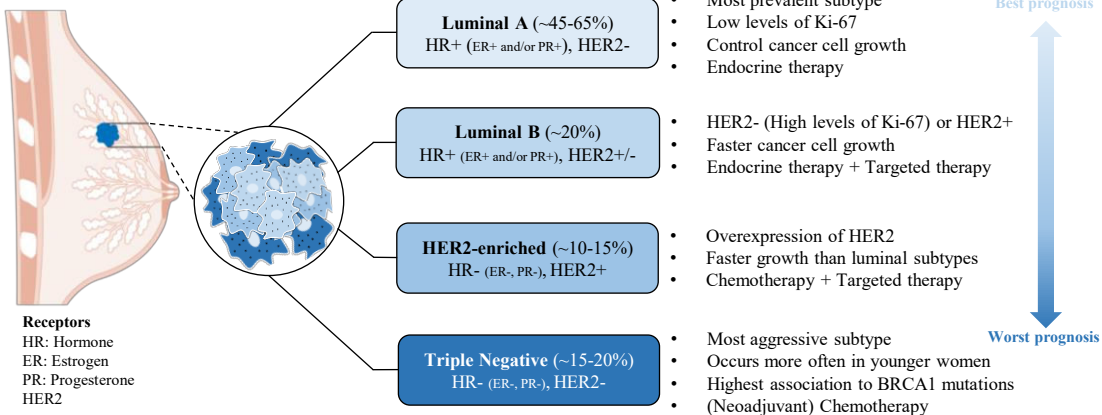


# Predicting molecular subtypes of breast cancer using multimodal deep learning and incorporation of the attention mechanism

## Introduction

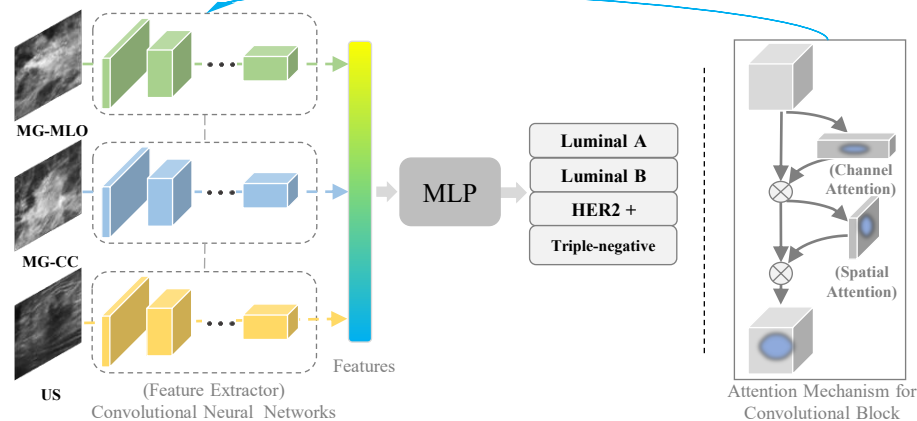
- Breast cancer can be divided into four molecular subtypes, which is usually based upon the expression levels of ER, PR, HER2 and Ki-67.



- The molecular subtypes of breast cancer is an important factor for the **prognosis** of breast cancer patients, and can **guide treatment selection**.

## Model

- The overall pipeline diagram of multimodal deep learning with attention mechanism (MDLA) model. The model was developed based on residual neural network.

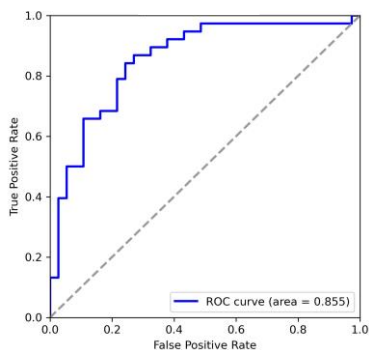


- Two views of mammography (MG, medio-lateral oblique -MLO- and cranio-caudal -CC-) and an ultrasound (US) image of each case (lesion location) were used as input.

## Results

- The confusion matrix (for 4-class molecular subtypes) and ROC curve (for Luminal vs Non-Luminal) for predicting molecular subtypes.

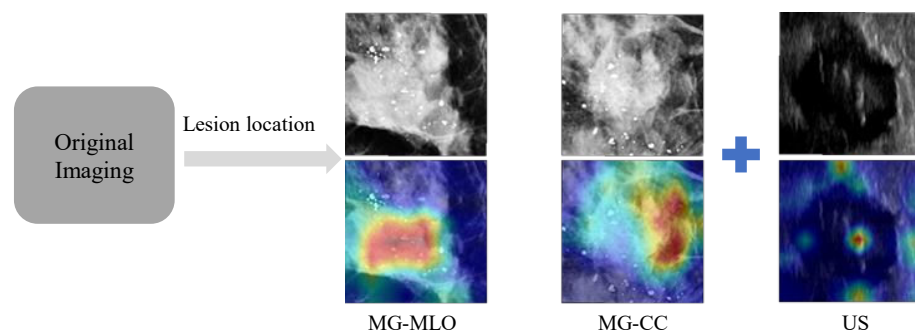
Ground true \ Predict	Luminal-A	Luminal-B	HER2+	Triple-negative
Luminal-A	93.2%	4.1%	2.7%	
Luminal-B	11.1%	87.0%	1.9%	
HER2+	5.9%	29.4%	64.7%	
Triple-negative	3.7%	18.5%		77.8%



- The MCC was **0.794** for predicting 4-class molecular subtypes of breast cancer, and the AUC was **0.855** for distinguishing between Luminal and Non-Luminal using MDLA model.

## Visualization

- The visualization of MDLA model in predicting 4-class molecular subtypes of breast cancer, including MG and corresponding US.



- The MDLA model focuses on the information of the lesion area of MG and US images, including masses, calcifications, lesion morphology and so on.



First Author: Tianyu Zhang



Corresponding Author:

Ritse M. Mann

[r.mann@nki.nl](mailto:r.mann@nki.nl)

[ritse.mann@radboudumc.nl](mailto:ritse.mann@radboudumc.nl)

## Co-authors:

Tianyu Zhang  
Luyi Han  
Yuan Gao  
Xin Wang  
Regina Beets-Tan  
Ritse M. Mann



MU & RUMC & NKI