MICRORNA EXPRESSION AND ASSESSMENT OF POTENTIAL
ROLE OF miR-181a IN HEAD AND NECK CANCER

NURUL SYAKIMA AB MUTALIB

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MICRORNA EXPRESSION AND ASSESSMENT OF POTENTIAL ROLE OF miR-181a IN HEAD AND NECK CANCER

By

NURUL SYAKIMA AB MUTALIB

Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia, in Fulfillment of the Requirements for the Degree of Doctor of Philosophy

May 2012
Dedicated to my parents
Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfillment of the requirement for the degree of Doctor of Philosophy

MICRORN A EXPRESSION AND ASSESSMENT OF POTENTIAL ROLE OF miR-181a IN HEAD AND NECK CANCER

By

NURUL SYAKIMA BINTI AB MUTALIB

May 2012

Chair: Cheah Yoke Kqueen, PhD

Faculty: Medicine and Health Sciences

MicroRNAs (miRNAs) represent a class of small non-coding RNAs that regulate gene expression by either inducing mRNA degradation or repressing mRNA translation. The involvements of miRNAs in various human cancer-related processes have been studied in recent years. The first objective of this study was to determine differentially expressed miRNAs in head and neck cancer. Global miRNA profiling was performed on 12 tissue samples from various head and neck cancers by using the microarray approach followed by real time RT-PCR validation. The microarray analyses identified 10 miRNAs that were able to distinguish malignant from normal tissues whereby seven miRNAs (hsa-miR-181a-2*, hsa-miR-29b-1*, hsa-miR-181a, hsa-miR-181b, hsa-miR-744, hsa-miR-1271 and hsa-miR-221*) showed up-regulation while three miRNAs (hsa-miR-141, hsa-miR-95 and hsa-miR-101) showed down-regulation. Therefore, these miRNAs may aid in simple profiling strategies to identify individuals at higher risk of developing head and neck cancers,
as well as elucidate the molecular mechanisms involved in head and neck cancers pathogenesis.

The second objective of this study was to identify the putative targets of miRNAs differentially expressed in head and neck cancers and the pathways involved, which was achieved through in silico analysis aided by online databases, whereby several cancer-associated genes and pathways were found to be targeted by miR-181a. The role of miR-181a in head and neck carcinogenesis was subsequently determined through functional analyses as the third objective of this study. It was found out that miR-181a regulates the proliferation, migration, invasion and colony-forming ability of head and neck cancer cell.

Fourth objective was achieved by using pathway analysis to profile changes in the activities of 10 signaling pathways related to cancer caused by miR-181a down-regulation. Six of these pathways, namely the p53/DNA damage, TGFβ, MAPK/ERK, MAPK/JNK, Wnt and NFκB pathways, were found to be significantly influenced, suggesting miR-181a may act as an oncomiR, and therefore its inhibition may be a potential therapeutic target for head and neck cancer patients. The fifth and final objective of this study involved visualizing miR-181a expression and localization in head and neck tissues, for which in situ hybridization was utilized. miR-181a is preferentially expressed in the cytoplasm of cancer cells, and its expression is significantly increased in malignant compared to benign tumors of the head and neck. Collectively, these findings provide basis for study into the role of miR-181a as a biomarker and/or therapeutic target in head and neck tumors.
Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

EKSPRESI MIKORRNA DAN PENILAIAN PERANAN POTENSI miR-181a DALAM KANSER KEPALA DAN LEHER

Oleh

NURUL SYAKIMA BINTI AB MUTALIB
Mei 2012

Pengerusi: Cheah Yoke Kqueen, PhD
Fakulti: Perubatan dan Sains Kesihatan

MikroRNA (miRNA) mewakili kelas RNA bukan pengekodan kecil yang mengawal selia gen sama ada secara menginduksi degradasi mRNA atau represi translasi. Penglibatan mereka dalam proses yang berkaitan dengan kanser telah dikaji dalam pelbagai jenis kanser manusia. Objektif yang hendak dicapai dalam kajian ini adalah pertamanya untuk menentukan pembezaan ekspresi miRNA dalam kanser kepala dan leher. miRNA profil global telah dilakukan ke atas 12 tisu sampel kanser kepala dan leher menggunakan pendekatan mikroarray diikuti oleh pengesahan menggunakan qRT-PCR. Analisis mikroarray mengenal pasti 10 miRNA yang boleh membezakan lesi kanser kepala dan leher dari tisu normal; 7 miRNA (hsa-miR-181a-2 *, hsa-miR-29b-1*, hsa-miR-181a, hsa-miR-181b, hsa-miR-744, hsa-miR-1271 dan hsa-miR-221*) dikawal selia naik manakala 3 miRNA (hsa-miR-141, hsa-miR-95 dan hsa-miR-101) dikawal selia turun. Kumpulan miRNA ini boleh menyumbang dalam strategi pemprofilan yang mudah untuk membantu dalam mengenal pasti individu berisiko tinggi untuk mendapat kanser kepala dan leher serta dapat membantu dalam
mengenal pasti mekanisma molekular yang terlibat dalam patogenesis kanser kepala dan leher.

