

Hippocampal microRNA-132 mediates stress-inducible cognitive deficits through its acetylcholinesterase target

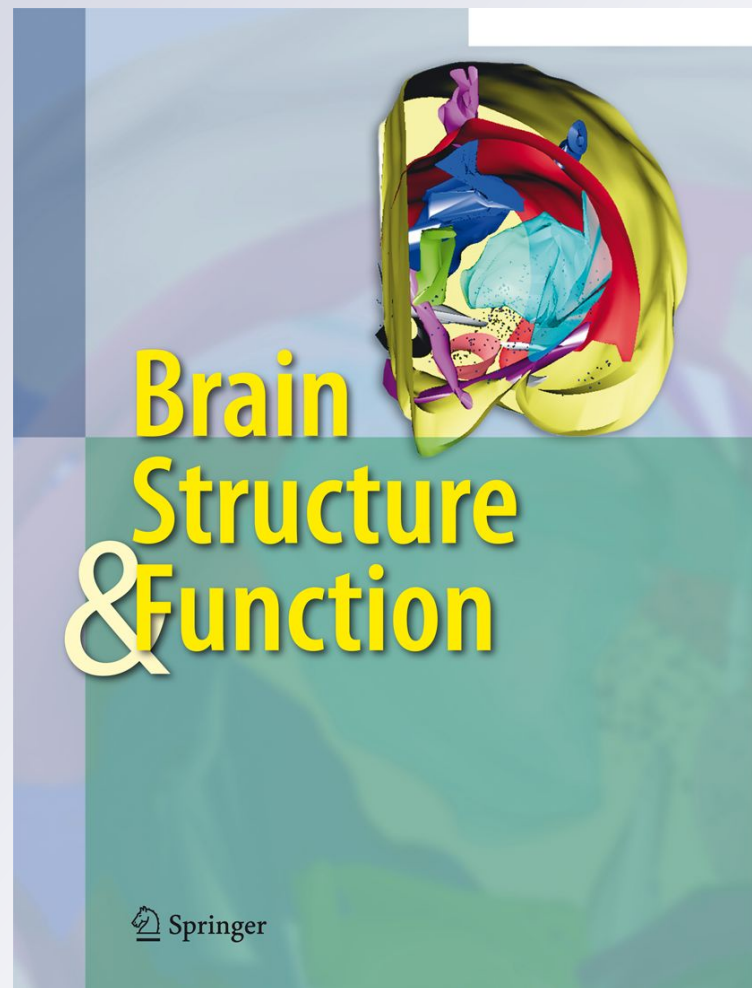
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Hippocampal microRNA-132 mediates stress-inducible cognitive deficits through its acetylcholinesterase target

G. Shaltiel · M. Hanan · Y. Wolf · S. Barbash · E. Kovalev · S. Shoham · H. Soreq

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Abstract

Abstract text describing the study's findings on hippocampal microRNA-132 and its target acetylcholinesterase.

3/4. hippocampal microRNA-132. acetylcholinesterase target. 250A. hippocampal microRNA-132A.

Keywords A 250A

Introduction

Introduction text discussing the role of microRNAs in cognitive function and the specific findings of microRNA-132.

Electronic supplementary material

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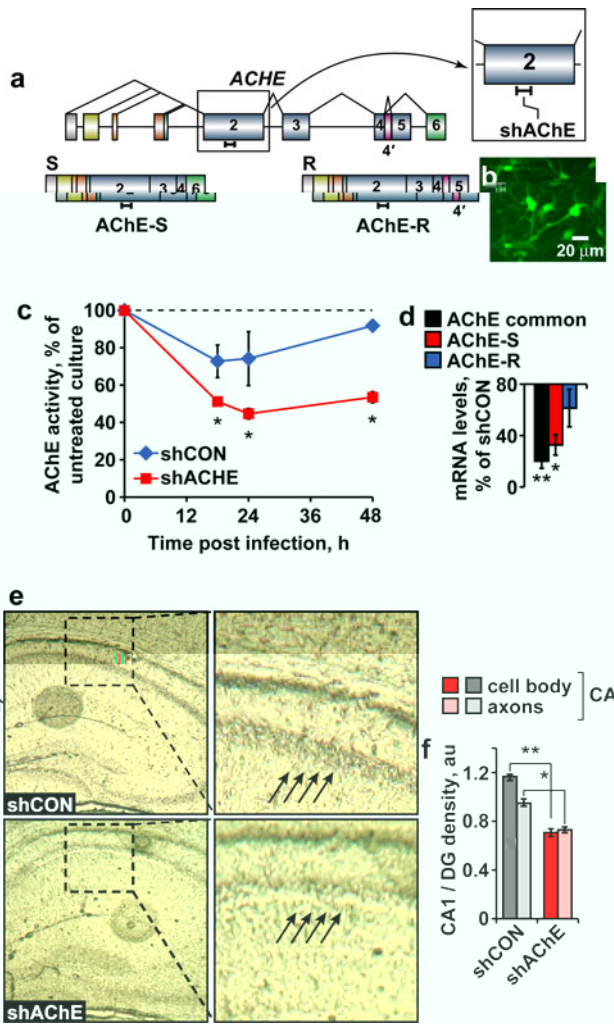


