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Expression of microRNA-96 and its relationship to matrix metalloproteinase regulator- RECK in Retinoblastoma

Thesis

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الخلوى RECK

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توطئة للحصول على درجة الماجستير

في الكيمياء الحيوية الطبية والبيولوجيا الجزيئية

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Abstract

Background: Retinoblastoma is the most common malignancy that develops in children's eyes. MiR- 96 was identified as an oncogenic micro-RNA. The RECK protein downregulates matrix metalloproteinases (MMP-2, MMP-9 and MT1-MMP) and suppresses tumor invasion, angiogenesis and metastasis. RECK mRNA was reported as a target of miR-96 in many cancers. This study aimed at evaluating the expression of miRNA-96 and the level of the matrix metalloproteinase regulator- RECK in 14 patients newly diagnosed with retinoblastoma before and after treatment, and in 14 controls, in order to give insights on their possible role as potential diagnostic and prognostic biomarkers in retinoblastoma.

Methods: MiR-96 and RECK protein were evaluated using sera of three groups of samples, retinoblastoma group before treatment, controls, and retinoblastoma group after treatment by quantitative reverse transcription real time polymerase chain reaction (qRT-PCR) and Enzyme Linked Immunosorbent Assay (ELISA), respectively.

Results: The expression of miR- 96 was significantly higher in the sera of patients than the control group ($P<0.05$). The concentration of RECK protein was significantly decreased in the patients compared to control group ($P<0.05$). Furthermore, the level of miR- 96 was significantly decreased ($P<0.05$), and concentration of RECK protein was significantly increased ($P<0.05$) in the sera after treatment in patients who responded to globe salvage treatment. Moreover the expression of RECK protein, and miR- 96 was negatively correlated with each other in the patients before treatment as well as after treatment and in the control group with r (-0.809), r (-0.926), r (-0.615), respectively.

Conclusion: Serum level of miR-96 and RECK protein concentration may serve as potential biomarkers in retinoblastoma.

KEYWORDS:

Retinoblastoma; MicroRNA; MiR-96; RECK.

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List of abbreviations

Ago	Argonaute protein
AML.....	Acute myeloid leukemia
ANT2.....	Adenine nucleotide translocase 2
ARE.....	AU-rich element
AUC	Area under the curve
c-Myc.....	Cellular Myelocytomatosis
Ct.....	Cycle threshold
CDK.....	Cyclin dependent kinase
CDKN1A.....	Cyclin dependent kinase inhibitor 1A
CNS.....	Central nervous system
CT.....	Computed tomography
DD.....	Disc diameters
DDR.....	DNA damage response
DEPC.....	Diethylpyrocarbonate
DNA.....	Deoxyribonucleic acid
ECM.....	Extra-cellular matrix
EGF	Epidermal growth factor
EGFR.....	Epidermal growth factor receptor
EGCG	Epigallocatechin 3 gallate
EMQN.....	European molecular quality network
FAK.....	Focal adhesion kinase
FOXO1.....	Forkhead box protein O1

FOXO3a	Forkhead box protein O3a
FXR1	Fragile-X-mental-retardation-related protein 1
HAT	Histone acetyltransferases
HbF	Fetal hemoglobin
HDAC1	Histone deacetylase 1
HCC	Hepatocellular carcinoma
HIF-1alpha	Hypoxia-inducible factor 1-alpha
HIF-2α	Hypoxia-Inducible Factor-2-alpha
HPV	Human papilloma virus
ICRB	International classification of retinoblastoma
ISH	In situ hybridization
JNK	c-Jun N-terminal kinase
Kda	Kilo Dalton
m7G	7-methylguanosine-cap
miR	Micro ribonucleic acid
miRNA	Micro ribonucleic acid
μg	Micro gram
mL	Mille liter
MMP	Matrix metalloproteinases
MRI	Magnetic resonance imaging
mRNA	Massager RNA
MTSS1	Metastasis suppressor-1
ng	Nano gram

nm..... Nano meter
NSCLCNon-small cell lung cancer
OD.....Optical density
ORFOpen reading frame
OS.....Overall survival
PACTProtein ACTivator of the
interferon- induced protein kinase
PFS.....Progression-free survival
pRb.....RB1 protein
pri-miRNA.....primary micro RNA
qRT-PCRQuantitative reverse transcription
Polymerase Chain Reaction
RB1.....Retinoblastoma gene
REReese-Ellsworth classification
RFS.....Relapse-free survival
RECK.....Reversion inducing cysteine rich
protein with kazal motifs
RIM.....RECK Intron-1 Methylation
RISC.....RNA induced silencing complex
RNase.....Ribonuclease
ROC.....Receiver-Operating Characteristic

RPM.....RECK Promoter/exon-1 Methylation
SACCSalivary adenoid cystic carcinoma
SAMD9.....Sterile alpha motif domain-