

Correction

Correction: Kędzierska, H., et al. Decreased Expression of SRSF2 Splicing Factor Inhibits Apoptotic Pathways in Renal Cancer. *Int. J. Mol. Sci.* **2016, 17, 1598**

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check for updates

The authors wish to make the following corrections to this paper [1]: in Figure 4 the same gel scans were mistakenly pasted to illustrate splicing changes of: i) BIM in KIJ-265T and KIJ308T cells, and ii) MCL-1 in UOK171 and KIJ-265T. In order to correct this mistake, we have provided the updated Figure 4 with the correct gel scans. Therefore, please replace the old Figure 4

		Caki-2		UOK171		KIJ-265T		KIJ-308T		
		siSRSF2	siControl	siSRSF2	siControl	siSRSF2	siControl	siSRSF2	siControl	
Caspase-8L antiapoptotic	~		and the second second	1000			1000	Arra book Arra		_386 bp
Caspase-8a proapoptotic	-									∼250 bp
Caspase-9a proapoptotic	-									742 bp
Caspase-9b antiapoptotic	/									∼292 bp
CFLAR NEW	~	Name and Address of the Owner,	and the local		No. of Concession, Name	-	100 C 100 C	-	And And And	_419 bp
CFLAR var: 1,2,4,5,6	1							-		⊃308 bp
DIABLO var ' proapoptotic	1	the loss limit	and the last	state later with	them in the surf	Intel State Party	state local state	124 1.14 200	And in case	535 bo
Smac3 proapoptotic	1						===			~403 bp
Surv-2B proapoptotic Survivin antiapoptotic	11						-			_440 bp
Surv-deltaEx3 antiapoptotic	3/	All and all a second	NAME AND ADDRESS.		And And Street					*255 bp
TRAIL a proapoptotic	~									_371 bp
non-active	_									*327 bp
BimEL + Bimα1	_									_∕656 bp
BimL + Bimα2 BimS + Bimα3	1		===		===		===	===	===	386 bp
										000.00
MCL-1L antiapoptotic	/									- 885 bp
MCL-1S proapoptotic	/			and set		tor and soal		and star the		- 637 bp
HPRT1	_									- 180 bp

Figure 4. The effect of SRSF2 silencing on splicing patterns of apoptotic genes. Electrophoretic analysis of PCR-amplified splicing variants of apoptotic genes in four renal cancer-derived cell lines transfected



with SRSF2-specific (siSRSF2) or control (siControl) siRNA. CFLAR NEW designates a new CFLAR splice variant, identified in this study. Primers used for amplification of BIM isoforms detected three major variants (BimEL, BimL, and BimS), as well as minor variants (Bim α 1, Bim α 2, and Bim α 3). HPRT1—Internal RT-PCR control.

with the new Figure 4.



Figure 4. The effect of SRSF2 silencing on splicing patterns of apoptotic genes. Electrophoretic analysis of PCR-amplified splicing variants of apoptotic genes in four renal cancer-derived cell lines transfected with SRSF2-specific (siSRSF2) or control (siControl) siRNA. CFLAR NEW designates a new CFLAR splice variant, identified in this study. Primers used for amplification of BIM isoforms detected three major variants (BimEL, BimL, and BimS), as well as minor variants (Bim α 1, Bim α 2, and Bim α 3). HPRT1—Internal RT-PCR control.

The article conclusions and findings reported are not affected by this correction. All authors approve this correction. The authors would like to apologize for any inconvenience caused to the readers by these changes.

Conflicts of Interest: The authors declare no conflict of interest.

Reference

 Kędzierska, H.; Popławski, P.; Hoser, G.; Rybicka, B.; Rodzik, K.; Sokół, E.; Bogusławska, J.; Tański, Z.; Fogtman, A.; Koblowska, M.; et al. Decreased Expression of SRSF2 Splicing Factor Inhibits Apoptotic Pathways in Renal Cancer. *Int. J. Mol. Sci.* 2016, *17*, 1598. [CrossRef] [PubMed]



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