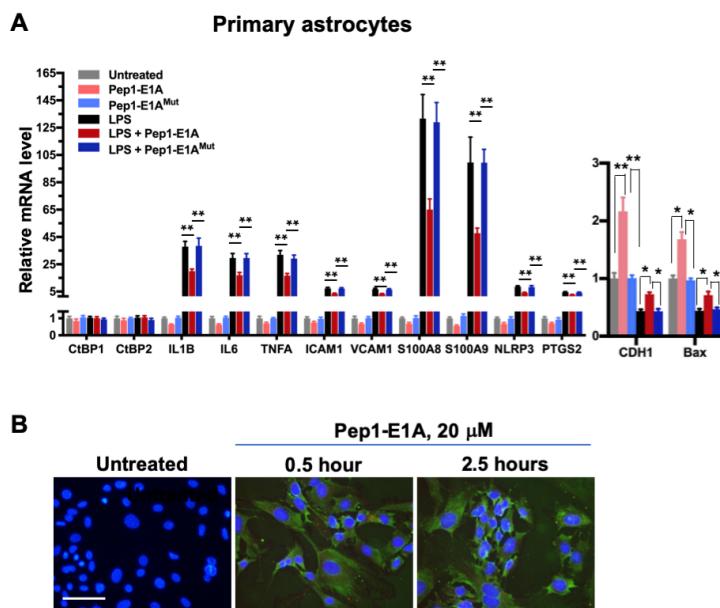


## **SUPPLEMENTARY MATERIALS**

### **C-terminal binding proteins 1 and 2 in traumatic brain injury-induced inflammation and their inhibition as an approach for anti-inflammatory treatment**

