



Corrigendum: Machine Learning Enables Accurate and Rapid Prediction of Active Molecules Against Breast Cancer Cells

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A Corrigendum on

Machine Learning Enables Accurate and Rapid Prediction of Active Molecules Against Breast Cancer Cells

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In the original article, there was a mistake in **Figures 4 and 5** as published. There are some errors in the figure insertion, **Figure 4** is repeated with **Figure 3**, and **Figure 5** is the result of **Figure 4**. The corrected **Figures 4 and 5** appear below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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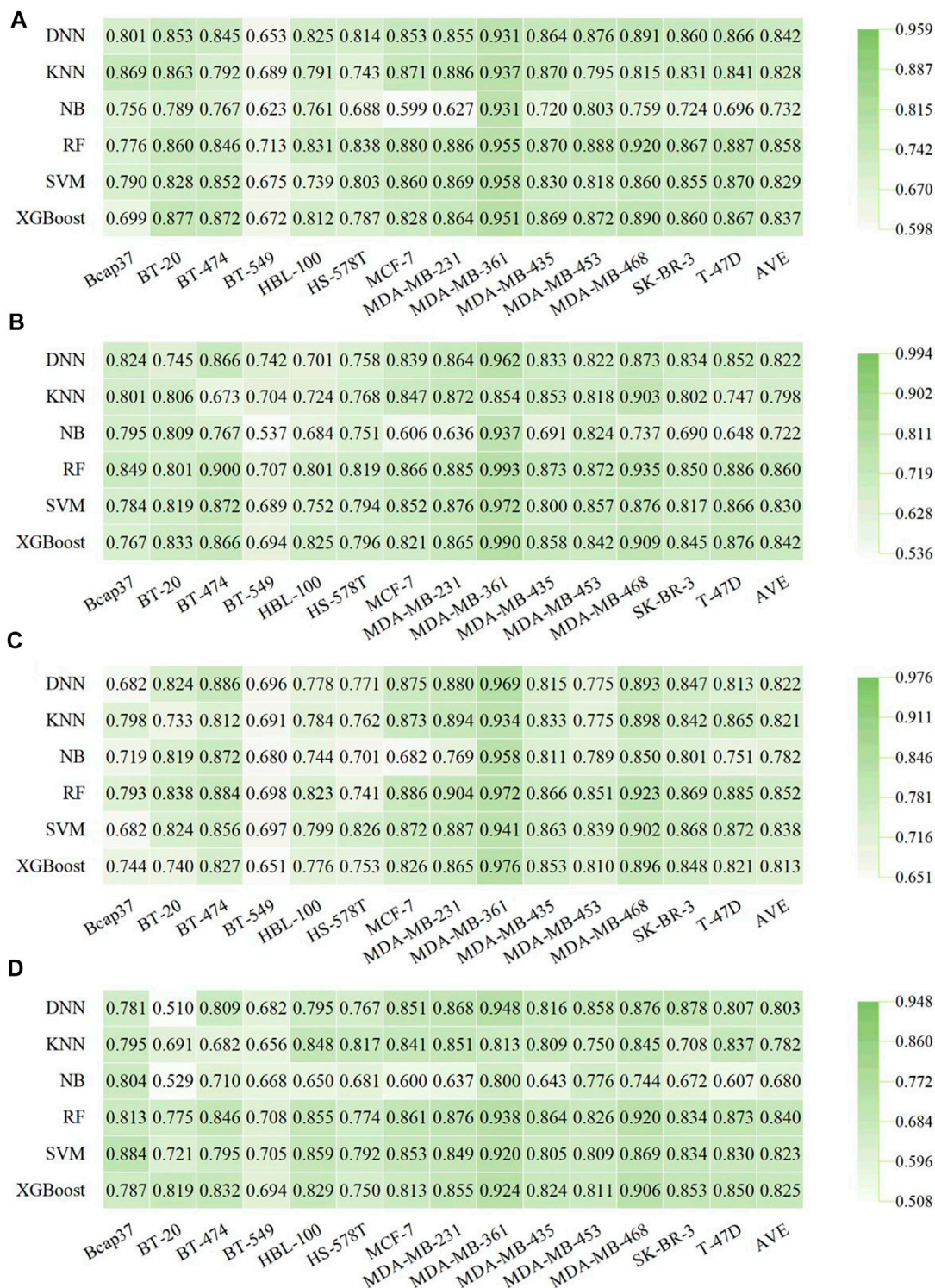


FIGURE 4 | Performance of fingerprint-based BC prediction models. **(A)** AUC results of the AtomPairs-based models. **(B)** AUC results of the MACCS-based models. **(C)** AUC results of the Morgan-based models. **(D)** AUC results of the PharmacofPF-based models.

