



Communication

The Effect of Silencing Fatty Acid Elongase 4 and 6 Genes on the Proliferation and Migration of Colorectal Cancer Cells

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Table S1. Relative expression level of selected fatty acid elongase in control conditions (basic culture medium), in the presence of positive siRNA control (siRNA GAPDH), negative siRNA control (siRNA(-)) and lipofectamine. Data presented as mean \pm SD. * $p < 0.05$ compared to control condition (base medium).

	ELOVL1	ELOVL2	ELOVL3	ELOVL4	ELOVL5	ELOVL6
CCD-841-CoN						
Control	1.01 \pm 0.09	1.03 \pm 0.15	1.13 \pm 0.05	1.02 \pm 0.12	1.05 \pm 0.19	1.02 \pm 0.11
siRNA(-)	1.00 \pm 0.04	1.15 \pm 0.18	1.06 \pm 0.15	1.22 \pm 0.13	0.99 \pm 0.11	0.92 \pm 0.05
siRNA GAPDH	0.97 \pm 0.13	1.00 \pm 0.19	1.05 \pm 0.05	1.08 \pm 0.14	1.08 \pm 0.07	1.10 \pm 0.11
Lipofectamine	1.04 \pm 0.09	1.07 \pm 0.17	1.06 \pm 0.13	1.03 \pm 0.05	0.99 \pm 0.10	0.97 \pm 0.08
HT-29						
Control	1.01 \pm 0.10	1.02 \pm 0.06	1.06 \pm 0.16	1.11 \pm 0.21	1.14 \pm 0.17	1.04 \pm 0.15
siRNA(-)	1.04 \pm 0.27	1.08 \pm 0.05	1.23 \pm 0.03	0.94 \pm 0.05	1.05 \pm 0.18	1.03 \pm 0.17
siRNA GAPDH	1.13 \pm 0.05	1.18 \pm 0.11	1.19 \pm 0.07	1.09 \pm 0.15	0.95 \pm 0.10	1.15 \pm 0.24
Lipofectamine	1.16 \pm 0.22	0.96 \pm 0.08	1.04 \pm 0.09	1.25 \pm 0.11	1.01 \pm 0.15	0.97 \pm 0.11
WiDr						
Control	1.01 \pm 0.04	1.02 \pm 0.03	1.01 \pm 0.02	1.01 \pm 0.08	1.00 \pm 0.04	1.00 \pm 0.10
siRNA(-)	1.12 \pm 0.12	0.99 \pm 0.22	0.95 \pm 0.11	1.10 \pm 0.12	0.98 \pm 0.06	1.16 \pm 0.06
siRNA GAPDH	1.13 \pm 0.03	1.14 \pm 0.06	1.06 \pm 0.07	1.13 \pm 0.06	1.15 \pm 0.12	1.11 \pm 0.06
Lipofectamine	0.99 \pm 0.01	1.12 \pm 0.17	1.20 \pm 0.14	1.17 \pm 0.11	1.04 \pm 0.14	0.98 \pm 0.03

Table S2. Cell viability (%) after 72h incubations with lipofectamine, negative control, and positive control. Data presented as mean \pm SD. * $p < 0.05$ compared to control condition (base medium).

Culture conditions	Cell line type		
	CCD-841-CoN	HT-29	WiDr
Control	100 \pm 4.58	100 \pm 8.87	100 \pm 8.25
Lipofectamine	98.69 \pm 7.45	95.05 \pm 8.86	99.15 \pm 4.45
siRNA(-)	98.46 \pm 7.84	92.14 \pm 1.31	99.75 \pm 2.95
siRNA GAPDH	42.20 \pm 7.82 *	27.25 \pm 7.11 *	23.54 \pm 3.64 *

Table S3. The migration rate [%] of CCD-841-CoN, HT-29, and WiDr at 0h, 24h, 48h and 72h incubation with negative control (siRNA(-)) or lipofectamine. * $p < 0.05$ compared to control condition (base medium).

	0h	24h	48h	72h
CCD-841-CoN				
Control	0.00	40.00 \pm 2.95	55.5 \pm 1.52	100.00 \pm 4.36
siRNA(-)	0.00	32.8 \pm 4.37	63.08 \pm 2.73	97.62 \pm 2.55
Lipofectamine	0.00	36.7 \pm 4.74	66.9 \pm 4.93	99.62 \pm 1.52
HT-29				
Control	0.00	40.02 \pm 2.36	63.00 \pm 4.76	100.00 \pm 3.65
siRNA(-)	0.00	44.3 \pm 7.81	66.1 \pm 2.91	97.4 \pm 5.58
Lipofectamine	0.00	46.84 \pm 2.45	62.62 \pm 4.84	96.39 \pm 4.32
WiDr				
Control	0.00	33.8 \pm 1.69	75.6 \pm 4.42	100.00 \pm 3.66
siRNA(-)	0.00	35.6 \pm 2.26	76.3 \pm 3.84	97.26 \pm 4.51
Lipofectamine	0.00	30.40 \pm 2.25	75.50 \pm 2.51	92.36 \pm 5.58