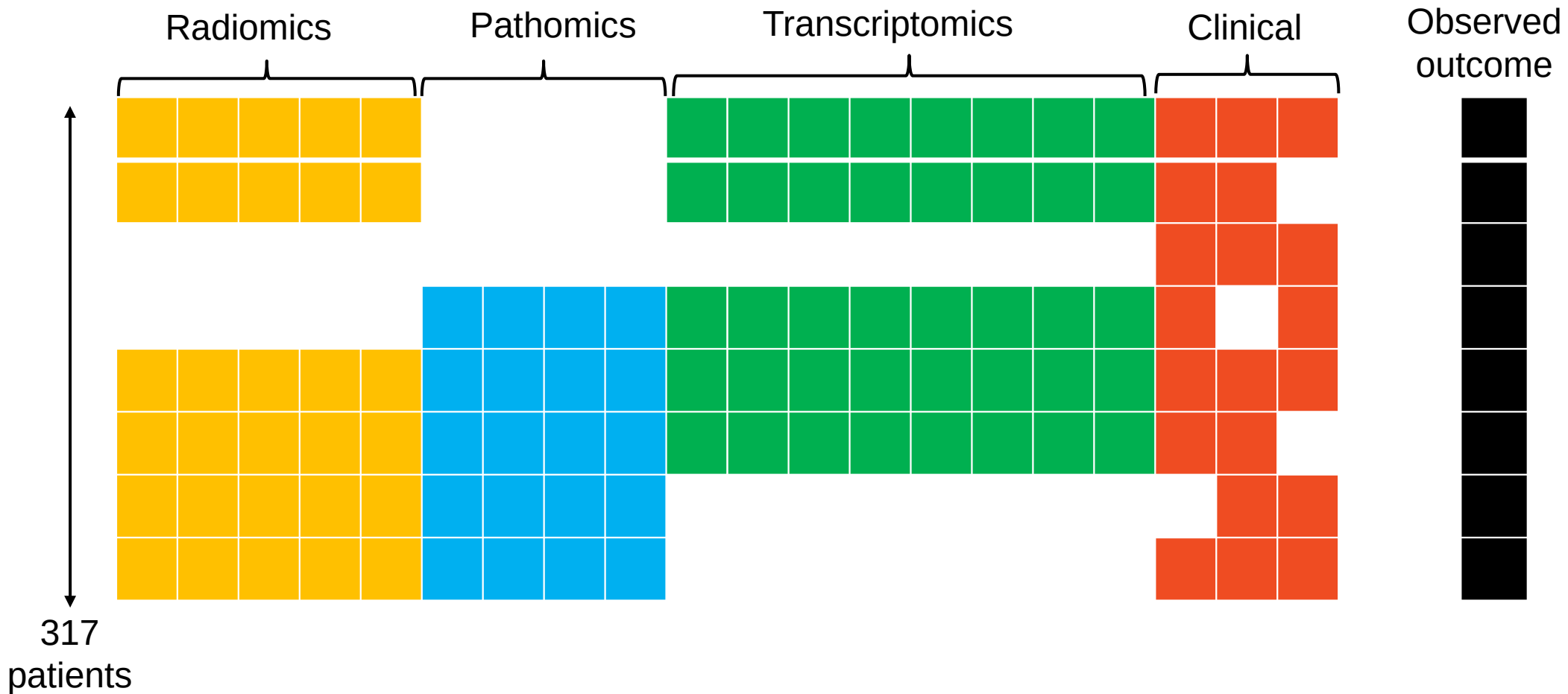


236 annotated/segmented  
digitized HE slides



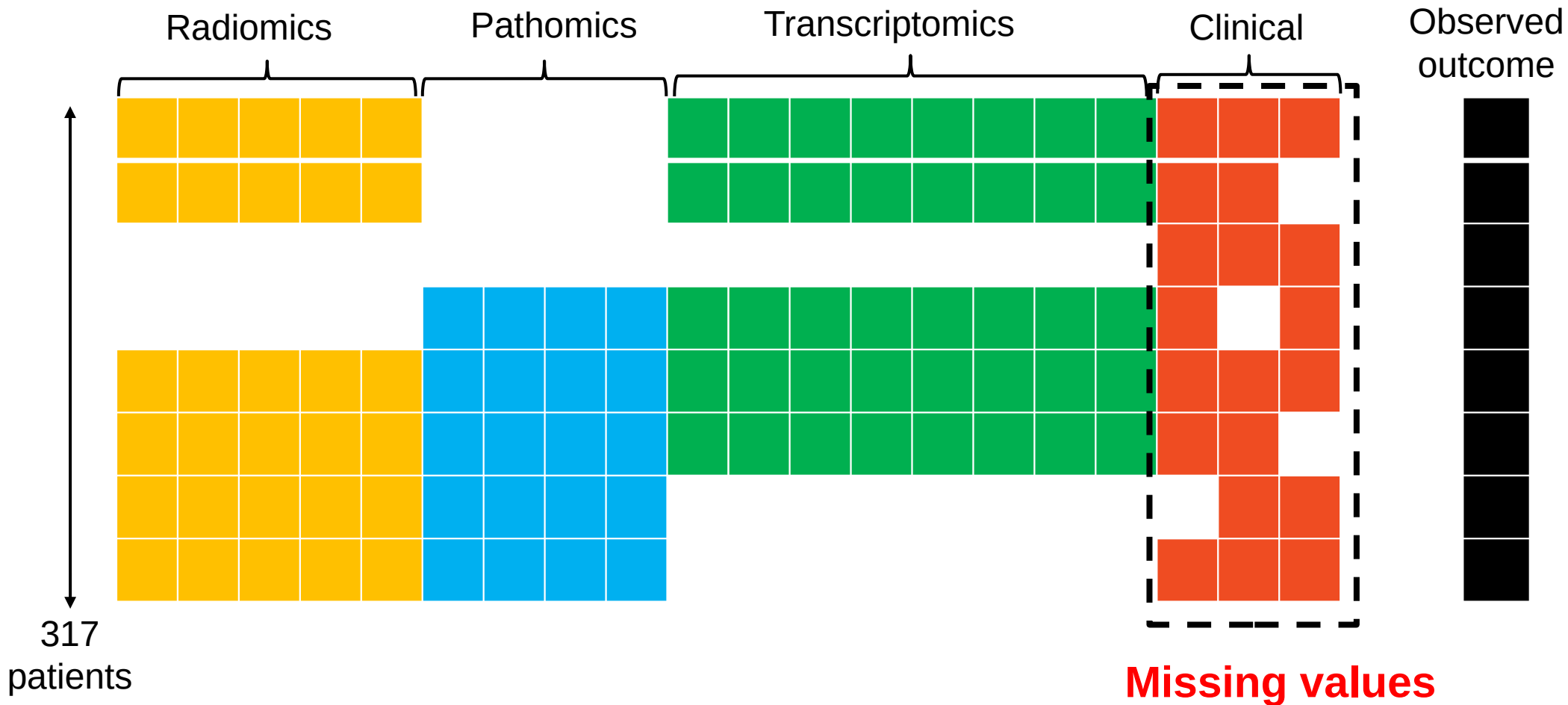
134 bulk RNA-seq profiles  
from solid biopsy

