

Interactions between breast cancer cells and their stromal component: An analysis of alterations in gene expression

Christos Sotiriou, Frank El Ouriaghli, Samira Majjaj, Benjamin Haibe-Kains, Christine Desmedt, Denis Larsimont, Martine Piccart. Translational Research and Medical Oncology Unit, Université Libre de Bruxelles (U.L.B.), Institut Jules Bordet, 121 Boulevard de Waterloo, 1000 Brussels, Belgium

Abstract

Background: Epithelial-stronal interactions are known to be important in normal mammary gland development and to play a role in breast carcinogenesis. The aim of this study was to explore the influence of breast tumor microenvironment on primary tumor growth, breast cancer sub-typing and prognosis.

Methods: Myo-fibroblast cells (Cb10) were isolated and purified from breast tumor (№ 28) and normal (№4) fissues. Gene expression analysis was performed using the Affymetrik GencChip@ Human Genome U133 Plus 2.0 arrays. Survival analysis was carried out using 12 publicly available microarray datasets including more than 1200 systemically untreated breast cancer patients.

Resulting Breast tumor myo-fibroblast stroma cells showed an alterced gave expression patterns to the ones isolated from ormal breast tissues. While some of the differentially expressed gaves are found to be associated with extracellular matrix formation/degradation and angiogenesis, the function of several other genes remains largely unknown. Unsupervised hierarchical clustering analysis clustered breast cancer based on ER, IRER status and tumor differentially, expression sportaries of breast cancer based on ER, IRER status and tumor differentiation, expression sportaries of breast cancer based on ER, IRER status and tumor differentiation, expression showed a statusfically significant expression is and tumor differentiation. As the status and tumor differentiation, status and tumor differentiation sports and breast tumors with expression levels of the stroma significant were significantly associated with weres progress (HER2 havens) with high and low acgression levels of the stroma significant based to the stroma significant were showed 40% and 80% distant metastasis free survival at 5-year follow-up respectively (IRE 2.32, (12):4.490, pp.529 10-).

Conclusions: Our results highlight the importance of tunor epithelial-stroma cell interactions in breast carcinogenesis and breast cancer sub-yoing. Moreover, it shows the road of stroma cells in tumor dissemiration particularly within the HER2+ subtype and provide basis for the development of novel therepartic strategies.















Differences in Gene Expression of CD10 (fibroblasts) According to Molecular Subtypes



Type 1: ER-/HER2- (basal-like) Type 2: HER2+ (HER2-like) Type 3: ER+/HER2- (luminal-like Ligrade 1; I:grade 2; H:grade 3

Clinical Relevance? All patients N≈1400 Systemically Untreated





Clinical Relevance? Molecular Subtypes





HR=3.076 CI 1.544-6.131 P=0.001



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