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MicroRNA-34a Acutely Regulates Synaptic Efficacy in the Adult Dentate Gyrus In Vivo

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Abstract

Activity-dependent synaptic plasticity involves rapid regulation of neuronal protein synthesis on a time-scale of minutes. miRNA function in synaptic plasticity and memory formation has been elucidated by stable experimental manipulation of miRNA expression and activity using transgenic approaches and viral vectors. However, the impact of rapid miRNA modulation on synaptic efficacy is unknown. Here, we examined the effect of acute (12 min), intrahippocampal infusion of a miR-34a antagonist (antimiR) on medial perforant path-evoked synaptic transmission in the dentate gyrus of adult anesthetised rats. AntimiR-34a infusion acutely depressed medial perforant path-evoked field excitatory post-synaptic potentials (fEPSPs). The fEPSP decrease was detected within 9 min of infusion, lasted for hours, and was associated with knockdown of antimir-34a levels. Antimir-34a-induced synaptic depression was sequence-specific; no changes were elicited by infusion of scrambled or mismatch control. The rapid modulation suggests that a target, or set of targets, is regulated by miR-34a. Western blot analysis of dentate gyrus lysates revealed enhanced expression of Arc, a known miR-34a target, and four novel predicted targets (Ctip2, PKI-1 α , TCF4 and Ube2g1). Remarkably, antimiR-34a had no effect when infused during the maintenance phase of long-term potentiation. We conclude that miR-34a regulates basal synaptic efficacy in the adult dentate gyrus in vivo. To our knowledge, these in vivo findings are the first to demonstrate acute (< 9 min) regulation of synaptic efficacy in the adult brain by a miRNA.

Keywords microRNA · miR-34a · Gene expression · Hippocampus · Protein synthesis · Synaptic plasticity · Synaptic efficacy

Introduction

The highly pleiotropic nature of miRNAs has changed our view of neuronal development, function and aging. Some 60% of human genes may be post-transcriptionally regulated by miRNAs [1, 2], and in the brain, the list of miRNAs implicated in neuronal plasticity paradigms is growing [3, 4]. Synaptic stimulation resulting in long-term potentiation (LTP) and long-term depression (LTD) have been

demonstrated to involve tight temporal control of many brain-specific miRNAs, suggesting that miRNAs coordinate protein expression underlying the plasticity of synaptic transmission [3, 5–10]. Causal roles for specific miRNAs in LTP and LTD mechanisms have been identified using viral vectors to chronically knockdown microRNA expression or inhibit miRNA binding to target mRNA [11, 12]. Rapid, neuronal activity-dependent regulation of the RNA-induced silencing complex (RISC) has also been demonstrated [7, 13, 14]. However, to date, no miRNAs have been demonstrated to play an acute role in regulating synaptic strength on the order of minutes.

miR-34a plays a critical regulatory role in neurodegenerative diseases such as Alzheimer's disease (AD) [15], neuropsychiatric disorders such as bipolar disorder [16, 17], and schizophrenia [18], in addition to being implicated in development and many forms of cancer [19–21]. miR-34a has emerged as a key regulator of cell proliferation, apoptosis, and differentiation, and at least 77 targets that have been experimentally validated across multiple cell types [19]. miR-34a is ubiquitously expressed with the highest abundance in

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57 the brain (and testes) [22]. In neurons, miR-34a has a
58 somatodendritic distribution and it is implicated in the regula-
59 tion of synaptic protein targets [23]. In the rat dentate gyrus
60 in vivo, induction of LTP has been associated with enhanced
61 miR-34a expression in the Argonaute 2/RISC [9] and altered
62 total miR-34a expression in lysates samples [5, 8, 9]. Our
63 previous work identified the immediate early gene Arc as a
64 potential target of miR-34a. In cultured hippocampal neurons,
65 overexpression of miR-34a resulted in downregulation of Arc
66 expression [5]. Intriguingly, no changes in miR-34a were ob-
67 served following brain-derived neurotrophic factor (BDNF)
68 treatment in primary hippocampal neuronal cultures in vitro
69 or following high-frequency stimulation (HFS)-induced LTP
70 in vivo, despite increased Arc expression [5]. In total lysates,
71 we have reported a small increase in miR-34a expression, and
72 a large increase in Ago2-associated miR-34a expression,
73 30 min following LTP induction, in vivo [9]. The work sug-
74 gested a preferential loading of miR-34a into the Ago2/RISC
75 following HFS-induced NMDA-receptor-dependent LTP. In
76 awake rats, miR-34a expression was shown to be reduced
77 20 min after LTP induction in the dentate gyrus, leading the
78 authors to suggest that NMDA receptor-mediated reduction of
79 miRNA levels out-competes an independent process working
80 to increase miRNA expression [24].

81 Given the key role of Arc in activity-dependent synaptic
82 plasticity and memory [25], we wanted to explore the possible
83 role of endogenous miR-34a in regulating Arc expression and
84 synaptic efficacy. Herein, we examined a possible role of en-
85 dogenous miR-34a in the acute regulation of synaptic efficacy
86 at medial perforant path (MPP)-dentate gyrus synapses
87 in vivo. Using acute intrahippocampal infusion of miR-34a-
88 targeting antimiR and corresponding mismatch and scrambled
89 controls, we show that acute inhibition of miR-34a profoundly
90 depresses MPP-evoked field EPSPs. Remarkably, antimiR-
91 34a had no effect on synaptic transmission when infused dur-
92 ing the maintenance phase of LTP. The results suggest that
93 miR-34a plays a central role in regulating basal synaptic
94 efficacy.

95 Materials and Methods

96 Reagents

97 Custom-made locked nucleic acid oligonucleotides antimiR-
98 34a, mismatch antimiR-34a, and scrambled-antimiR-34a were
99 obtained from Exiqon, Qiagen, (1 mM prepared in 1 × PBS).
100 Sequences shown in Table S1. HiPerFect Transfection reagent
101 13.5% (Mat. No. 1029975; Lot No. 139311238; Qiagen) was
102 diluted in PBS and added prior infusion to yield 1 mM and
103 100 μM concentrations. Custom-made 6-FAMTM Fluorescein
104 antimiR-34a and scrambled-miR-34a (1 mM prepared in 1 ×
105 PBS; Exiqon) was used for evaluation of neuronal uptake,

106 distribution and localisation. Antibodies used for western blot
107 analysis were Arc C7 mouse monoclonal (1:500, sc-17839),
108 GAPDH mouse monoclonal (1:1000, sc-32233), PKI-1α goat
109 polyclonal (1:1000, sc-1944), from Santa Cruz Biotech, USA
110 and BCL11B/Ctip2 rabbit polyclonal (1:200, ABIN487046)
111 Aviva Systems Biology, USA, TCF4; TCF7L2 rabbit poly-
112 clonal (1:1000, ABIN487067) Aviva Systems Biology,
113 USA, UBE2G1 rabbit polyclonal (1:200, ABIN1385714),
114 Bioss, USA, and Gnao1 rabbit polyclonal antibody from
115 Aviva Systems Biology, USA.