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**Figure S1**

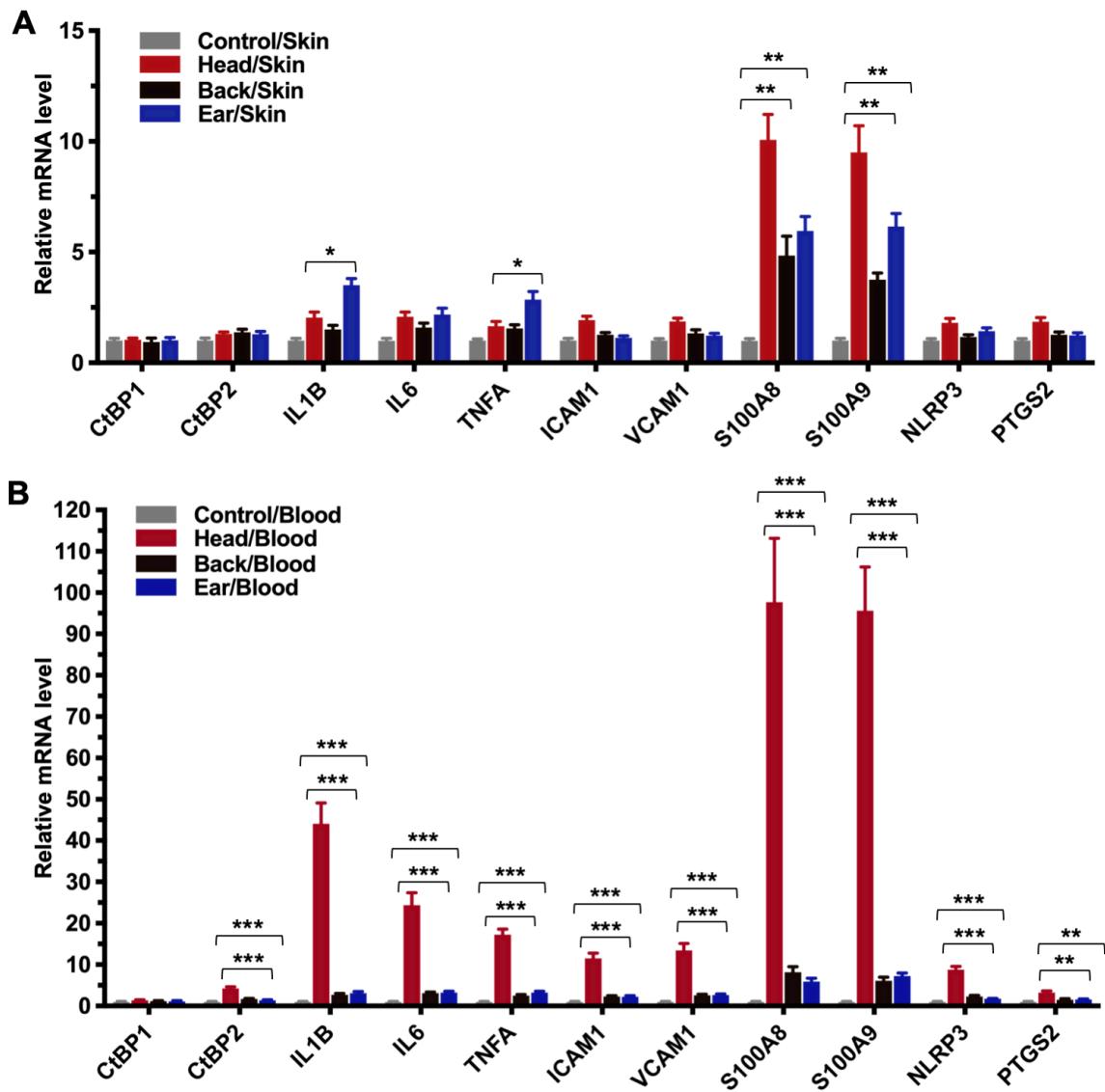


**Figure S1. Pep1-E1A suppresses LPS-induced inflammatory gene expression in mouse primary astrocytes.**

(A) Effects of pretreatment with 20  $\mu$ M Pep1-E1A or Pep1-E1A<sup>Mut</sup> on the basal and LPS-induced mRNA expression of CtBP target genes in mouse primary astrocytes.  $n = 3$ ; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

(B) Representative immunostaining images of Pep1-E1A internalization into cultured astrocytes after 0.5 and 2.5 h incubation. Scale bar, 50  $\mu$ m.

**Figure S2**



**Figure S2. CHIMERA-delivered mild TBI causes systemic inflammatory response.**

Mice received sham procedure (control) or a single impact of 0.7 J energy to the head, the back, or the ear. Skin tissues containing the site of the impact (**A**) and peripheral blood leukocytes (**B**) and were collected 24 h postinjury for total RNA extraction and RT-qPCR analysis.  $n = 4$ ; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

**Table S1. Sequences of primers used in this study.**

<b>ChIP-qPCR</b>	<b>Forward (5' to 3')</b>	<b>Reverse (5' to 3')</b>
mIL1B	ACATGAGACTGGCTATGGTATT	TTATCCCTTTCCAGGTCTCCC
mIL6	AGGGCTAGCCTCAAGGATGACT	GGAGTCAACTCTCTAATTTGA
mTNFA	CAGCCCTCCCAAAGCCCATGCA	GTGCTTCTGAAAGCTGGGTGCA
mS100A8	GCCTAGACATGGACTTATTGCC	AGCATTCTGAGTCTGAGGAAG
<b>RT-qPCR</b>		
mCtBP1	TAGGCGGGGCAAGAGGAAGC	CGAGGAACGCAAAGGACACAGG
mCtBP2	GGCACACCCCTCCCAAGCTC	GGCACACCCCTCCCAAGCTC
mIL1B	GCCCCATCAGAGGCAAGGAGGA	CAGGTCGCTCAGGGTCACAAGA
mIL6	TGGCTAAGGACCAAGACCATTCA	CATAACGCACTAGGTTGCCGAGTAGA
mTNFA	CCCACACCGTCAGCCGATTT	CCTTGGGGCAGGGCTCTT
mICAM1	TGGTACATACGTGTGCCATGCCT	ATCCTGATCTTCTCTGGCGGT
mVCAM1	TCCGCCAGGCACAGCTGCAGGA	ATTACCAAGGAAGATGCGCAG
mS100A8	TCGAGGAGTTCTTGCGATGG	ACATATCCAGGGACCCAGCCCTA
mS100A9	TGGAGGACCTGGACACAAACCAG	TTCCACAGCCTTGCCATGA
mNLRP3	CCCCCTCCCTCCGCTTCT	CTAGCCCACGGGCAGCTCCT
mPTGS2	GGACTGCAGAAGGCCCATGT	GGAACACAGCTACGAAAACCAATCA
mCDH1	CCCCTCCAATGCCTGCTTTG	TCTGACTGCCTCTGCCTCTGA
mBax	AGTCCTCACCGCCTCGCTCA	GCCTTCCCCTTCCCCATT