# A retrospective multimodal cohort - III

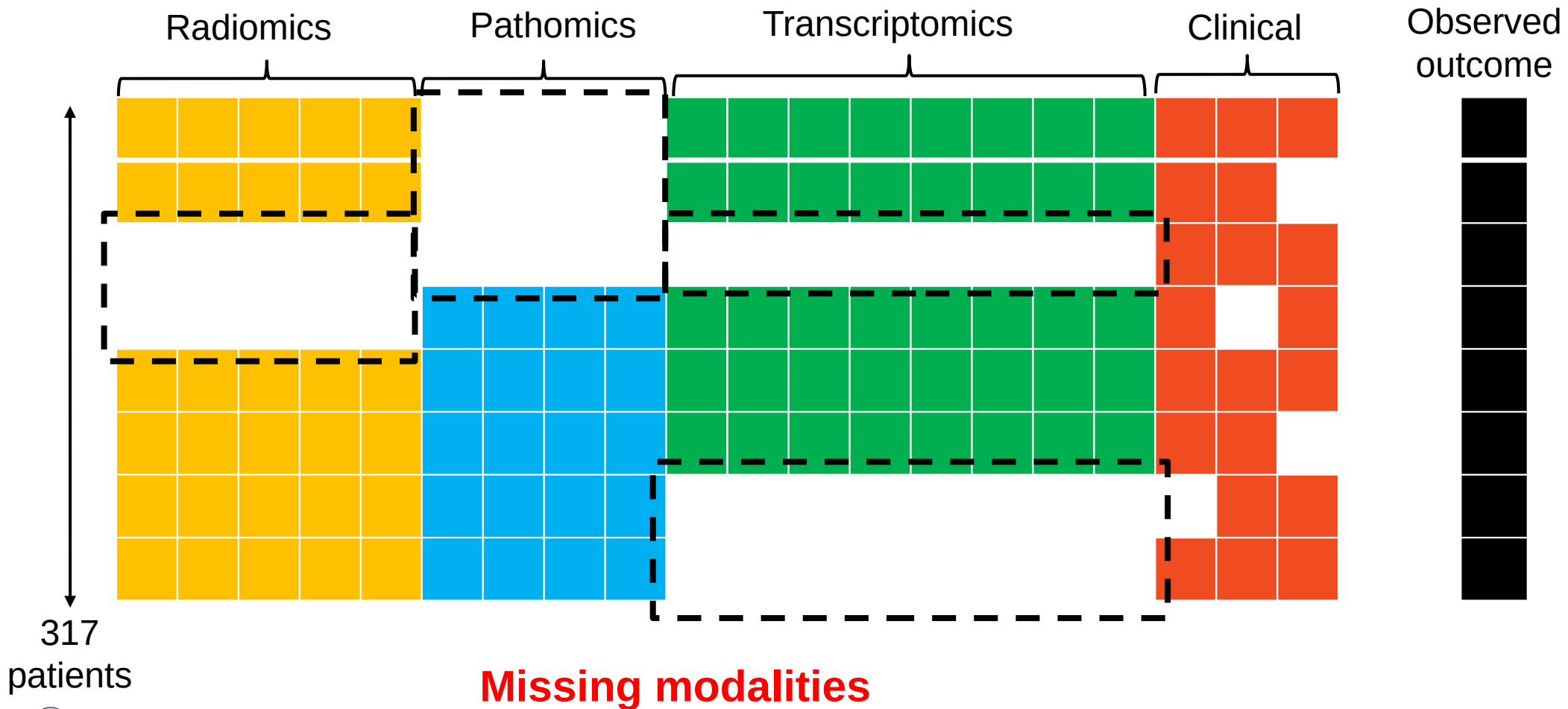




# A retrospective multimodal cohort - III



# A retrospective multimodal cohort - III



# Investigate multiple learning tasks to extract consensus trends

## Multiple outcomes

### Survival outcomes:

Overall Survival (OS)

Progression-Free Survival  
(PFS)

### Binary outcomes:

Death at 1 year

Progression at 6 months



# Investigate multiple learning tasks to extract consensus trends

| Multiple outcomes                                  | Multiple algorithms   |
|--|---|
| <b>Survival outcomes:</b><br>Overall Survival (OS) | <b>Linear algorithms:</b><br>Logistic regression with elastic net penalty |
| Progression-Free Survival (PFS)                    | Cox model with elastic net penalty  |
| <b>Binary outcomes:</b><br>Death at 1 year         | <b>Tree ensemble algorithms:</b><br>XGBoost                               |
| Progression at 6 months                            | Random Survival Forest  |

# Investigate multiple learning tasks to extract consensus trends

| Multiple outcomes                                  | Multiple algorithms   | Multiple fusion strategies                                     |
|--|---|--|
| <b>Survival outcomes:</b><br>Overall Survival (OS) | <b>Linear algorithms:</b><br>Logistic regression with elastic net penalty | <u>Late fusion</u>   |
| Progression-Free Survival (PFS)                    | Cox model with elastic net penalty  | Early fusion<br>(without or with univariate feature selection) |
| <b>Binary outcomes:</b><br>Death at 1 year         | <b>Tree ensemble algorithms:</b><br>XGBoost                               | Fusion with attention weights<br>(DyAM)                        |
| Progression at 6 months                            | Random Survival Forest  |  |



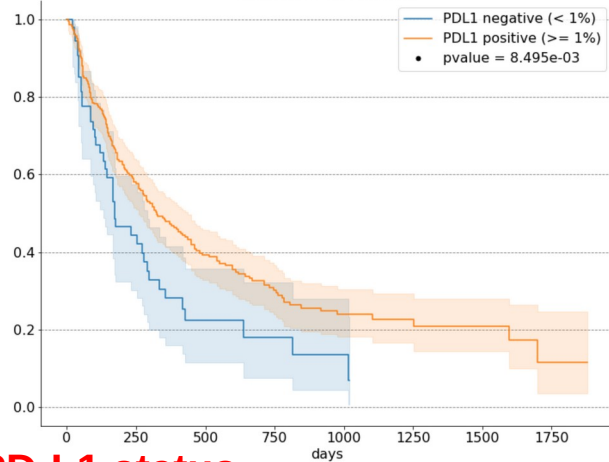
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# Univariate established biomarkers show limited performance

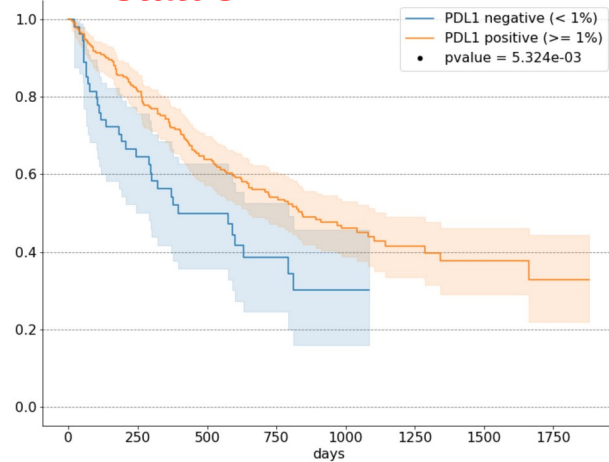
Progression-Free Survival



— PDL1 negative (< 1%)  
— PDL1 positive ( $\geq 1\%$ )  
• pvalue =  $8.495e-03$

**PD-L1 status**

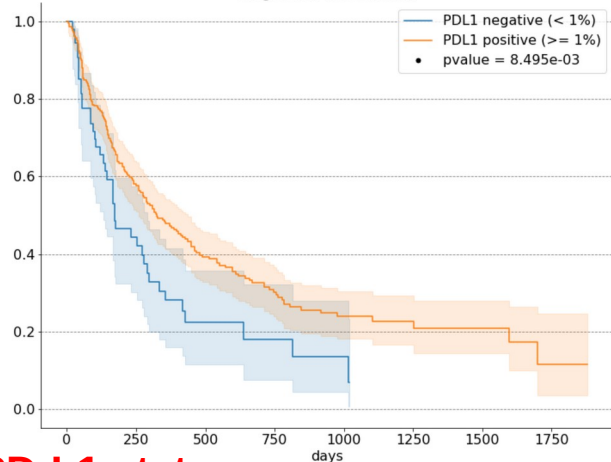
Overall Survival



— PDL1 negative (< 1%)  
— PDL1 positive ( $\geq 1\%$ )  
• pvalue =  $5.324e-03$

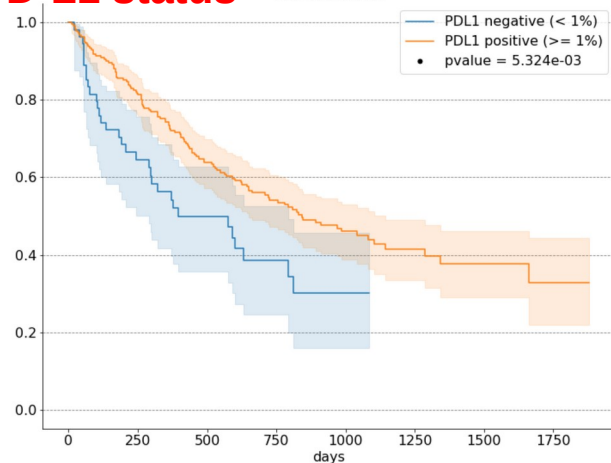
# Univariate established biomarkers show limited performance