116 Electrophysiology and Intrahippocampal Infusion 117 in Anesthetized Rats

118 Intrahippocampal drug infusion was performed as described
119 previously [26]. All experiments were carried out under ethi-
120 cal standards approved by the Norwegian Committee for
121 Experiments on Animals. The experiments were carried out
122 on 57 adult male rats of the Sprague-Dawley strain (Taconic,
123 Denmark), weighing 250–350 g. Rats were anesthetized with
124 urethane (1.5 mg/kg, i.p.), positioned in a stereotaxic frame,
125 and body temperature was maintained at 37 °C with a thermo-
126 statically controlled electric heating pad. A concentric bipolar
127 stimulating electrode (tip separation 500 μm; SNEX 100;
128 Rhodes Medical Instruments, Woodland Hills, CA) was
129 lowered into the angular bundle for stimulation of the medial
130 perforant path. Stereotaxic coordinates for stimulation were as
131 follows (in mm): 7.9 posterior to bregma, 4.2 lateral to the
132 midline and 2.5 below the dura.

133 A Teflon-coated tungsten wire recording electrode (outer
134 diameter of 0.075 mm; A-M Systems #7960) was glued to
135 the infusion cannula (30 gauge). The electrode was then cut
136 so that it extended 800 μm from the end of the cannula.
137 Stereotaxic coordinates for recording in the dentate hilus were
138 as follows (in mm): 3.9 posterior to bregma, 2.3 lateral and
139 2.8–3.1 below the dura. The recording electrode was slowly
140 lowered into the dorsal hippocampus until a positive-going
141 field EPSP (fEPSP) of maximum slope was obtained in the
142 dentate hilus. The tip of the infusion cannula was located in
143 the deep striatum lacunosum-moleculare of field CA1,
144 800 μm above the hilar recording site and 300–400 μm above
145 the medial perforant synapse. The infusion cannula was con-
146 nected via a polyethylene (PE50) tube to a 10-μl Hamilton
147 syringe (Reno, NV) and infusion pump. After baseline record-
148 ing for 20 min, 1 μl drug (100 μM or 1 mM in PBS, 13.5% HP
149 transfection reagent, lot no. 139311238, Qiagen) was infused
150 for 12 min at a rate of 0.085 μl/min. Evoked responses were
151 recorded for 140 min after infusion. Biphasic rectangular test
152 pulses of 150-μs duration were applied every 30 s throughout
153 the experiment (0.033 Hz). Responses were allowed to stabi-
154 lize for 1 h at a stimulation intensity that produced a popula-
155 tion spike 30% of maximum. A stable 20-min baseline of
156 evoked potentials was recorded (pulse-width 0.15 ms, at

157 0.033 Hz) before intra-hippocampal drug infusion. Evoked
 158 responses were recorded for 120 min post-infusion. Signals
 159 from the dentate hilus were amplified, filtered (0.1 Hz to 10
 160 kHz), and digitized (25 kHz). Acquisition and analysis of field
 161 potentials were accomplished using Datawave Technologies
 162 Software. The maximum slope of the fEPSP was measured,
 163 and averages of four consecutive responses were obtained.
 164 Changes in the fEPSP slope were expressed in percent of
 165 baseline (20-min preceding infusion). Responses for input-
 166 output (I/O) curves were collected immediately before base-
 167 line recording and at 30 min and 150 min post-infusion. Seven
 168 stimulus intensities ranging from 100 to 1000 μ A were applied
 169 in randomized sequence.

170 **Dissection of the Adult Rat Hippocampal Dentate**
 171 **Gyrus**

172 After recordings were completed, the electrodes were re-
 173 moved, rats were decapitated and DG microdissection was
 174 carried out within 2–3 min. The brain was removed from the
 175 skull and placed onto an ice-cold glass slide where it was
 176 continuously rinsed with ice-cold PBS. The cerebellum was
 177 removed and hemispheres were separated using a scalpel
 178 along the cerebral longitudinal fissure. The hemispheres were
 179 placed dorsal side down. Forceps were used to lift the brain
 180 stem exposing the corpus callosum and the medial side of the
 181 hippocampus. The hippocampus was tilted out from the tem-
 182 poral cortical fold. Fimbria and blood vessels were removed.
 183 Forceps were carefully inserted into the hippocampal fissure
 184 isolating the DG from the region of Cornu Ammonis, along its
 185 septotemporal axis. The microdissected DGs were immedi-
 186 ately frozen on dry ice and stored at -80°C until use.

187 **Microscopy**

188 FAM-labelled antimiR-34a and SC-miR-34a expression in
 189 hippocampal neurons was imaged on Axio Imager 2 Zeiss
 190 light fluorescent microscope.

191 **Cell Culture and antimiR-34a Transfection**

192 Normal rat kidney cells (NRK) were maintained in
 193 Dulbecco's modified Eagle medium (DMEM, Sigma) supple-
 194 mented with 10% heat-inactivated foetal calf serum (Sigma),
 195 100 U/ml penicillin (Sigma), 100 μ g/ml streptomycin (Sigma)
 196 and 1 mM L-glutamine at 37°C and 5% CO_2 . Cells were
 197 seeded in 12-well plates at a density of 8000 cells/ml.
 198 Hippocampal neuronal cultures were prepared with slight
 199 modifications from the protocol originally described by
 200 Banker and co-workers [27, 28]. Hippocampi of Wistar rat
 201 embryos (E18) were dissected and dissociated by trypsin treat-
 202 ment followed by trituration. After removal of trypsin, neu-
 203 rons were plated at a density of 200,000 cells/well in a 12-well

plate for biochemical analysis. Plates were precoated with
 poly-D-lysine (Sigma). The cultures were maintained in
 MEM (M2279, Sigma) growth medium supplemented with
 B-27 supplement (B-27® Supplement, Invitrogen), sodium
 bicarbonate, glucose, and pyruvate, that had been conditioned
 on astrocyte cultures for 3 days. Half of the neuronal growth
 medium was replaced with fresh growth medium twice a
 week. Hippocampal neurons were transfected at 8 DIV.
 AntimiR-34a and SC-miR-34a were transfected at a final con-
 centration of 1 mM using HiPerFect Transfection reagent di-
 luted to 13.5% in PBS.

The transfection mix was replaced with conditioned growth
 medium (neurons) or complete DMEM medium (NRK cells)
 after 3 h. Cells were washed with once with PBS and harvest-
 ed in RNA lysis buffer (PureLink® miRNA Isolation Kit (Life
 technologies, Invitrogen, Carlsbad, CA, USA) after 48 h.

Cell Culture and Fluorescent-Labelled antimiR-34a

Primary hippocampal neuronal cultures were prepared from
 embryonic day 18 (E18) Sprague-Dawley rat brains [29, 30].
 Cells were plated on coverslips coated with poly-L-lysine (100
 μ g/ml) and laminin (2 μ g/ml) at a density of 30,000/well.
 Hippocampal cultures were grown in Minimum Essential
 Medium (MEM) supplemented with B27, glutamate, sodium
 bicarbonate, glucose, glutamate, pyruvate and antibiotic (peni-
 cillin/streptomycin/neomycin) and added to astrocyte growing
 plate. Fifty percent of the MEM was replaced every 3 days.
 Fourteen DIV neurons were incubated for 2 h with 1 μ l (1
 mM) fluorescent-labelled antimiR-34a or SC-miR-34a diluted
 in 400-ml medium, followed by 10-min fixation by 4% para-
 formaldehyde/4% sucrose in PBS at room temperature. After
 fixation cells were washed three times in PBS for 30 min at
 room temperature. Slides were mounted using FluoroGold
 mounting medium with DAPI (Invitrogen) and image acquired
 on Axio imager 2 Zeiss light fluorescent microscope.

