* **Title**

**Identification of breast cancer metastasis markers using machine learning approaches with gene expression profiles**

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텍스트, 도표, 스크린샷, 라인이(가) 표시된 사진

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**Figure S1. Boxplots of the preprocessed gene expressions for randomly selected participants (left) and genes (right)**

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**Figure S2. The box plots of the 50 AUCs obtained from the 50 trained XGBoost models.**