Progression-Free Survival



**PD-L1 status**

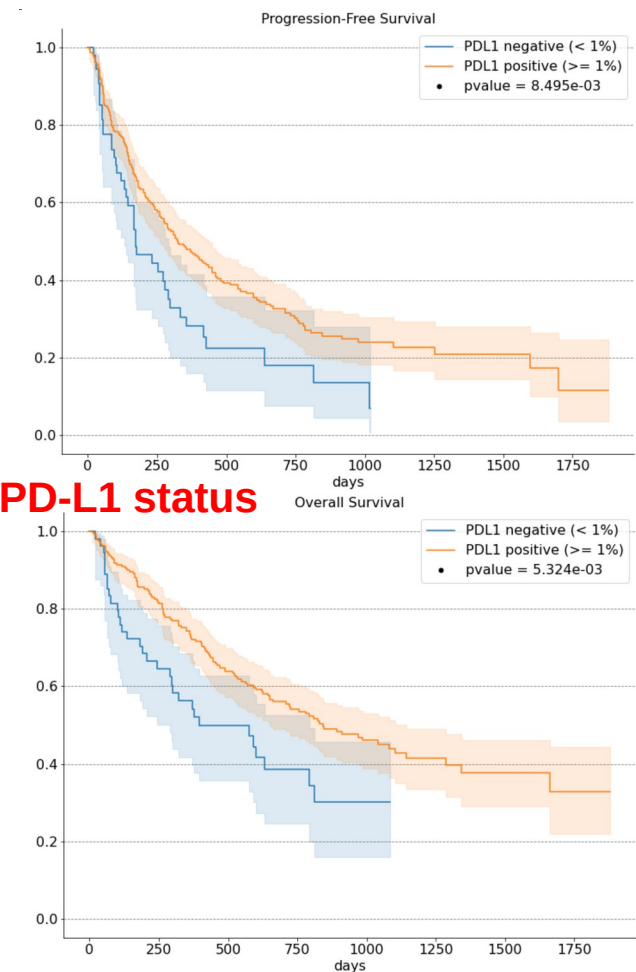
Overall Survival



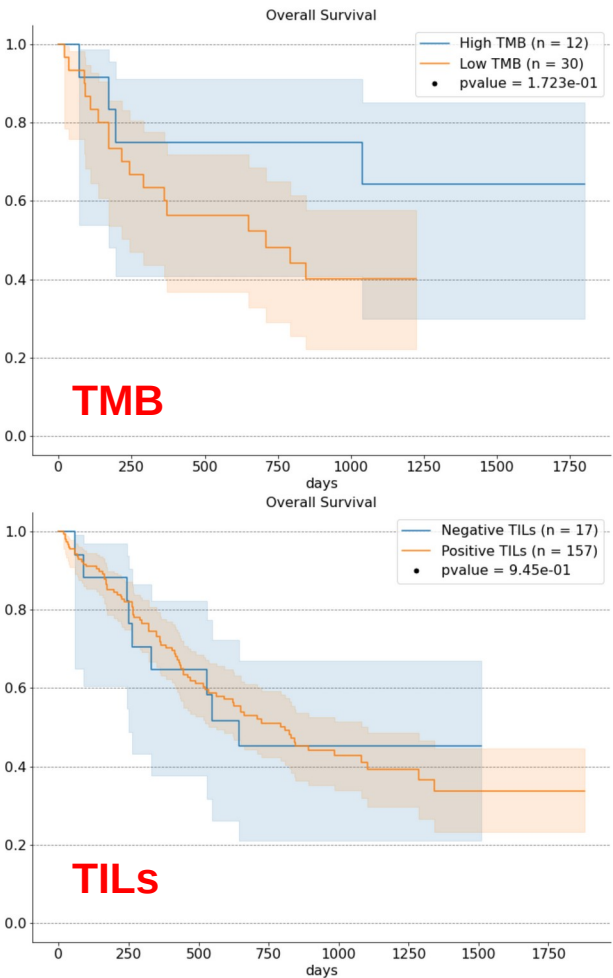
|                            | C-index OS<br>(whole cohort)     |
|----------------------------|----------------------------------|
| PD-L1<br>binary status     | 0.54 [0.51 – 0.57]<br>pval=0.014 |
| PD-L1 score<br>(100 – TPS) | 0.53 [0.48 – 0.58]<br>pval=0.104 |



# Univariate established biomarkers show limited performance



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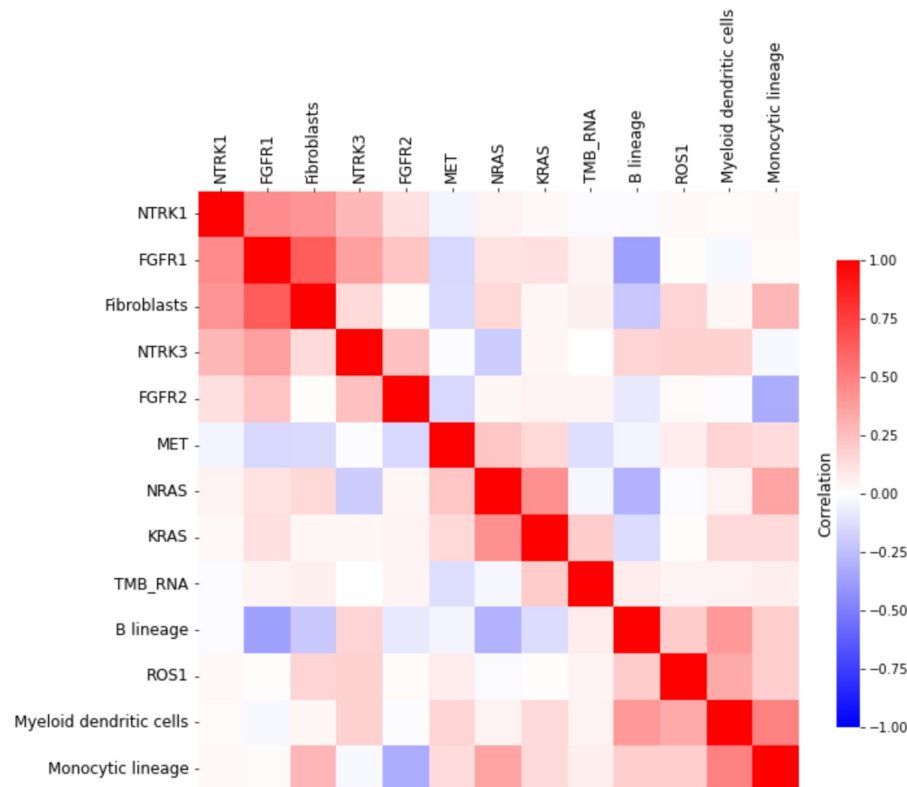
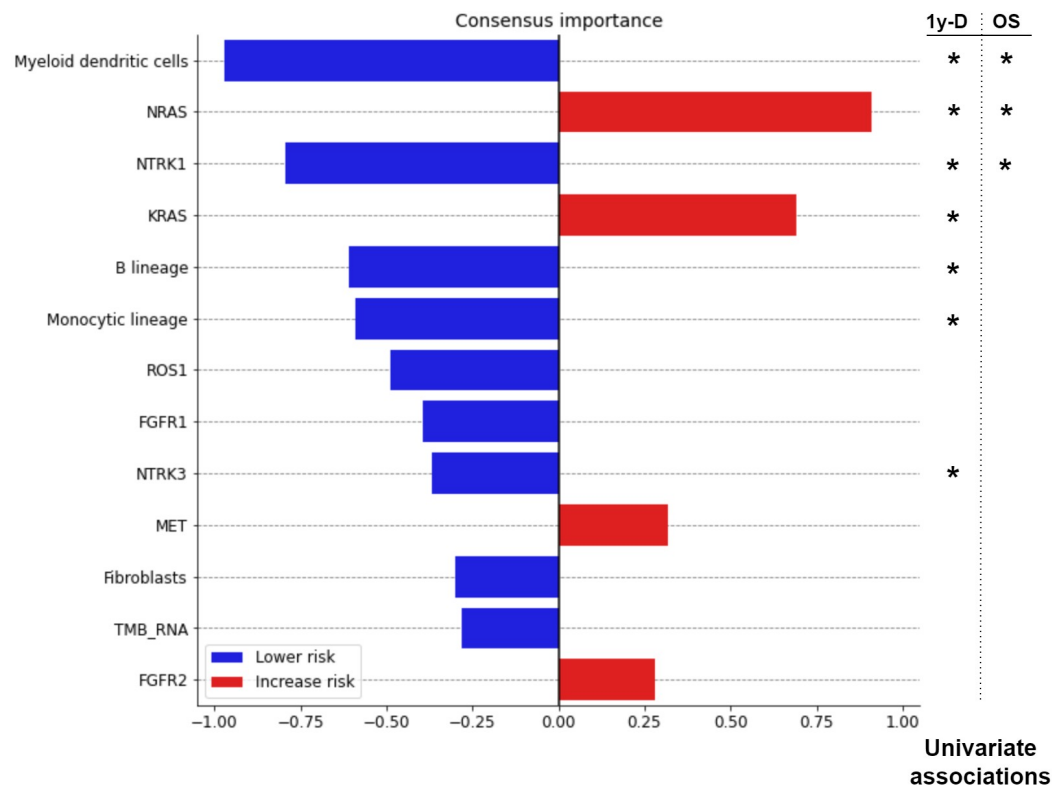
# A benchmark of unimodal predictors

| Target (number of patients) |                | OS<br>(n=79)       | 1-year death<br>(n=77) | PFS<br>(n=80)      | 6-months progression<br>(n=75) |
|-----------------------------|----------------|--------------------|------------------------|--------------------|--------------------------------|
| Metric                      |                | C-index            | AUC                    | C-index            | AUC                            |
| Clinical                    | Tree ensembles | 0.67±0.01 *        | 0.59±0.05              | 0.56±0.02          | 0.58±0.04                      |
|                             | Linear         | 0.60±0.02 *        | 0.73±0.02 *            | 0.53±0.03          | <b>0.61±0.03 *</b>             |
| Radiomics                   | Tree ensembles | 0.61±0.02 *        | 0.62±0.04              | 0.57±0.01          | 0.56±0.05                      |
|                             | Linear         | 0.61±0.02 *        | 0.47±0.03              | 0.55±0.02          | 0.48±0.04                      |
| Pathomics                   | Tree ensembles | 0.59±0.02          | 0.54±0.05              | 0.56±0.02          | 0.58±0.06 *                    |
|                             | Linear         | 0.58±0.02          | 0.56±0.03              | 0.51±0.02          | <b>0.61±0.03 *</b>             |
| RNA                         | Tree ensembles | <b>0.69±0.02 *</b> | <b>0.75±0.04 *</b>     | 0.57±0.02          | 0.60±0.04 *                    |
|                             | Linear         | 0.58±0.02          | 0.65±0.03              | <b>0.59±0.02 *</b> | <b>0.61±0.03</b>               |

