

ERRATUM

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# Erratum to: Next-generation sequencing reveals novel differentially regulated mRNAs, lncRNAs, miRNAs, sdRNAs and a piRNA in pancreatic cancer

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## Erratum

Due to a publisher error, this article [1] was originally published with an incorrect table 3. The correct table is provided here Table 1.

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## Reference

1. Müller S, Raulefs S, Bruns P, Grunz FA, Plötner A, Thermann R, et al. Kahl G1, Koch I, Theis FJ, Kleeff J, Winte P, Michalsk CW. Next-generation sequencing reveals novel differentially regulated mRNAs, lncRNAs, miRNAs, sdRNAs and a piRNA in pancreatic cancer. *Mol Cancer*. 2015;14:94.

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**Table 1 Differentially expressed lincRNAs**

lincRNA	NEV normal	NEV PDAC	Log2 FC	FDR	Class	Regulation	PED	other
LINC00671	81.13	0.47	7.45	0.0000000	lincRNA	Down		
DGCR5	83.72	2.89	4.86	0.0000000	lincRNA	Down		
LINC00261	278.16	7.04	5.30	0.0000000	lincRNA	Down		D [24]
TRHDE-AS1	24.54	1.05	4.54	0.0000015	NAT	Down		
PITPNA-AS1	94.22	16.50	2.51	0.0000085	NAT	Down		
SNHG8	820.02	190.34	2.11	0.0000189	lincRNA	Down	D	
SNHG9	595.61	149.36	2.00	0.0000217	lincRNA	Down		
LINC00152	27.79	134.36	-2.27	0.0000230	lincRNA	Up		U [24]
CEBPA-AS1	37.03	5.21	2.83	0.0000629	NAT	Down		
DANCR	357.77	90.05	1.99	0.0000661	lincRNA	Down		
SCARNA22	56.28	7.07	2.99	0.0001779	ncRNA	Down		
SCARNA2	27.72	4.21	2.72	0.0002533	ncRNA	Down		
HNF1A-AS1	72.29	10.99	2.72	0.0002943	NAT	Down		U [26]
SDCBP2-AS1	18.87	1.90	3.31	0.0002948	NAT	Down		
CECR5-AS1	23.83	3.16	2.91	0.0003537	NAT	Down		
HOTAIRM1	11.60	56.22	-2.28	0.0003752	NAT	Up	U	
LINC00339	355.98	10.67	5.06	0.0004037	lincRNA	Down		
TMEM44-AS1	36.53	7.99	2.19	0.0009294	NAT	Down		
LINC00263	44.69	10.62	2.07	0.0010579	lincRNA	Down		
MIAT	10.01	46.87	-2.23	0.00119361	lincRNA	Up	U	
NAPA-AS1	25.94	4.71	2.46	0.0012669	NAT	Down		
LINC00340	4.16	28.42	-2.77	0.0013636	lincRNA	Up		
EPB41L4A-AS1	303.58	97.66	1.64	0.0015839	NAT	Down		
SERTAD4-AS1	3.19	22.21	-2.80	0.0016944	NAT	Up		
LHFPL3-AS2	20.61	3.08	2.74	0.0023207	NAT	Down		
LINC00673	8.59	42.30	-2.30	0.0023447	lincRNA	Up		
DNAJC27-AS1	20.43	3.65	2.49	0.0030115	NAT	Down		
PVT1	6.87	43.83	-2.67	0.0056448	lincRNA	Up	U	
BOLA3-AS1	34.96	9.42	1.89	0.0074620	NAT	Down		
ZNF503-AS2	21.89	4.65	2.24	0.0079046	NAT	Down		
ACVR2B-AS1	13.06	1.46	3.16	0.0095731	NAT	Down		
PRRT3-AS1	28.36	7.04	2.01	0.0095911	NAT	Down		
LINC00346	1.41	13.14	-3.22	0.0121525	lincRNA	Up		
LINC00086	11.86	1.76	2.76	0.0132691	lincRNA	Down		
RUNX1-IT1	2.57	30.23	-3.56	0.0136106	ncRNA	Up		
THAP7-AS1	29.29	8.11	1.85	0.0198907	NAT	Down		
ZNF503-AS1	11.56	1.93	2.58	0.0240067	NAT	Down		
RNF157-AS1	10.82	1.91	2.50	0.0245309	NAT	Down		
AFAP1-AS1	17.34	153.08	-3.14	0.0265179	NAT	Up		U [25]
PSMG3-AS1	24.17	6.92	1.80	0.0291786	NAT	Down		

**Table 1 Differentially expressed lncRNAs (Continued)**

LINC00578	2.50	13.79	-2.46	0.0404404	lincRNA	Up	
H19	2,303.62	721.52	1.67	0.0452843	lincRNA	Down	D
GASS	964.77	471.70	1.03	0.0464071	lncRNA	Down	

For each differentially expressed lncRNA (FDR < 0.05) the mean normalized expression values (NEV) for each group is provided together with the ratio between normal and PDAC expression in log2 scale (log2 FC). Furthermore, "Class" indicates the lncRNA type, "PED" the expression as provided by the pancreatic cancer expression database (U: Upregulated in PDAC, D: Downregulated in PDAC) and the last column presents information about the expression level of the RNA in other types of cancer