

MiR-214-3p Regulates Apoptosis of Ameloblasts under Excessive Fluoride via PI3K/AKT Signaling Pathway



Xia Liu^{1,2,#}, Nanqing Fu^{1,2,#}, Bin Chen¹, Yueyue Wang^{1,2}, Xin Shu¹, Jianguo Liu¹, Guohui Bai^{1,2,*} and Yuan Tian^{1,2,#}

¹Key Laboratory of Oral Disease Research, School of Stomatology, Zunyi Medical University, Zunyi 563000, China

²Hospital of Stomatology, Zunyi Medical University, Zunyi 563000, China

© 2024 The Author(s). Published by Bentham Open.

This is an open access article distributed under the terms of the Creative Commons Attribution 4.0 International Public License (CC-BY 4.0), a copy of which is available at: <https://creativecommons.org/licenses/by/4.0/legalcode>. This license permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.



*Address correspondence to this author at the Key Laboratory of Oral Disease Research, School of Stomatology, Zunyi Medical University, Zunyi 563000, China; E-mail: baiguohui1228@126.com

#These authors contributed equally to this work

Published: March 21, 2024

Cite as: Liu X, Fu N, Chen B, Wang Y, Shu X, Liu J, Bai G, Tian Y. MiR-214-3p Regulates Apoptosis of Ameloblasts under Excessive Fluoride via PI3K/AKT Signaling Pathway. Open Dent J, 2024; 18: e18742106290670.

<http://dx.doi.org/10.2174/0118742106290670240314063143>



Send Orders for Reprints to reprints@benthamscience.net

Table S1. Cell viability after treatment with different concentrations of NaF (%).

Time	Blank Group	0.4 mmol/L	0.8 mmol/L	1.6 mmol/L	3.2 mmol/L	6.4 mmol/L
24 h	100	91.60 ± 0.33	88.26 ± 1.01	57.97 ± 0.63	39.02 ± 0.36	21.56 ± 1.15
48 h	100	87.73 ± 0.92	76.16 ± 0.57	37.61 ± 0.35	26.67 ± 0.59	12.62 ± 0.33
72 h	100	79.11 ± 0.73	56.21 ± 0.94	28.56 ± 0.42	12.44 ± 1.06	3.32 ± 0.35

Table S2. Influences of different concentrations of NaF on apoptotic proteins.

Apoptosis-related Proteins	Blank Group	0.4 mmol/L	0.8 mmol/L	1.6 mmol/L	3.2 mmol/L	6.4 mmol/L
Caspase 3	0.42 ± 0.04	0.45 ± 0.04	0.51 ± 0.02	0.62 ± 0.04	0.73 ± 0.09	0.85 ± 0.10
Bax	3.74 ± 0.45	4.21 ± 0.29	5.37 ± 0.40	6.16 ± 0.21	6.40 ± 0.38	7.80 ± 0.24
Bcl-2	2.83 ± 0.09	2.46 ± 0.13	2.55 ± 0.15	2.52 ± 0.27	2.14 ± 0.21	2.06 ± 0.14
Bax/Bcl-2	1.32 ± 0.15	1.71 ± 0.11	2.10 ± 0.11	2.46 ± 0.24	3.00 ± 0.15	3.83 ± 0.33

Table S3. Statistical result of sequencing data.

Sample	Raw Reads	Clean Reads	Percentage of Clean Reads (%)
C-1	24,417,752	20,759,412	85.02
C-2	25,143,022	21,284,681	84.65
C-3	25,152,901	21,394,616	85.06
T1-1	24,202,232	21,389,076	88.38
T1-2	23,614,695	21,754,713	92.12
T1-3	24,997,385	21,949,984	87.81
T2-1	24,025,453	21,058,801	87.65

Sample	Raw Reads	Clean Reads	Percentage of Clean Reads (%)
T2-2	24,335,049	21,641,727	88.93
T2-3	23,090,840	22,090,808	95.67

Note: C (1-3) are three replicates of the control group, T1 (1-3) are three replicates of the 1.6 mmol/L NaF group and T2 (1-3) are three replicates of the 3.2 mmol/L NaF group.

Table S4. Statistical result of miRNA relative expression levels.

miRNAs	Blank Group	1.6 mmol/L	3.2 mmol/L
miR-3110-5p	1.06 ± 0.02	2.35 ± 0.09	3.06 ± 0.18
miR-324-5p	1.02 ± 0.03	2.63 ± 0.32	2.61 ± 0.25
miR-21a-3p	1.01 ± 0.01	1.65 ± 0.28	1.60 ± 0.28
miR-214-3p	1.01 ± 0.01	0.19 ± 0.07	0.36 ± 0.07
miR-210-5p	1.02 ± 0.02	0.16 ± 0.08	0.26 ± 0.06
miR-574-5p	1.01 ± 0.01	0.09 ± 0.04	0.15 ± 0.06

Table S5. Statistical result of relative expression of pathway proteins.

Proteins	Blank Group	NaF Group	NaF+miR-214-3p Mimic Group	NaF+miR-214-3p Mimic+LY294002 Group
PI3K	0.29±0.02	0.44±0.10	0.13±0.05	0.31±0.08
p-PI3K	0.45±0.07	0.25±0.06	0.40±0.03	0.18±0.07
AKT	0.31±0.03	0.65±0.08	0.37±0.05	0.52±0.07
P-AKT	0.44±0.09	0.25±0.06	0.38±0.06	0.25±0.05
Bcl-2	2.51±0.14	1.99±0.20	2.51±0.25	1.51±0.25
Bax	1.44±0.24	2.43±0.21	1.56±0.17	2.14±0.20