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Expression and predictive value of mir-489 and mir-21 in melanoma metastasis



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### Serum microRNA expression levels can predict lymph node ...

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3782554>

Serum microRNA **expression** levels can predict lymph node **metastasis** in patients with early-stage cervical squamous cell carcinoma JUNYING CHEN , 1, 2 DESHENG YAO , 1 YUE LI , 3 HONG CHEN , 2 CHANJUAN HE , 1 NAN DING , 1 YAN LU , 1 TINGYU OU , 1 SHAN ZHAO , ...

**Cited by:** 103

**Author:** Junying Chen, Desheng Yao, Yue Li, Ho...

**Publish Year:** 2013

### MicroRNA and mRNA expression profiling in metastatic ...

<https://onlinelibrary.wiley.com/doi/10.1111/pcmr.12343>

The role of microRNAs (miRNAs) in **melanoma** is unclear. We examined global miRNA **expression** profiles in fresh-frozen metastatic melanomas in relation to clinical outcome and BRAF mutation, with validation in independent cohorts of tumours and sera. We integrated miRNA and mRNA information from the same samples and elucidated networks associated with outcome and mutation.

**Cited by:** 30

**Author:** Varsha Tembe, Sarah-Jane Schramm, M...

**Publish Year:** 2015

### A 4-microRNA signature predicts lymph node metastasis and ...

<https://www.sciencedirect.com/science/article/pii/S004681771830087X>

Decreased miR-**489 expression** was associated with LNM, advanced pN stage and pTNM stage, and positive PR status. In the combined cohort, all the 4 miRNAs were associated with LNM status. High **expression** of miR-191-5p and low **expression** of miR-451a and miR-**489** were associated with larger tumor size and advanced pT stage, pN stage, and pTNM stage.

**Cited by:** 3

**Author:** Xu Chen, Ya-Wen Wang, Wen-Jie Zhu, ...

**Publish Year:** 2018

### MiR-221, a potential prognostic biomarker for recurrence ...

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5219708>

Background. Thyroid cancer is the most prevalent and rapidly increasing endocrine neoplasm. The number of new cases is almost 300,000 worldwide with a median age at diagnosis of 50 years and nearly 40,000 deaths per year [].The most common histological type of all thyroid cancers is papillary thyroid cancer (PTC), which is defined as a differentiated neoplasia and accounts for approximately 80%.

**Name of Journal:** *World Journal of Clinical Cases*

**Manuscript NO:** 50275

**Manuscript Type:** ORIGINAL ARTICLE

*Case Control Study*

Expression and predictive value of miR-489 and miR-21 in melanoma metastasis

Mo H *et al.* The role of miR-489 and miR-21 in metastasis of melanoma

Hao Mo, Jian Guan, Zhen-Chao Yuan, Xiang Lin, Zhen-Jie Wu, Bin Liu, Ju-Liang He

**Abstract**

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### microRNA-10b is a prognostic biomarker for melanoma ...

<https://www.nature.com/articles/modpathol2015149>

Melanoma is an aggressive **skin** cancer. Most tumors can be cured by local surgical excision but once spread to distant sites occurs the prognosis is poor. However, recent clinical trials have ...

### A 4-microRNA signature predicts lymph node metastasis and ...

<https://www.sciencedirect.com/science/article/pii/S004681771830087X>

Decreased miR-489 **expression** was associated with LNM, advanced pN stage and pTNM stage, and positive PR status. In the combined cohort, all the 4 miRNAs were associated with LNM status. High **expression** of miR-191-5p and low **expression** of miR-451a and miR-489 were associated with larger tumor size and advanced pT stage, pN stage, and pTNM stage.

Cited by: 3

Author: Xu Chen, Ya-Wen Wang, Wen-Jie Zhu, Yan L...

Publish Year: 2018

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Cited by: 6

Author: Lei Dai, Yaozong Wang, Liangliang Chen, J...

Publish Year: 2017

### Melanoma MicroRNA Signature Predicts Post-Recurrence ...

[clincancerres.aacrjournals.org/content/16/5/1577](http://clincancerres.aacrjournals.org/content/16/5/1577) ▼

Mar 01, 2010 · Purpose: To identify a melanoma microRNA (miRNA) **expression** signature that is **predictive** of outcome and then evaluate its potential to improve risk stratification when added to the standard-of-care staging criteria. Experimental Design: Total RNA was extracted from 59 formalin-fixed paraffin-embedded melanoma metastases and hybridized to miRNA arrays containing 911 probes.

Published in: Clinical Cancer Research · 2010

Authors: Miguel F Segura · Ilana Belitskayalevy · Amy E Rose · Jan Zakrzewski · Avital Gaziel · D...

About: microRNA · RNA interference · Gene silencing · Melanoma · Cancer · Gene **expression** ...

### Serum microRNA expression levels can predict lymph node ...

[https://www.academia.edu/20644665/Serum\\_microRNA...](https://www.academia.edu/20644665/Serum_microRNA...) ▼

After confirmation and in order to access the this manner, if the microRNA is a qualified LNM marker in