Q-PCR

To validate the inhibition of microRNAs after antimiR-34a
 transfection, RNA enriched in small RNAs was purified using
 PureLink® miRNA Isolation Kit (Life technologies, Invitrogen,
 Carlsbad, CA, USA). Changes in mature microRNA levels were
 determined using the TaqMan® MicroRNA Reverse
 Transcription Kit and TaqMan® microRNA Assays (Applied
 Biosystems, Foster City, CA) according to the manufacturer's
 protocol. Fifteen microliters of cDNA was generated from 30 ng
 of total RNA, and 3 μ l of a 15-fold dilution was used for real-
 time PCR reactions. The data was normalized to Y1. Changes in
 relative concentration were calculated with the second derivative
 maximum method $2^{-\Delta\text{CT}}$. ΔCT was calculated by subtracting the
 CT of the geometric mean of the two housekeeping genes from
 the CT of the gene of interest.

253	RNA Preparation		
254	Micro-dissected dentate gyri were homogenized in a	Given that miR-34a is implicated in the regulation of synaptic	302
255	Dounce homogenizer on ice in lysis buffer (20 mM	protein targets, it presents as an attractive candidate to manip-	303
256	Tris, pH 7.5, 150 mM NaCl, 2 mM MgCl ₂ , 1 mM	ulate for investigation of synaptic function and plasticity. In	304
257	NaF, 0.5% DTT, 2 mM EDTA, protease inhibitor,	order to investigate the function of miR-34a the dentate gyrus	305
258	RNAse inhibitor, DEPC water), and protein concentra-	in vivo, initial experiments were carried out in vitro for veri-	306
259	tion was determined using the Pierce BCA protein assay	fication of anti-miR uptake in cells, cellular distribution and	307
260	reagent (Thermo Scientific, Pierce).	efficiency of miRNA knockdown.	308
261	SDS-PAGE and Western Blotting		
262	Equal protein amounts (40 µg) were loaded onto 8% or	AntimiR Downregulates Endogenous miRNA 34a	309
263	10% SDS-PAGE gels and run for 2 h at a constant	First, we tested endogenous knockdown of miR-34a in NRK	310
264	voltage of 100 V. Separated proteins were subjected to	cells. RNA enriched for small RNAs was isolated 48 h after	311
265	western blotting and transferred to HyBond ECL nitro-	transfection of anti-miR-34a and scrambled-miR-34a (SC-	312
266	cellulose membrane (Amersham, Little Chalfont, UK) at	miR-34a) control, and the level of unbound microRNA was	313
267	a constant voltage of 100 V for 1.5 h. Membranes were	assayed by RT-PCR. Approximately 8% of miR-34a remained	314
268	stained with Ponceau to check for proper transfer	unblocked after specific anti-miR-34a transfection compared	315
269	followed by blocking with buffer (5% BSA, 0.1%	to control (Fig. 1a), demonstrating a 92% knockdown of en-	316
270	Tween and Tris-buffered saline (TBST) for 1 h on a	dogenous miR-34a in vitro. Y1 RNA was used for normaliza-	317
271	gyro-rocker at room temperature. The primary and sec-	tion. Functional miRNA inhibition was also assessed in den-	318
272	ondary antibodies were diluted in 5% BSA in 0.1%	tate gyrus in vivo. Taqman qPCR was performed on dentate	319
273	TBST, and 0.1% TBST, respectively. The membranes	gyrus total homogenates 2 h after intrahippocampal infusion	320
274	were incubated with primary antibody over night at 4	of SC-miR-34a or anti-miR-34a (sequences listed in	321
275	°C on a gyro-rocker. Following three washes of TBST,	Supplementary data Table S1). Results revealed that	322
276	blots were incubated for 1 h in horseradish peroxidase-	anti-miR-34a downregulated endogenous miR-34a 0.5-fold	323
277	conjugated secondary antibody dissolved in TBST at	compared to SC-miR-34a control (Fig. 1b). Significance was	324
278	room temperature. The blots were washed three times	tested by one-way ANOVA (* <i>p</i> < 0.05). These results indicate	325
279	with TBST, and proteins were visualized using en-	that anti-miR-34a is a potent inhibitor of endogenous miR-34a	326
280	hanced chemiluminescence (ECL Western Blotting	in cell cultures in vitro and in the hippocampus of live rats.	327
281	Analysis System, Amersham Pharmacia Biotech,		
282	Norway). The blots were scanned using Gel DOC	AntimiR-34a Is Detected in Cell Bodies and Dendrites	328
283	XRS+ (BIO RAD), and densitometric analyses were	of Hippocampal Neurons	329
284	performed with ImageJ software (NIH, Bethesda, MD).	To evaluate neuronal uptake, distribution and localization, an	330
285	miRNA Target Prediction	anti-miR-34a uptake experiment was conducted using cultured	331
286	miRNA target prediction sources have been described	primary hippocampal neurons. Anti-miR-34a and scrambled-	332
287	elsewhere [9]. To identify the putative target genes of	miR-34a were labelled with green fluorescent dye (FAM-6,	333
288	each miRNA, we first queried four of the most widely	Exiqon) and added (1 µl, 1 mM) to the culture medium (400	334
289	used target prediction sources: DIANA [31], miRanda	ml) followed by an incubation period of 2 h. Fluorescence	335
290	[32], TargetScan [33] and PicTar [34]. We quantified	microscopy showed uptake of FAM-labelled anti-miR-34a	336
291	the agreement between predicted target lists using	and SC-miR-34a by primary hippocampal neurons. FAM-6-	337
292	Rank Product (RP) analysis [35]. Each gene was or-	anti-miR-34a distribution was somatodendritic similar to en-	338
293	dered by quality score and the geometric mean of the	dogenous miR34a (Fig. 2), whereas FAM labelled SC-miR-	339
294	gene rank calculated across prediction sources. Missing	34a was more soma restricted. These results indicate effective	340
295	ranks were imputed for target genes missing only one	neuronal uptake and somatodendritic distribution of anti-miR-	341
296	rank value, and genes missing more source values were	34a.	342
297	discarded. To assess the robustness of the computed		
298	ranks, we performed a bootstrap analysis with 1000	miRNA-34a Acutely Regulates Synaptic Efficacy	343
299	permutations of rank order using the Bioconductor	We asked whether miR-34a is involved in the regulation of	344
300	RankProd package [36].	basal synaptic efficacy in the dentate gyrus of the adult anes-	345
		thetized rat. Anti-miR-34a and scrambled anti-miR-34a were	346