Fig. 3 Schematic of AChE-S and AChE-R. **a** Schematic of AChE-S and AChE-R. AChE-S (S) and AChE-R (R) are shown with domains 2, 3, 4, 5, and 6. AChE-R has a 4' tail. shAChE is a shRNA targeting domain 2. **b** Fluorescence microscopy image of neurons showing green fluorescence. Scale bar = 20 μm. **c** Line graph of AChE activity (% of untreated culture) over 48 hours. **d** Bar graph of mRNA levels (% of shCON) for AChE common, AChE-S, and AChE-R. **e** Histology images of CA1 and DG regions in shCON and shAChE groups. **f** Bar graph of CA1/DG density (au) for shCON and shAChE groups.

$(F_{1,10} = 18.97, *p < 0.01)$, $(F_{2,10} = 1.39, p = \dots)$, $(F_{2,10} = 0.46, p = \dots)$. **d** $(F_{1,10} = 18.97, *p < 0.01)$, $(F_{2,10} = 1.39, p = \dots)$, $(F_{2,10} = 0.46, p = \dots)$. **e** $(t_{13} = 5.24, **p < 0.001)$, $(t_{13} = 18.5, **p < 0.001)$. **f** $(t_{13} = 5.24, **p < 0.001)$, $(t_{13} = 18.5, **p < 0.001)$.

AChE-R (R) is shown with domains 2, 3, 4, 5, and 6. AChE-R has a 4' tail. shAChE is a shRNA targeting domain 2. **b** Fluorescence microscopy image of neurons showing green fluorescence. Scale bar = 20 μm. **c** Line graph of AChE activity (% of untreated culture) over 48 hours. **d** Bar graph of mRNA levels (% of shCON) for AChE common, AChE-S, and AChE-R. **e** Histology images of CA1 and DG regions in shCON and shAChE groups. **f** Bar graph of CA1/DG density (au) for shCON and shAChE groups.

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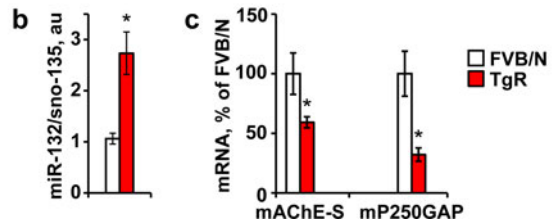
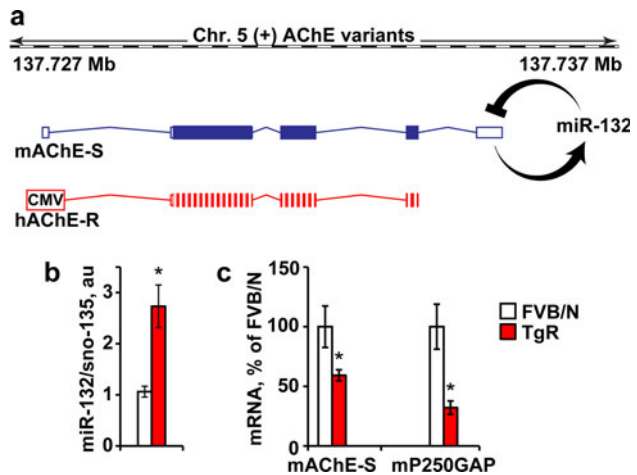


Fig. 6 miR-132 expression and its effect on AChE mRNA levels. **a**, Schematic of the AChE gene structure on Chromosome 5 (+). **b**, miR-132/sno-135 levels in FVB/N (white) and TgR (red) mice. **c**, mRNA levels of mAChE-S and mP250GAP in FVB/N (white) and TgR (red) mice. *p < 0.05.