\*: permutation p-value  $\leq 0.05$

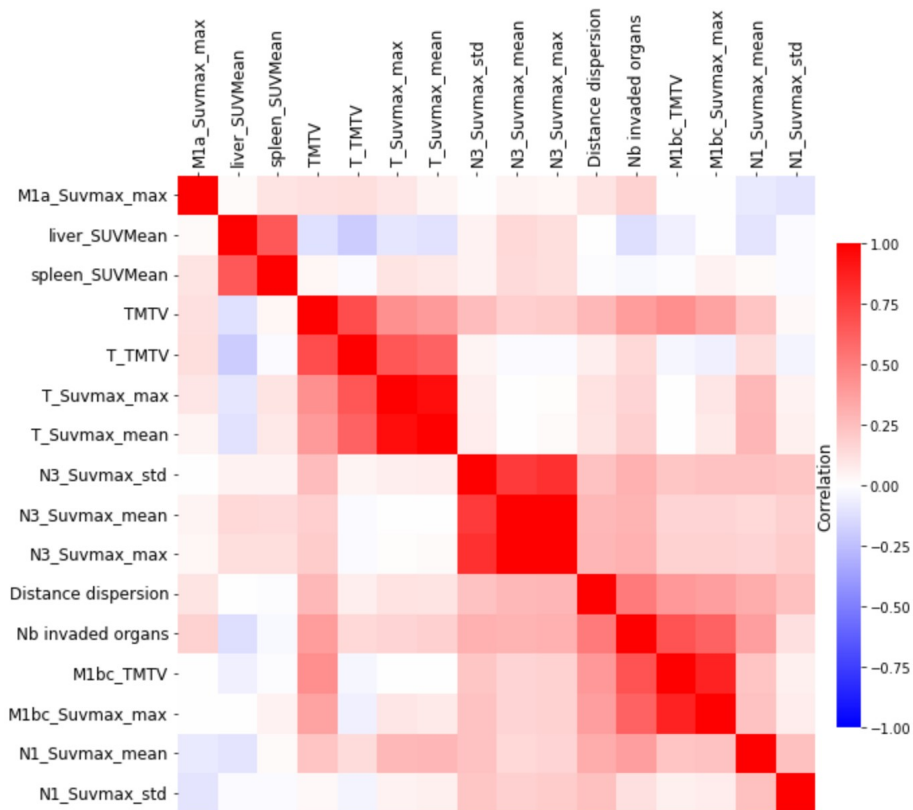
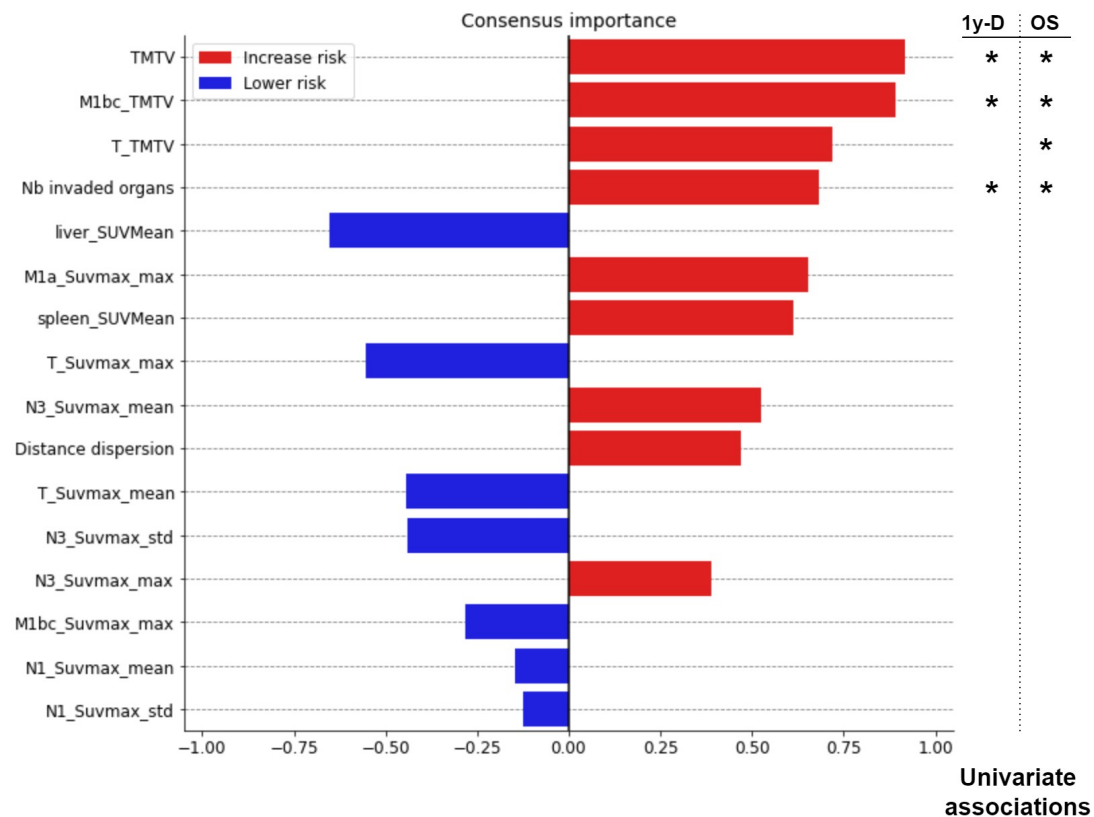
# Robust feature importance – RNA modality

Feature importance ranking aggregated over both tasks (OS and 1 year death) and both algorithms (linear and tree ensemble)

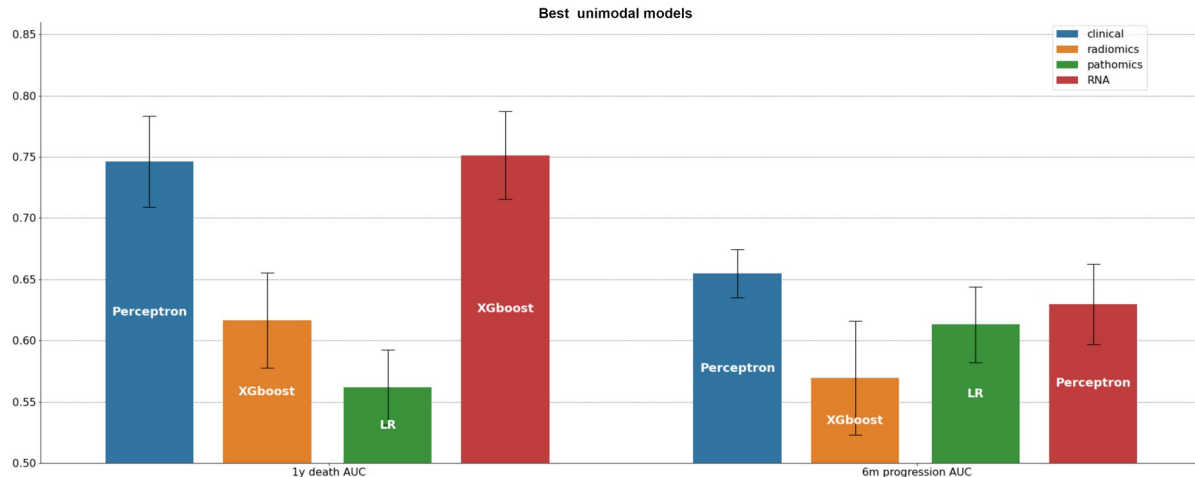
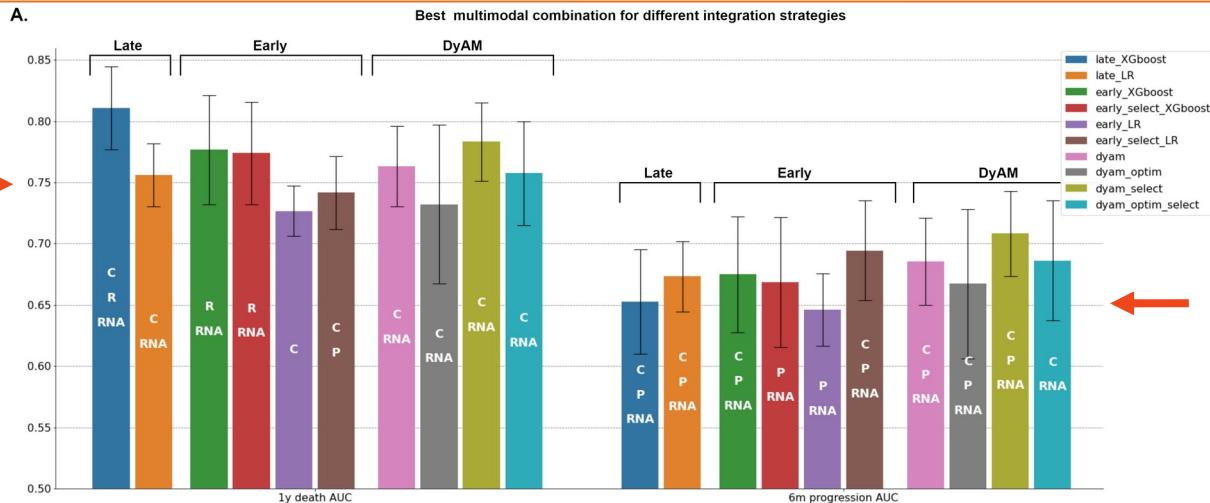


