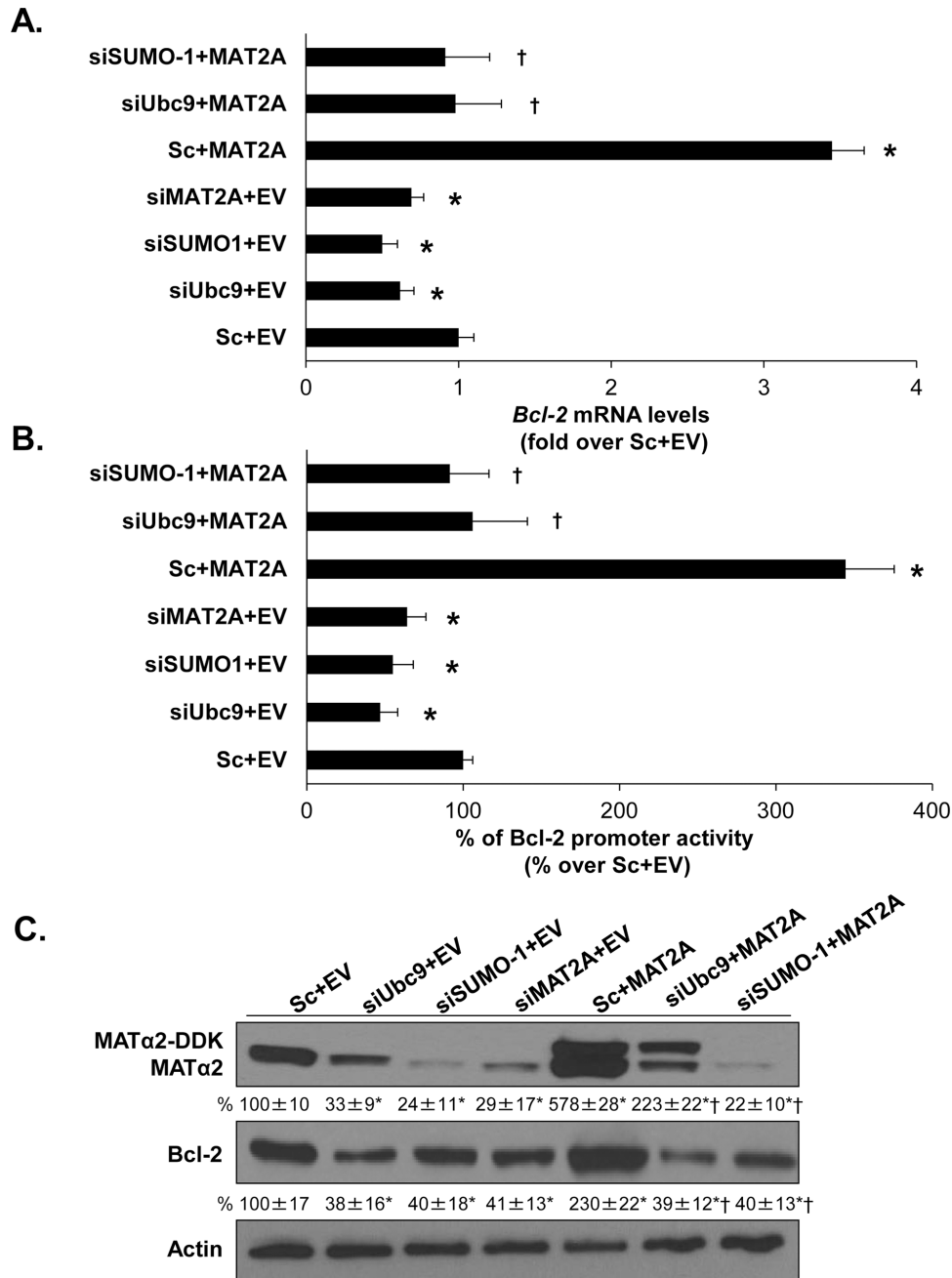
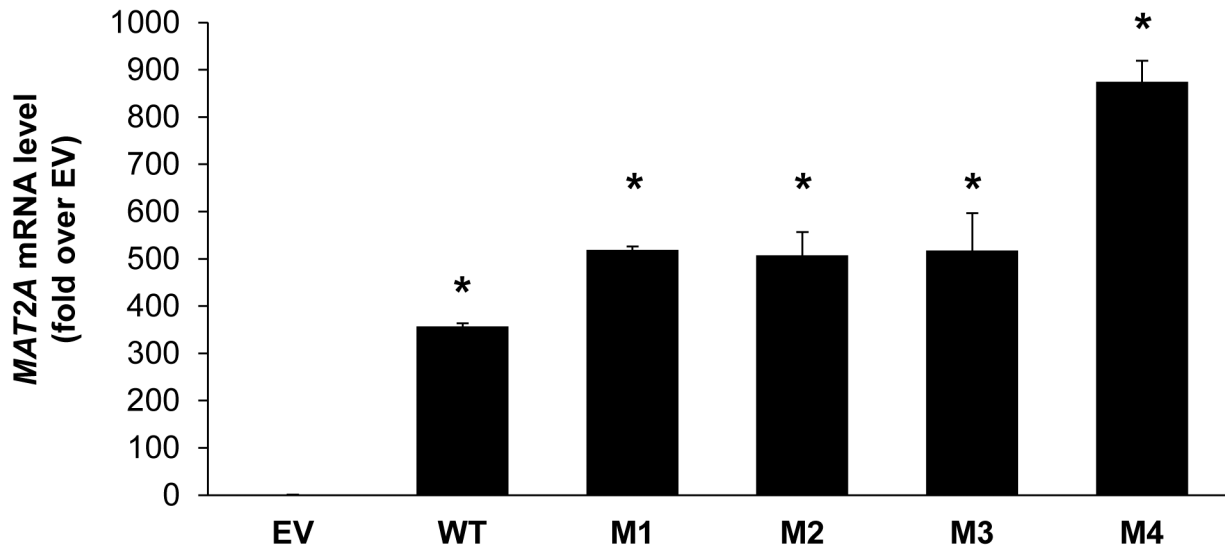