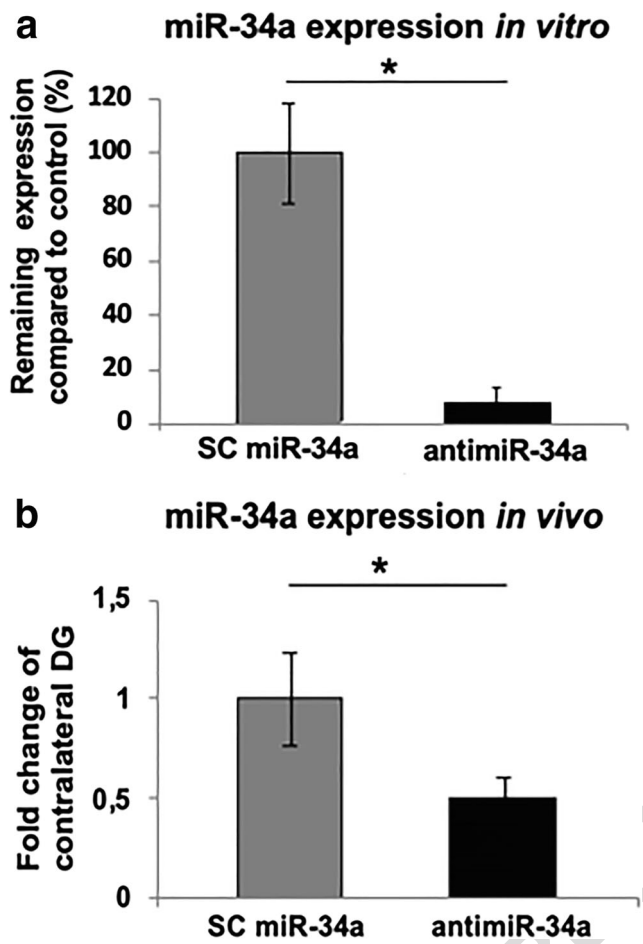


Fig. 1 AntimiR downregulates endogenous microRNA. **a** RNA enriched for small RNAs was isolated 48 h after transfection of antimiR and scrambled control, and the level of unbound microRNAs was assayed by real-time PCR. Approximately 8% of miR-34a remained unblocked after specific antimiR-34a transfection compared to scrambled control. Y1 was used for normalization. **b** Taqman qPCR was performed on dentate gyrus total homogenates 2 h after infusion of scrambled antimiR-34a or antimiR-34a. Bar graphs shows relative miRNA-34a expression levels post-antimiR-34a ($n = 6$) and SC-antimiR-34a ($n = 4$) treatment. AntimiR-34a reduced endogenous miR-34a levels by 50% compared to SC-miR-34a control. Normalised to Y1 and SNO. Values are means of \pm S.E.M. One-way ANOVA analysis was used to test significance between groups ($*p < 0.05$)

347 infused into deep stratum lacunosum-moleculare of dorsal
 348 CA1, immediately above the dentate gyrus (1 μ l, 12 min).
 349 Infusion of SC-antimiR-34a (grey circles, $n = 13$) served as
 350 a control to distinguish sequence-specific silencing from non-
 351 specific effects. Figure 3a shows the experimental design. No
 352 effects were observed on MPP-evoked responses post-
 353 infusion of SC-antimiR-34a throughout the recording
 354 (Fig. 3b). Strikingly, infusion of antimiR-34a (black circles,
 355 $n = 12$) elicited a decrease of the MPP-evoked
 356 fEPSP. Already ~ 9 min after completing antimiR-34a infusion,
 357 the fEPSP slope started to progressively decrease. During the
 358 140-min recording post-infusion, the fEPSP decreased to \sim
 359 60% below baseline (Fig. 3b, c). Input-output curves obtained

during baseline (1), 30 min post-infusion (2) and 150 min after
 360 time point of infusion (3) (Fig. 3e) show a decrease in synaptic
 361 transmission across a range of stimulus intensities. Figure 3d
 362 shows representative sweeps collected at five time points be-
 363 fore and after infusion (as labelled in Fig. 3b).
 364

To further assess the specificity of the antimiR-34a effect,
 365 rats were infused with mismatch-antimiRNA-34a (MM-
 366 antimiR-34a, striped triangles, $n = 6$) harbouring nucleotide
 367 mismatches at nucleotides 2, 6, and 10 in the 15-nt sequence
 368 (Supplementary S1). Similar to scrambled miR-34a, infusion
 369 of MM-antimiR-34a did not impact field EPSP slope values
 370 across the duration of recording (Fig. 3b), and no difference
 371 was found between pre and post-infusion input-output curves
 372 (Fig. 3e) ($p > 0.05$). These results indicate that perfect base pair
 373 complementarity at miR-34a nucleotides 2, 6 and 10 are crucial
 374 for inhibition of endogenous miR-34a and acute downregulation
 375 of synaptic transmission. Previously, we identified Arc-targeting
 376 miRNAs and showed that overexpression of both miR-34a and
 377 miR-193 inhibits Arc expression in cultured hippocampal neu-
 378 rons. In adult dentate gyrus, miR-193 has a somatodendritic
 379 expression profile similar to miR-34a. However, as shown in
 380 Fig. 3a, infusion of antimiR-193 (white circles, $n = 6$) did not
 381 elicit a change in MPP-evoked fEPSPs. The robust effect of
 382 antimiR-34a and complete lack of effect of SC-miR-34a, MM-
 383 antimiR-34a, and antimiR-193 indicates a profound role of en-
 384 dogenous miR-34a in regulation of basal synaptic efficacy
 385 in vivo. Western blots on total homogenates showed that
 386 antimiR-34a enhances expression of Arc 3.1-fold in the ipsilat-
 387 eral infused dentate gyrus relative to control (Fig. 5a, b), whereas
 388 scrambled miR-34a and mismatched antimiR-34a controls had
 389 no significant effect ($n = 6$, $p > 0.05$).
 390

LTP Induction Blocks Antimir-34a-Mediated Depression

Next, we sought to inhibit endogenous miR-34a during LTP in
 393 the dentate gyrus of the adult anaesthetised rat in vivo. Our
 394 previous work has shown that Arc translation is necessary for
 395 LTP consolidation [26] and that miR-34a downregulates Arc
 396 protein in vitro [5]. Given the profound depression of synaptic
 397 transmission elicited by antimiR-34a at baseline, we wanted to
 398 assess the effect of antimiR-34a infusion during the mainte-
 399 nance phase of LTP. We speculated that if miR-34a is derepressed
 400 following high-frequency stimulation (HFS)-induced LTP, the
 401 effect of antimiR-34a infusion during the LTP maintenance
 402 phase may be attenuated. Figure 4a shows the experimental
 403 design. HFS (400 Hz, 8-pulse bursts) of the medial perforant
 404 path (MPP) generated a lasting increase in the slope of the
 405 fEPSP. AntimiR-34a or SC-miR-34a was infused (1 μ l/12 min)
 406 at 2 h post-HFS, and recording was continued to 4 h post-HFS.
 407 Time course plots are shown in Fig 4c. Remarkably, infusion
 408 of antimiR-34a during LTP maintenance had no effect on fEPSPs
 409 recording over the 2-h post-
 410

