

[illegible]

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Figure S1: Gene sequences are shown. Sequences of capture extenders (blue), label extenders (red), blocking probes (green) and flanking sequences (black) are indicated.

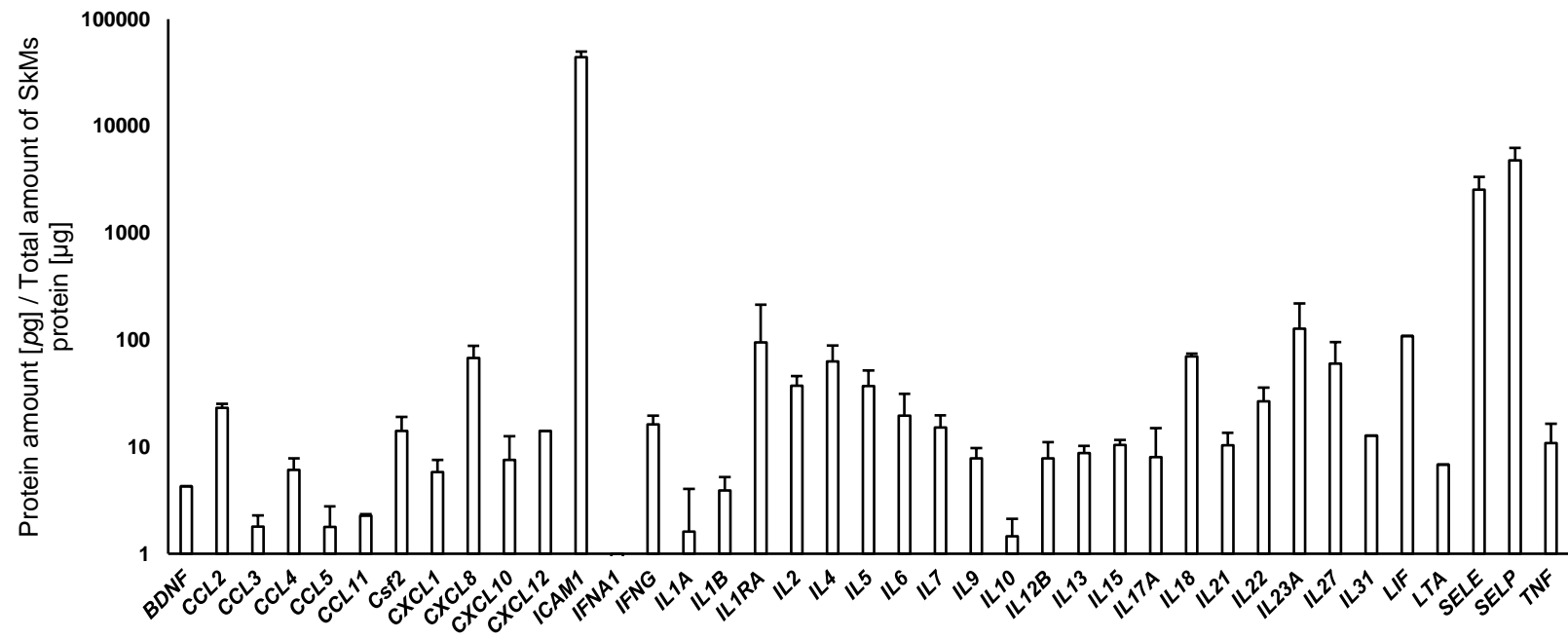


Figure S2: Constitutive expression of proteins in untreated SkMs. SkM cell extracts were subjected to multiplex protein quantification three times to precisely determine the concentration of inflammatory markers in total cellular proteins. The results are presented as the mean \pm SD from three independent experiments.

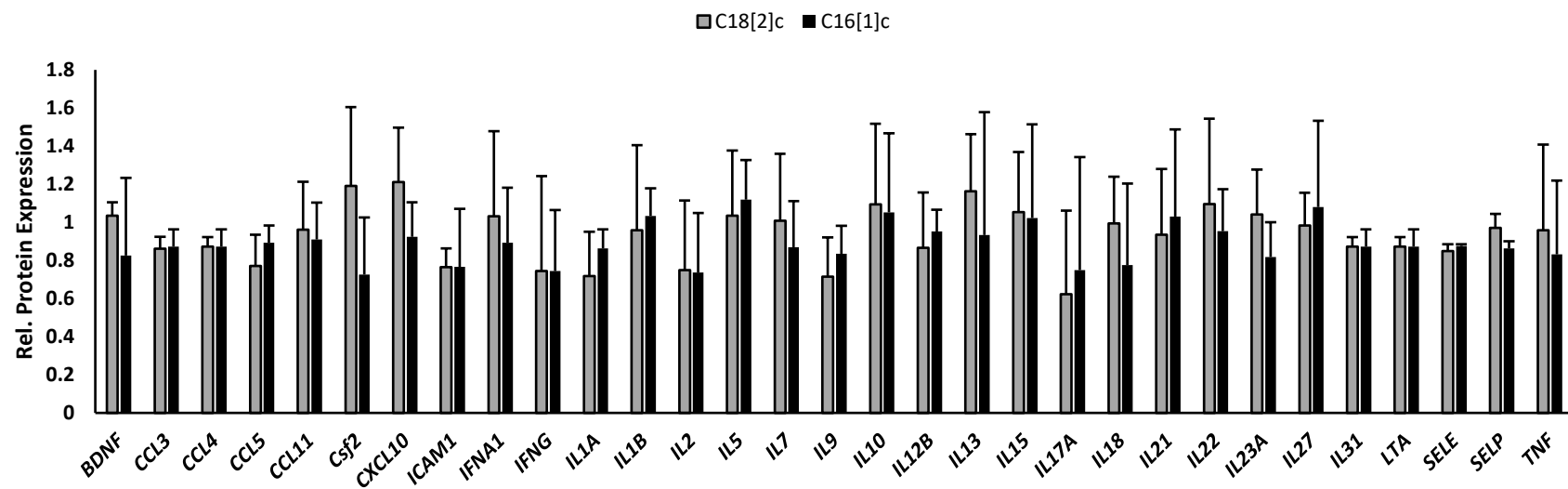


Figure S3: CC18[2]c and C16[1]c mediated relative protein expression in SkMs. The experiments and data analysis were performed as described in the legend to Fig. 1a-h.

