

Targeting CDC7 sensitizes resistance melanoma cells to BRAF^{V600E}-specific inhibitor by blocking the CDC7/MCM2-7 pathway

Shaimaa A. Gad^{1,4}, Hamdy EA Ali¹, Rofaida Gaballa¹, Rania M. Abdelsalam², Mourad Zerfaoui³, Hamed I. Ali¹, Salwa H. Salama⁴, Sanaa A. Kenawy², Emad Kandil³, Zakaria Y. Abd Elmageed¹

¹Department of Pharmaceutical Sciences, Rangel College of Pharmacy, Texas A&M Health Science Center, Kingsville, TX 78363, ²Department of Pharmacology and Toxicology, Faculty of Pharmacy, Cairo University, ³Department of Surgery, Tulane University School of Medicine, New Orleans, LA 70118, ⁴Department of Pharmacology, Medical Division, National Research Center, Cairo.

Supplementary Information

Supplementary Table 1. Clinical information of melanoma patients used in the study

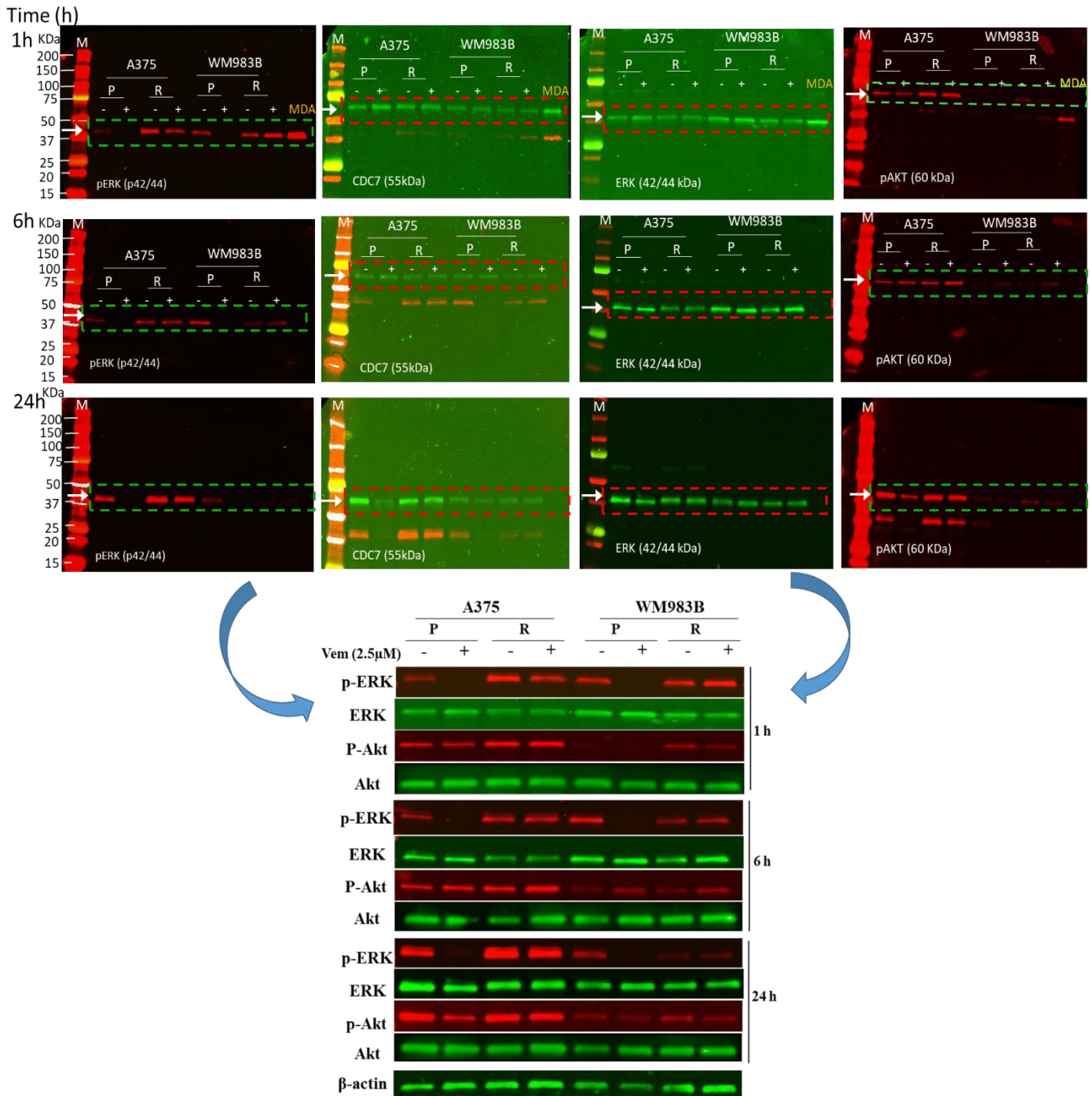
Parameter	Description	Mean (%)
Age (Mean±SD)		54.4 ± 14.2
Gender	Male	50 (55%)
	Female	40 (45%)
Organ	Skin	70 (77.7)
	Rectum	10 (11.1)
	Vulva	6 (6.6)
	Others	4 (4.4)
T-stage	T1+2	10 (11.1)
	T3+4	70 (77.7)
Pathological stage	I	6 (6.6)
	II	62 (68.9)
	III+IV	12 (13.3)

