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Tumor Biology

Abstract 1163: Microenvironment targets in KRAS-mutated metastatic colorectal cancer

Serena Marchio, Alice Bartolini, Sabrina Cardaci, Marco Soster, Giorgio Corti, Simona Lamba, Federico Bussolino, Davide Cora', and Federica D. Nicolantonio

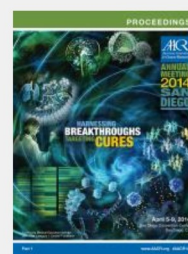
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Abstract

The introduction of biodrugs, e.g. the anti-EGFR antibodies, was initially seen as a promise to radically change the landscape for patients with metastatic colorectal cancer. However, although EGFR-targeted therapies, combined with chemotherapy, have prolonged survival expectancy to 24 months, cure remains anecdotal. Target therapies suffer of high costs, important side effects, and low response rates: it is now clear that this approach as it was originally conceived failed to meet the majority of its expectations. Importantly, because of novel prescription guidelines, patients with KRAS-mutated tumors are excluded from



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[Table of Contents](#)
[Index by Author](#)

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EGFR-targeted therapies: for these patients alternative treatments are urgently needed.

We here propose an innovative strategy based on the identification of molecular targets specifically associated with the microenvironment of metastatic colorectal cancer in patients carrying oncogenic KRAS. For this purpose, we set up mice models of metastatic colorectal cancer by intrasplenic (to evaluate liver homing) and intrahepatic (to investigate liver colonization) implantation of human colorectal cancer cell lines (SW48 and LIM1215) in which oncogenic KRAS (G12D, G12V, G13D) variants have been somatically knocked-in. We based our “target fishing” strategy on high-throughput, phage display-mediated proteomic screenings of deriving tumor samples *ex vivo*. Proteome signatures from the cognate cell lines *in vitro* served as a subtractive reference for the *ex vivo* biopanning, aimed at the identification of peptide ligands specific for non-epithelial components. By this combined biological-genetic-bioinformatics approach, we identified peptide/protein signatures selectively associated to metastasis microenvironments with controlled genetic backgrounds *in vitro* and *in vivo*.

We selected 2 target proteins and 2 targeting peptides, which were exploited for diagnostic and therapeutic purposes. First, we evaluated by IHC the presence and localization of the target proteins in samples (tumor, healthy liver) from a panel of human biopsies and from the described *in vivo* models. This analysis confirmed a specific enrichment in KRAS-mutated microenvironments. Second, we tested the capability of dye-conjugated targeting peptides to home to these same tumor microenvironments, observing a specific accumulation in target tissues, compared to their scrambled versions and to control tissues. Third, we investigated a potential anti-metastatic effect of these peptides when orthotopically co-injected with colorectal cancer cell lines; preliminary experiments revealed that

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the targeting peptides, but not the scrambled variants, inhibit the onset of liver metastases from KRAS-mutated cell lines.

In summary, we obtained proof-of-concept results in preclinical studies and produced prototype compounds to provide innovative tools that can be translated into the clinical practice with a mid-term perspective.

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