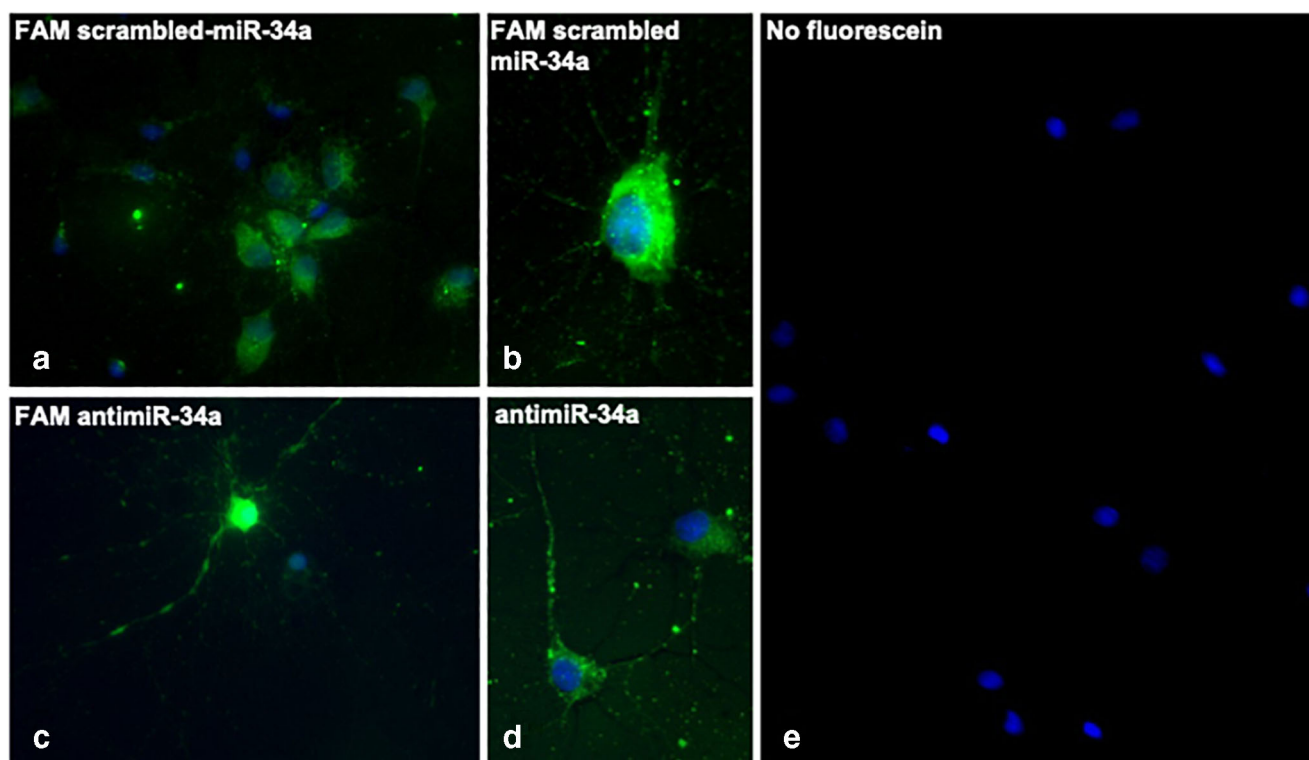


Fig. 2 AntimiR-34a is detected in cell bodies and dendrites of hippocampal neurons. Cultured hippocampal neurons (DIV 14) were treated with fluorescent antimiR-34a or scrambled antimiR-34a for 2 h. Expression was assessed by light microscopy. **a, b** Representative image of cells treated with fluorescent scrambled miR-34a. **c, d** Representative images of cells treated with antimiR-34a. AntimiR-34a is detected in the somata and the dendrites. **e** Dapi stained control, no fluorescence

411 infusion period, and there was no difference between antimiR-
 412 34a and SC-miR-34a groups in LTP magnitude recorded 4 h
 413 post-HFS. Figure 4b shows representative sweeps collected at
 414 five different time points before and after HFS and drug infu-
 415 sion (indicated in Fig. 4c). Western blot analysis performed on
 416 whole dentate gyrus lysates confirmed upregulation of Arc
 417 protein 4 h post-HFS, but there was no significant dif-
 418 ferences in Arc levels between antimiR-34a ($n = 5$) and
 419 SC-miR-34a ($n = 6$) infused rats (Fig. 4d). Significance
 420 was tested by one-way ANOVA ($*p < 0.05$).

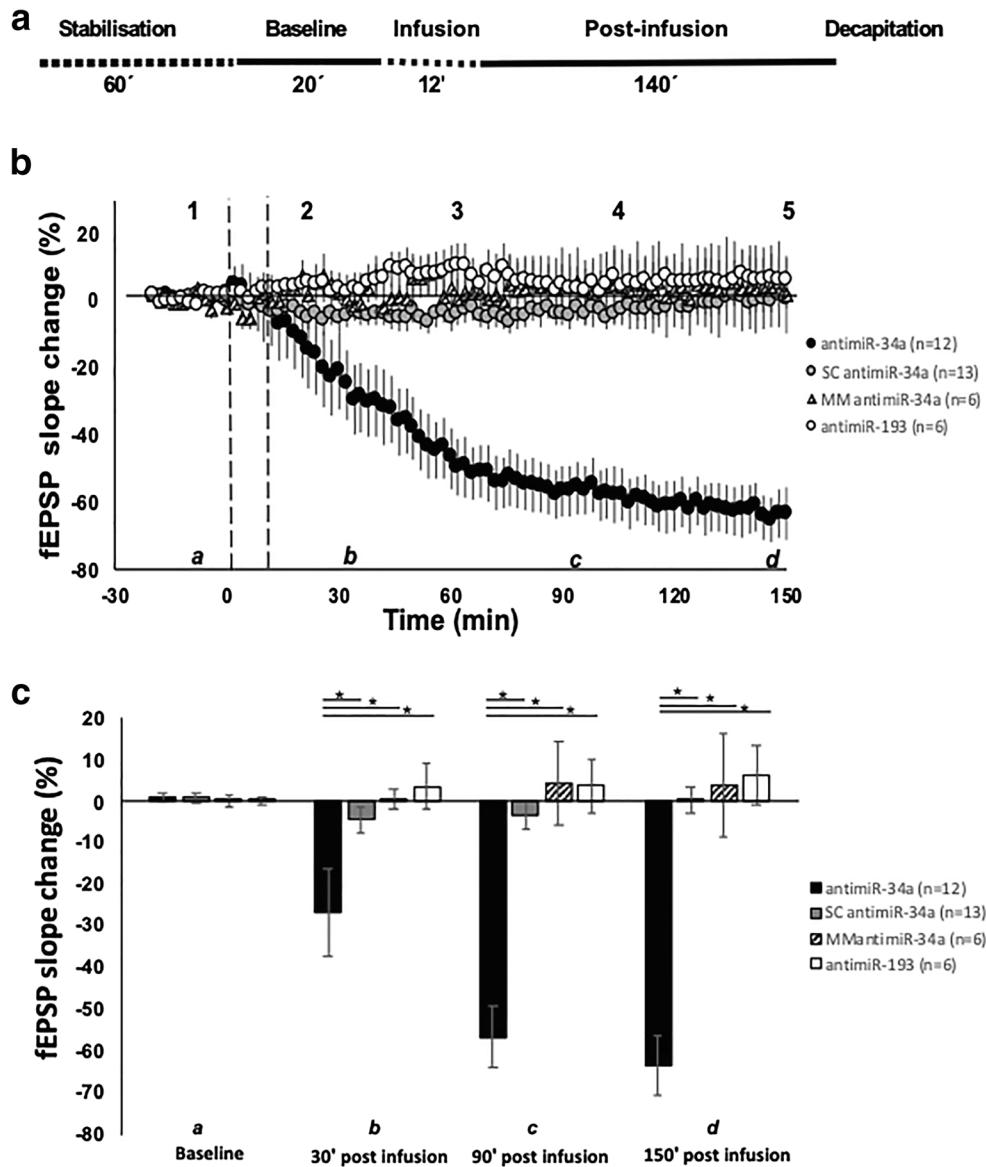
421 mRNA Target Prediction

422 To better understand the downstream effects of antimiR infu-
 423 sion, we integrated predictions from four miRNA target pre-
 424 diction resources: PicTar [34], TargetScan [33], Miranda [32]
 425 and DIANA [31]. We used the rank product method [35, 36]
 426 where ranks are aggregated across prediction resources using
 427 a geometric mean where a gene must appear in at least two
 428 sources. This allowed us to mitigate for the poor agreement
 429 normally found between different target gene prediction algo-
 430 rithms. miR34a was predicted to target 397 genes
 431 (Supplementary S2). Using the Bioconductor KEGG profile
 432 package, the common targets were predicted to be involved in
 433 96 pathways (Supplementary S3), including regulation of the