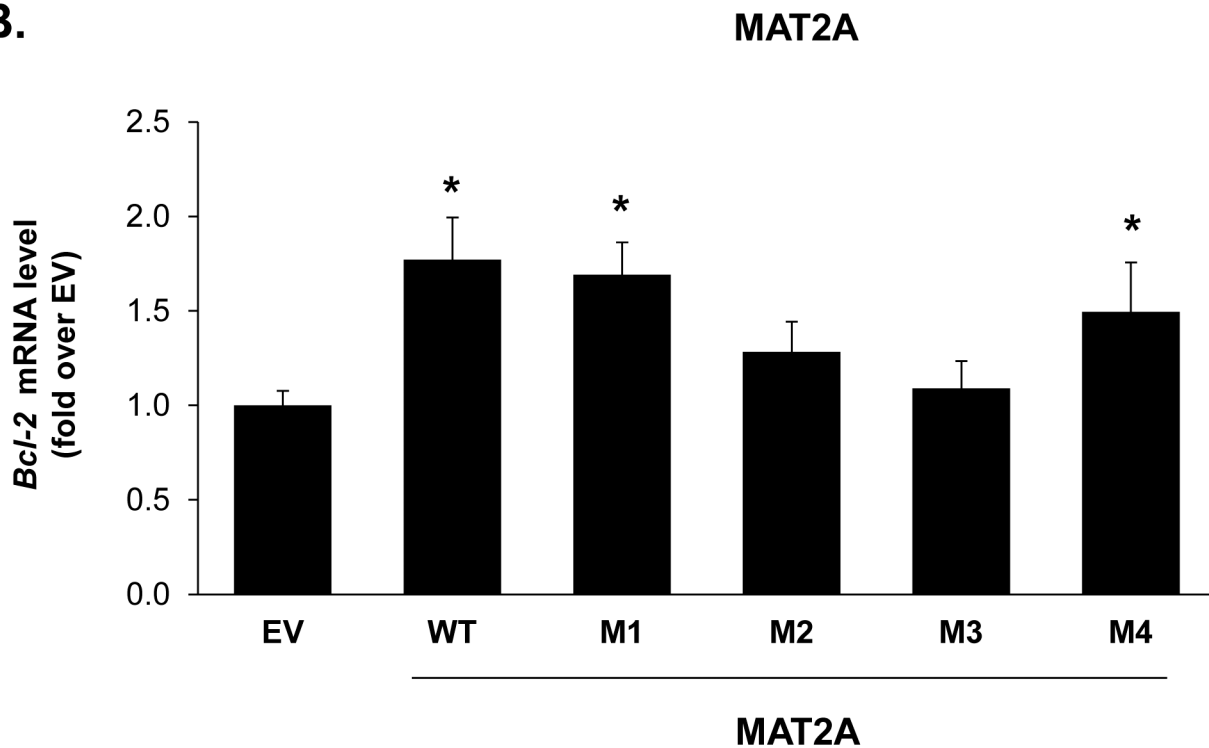


