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What could microRNA expression tell us more about colorectal serrated pathway carcinogenesis?

Milena Peruhova, Monika Peshevsk-Sekulovska, Boris Krastev, Gabriela Panayotova, Viktoriya Georgieva, Rossitza Konakchieva, Georgi Nikolaev, Tsvetelina Veselinova Velikova

Abstract

In the last two decades, the vision of a unique carcinogenesis model for colorectal carcinoma (CRC) has completely changed. In addition to the adenoma to carcinoma transition, colorectal carcinogenesis can also occur *via* the serrated pathway. Small non-

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Introduction. MiRNAs are small, non-protein-coding RNA molecules that regulate gene **expression** either by post-transcriptionally suppressing mRNA translation or by causing mRNA degradation 1–6. We know that miRNAs play a critical role in regulation of proliferation, differentiation, apoptosis, and stress response and are involved in the majority of physiological processes 7, 8.

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BACKGROUND. **MicroRNA** (miR)-320a, miR-145, and miR-192 have been shown to play a role in **colorectal carcinogenesis** and metastasis. We examined if there is a difference in **expression** during the histologic progression from normal mucosa (NM) to high-grade dysplastic adenomas (HG).

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