# Robust feature importance – radiomic modality

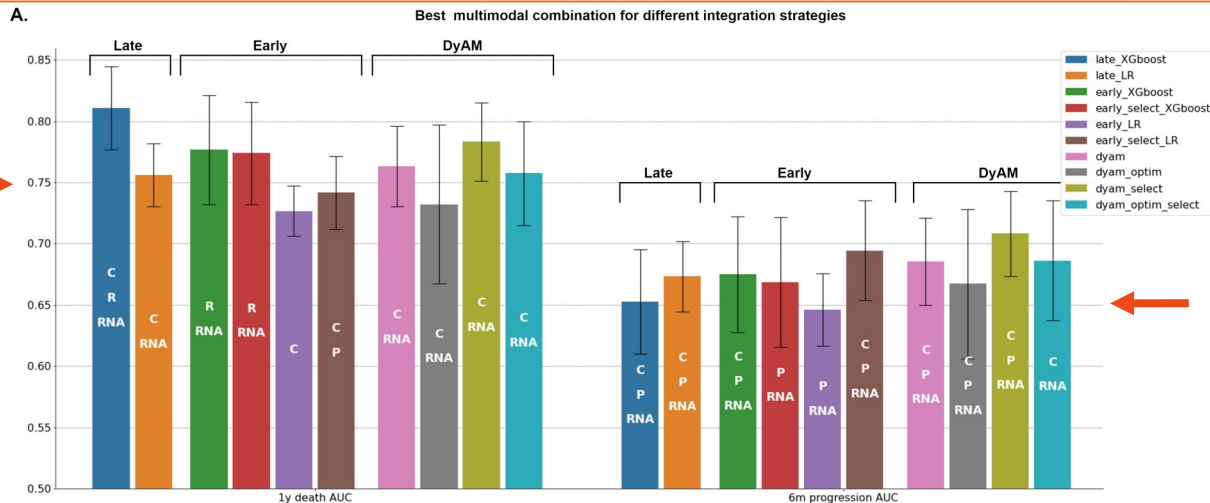
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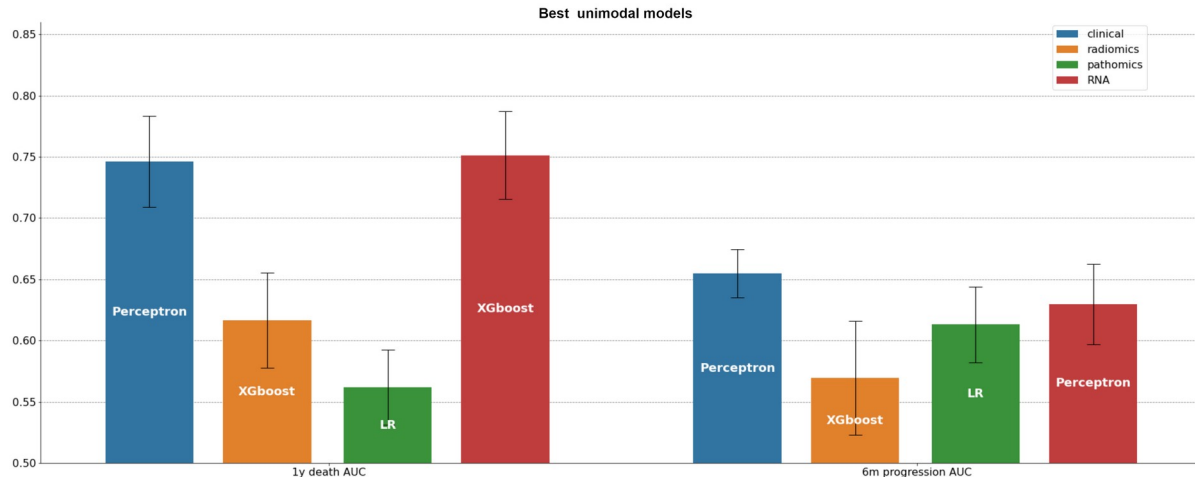
# Benchmark highlights benefits of multimodal approaches



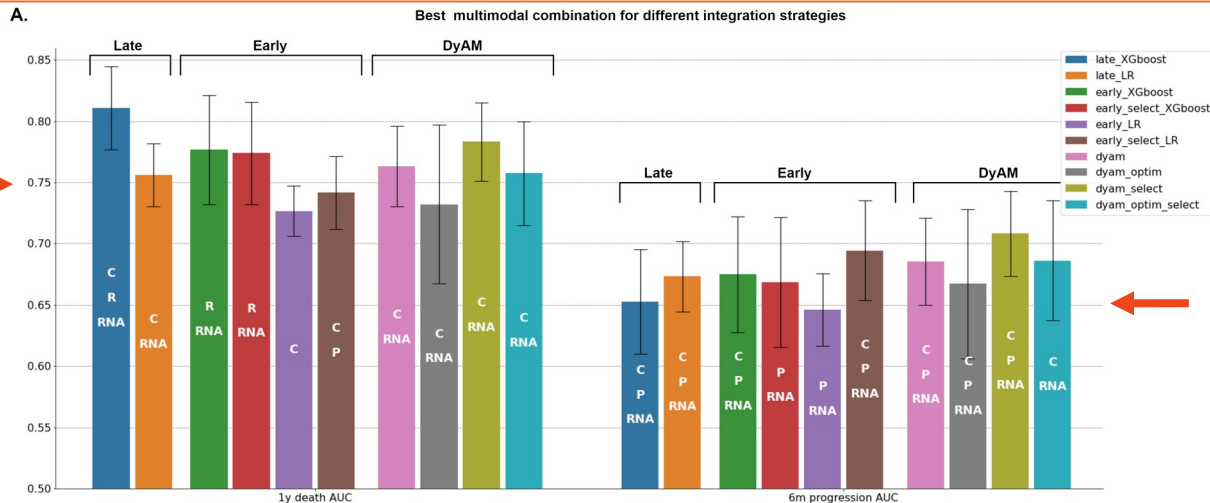
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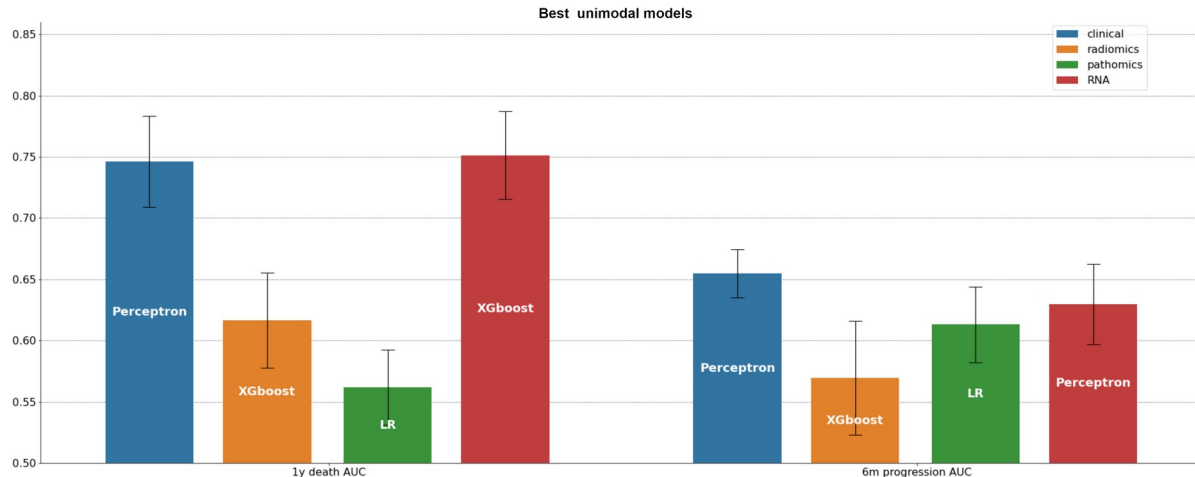
- The majority of multimodal models outperformed the best unimodal models