Supplementary Table 2. Correlation of CDC7 expression with clinical outcomes of melanoma patients

Parameter		Cyto.	Nucl.	Age	Gender	TNM	Stage
Cyto.	<i>r</i>	1.0000	0.0048	0.3195	0.2547	0.1463	0.2810
	<i>p</i>		0.9656	0.0034	0.0209	0.2200	0.0167
Nucl.	<i>r</i>		1.0000	-0.0623	-0.0764	0.1869	0.1016
	<i>p</i>			0.5777	0.4948	0.1157	0.0167
Age	<i>r</i>			1.0000	-0.0103	0.1498	0.0735
	<i>p</i>				0.9229	0.1845	0.5224
Gender	<i>r</i>				1.0000	-0.1744	0.2059
	<i>p</i>					0.1216	0.0704
TNM	<i>r</i>					1.0000	0.6348
	<i>p</i>						0.0004
Stage	<i>r</i>						1.0000
	<i>p</i>						

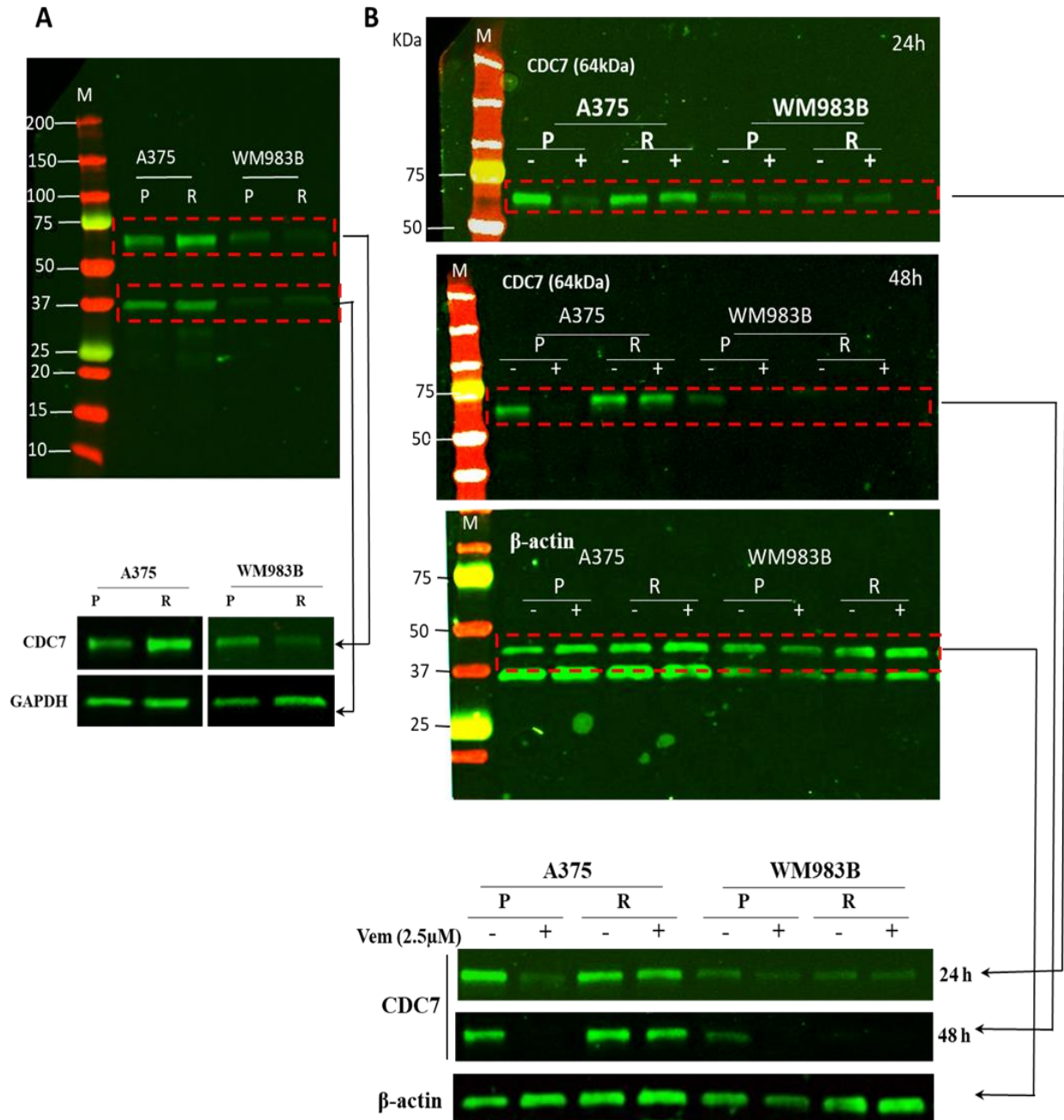
Correlation coefficient (*r*) was calculated among different groups and significant data considered at $p < 0.05$ (bold numbers).

