

Background:

In the past few years, it has been observed, that the number of registered neuroendocrine tumors (NET) considerably increased (from 3,9 (2001) to 7,9 (2012) / 100000 people) (1,2,3). We have focused on NETs of the pancreas (pNETs). The therapy of pNETs is principally influenced by their histological grade, which can change through the progression of the tumor. Currently, to determine the grade, it is inevitable to use invasive methods, such as biopsy and surgery, and it requires an experienced pathologist. According to the WHO classification pNETs are sorted into 3 grades based on their KI-67 index, mitotic index, and differentiation. MicroRNAs are short, stable, non-coding RNAs that play a role in the posttranscriptional regulation of gene expression. Their expression is tissue- and tumor-specific and may be characteristic of the grade. MicroRNAs are actively secreted by cells into their environment, thus can be detected in body fluids as well. In this study we selected 4 microRNAs based on literature data (4,5), and measured their expression in FFPE (formalin-fixed paraffin-embedded) blocks of G1 and G2 samples.

Aims:

Our aim was to find miRNAs that show significant differences in expression according to grade, thus could be suitable for grade determination. The significant miRNAs could be used to complement the histological approach.

Selected microRNAs for this study:

Based on literature search we have selected four miRNAs (hsa-miR-96-5p, hsa-miR-130b-3p, hsa-miR-106b and hsa-miR-194-5p). These miRNAs, were examined in pNETs from FFPE samples. The expression of hsa-miR-106b was shown to have prognostic relevance. Its expression is was found to be upregulated in G3 tumors compared to G1 and G2 tumors. (4,6) The hsa-miR-96-5p is an oncogenic microRNA. Its expression increased with PNET grade and in other tumors it is associated with tumor progression. It inhibits FOXO1 and activates AKT/GSK-3 β / β -catenin signaling pathway. (4,5) The expression of hsa-miR-130b-3p was found to increase with the grade, whereas the expression of hsa-miR-194-5p decreased with grade. For these miRNAs no specific function in PNET is determined yet. (4,5)

Materials and Methods:

We've collected 33 FFPE samples (16 G1, 17 G2) from the two Departments of Pathology of Semmelweis University. From 80-80 μ m sections of the FFPE blocks we isolated total RNA with Recover ALL Kit (Thermo Fisher Scientific). The sections were from tumor tissue only, marked by a pathologist. The expression of hsa-miR-96-5p, hsa-miR-130b-3p, hsa-miR-106b and hsa-miR-194-5p was determined by RT-qPCR, using TaqMan MicroRNA Assays. We calculated the Δ CT values using cel-miR-39 spike-in external control and RNU48 internal control, and analyzed them with unpaired t-test, using GraphPad Prism.

The miR-130b and miR-194 have shown significant difference in expression in grade 1 and grade 2 pNETs.

pNET, grade1, grade2, FFPE, miRNA, miR-130b, miR-194, miR-96, miR-106b

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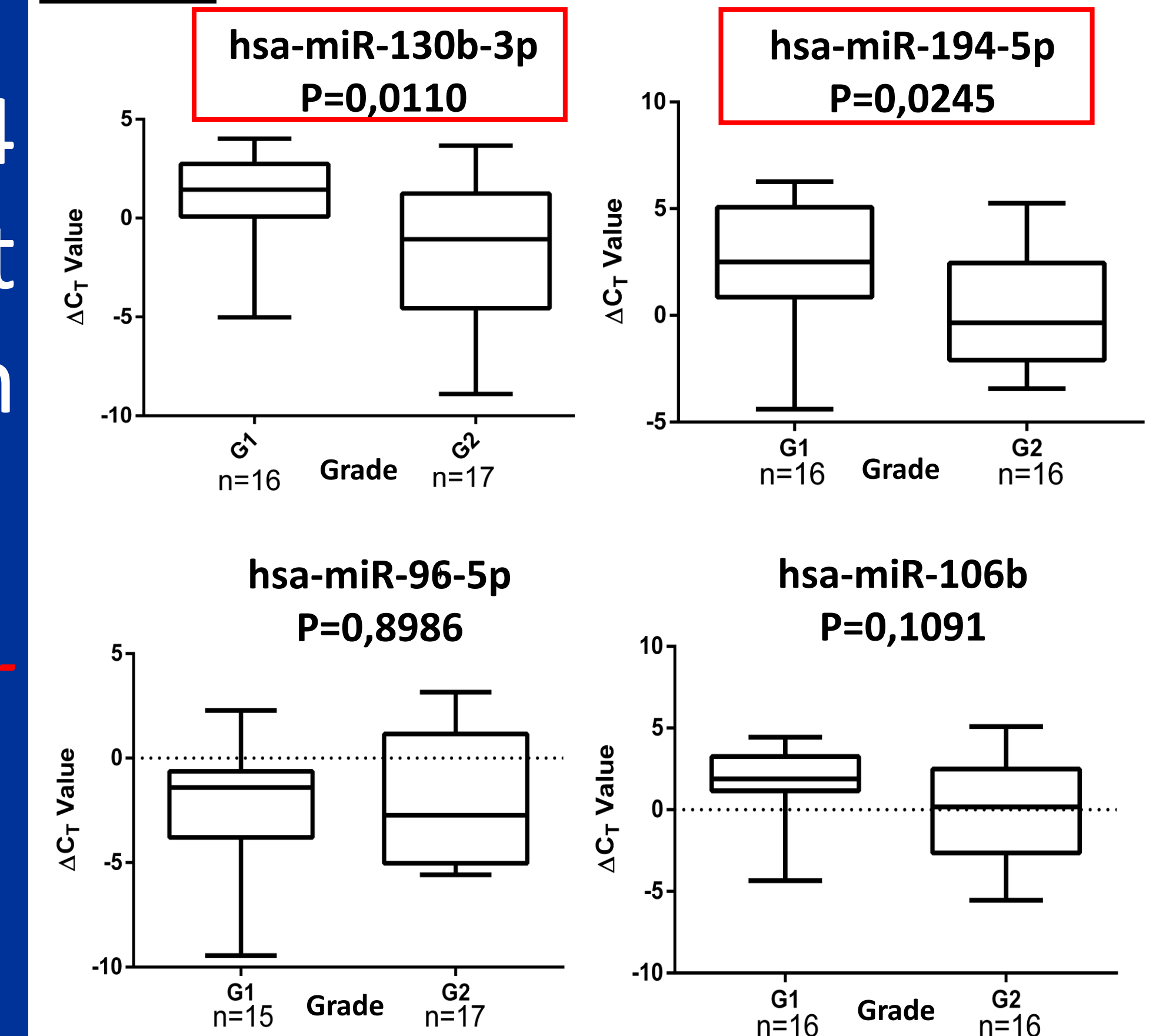
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Results:



Conclusions:

In conclusion, two of the four miRNAs (*hsa-miR-130b-3p*, *hsa-miR-194-5p*) have shown significant difference in expression in grade 1 and grade 2 pNETs. Both microRNAs showed lower expression in G2 compared to G1 pNETs.

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