MiRNA profile associated with invasiveness in non-functioning pituitary adenomas



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Background

Non-functioning pituitary adenomas (NFPAs) are not associated with hormonal hypersecretion and are among the most common pituitary tumours. In some patients tumour recurrence and invasion have been observed. So far, the diagnosis and monitoring of NFPAs remains difficult due to the lack of clinical biomarkers. In the present study, microRNAs (miRs) expression profiles were analysed in patients with invasive and non-invasive NFPAs and their biomarker potential was evaluated.

Methods

MiRs profiles were analysed in 12 patients with non-invasive and 8 patients with invasive NFPAs using miRCURY LNA miRNA Cancer Focus PCR Panel (Qiagen). ROC curve analysis evaluated the diagnostic ability of the selected miRNAs. P<0.05 was assumed for statistically significant and was calculated using un-paired. 2-tailed Student's t-test.

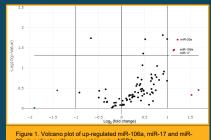






Results

Four miRs were found differentially expressed related to invasion and recurrence of NFPAs. MiR-106a, miR-17 and miR-20a were up-regulated in patients with recurrent invasive NFPAs (fig. 1), while miR-210-3p was down-regulated in invasive NFPAs compared to patients with noninvasive NFPAs (fig. 2).





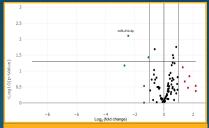
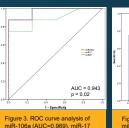


Figure 2. Volcano plot of down-regulated miR-210-3p in invasive NFPAs compared to non-invasive NFPAs

The average AUC was 0.943 (95% CI = 0.876-1) for miR-106a, miR-17 and miR-20a, and AUC = 0.915 (95% CI 0.702-1)as markers differentiating invasiveness.



(AUC=0.891) and miR-20a



Conclusion

The ability to predict tumour invasion and recurrence after the initial surgery will decrease morbidity and mortality rate in patients with NFPAs. The selected profile of miR-106a, miR-17, miR-20 and miR-210-3p showed biomarker potential. The next step is confirmation of the results in a larger patient cohort.

Funding

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