

# Stabilization of Pim-3 by TCTP

Zhang et al. Page 1508

Pim-3, a proto-oncogene with serine/ threonine kinase activity, is aberrantly expressed in cancerous tissues and is known to play a critical role in tumor development and progression. In order to identify potential novel regulators of Pim-3, Zhang and colleagues performed a yeast two-hybrid screen using a human HeLa matchmaker cDNA library and revealed that translationally controlled tumor protein (TCTP/TPT1) specifically interacts with and enhances Pim-3 protein stability by blocking its ubiquitin-proteasome-mediated degradation, thereby promoting tumor growth in vitro and in vivo. These findings recognize the TCTP/Pim-3 pathway as a new therapeutic target in human pancreatic cancer.

# miR-103 and miR-107 Inhibit Homologous Recombination

Huang et al. Page 1564

Error-free repair of DNA double-strand breaks by homologous recombination is essential for maintaining genomic stability and promoting resistance to DNA-damaging chemotherapies, but its regulation by microRNAs is poorly characterized. Huang and colleagues used a cell-based screening approach to identify miR-103 and miR-107 as potent inhibitors of homologous recombination. Mechanistically, miR-103 and miR-107 directly targeted RAD51 and RAD51D to mediate chemosensitivity to cisplatin and PARP inhibitors and are implicated in the endogenous regulation of RAD51D in several tumor subtypes. These findings not only demonstrate that miR-103 and miR-107 promote genomic instability and chemosensitivity but also show their utility as therapeutic agents for the chemosensitization of tumors.

#### Angiogenesis Genes Induce FGFmediated Resistance in HNSCC

Gyanchandani et al. Page 1585

Increased resistance to antiangiogenic therapies underscores the need for relevant preclinical models to gain insight into the mechanisms that drive refractoriness. Gyanchandani and colleagues developed a novel head and neck squamous cell carcinoma (HNSCC) xenograft model of acquired resistance to the VEGF inhibitor bevacizumab and uncovered an FGF signaling network in resistant tumors. Increased expression of FGF2 was regulated by overexpression of angiogenesis-related genes, including PLCG2, FZD4, CX3CL1, and CCL5, via increased ERK signaling. Inhibition of FGFR in resistant tumors led to restoration of sensitivity to bevacizumab, suggesting that cotargeting of VEGF and FGF overcomes resistance to antiangiogenic therapy in HNSCC.



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