Supplementary Figure 1. Representative photomicrograph showing the effect of Vemurafenib treatment on parental (P) and Vemurafenib resistant (R) melanoma cells. Total cell lysate collected at different time points (from 1-24h) of 2.5 μ M Vemurafenib treatment. Western blot analysis was performed and membranes were probed with the designated antibodies as shown. B-actin was used as a loading control. Protein signals were detected by Odyssey CLX Imaging System. M: protein marker

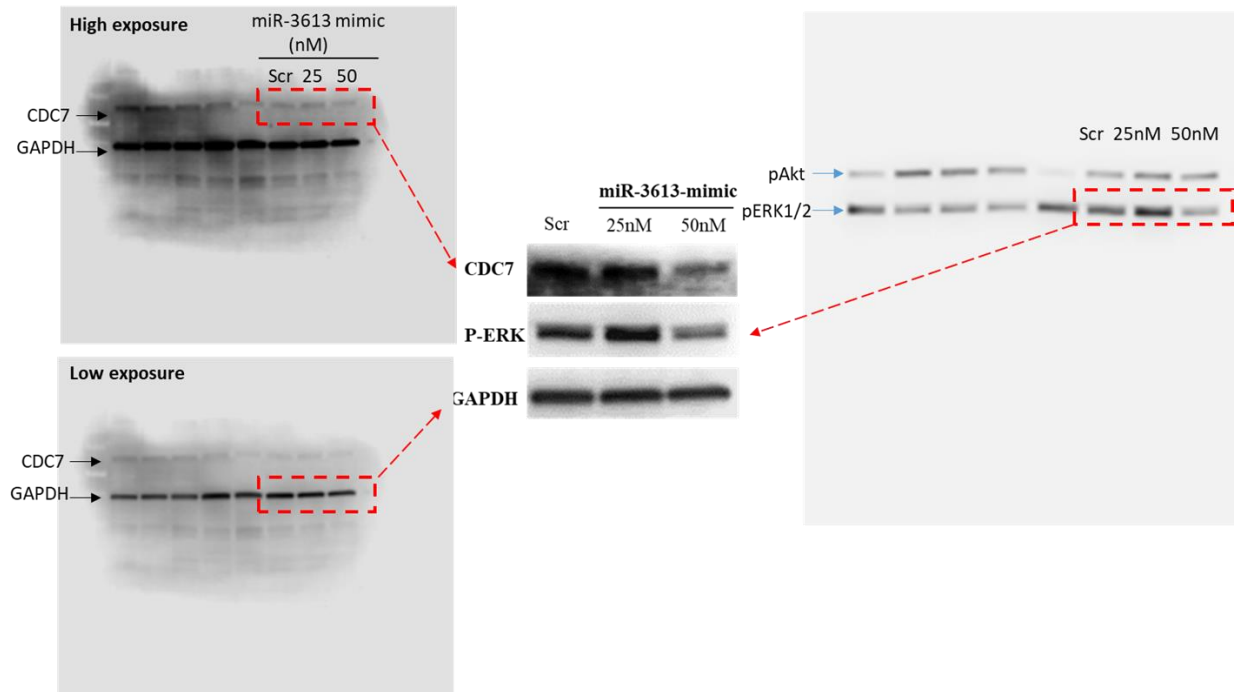


Supplementary Figure 2. Representative photomicrograph of endogenous expression of CDC7 in sensitive (P) and resistant (R) cells before and after treatment with Vemurafenib. Total cell lysate collected before (A) or after (B) treatment of melanoma cells with 2.5 μ Vemurafenib at different time points (24 & 48h). Western blot analysis was performed and membranes were probed with the designated antibodies. GAPDH or β -actin was used as loading control.

M: protein marker



Supplementary Figure 3. Representative photomicrograph showing targeting of CDC7 and phosphorylated ERK1/2 by miR-3613 mimic. Total cell lysate collected after transfection of A375 cells with 25 & 50nM miR-3613-3p mimic in addition control cells transfected with non-specific siRNA (scrambled). Western blot analysis was performed and membranes were probed with the designated antibodies. GAPDH was used as a loading control.



Supplementary Figure 4. Representative photomicrograph after treatment of A375-R cells with Vemurafenib or CDC7 inhibitor TAK-931. Total cell lysate collected after treatment of A375-R cells with either 2.5 μ Vemurafenib or 500nM or 1000nM TAK-931 at different time points. Western blot analysis was performed and membranes were probed with the designated antibodies.

