

RNA-seq transcriptome analysis of differentiated human oligodendrocytic MO3.13 cells shows up-regulation of genes involved in myogenesis

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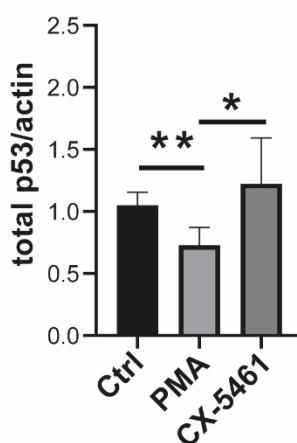


Figure S1. The level of total p53 in control (Ctrl), PMA or CX-5461 MO3.13 treated cells. Results are presented as a ratio of total p53 normalized to actin (loading control). Densitometric analysis of results (n=6) was performed with the use of one sample *t* test. Results are presented as means \pm standard error. The level of statistical significance is indicated using $*p \leq 0.05$, $**p \leq 0.01$.

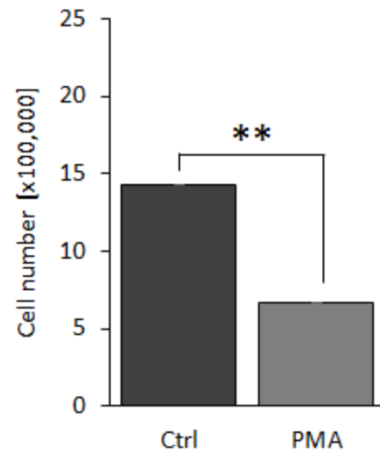


Figure S2. Analysis of cell proliferation. MO3.13 cells were cultured for 72 h in the absence (control; Ctrl) or presence of PMA (n=4), ** $p \leq 0.01$.

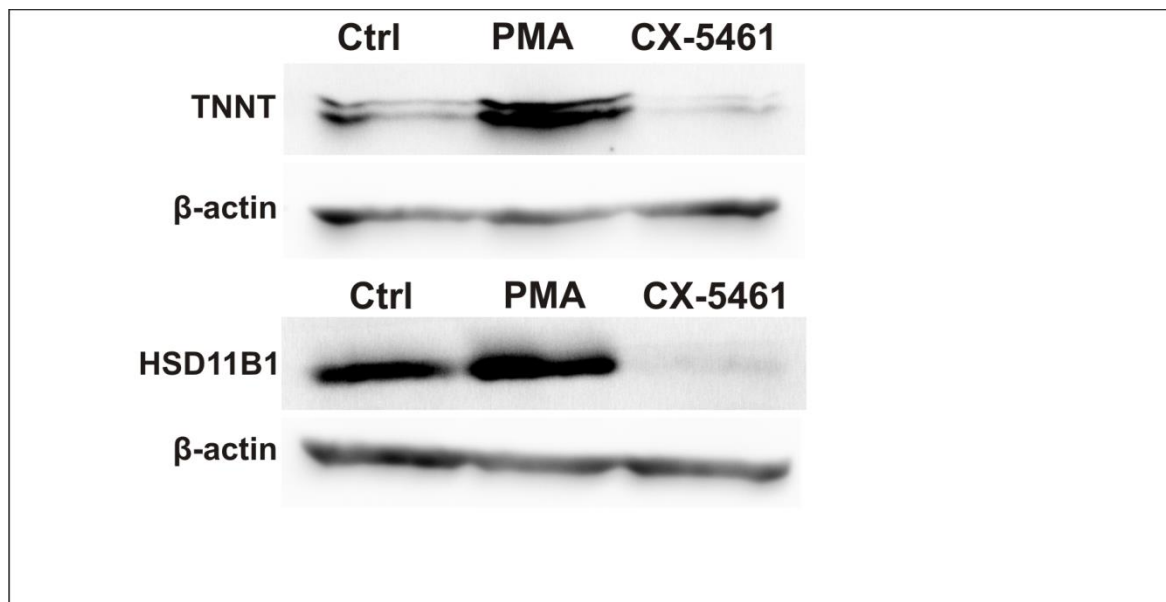


Figure S3. The protein level of TNNT and HSD11B1 in MO3.13 cells, control (Ctrl) and treated with PMA or CX-5461. Actin served as loading control.

Table S1. Sequences of primers used in RT-qPCR.

Gene	Forward Primer (5'->3')	Reverse Primer (5'->3')
<i>18S</i>	CGCCGCTAGAGGTGAAATTC	TTGGCAAATGCTTTCGCTC
<i>CASQ2</i>	AGCTTGTGGAGTTTGTGAAG	GGATTGTCAGTGTTGTCCC
<i>HSD11B1</i>	GACAGCGAGGTCAAAAGAAA	GTCCTCCCATGAGCTTTCCTG
<i>MYH *</i>	GAGGACATGGCCATGATGAC	GGCATTGTCAGAGATGGAGAAG
<i>TNNT3</i>	AGGAGGAGGATGCCAAGAGGA	TCCTTGGCCTTGTCCTCAGTTT
<i>GPR17</i>	GAGAGATGCTGAAACTCTCAGC	CAGGGAGAAGTTGGTGATCAGAC
<i>MBP</i>	ACCCAAGATGAAAACCCCGTA	TCCGTAGCCAAATCCTGGTCT
<i>MOG</i>	TTTTGATCCCCACTTTCTGAGG	CGTAGCTCTTCAAGGAATTGCC

Table S2. MO3.13 genes with statistically significant change in expression after PMA treatment. 6769 genes were up-regulated and 5856 - down-regulated.

(Table S2, please see the excel file)