actin cytoskeleton, LTD, MAPK signalling, axon guidance 434
 and the calcium signalling pathway. Five predicted targets 435
 were selected as candidates for Western blot analysis: 436
 Bcl11b/Ctip2 (from now on referred to as Ctip2), Ube2g-11, 437
 TCF4, PKI-1 α and Gnao1. Ctip2, COUP-TF interacting pro- 438
 tein 2, is expressed predominantly in central nervous system 439
 (CNS). In the dentate gyrus, loss of Ctip2 expression selectiv- 440
 ity impairs spatial working memory [37]. TCF4 is a transcrip- 441
 tion factor known to regulate synaptic plasticity and memory 442
 function [38]. Ube2g1 is involved in mechanisms targeting 443
 abnormal or short-lived proteins for degradation [39]. PKI- 444
 1 α negatively modulates synaptic activity [40]. Gnao1 is 445
 brain-enriched and mutations in the gene are associated with 446
 epileptic encephalopathy [41]. 447

448 miRNA-34a Regulates Multiple Targets in Addition 449 to Arc in the Dentate Gyrus In vivo

450 Next, we investigated the effects of acute intrahippocampal 450
 infusion of antimiR-34a on Arc protein expression along with 451
 the five new predicted miR-34a targets Ctip2, PKI-1 α , Ube2g- 452
 11, TCF4 and Gnao1 protein expression. Dentate gyri were 453
 micro-dissected 140 min after infusion of antimiR-34a or con- 454
 trol sequences. Replicating results from Fig. 3, antimiR-34a 455
 infusion elicited a stable decrease of the fEPSP slope to ~ 456



Q3 **Fig. 3** miRNA-34a regulates synaptic efficacy, and variation in sequence alters target recognition. **a** Experimental design. **b** Time course plots show changes in the medial perforant path-evoked fEPSP slope expressed in percentage change from baseline. Values are means \pm S.E.M. Infusion of scrambled anti-miR-34a and anti-miR-34a is indicated by vertical, dotted lines. Synaptic efficacy is rapidly and gradually reduced to $\sim 60\%$ after local infusion (1 μ l, 1 mM) of anti-miR-34a (black circles, $n = 12$). Infusion of the multiple controls scrambled anti-miR-34a (grey circles, $n = 13$), or mismatched anti-miR-34a (striped triangles, $n = 6$) or anti-miR-193 (white circles, $n = 6$), had no effect on synaptic efficacy. **c** Bar graph of % fEPSP slope change from baseline. **d** Representative progression sweeps of scrambled anti-miR-34a, anti-miR-

34a, anti-miR-193, mismatch anti-miR-34a during baseline (1), 20 min post-infusion (2), 60 min post-infusion (3), 90 min post-infusion (4), and 120 min post-infusion (5). Averaged field potentials traces (20 sweeps) collected at the beginning of baseline recording (1), 20 min after infusion (2), 60 min post-infusion (3), 90 min post-infusion (4), and 150 min post-infusion (5). Scale bars 5 mV, 2 ms. **e** Input-output curves (average of 4 sweeps) collected at baseline (20 min), post-infusion (30 min) and at the end of the experiment (150 min). Significance was tested by factorial ANOVA, and post hoc tests. A probability level of $p < 0.05$ was considered statistically significant ($b = *p < 0.05$, $c = **p < 0.01$, and $d = ***p < 0.001$)

457 60% of baseline. Western blot analysis of dentate gyrus homog-
 458 enates showed significantly enhanced expression of Arc (3.1-
 459 fold, $n = 6$, $p > 0.05$), Ctip2 (1.6-fold, $n = 13$, $p > 0.05$), PKI-1 α
 460 (2.2-fold, $n = 10$, $p > 0.05$), Ube2g-1 (1.5-fold, $n = 10$, $p > 0.05$)
 461 and TCF4 (2.6-fold, $n = 6$, $p > 0.05$) in the anti-miR-infused
 462 dentate gyrus relative to the contralateral dentate gyrus, which

was significantly different from the MM-anti-miR-34a and SC-
 463 miR-34a controls (Fig. 5a, b). There was no statistically signif-
 464 icant alteration in Gnao1 expression ($n = 12$, $p < 0.05$), or
 465 difference between groups indicating that Gnao1 is not regulat-
 466 ed by miR-34a under the present conditions. GAPDH was used
 467 as a loading control and for normalisation. 468