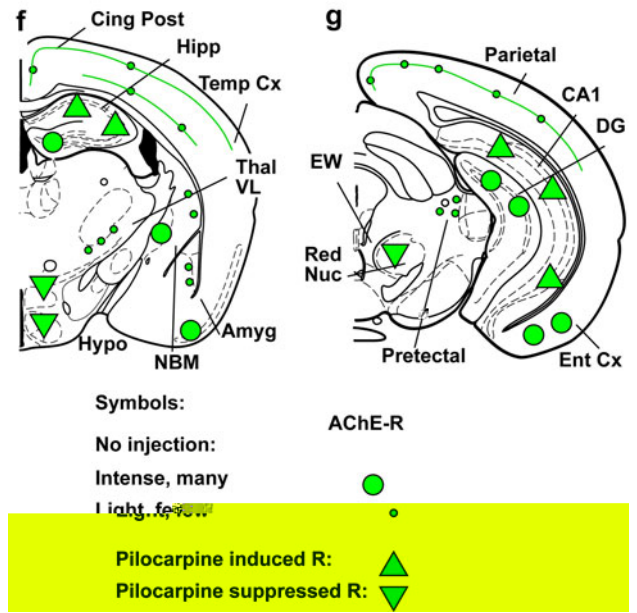
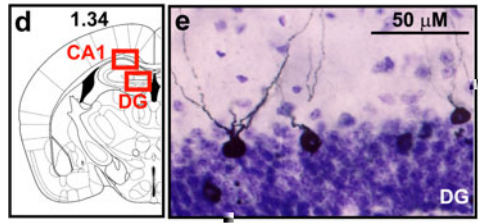
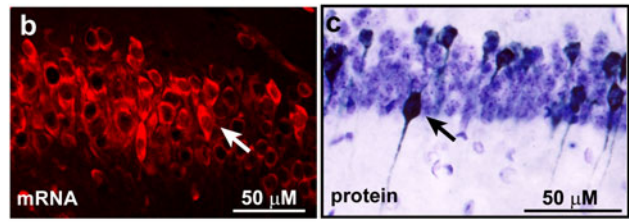
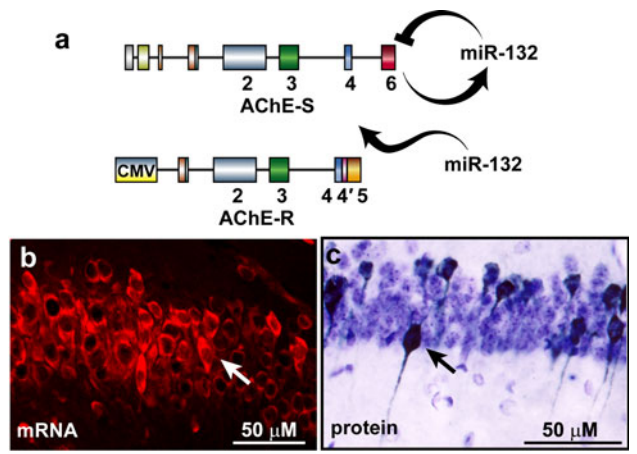


Fig. 7 miR-132 expression and its effect on AChE-R protein levels. **a**, Schematic of the AChE-R gene structure. **b**, miR-132/sno-135 levels in FVB/N (white) and TgR (red) mice. **c**, mRNA levels of AChE-R and mP250GAP in FVB/N (white) and TgR (red) mice. **d**, Schematic of the brain showing the location of the CA1 and DG regions. **e**, Histology of the brain showing the location of the CA1 and DG regions. **f**, Schematic of the brain showing the distribution of AChE-R in various regions. **g**, Schematic of the brain showing the distribution of AChE-R in various regions. *p < 0.05.

... (7). ...
 ... (8). ...
 ... (8). ...