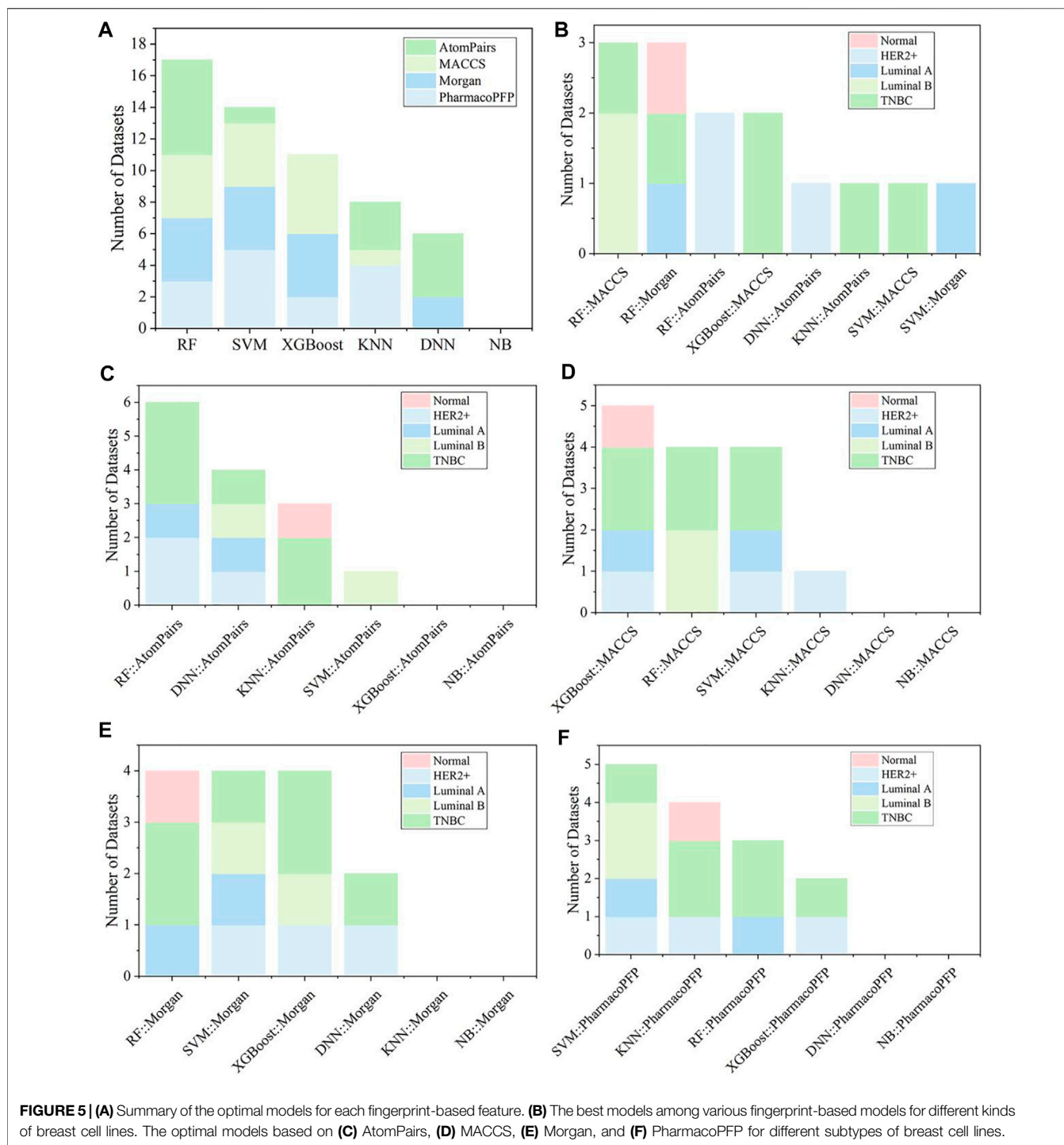


FIGURE 5 | (A) Summary of the optimal models for each fingerprint-based feature. **(B)** The best models among various fingerprint-based models for different kinds of breast cell lines. The optimal models based on **(C)** AtomPairs, **(D)** MACCS, **(E)** Morgan, and **(F)** PharmacoPFP for different subtypes of breast cell lines.