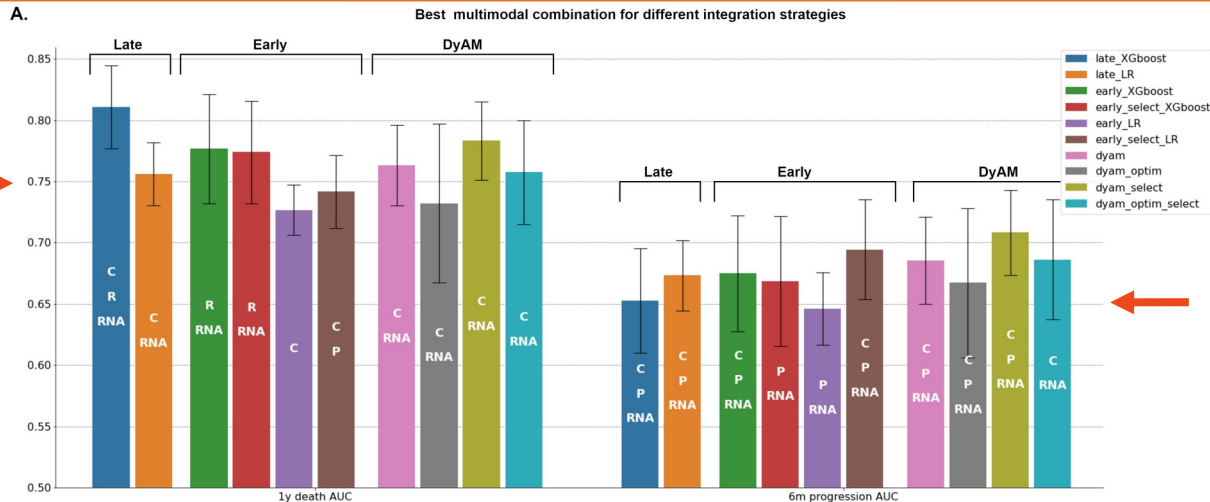
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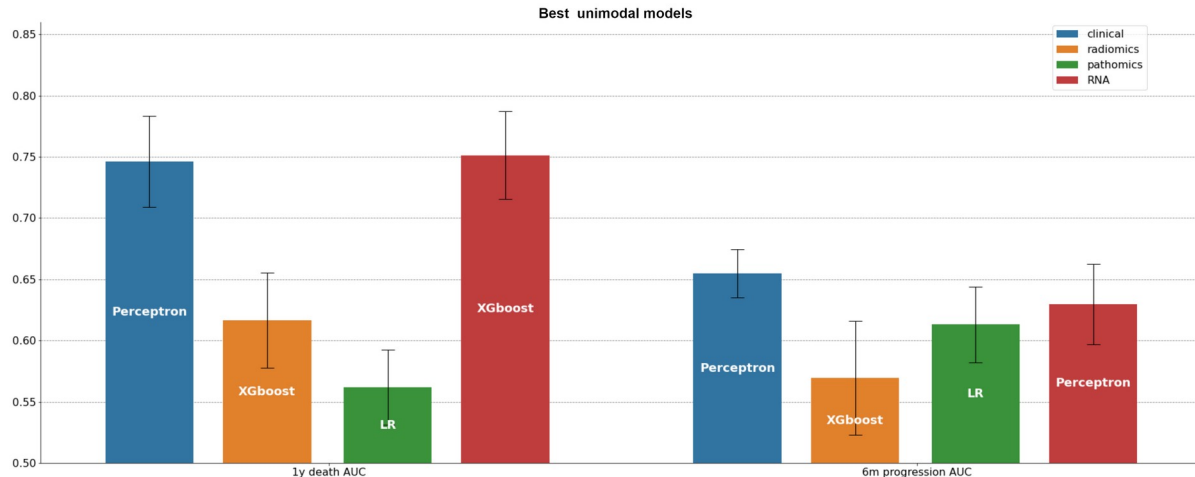
- The majority of multimodal models outperformed the best unimodal models
- Late fusion performed the best for 1 year death prediction



# Benchmark highlights benefits of multimodal approaches

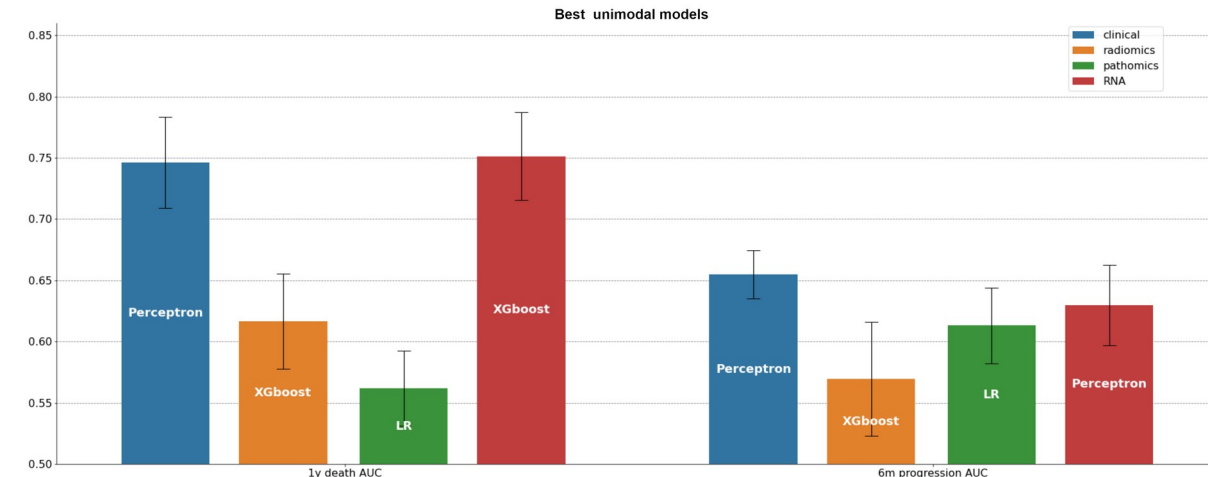
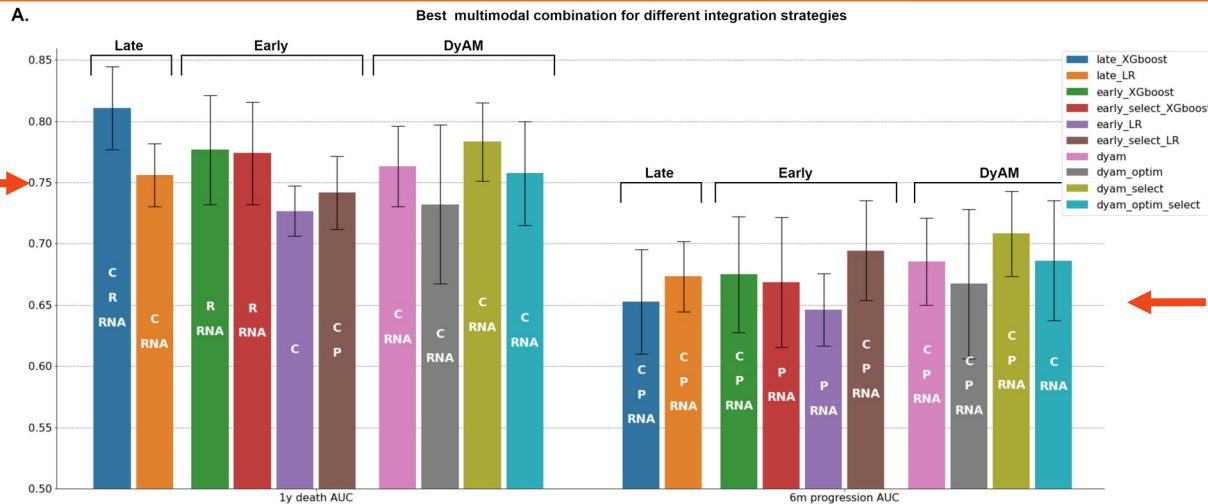


- The majority of multimodal models outperformed the best unimodal models
- Late fusion performed the best for 1 year death prediction
- Clinical, RNA and radiomic consistently involved in the best multimodal models for 1y death prediction





# Benchmark highlights benefits of multimodal approaches



- The majority of multimodal models outperformed the best unimodal models
- Late fusion performed the best for 1 year death prediction
- Clinical, RNA and radiomic consistently involved in the best multimodal models for 1y death prediction

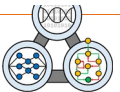
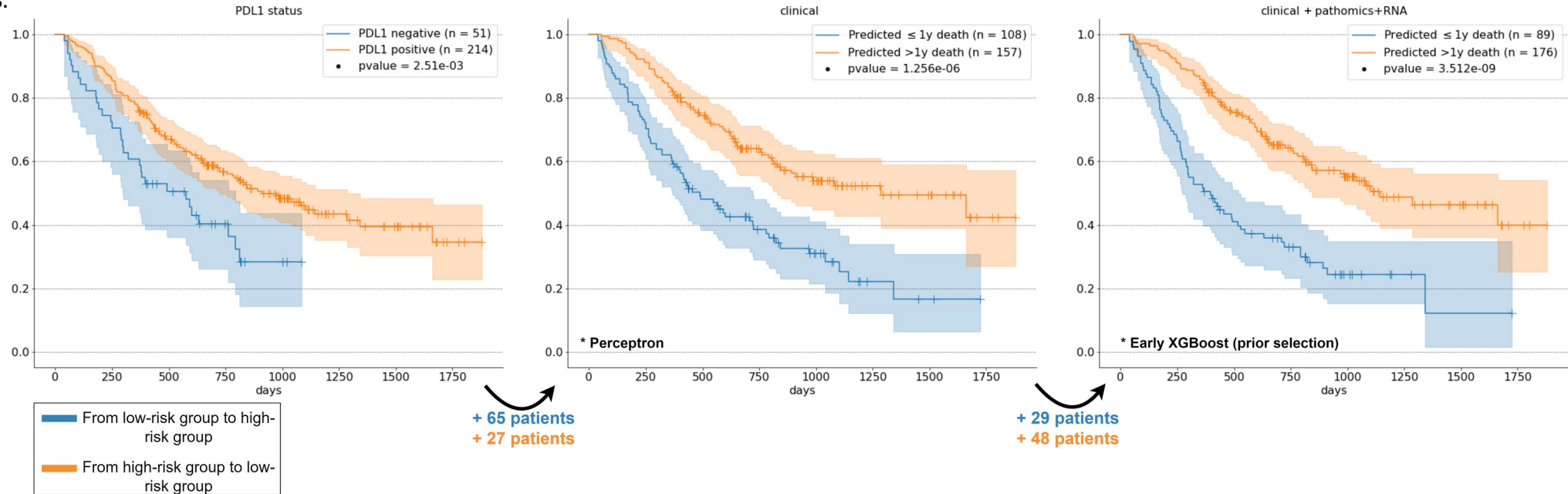
No integration strategy performed the best for all the tasks



