

SUPPLEMENTARY TABLE S5. NOVEL miRNAs ASSOCIATED WITH TBI

<i>Novel miRNA</i>	<i>Sequence</i>	<i>EV mTBI vs. CC Log2 fold change</i>	<i>p value</i>	<i>EV mTBI vs. DC Log2 fold change</i>	<i>p value</i>
Novel:chr1_407-5p	GAUGAUAGGCUGGGUGUG	-1.28	0.00434	-1.34	0.00960
Novel:chr10_425-3p	AAACAGUCAGGAUGUUGCCU	-1.01	0.02101	-0.67	0.36737
Novel:chr18_550-5p	UGACAGCAGGACGGAGGU	1.21	0.00353	1.61	0.00044
Novel:chr3_5-3p	UGGGACUGAGACACUGCC	-0.54	0.14533	-1.42	0.02593
Novel:chr5_575-3p	CGCGGGUGCUUCCAGCUCUUU	-2.97	0.00043	-0.59	0.37790
Novel:chr7_33-5p	AGUUGGUCCGAGUGUUGUG	-1.89	0.00186	-1.66	0.01809

CC, community controls; DC, deployed controls; EV, extracellular vesicle; miRNA, microRNA; mTBI, chronic mild traumatic brain injury.