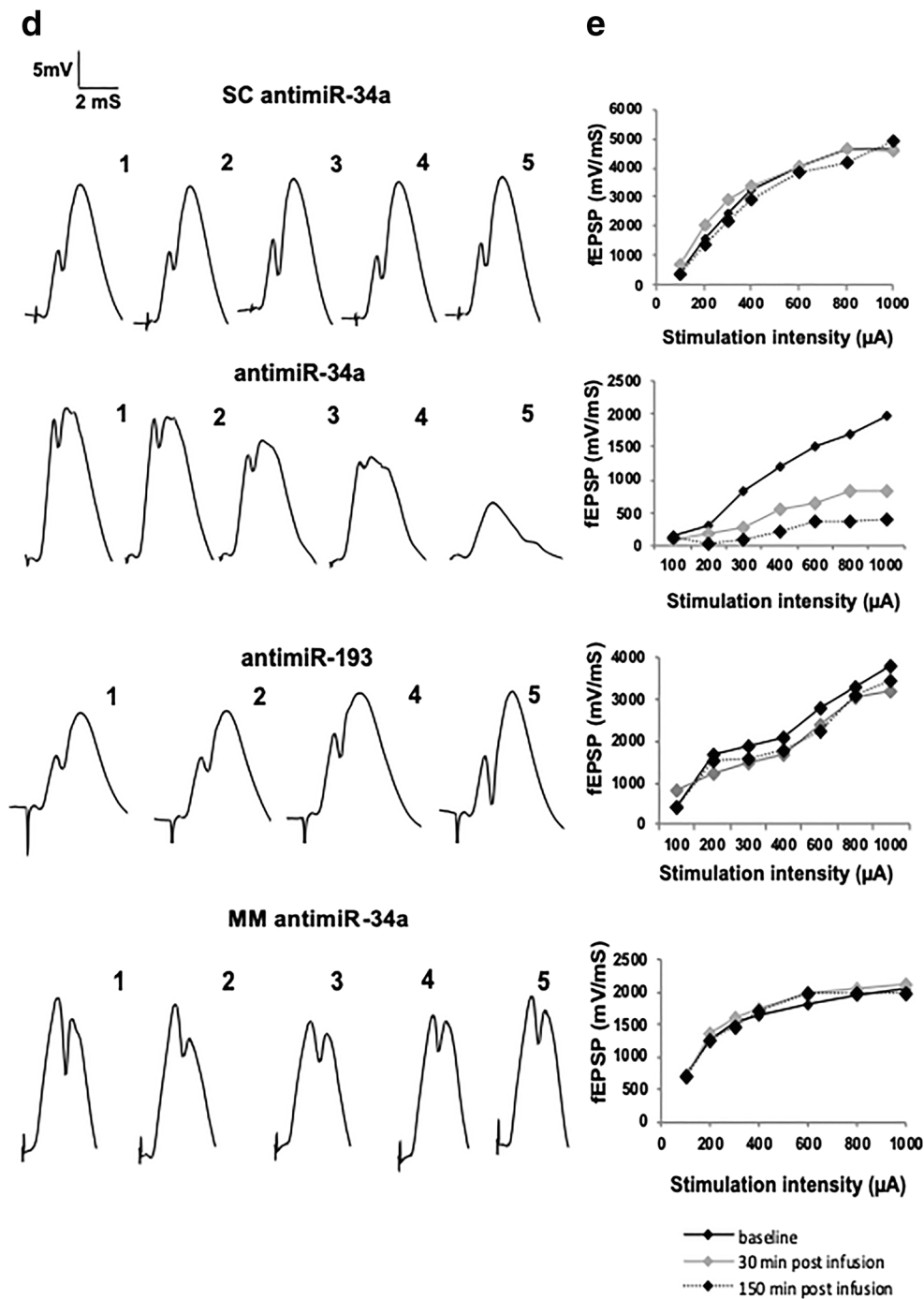


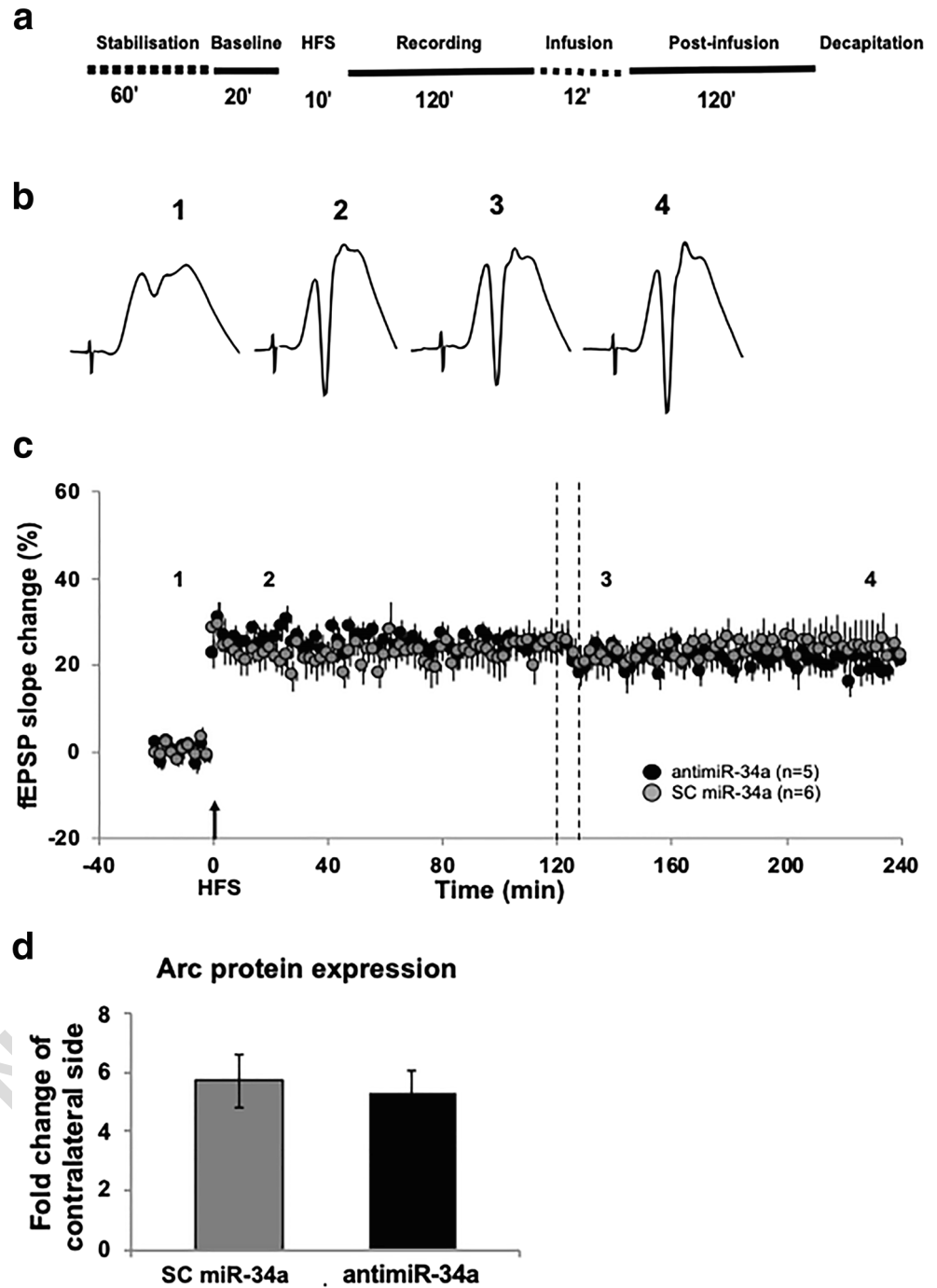
Fig. 3 (Continued)

469 **Discussion**

470 Synaptic plasticity such as LTP, LTD and homeostatic synap-
 471 tic scaling are alterations of synaptic transmission and efficacy
 472 in response to neural activity. Long-lasting activity-dependent
 473 modifications of synapses typically require de novo protein
 474 synthesis. In addition to somatic protein synthesis, translation
 475 of mRNA in dendritic processes is important for synaptic

homeostasis and plasticity [42, 43]. Many miRNAs are involved
 476 in the spatial-temporal control of neuronal protein syn-
 477 thesis and regulation of synaptic plasticity. MicroRNA func-
 478 tion has been elucidated primarily through loss-of-function
 479 approaches in which a specific miRNA is deleted or chroni-
 480 cally inhibited, and by viral vector-mediated miRNA overex-
 481 pression [7]. However, the regulation of protein synthesis dur-
 482 ing activity-dependent synaptic plasticity operates on a
 483

Fig. 4 LTP induction blocks miRNA-34a-mediated depression of synaptic transmission. **a** Experimental timeline. **b** Representative sweeps collected at baseline (20 min) (1), 5 min post-HFS (2), 130 min post-HFS (3) and 240 min post-HFS (4). **c** Time course plots show changes in the medial perforant path-evoked fEPSP slope expressed in percentage of baseline. Values are means \pm S.E.M. Scale bars 5 mV, 2 ms. Vertical dotted lines indicate infusion (1 μ L/12 min). HFS is indicated by black arrow (antimiR-34a, $n = 5$; SC-miR-34a, $n = 6$). Dentate gyrus tissue was collected 2 h post-infusion. **d** Western blots were performed in dentate gyrus homogenates 4 h after HFS and 2 h after infusion of antimiR-34a or SC-miR-34a during LTP. Arc protein is significantly enhanced during LTP, but there is no significant difference between the two groups ($n = 4$ in both groups)

