

**Identification of non-intuitive biomarkers predicting dependency on AKT vs. MAPK signaling of ERBB2+ cancer cells using an integrated experimental and computational framework****Jinyan Du**

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Understanding the molecular pathways by which oncogenes drive cancerous cell growth and how dependence on such pathways varies between tumors, could be highly valuable for the design of anti-cancer treatment strategies. In this work, we study how dependence upon the canonical PI3K and MAPK cascades varies across HER2+ cancers and define biomarkers predictive of pathway dependencies. A panel of 18 HER2+ (ERBB2-amplified) cell lines representing a variety of tumor types was used to characterize the functional and molecular diversity within this oncogene-defined cancer. PI3K and MAPK-pathway dependencies were quantified by measuring *in vitro* cell growth responses to combinations of AKT (MK2206) and MEK (GSK1120212; trametinib) inhibitors, in the presence and absence of the ERBB3 ligand heregulin (NRG1). A combination of three protein measurements comprising the receptors EGFR, ERBB3 (HER3) and the cyclin-dependent kinase inhibitor p27 (CDKN1B) was found to accurately predict dependence on PI3K/AKT vs. MAPK/ERK signaling axes. Notably, this multivariate classifier outperformed the more intuitive and clinically employed metrics, such as expression of phospho-AKT and phospho-ERK and PI3K pathway mutations (PIK3CA, PTEN and PIK3R1). The predictability of the three protein biomarkers for differentiating PI3K/AKT vs. MAPK dependence in HER2+ cancers was confirmed using external datasets (Project Achilles and GDSC), again out-performing clinically used genetic markers. Measurement of this minimal set of three protein biomarkers could thus inform treatment and predict mechanisms of drug resistance in HER2+ cancers. More generally, our study provides an integrated framework for identifying non-intuitive biomarkers in cancer patients.

**Biography**

Jinyan Du has completed her PhD at Harvard University and Postdoctoral training at the Broad Institute. She is currently a Principal Scientist at Merrimack Pharmaceuticals, a fully integrated biopharmaceutical company that is building one of the most robust oncology pipelines in the industry. She has published over 30 papers in reputed journals.

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