SUPPLEMENTARY FIGURES AND TABLES



Supplementary Figure S1: MAT2A and sumoylation machinery regulate Bcl-2 expression in RKO cells. **A.** RKO cells were transfected with siMAT2A, siUbc9 and siSUMO-1 (20 nM) or scrambled siRNA (Sc) for 48 hours and/or MAT2A overexpression vector or empty vector (EV) for 24 hours. The mRNA levels of Bcl-2 were compared to Sc+EV using real-time PCR. Results represent mean \pm SEM from 3 experiments in duplicates, * p < 0.04 vs. Sc+EV control; † p < 0.04 vs. Sc+MAT2A overexpression vector. **B.** Following the same treatments as in “A”, RKO cells were transfected with the human *Bcl-2* promoter or pBS-basic as described in Methods. The luciferase activity driven by *Bcl-2* promoter was normalized to that of pBS-Basic and expressed as % over Sc+EV. Results represent mean \pm SEM from 4 experiments in duplicate, * p < 0.04 vs. Sc+EV control; † p < 0.03 vs. Sc+MAT2A overexpression vector. **C.** Cells were treated as in “A” and total cellular protein was subjected to Western blotting with antibodies against MAT α 2 or Bcl-2. Densitometric changes were normalized to actin. Representative images and densitometric analysis (% mean \pm SEM of Sc+EV, indicated below the blots) from 3 experiments in RKO cells are shown. * p < 0.03 vs. Sc+EV, † p < 0.001 vs. Sc+MAT2A overexpression vector.

A.**B.**

Supplementary Figure S2: Effects of wild-type or mutant MAT2A overexpression on MAT2A and Bcl-2 mRNA levels in RKO cells. RKO cells were transfected with empty vector (EV), hMAT2A-WT, hMAT2A-M1, and hMAT2A-M2, hMAT2A-M3 and hMAT2A-M4 for 24 hours as described in Methods. **A–B.** MAT2A and Bcl-2 expression was determined by real-time PCR. mRNA results are expressed as mean % of EV \pm SEM from 3 independent experiments. * $p < 0.01$ vs. EV control for (A), * $p < 0.03$ for (B).

Supplementary Table S1: EMSA primers of Bcl-2 P2 promoter

Probe 1 Fw	TGCGGATTGACATTTCTGTGAA
Probe 1 Rev	TTCACAGAAATGTCAATCCGCA
Probe 2 Fw	GGTTTAATGTAAC TTCAATGG
Probe 2 Rev	CCATTGAAAGTTACATTA AAC
Probe 3 Fw	GGCAGCTTAATACATTCTTTT TAGCCG
Probe 3 Rev	CGGCTAAAAAGAATGTATTAAGCTGCC
<p>BCL-2 P2 PROMOTER SEQUENCE:</p> <p>GGTTGGGATTCCTGCGGATTGACA TTTCTGTGAAGCAGAAGTCTGGGAATCGATCTGGAAATCCTCCTAATTTT TACTCCCTCTCCCCGCGACTCCTGATTCATTGGGAAGTTTCAAATCAGCT ATAACTGGAGAGTGCTGAAGATTGATGGGATCGTTGCCTTATGCATTTGT TTTGGTTTTACAAAAAGGAACTTGACAGAGGATCATGCTGTACTTAAAA AATACAAGTAAGTTCTCTGCACAGGAAATTGGTTTAATGTAAC TTCAAT GGAAACCTTTGAGATTTTTTACTTAAAGTGCATTCGAGTAAATTTAATTT CCAGGCAGCTTAATACATTCTTTT TAGCCGTGTTACTTGTAGTGTGTATG CCCTGCTTTCACTCAGTGTGTACAGGGAAACGCACCTGATTTTTTACTTA TTAGTTTGTTTTTTCTTTAACCTTTCAGCATCACAGAGGAAGTAGACTGA TATTAACAATACTTACTAATAATAACGTGCCTCATGAAATAAAGATCCGA AAGGAATTGGAATAAAAATTTCTGCATCTCATGCCAAGGGGAAACACC AGAATCAAGTGTTCCGCGTGATTGAAGACACCCCTCGTCCAAGAATGCA AAGCACATCCAATAAATAAGCTGGATTATAACTCCTCTTCTTTCTCTGGG GGCCGTGGGGTGGGAGCTGGGGCGAGAGGTGCCGTTGGCCCCCGTTGCTT TTCCTCTGGGAAGGATG</p>	

Supplementary Table S2: Site-directed mutagenesis primers of MAT α 2 sumoylation sites

MAT2A mut 1 Fw	TTGGGTGGCAAATCCCTTGTTAGAGGAGGTCTGTG
MAT2A mut 1 Rev	CACAGACCTCCTCTAACAAGGATTTTGCCACCCAA
MAT2A mut 2 Fw	CATTATGGTACCTCTCAGAGGAGTGAGAGAGAGCTATTA
MAT2A mut 2 Rev	TAATAGCTCTCTCTCACTCCTCTGAGAGGTACCATAATG
MAT2A mut 3 Fw	TCAGGGATCTGGATCTGAGGAAGCCAATTTATCAGAG
MAT2A mut 3 Rev	CTCTGATAAATTGGCTTCCTCAGATCCAGATCCCTGA
MAT2A mut 4 Fw	AGTGCCCAAAAAGCTTAGATATACGCGTACGCGGC
MAT2A mut 4 Rev	GCCGCGTACGCGTATATCTAAGCTTTTTGGGCACT