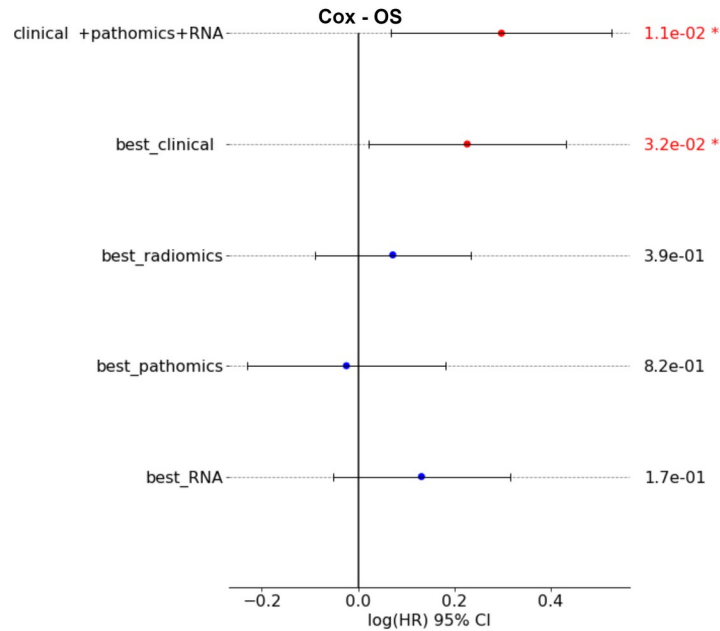
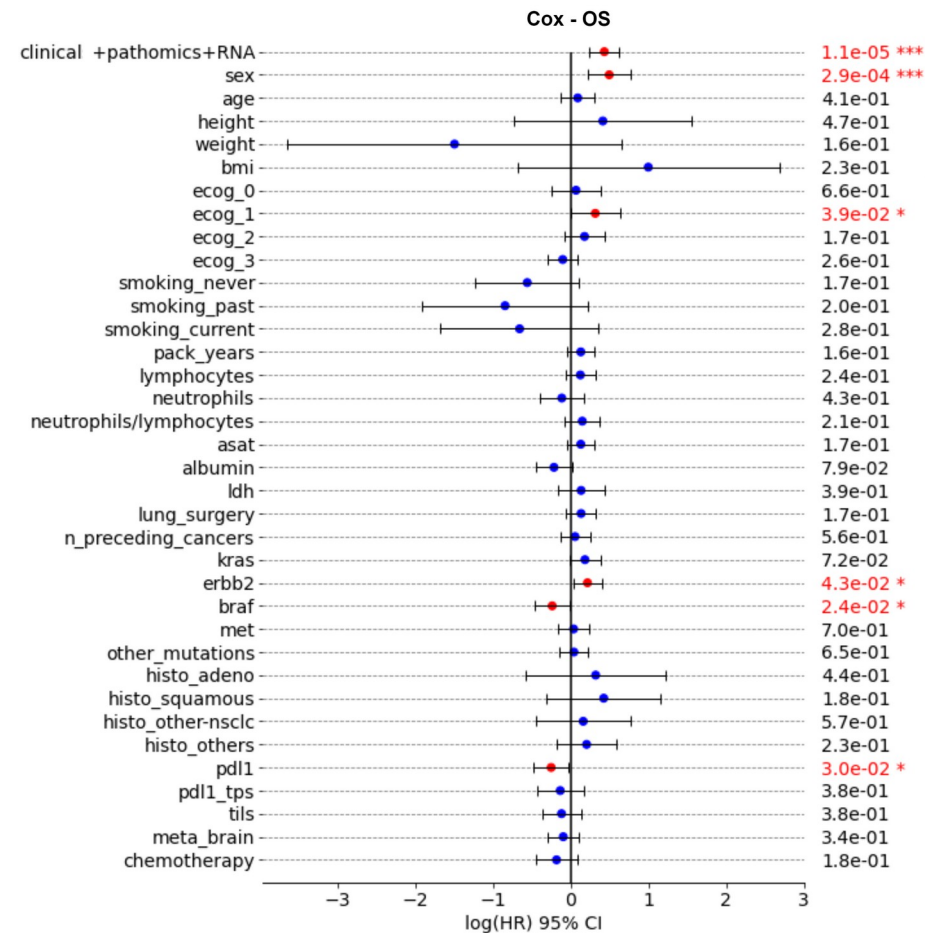
# Multimodal score improves patient risk stratification

- A threshold is learnt on the training set of each fold of the cross-validation scheme and applied to the test set.
- Group membership is defined as the most frequently attributed group (low risk vs high risk) across the 100 repeats.

B.



# Multimodal score brings additional predictive information



Likelihood-ratio tests show a significant effect of the multimodal score wrt to routine clinical information



# Conclusions

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1. Late fusion is a relevant baseline strategy/starting point for further multimodal studies (simplicity, handle missing modalities easily...)
2. We provided new evidence of the relevance of multimodal approach for building powerful predictors □ should motivate others to collect new and larger multimodal cohorts.
3. Simple signatures of the Tumor MicroEnvironment seems to predict well NSLC outcome and lead to relevant hypotheses.
4. The results need to be further validated on larger an external data sets. More complex (e.g. end-to-end strategies) should be investigated.

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# Project 1: Study the link between radiomics and RNA expression

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- 93+ patients with RNAseq and PET images (biopsy site is missing for some of them)
- Are different clusters based on radiomics/imaging characteristics associated with differentially expressed genes or biological pathways ?
- Can we identify/probe biological pathways deregulation with radiomic characteristics ?
- Can we use RNAseq data to understand better the radiomic phenotype of tumors ?

## Project 2: Multiple Instance Learning for metastatic disease

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- Use each metastasis as an instance with a hidden outcome to predict patient's outcome

# Project 3: Improve PET representation with supervised learning

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- Predict the total tumor volume in lymph nodes, distant regions... from unannotated MIP images

