

PROPOSTA PROGETTUALE
DOTTORATO IN RNA THERAPEUTICS AND GENE THERAPY
(Mechanisms of diseases and Drug Target identification)
CICLO XLI

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PROJECT TITLE

Bile Acid-CoA:Amino Acid N-Acyltransferase as a Novel Target for RNA-Based Therapies in Colon Carcinoma

Project description (max 300 words)

Colorectal cancer (CRC) is the second leading cause of cancer-related deaths worldwide. Currently, the 5-year survival rate for patients in advanced stages remains only 14%¹. This poor prognosis is largely attributed to the emergence of an immunosuppressive tumor microenvironment (TME), which can severely limit the efficacy of advanced immunotherapies, especially immune checkpoint inhibitors². Preclinical in vitro and in vivo studies have shown that polyunsaturated fatty acids (PUFAs) are abundant in the TME and that their dysregulation contributes to cancer development and progression³. N-acyl taurines (NATs) are a new class of endogenous PUFAs, composed of long-chain fatty acids conjugated to the amino acid taurine. In humans, NATs are synthesized in peroxisomes by the enzyme bile acid-CoA:amino acid N-acyltransferase (BAAT)⁴.

Based on these premises, this project aims to provide unprecedented insights into the role of the enzyme BAAT enzyme and NATs in CRC development and progression, with a particular focus on tumor-stroma interactions. To achieve this goal, a multidisciplinary approach will be employed, combining multi-omics analyses and functional assays across a range of experimental models: human colon biopsies, patient-derived organoids (PDOs), genetically modified mouse models of CRC with high endogenous NAT levels, and 3D co-culture systems (PDOs/immune cells) to better mimic the complexity of the TME.

The findings of this project are expected to deepen the understanding of a novel molecular pathway involved in CRC and may pave the way for the identification of new therapeutic targets or strategies aimed at modulating the tumor microenvironment.

REFERENCES

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FUNDS

PRIN ,JAZZ PHARMACEUTICALS