Kemudian, sasaran diduga miRNA yang diekspresi secara berbeza dalam kanser kepala dan leher serta laluan kanser yang terlibat telah dikenal pasti melalui analisis silico menggunakan pangkalan data dalam talian. Beberapa gen dan laluan berkaitan dengan kanser didapat disasarkan oleh kumpulan miRNA tersebut. Fungsi miRNA yang dipilih (miR-181a) dalam karsinogenesis kepala dan leher ditentukan melalui analisis fungsi. Kajian ini mendapati bahawa miR-181a mengawal proliferasi, migrasi, pencerobohan dan keupayaan membentuk koloni dalam kanser kepala dan leher.

Analisis laluan dilakukan untuk memprofil perubahan dalam aktiviti 10 laluan isyarat kanser yang disebabkan oleh penurunan paras miR-181a. Laluan p53/kerosakan DNA, TGFβ, MAPK / ERK, MAPK / JNK, Wnt dan NFκB didapati banyak dipengaruhi oleh penurunan paras miR-181a. Keputusan ini mencadangkan bahawa miR-181a boleh dicalonkan sebagai oncomiR dan seterusnya boleh dijadikan sebagai sasaran potensi terapeutik untuk pesakit kanser kepala dan leher. Akhir sekali, untuk menggambarkan ekspresi miR-181a dan penyetempatan di dalam tisu kepala dan leher, hibridisasi in situ telah digunakan. miR-181a terzahir dalam sitoplasma sel-sel kanser, dan ekspresinya meningkat dengan ketara dalam tumor malignan berbanding dengan tumor benigna kepala dan leher. Secara kolektif, penemuan ini menyediakan asas kepada kajian terhadap peranan miR-181a sebagai biopenanda dan / atau sasaran terapeutik tumor di kepala dan leher.
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I certify that a Thesis Examination Committee has met on 7 May 2012 to conduct the final examination of Nurul Syakima binti Ab Mutalib on her thesis entitled “MicroRNA Expression and Assessment of Potential Role of miR-181a in Head and Neck Cancer” in accordance with the Universities and University Colleges Act 1971 and the Constitution of the Universiti Putra Malaysia [P.U.(A) 106] 15 March 1998. The Committee recommends that the student be awarded the Doctor of Philosophy.

Members of the Thesis Examination Committee were as follows:

**Sabrina binti Sukardi, PhD**  
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Chairman)

**Noorjahan Banu binti Mohammed Alitheen, PhD**  
Associate Professor  
Faculty of Biotechnology and Biomolecular Sciences  
Universiti Putra Malaysia  
(Internal Examiner)

**Tan Soon Guan, PhD**  
Professor  
Faculty of Biotechnology and Biomolecular Sciences  
Universiti Putra Malaysia  
(Internal Examiner)

**Chin-Yuan Tzen, PhD**  
Professor  
Fu Jen Catholic University  
Taiwan  
(External Examiner)

---

**SEOW HENG FONG, PhD**  
Professor and Deputy Dean  
School of Graduate Studies  
Universiti Putra Malaysia

Date: 23 July 2012
The thesis was submitted to the Senate of Universiti Putra Malaysia and has been accepted as fulfillment of the requirement for the degree of Doctor of Philosophy. The members of the Supervisory Committee were as follows:

**Cheah Yoke Kqueen, PhD**  
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Chairman)

**Shiran Mohd Sidik, MBBS, MPath**  
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Member)

**Sabariah Abdul Rahman, MBBS, MPath**  
Professor  
Faculty of Medicine  
Universiti Institute Teknologi Mara  
(Member)

**Avatar Singh Mohan Singh, MBBS, FAGE, MS (ORL-UKM)**  
Dr.  
Head of ENT Head-Neck Surgery Unit  
Taiping General Hospital  
(Member)

---

**BUJANG BIN KIM HUAT, PhD**  
Professor and Dean  
School of Graduate Studies  
Universiti Putra Malaysia  

Date:
DECLARATION

I declare that the thesis is my original work except for quotations and citations which have been duly acknowledged. I also declare that it has not been previously, and is not concurrently, submitted for any other degree at Universiti Putra Malaysia or at any other institution.

__________________________

NURUL SYAKIMA BINTI AB MUTALIB

Date: 7 May 2012
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