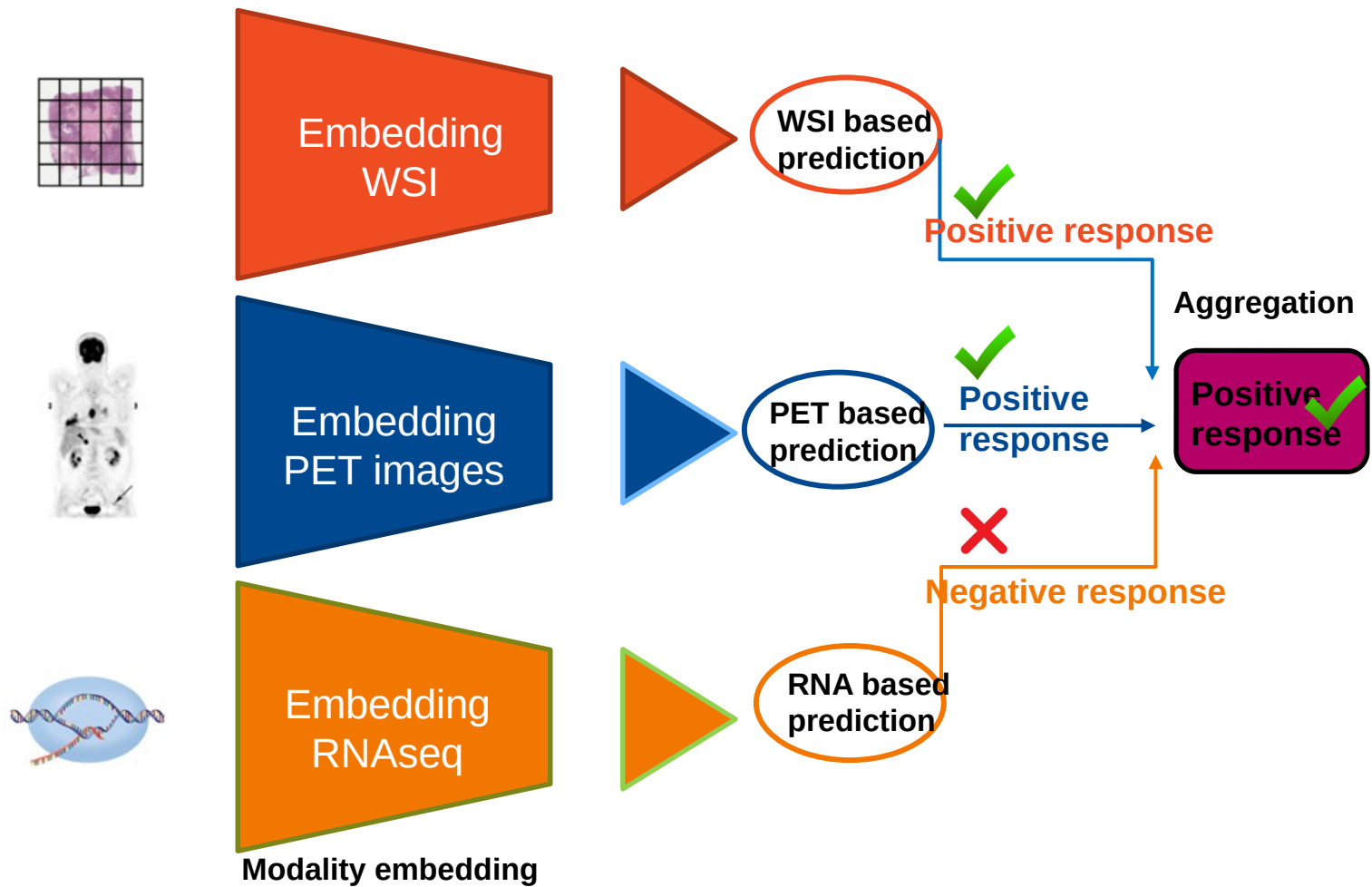


# Acknowledgements

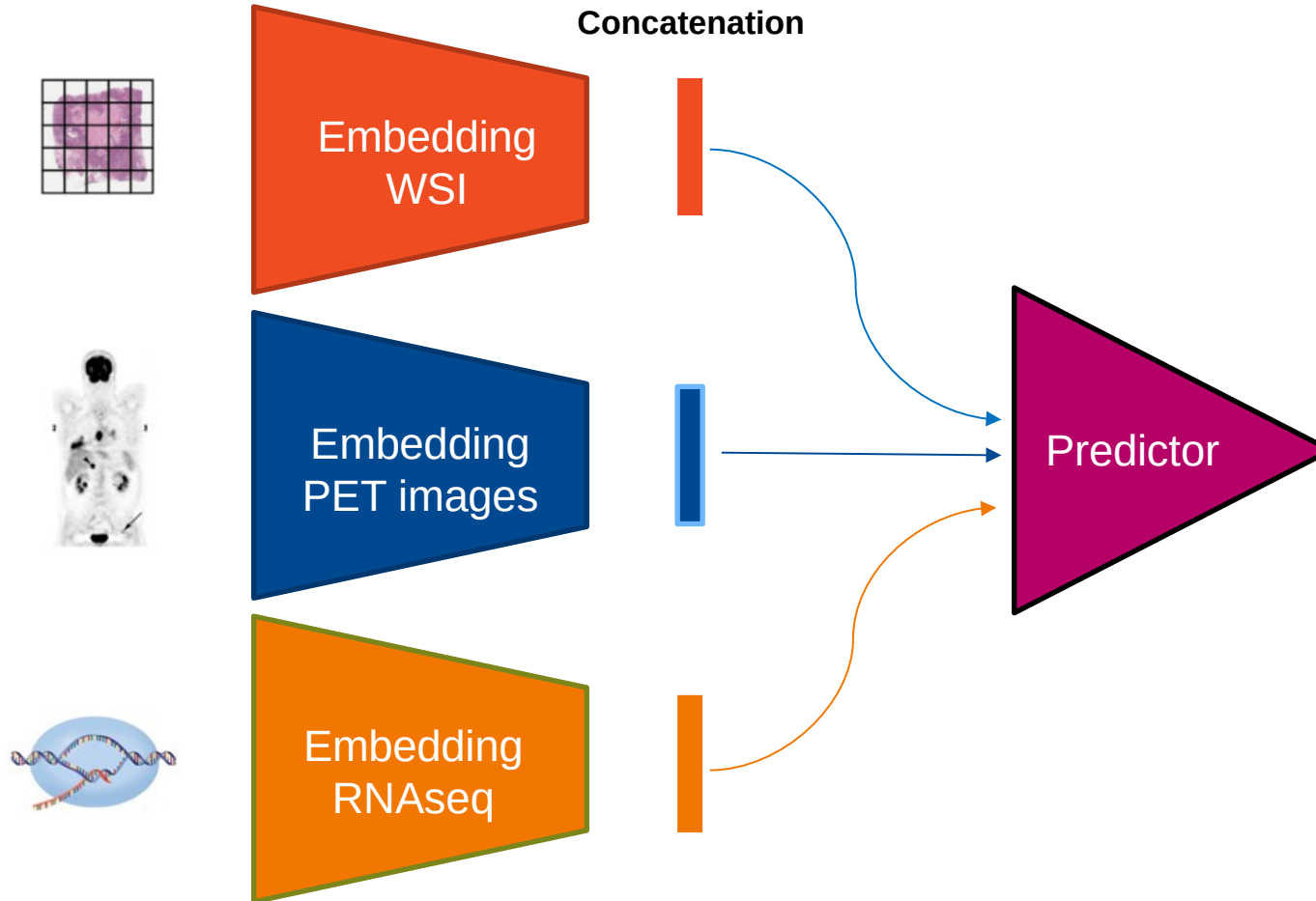
- **Curie-Montsouris Chest center:** Nicolas Girard, Sarah Lagha, Anne-Sophie Tedesco
- **Imaging department Curie:** Hervé Brisse, Marie Luporsi, Toulis Ramtohol
- **Pathology department Curie:** Clément Beaulaton
- **Data department Curie:** Paulette Salamoun Feghali
- **LITO Curie:** Irène Buvat, Fanny Orlhac, Narinée Hovhannisyan, Nicolas Captier, Erwin Woff
- **SysBIO U900 Curie:** Emmanuel Barillot, Andrei Zinovyev, Christine Lonjou, Nicolas Captier
- **CBIO U900 Curie-Mines:** Thomas Walter, Marvin Lerousseau
- \* **Immunity and Cancer:** Hélène Salmon



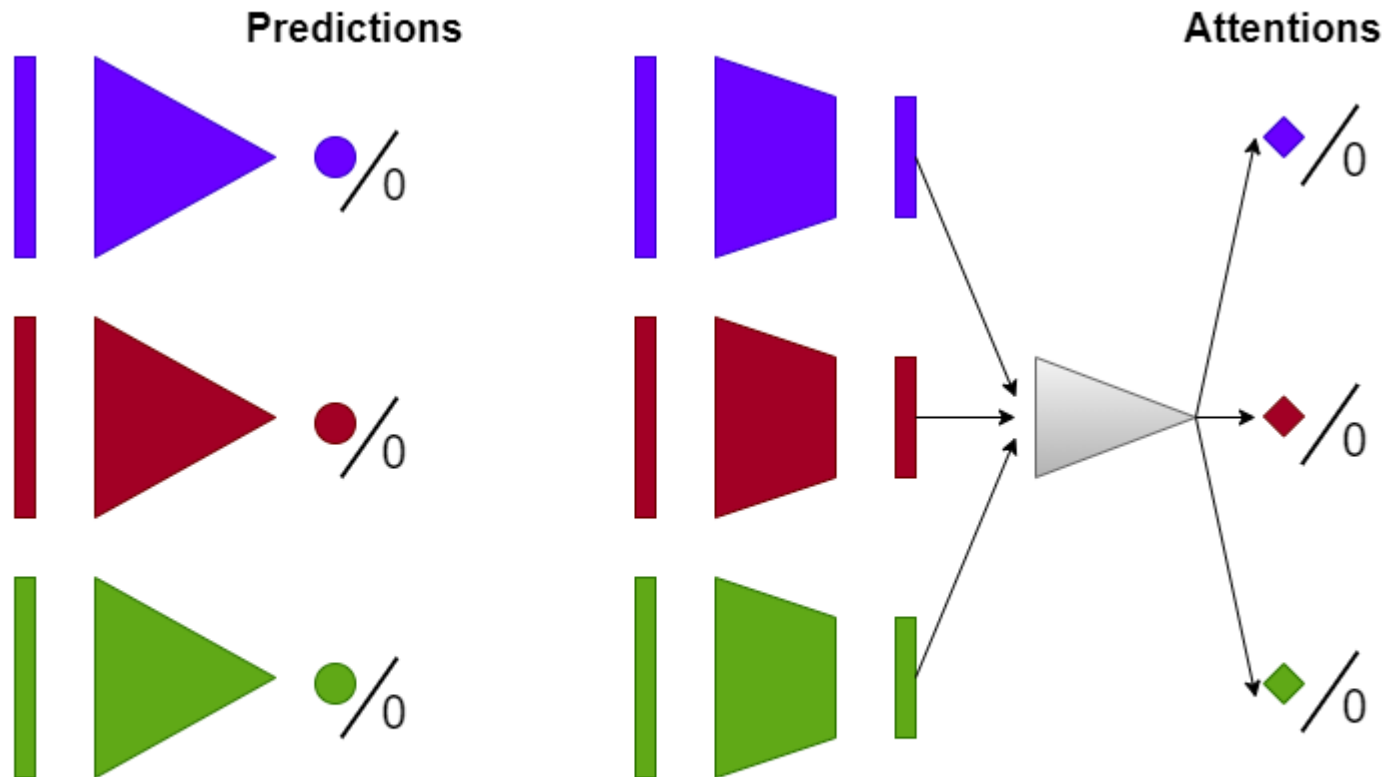
# Late fusion strategy



# Early fusion strategy



# DyAM strategies



- Attention mechanism weighs each modality for each patient
- The model learns to combine different modalities through normalization and grey network

Final prediction:  $\text{blue circle} \times \text{blue diamond} + \text{red circle} \times \text{red diamond} + \text{green circle} \times \text{green diamond}$