Supplementary information

Enlisting the *Ixodes scapularis* Embryonic ISE6 Cell Line to Investigate the Neuronal Basis of Tick-Pathogen Interactions

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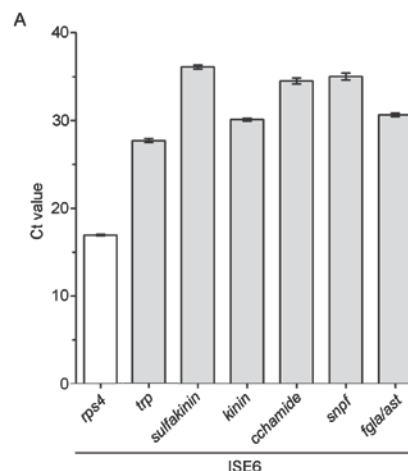
Supplementary figure S1. (A) Relative expression of ribosomal protein S4 (*rps4*) transcript and six different neuropeptide transcripts in uninfected ISE6 cells. The Ct values (i.e., the mean-standard error) for the two biological replicates are shown.

Supplementary figure S2. Protein (A) and nucleotide (B) alignment of *Ixodes scapularis* and *Ixodes ricinus* TRP. ISCI008383 – *I. scapularis* genome-predicted transcript; EL516783 - *I. scapularis* EST sequence; MW082607 – *I. ricinus* transcript (identified in this study). Please note the incorrect N-terminal sequence of ISCI008383 (A) caused by the incomplete 5'-end (B). In addition the shift in the 3'-end (B) reading frame caused an improper conceptual TRP translation (A) of ISCI008383.

Name	Forward primer (5'-3')	Reverse primer (5'-3')
fgla/ast	CACCATGCTGCTGAGCAA	CTACTTTCTCCGGAAGCAGG
cchamide	GTGTCCTCTGACGTGATGC	GTCTATCAAGCCTCGCCTG
short neuropeptide f	GCCTGCATCCTTCTCGATAG	CTTCCGTTCCATGGTGTGG
sulfakinin	CTTTGTCTTCTTGCAGCACTC	GACGGGATCGATCATGTCC
trp	CTTGCGGATGGCATTGGTTC	CACCTATTGGCGTAACCCG
kinin	GAGGCAGGACAAAGAGAGC	CGAGAAGGTGCTGTCCTTAG
natalisin (1st set)	AACACTCGGATCACAGACCC	CATCGTGACGTCGAAAGAGT
natalisin (2nd set)	AGGTAGTATGTCCGACCAAC	ACAGGTGACTACTGCGACAA
natalisin (3rd set)	CTTGCGGATGGCATTGGTTC	CACCTATTGGCGTAACCCG

Supplementary table S1. List of primers used in our qRT-PCR experiments.