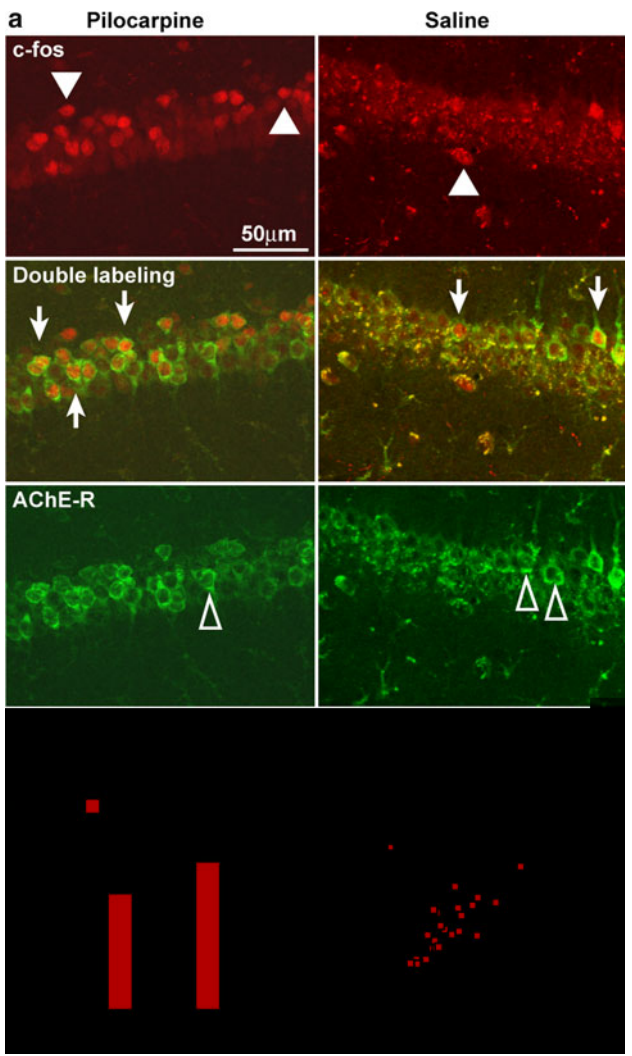


Fig. 8 ...
 ... (F_{1,19} = 226, p < 0.0001)
 ... (F_{2,19} = 5.88, p < 0.05). ...
 ... r = 0.86, p < 0.05 ...
 ... r = 0.77, p < 0.05

... (2004)
 ... r = 0.86, p < 0.05, ... r = 0.77, p < 0.05/ ... (8). ...
 ... p < 0.001).
 ... (2002),
 ... 55% ... (t ... p < 0.01/ ... (9)).

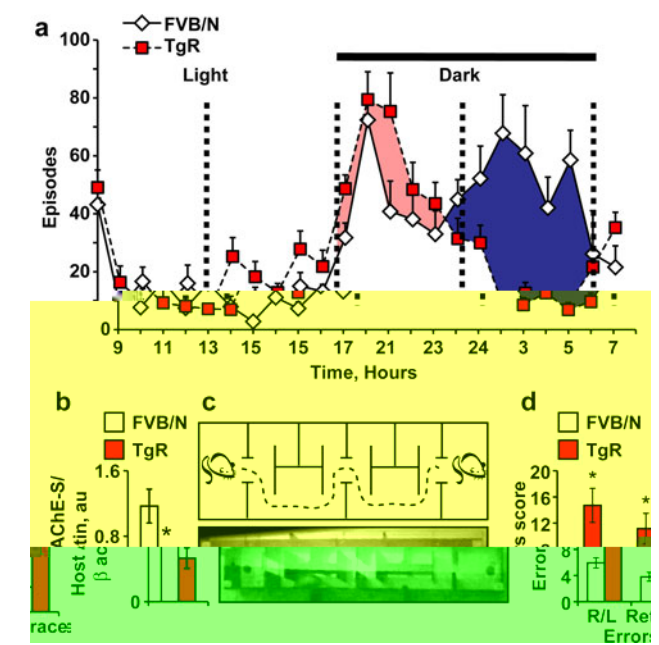


Fig. 9 ...
 ... Light F_{1,22} = 0.4, p = ... / Dark F_{1,22} = 79.02, p < 0.001. ...
 ... t₁₃ = 2.71, *p < 0.01). ...
 ... F_{1,33} = 9.37, *p < 0.005 ... F_{1,33} = 9.97/

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