

Figure S1. Canonical pathway analysis for the target-mRNAs. The negative logarithmic P-value is presented. The darker purple color tones correspond to comparably more significant regulation. However, the direction of regulation cannot be deduced from it. The asterisk (\*) symbol indicates pathways that were not considered relevant for colorectal cancer due to their sole function in the neuronal system. TIVA, total intravenous anesthesia; VAG, volatile anesthetic gas.

Canonical pathways	-Log (p-value)		Regulatory protein	
	TIVA	VAG	TIVA	VAG
p53_signaling	0.3	2.8	HIPK2	BCL2
Autophagy			-	BCL2
4-Aminobutyrate degradation I*			-	ALDH5A1
Cell cycle: G2/M DNA damage checkpoint regulation	0.3		HIPK2	-
Glutamate degradation III (via 4-aminobutyrate)*			-	ALDH5A1
Molecular mechanisms of cancer	0.3		HIPK2	-
Inflammasome pathway			-	P2RX7 (P2X7)
Senescence pathway	0.3		HIPK2	-
Cytotoxic T lymphocyte-mediated apoptosis of target cells			-	BCL2
B cell development			-	SPN (CD43)
Interferon signaling			-	BCL2
Neuroinflammation signaling pathway*			-	BCL2
Docosahexaenoic acid (DHA) signaling			-	BCL2
Myc mediated apoptosis signaling			-	BCL2