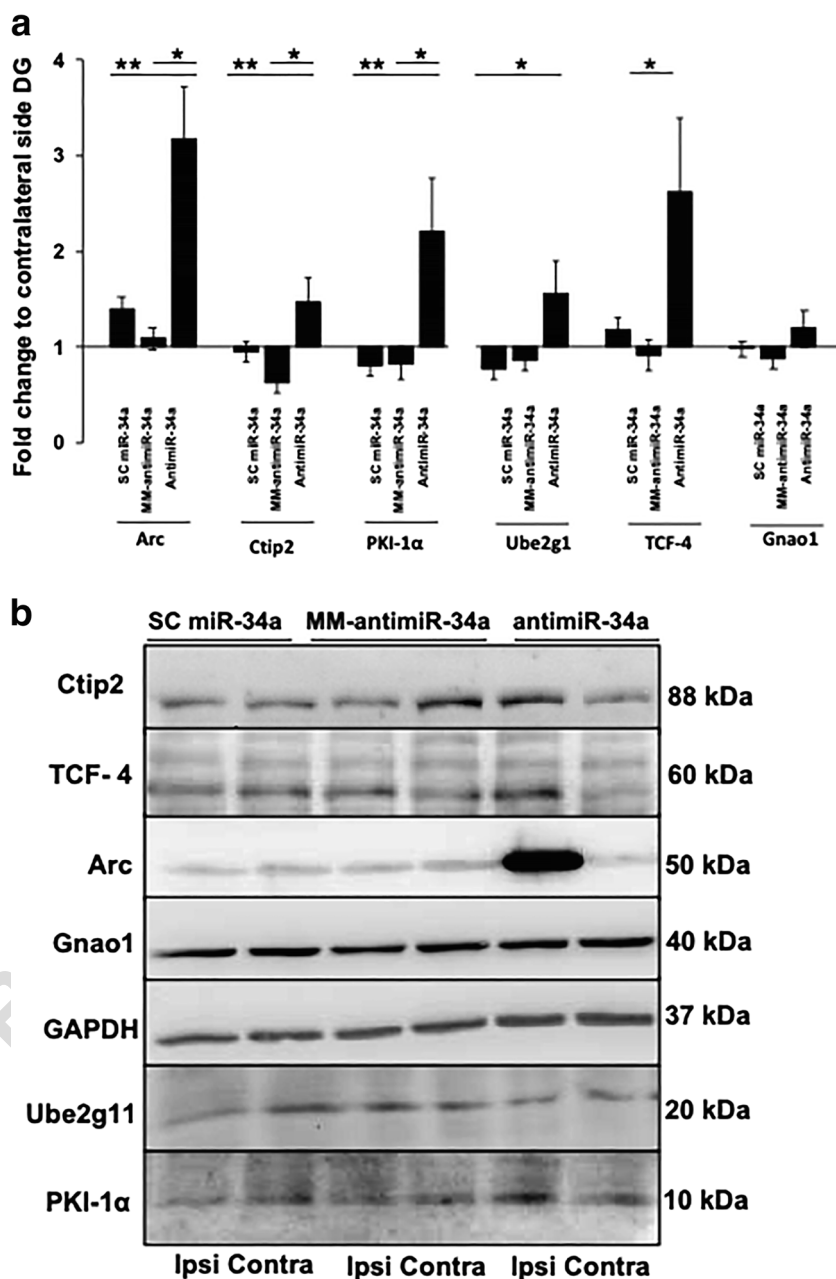


484 timescale of minutes. To our knowledge, the present study is
 485 the first to report acute regulation of synaptic efficacy in the
 486 adult brain by a miRNA.

487 We found that brief (12 min) intrahippocampal infusion of
 488 antimiR-34a results in a rapid and persistent decrease in
 489 MMP-evoked synaptic transmission in the dentate gyrus.
 490 The decrease in fEPSP started at approximately 9 min after
 491 completing the infusion and developed gradually, reaching a
 492 stable 60% decrease relative to pre-infusion baseline. This
 493 modulation was absent when a scrambled or mismatched

miR-34a sequence was introduced. Levels of endogenous
 494 antimiR-34a in the dentate gyrus were downregulated by
 495 antimiR-34a, but not by infusion of scrambled control
 496 sequence. Furthermore, antimiR-34a but not control
 497 sequences resulted in upregulation of Arc, a known
 498 miR-34a target and major regulator of synaptic
 499 plasticity. AntimiR-34a infusion similarly resulted in
 500 sequence-specific upregulation of several novel
 501 predicted miR34a targets (Ctip2, PKI-1 α , TCF4 and
 502 Ube2g) with roles in neuronal and synaptic function.
 503 This suggests that endogenous miR-34a potentially regulates synaptic

Fig. 5 AntimiR-34a infusion enhances synaptic protein expression during basal conditions. Western blots were performed in dentate gyrus homogenates 2 h after infusion of scrambled antimiR-34a, antimiR-34a, and mismatched antimiR-34a during baseline conditions. Immunoblots on total homogenates (left/right hemisphere ratio) showed that antimiR-34a enhances expression of different protein. **a** Mean and standard deviation of the protein levels detected in western blotting. One-way ANOVA with Tukey's post hoc analysis was used to test significance between groups compared to control ($*p < 0.05$, $**p < 0.01$). **b** Representative blots of dentate gyrus protein expression 2 h post-infusion of scrambled antimiR-34a, antimiR-34a, and mismatched antimiR-34a, relative to right hemisphere control, during baseline conditions



504 transmission through sequence-specific inhibition of one or
 505 more target RNAs. Release of the tonic inhibition mediated
 506 by miR-34a resulted in profound depression of synaptic trans-
 507 mission. Remarkably, antimiR-34a had no effect on synaptic
 508 efficacy when infused at 2 h post-HFS, during the process of
 509 Arc-dependent LTP consolidation.

510 **MiR-34a Regulation of Arc and Synaptic Efficacy**
 511 **In vivo**

512 The speed at which antimiR-34a impacts synaptic transmis-
 513 sion suggests that the effective target mRNAs are located near
 514 synapses, such that newly synthesized proteins can exert a

nearly immediate effect on synaptic efficacy. Arc is a miR-34a target that may fit such a role.

The diversity of Arc function in synaptic plasticity is a truly remarkable phenomenon. Arc scales neuronal action and controls excitation and inhibition through bidirectional regulation of synaptic strength. Synaptic activity can converge to alter Arc transcription and then diverge to induce different plasticity outcomes, such as AMPA-receptor endocytosis promoting LTD or actin cytoskeletal modulation promoting LTP [26, 42, 43]. In this study, we initially chose to explore miR-34a function in vivo due to its Arc-targeting properties shown in vitro [5]. We hypothesised that knockdown of miR-34a would enhance Arc expression and alter synaptic efficacy. Here,

528 infusion of anti-miR-34a increased Arc expression 3-fold and
529 acutely depressed MMP-evoked fEPSPs.

530 Arc is an immediate early gene with low levels of basal
531 expression [44]. Stimulus-evoked Arc transcription may occur
532 within minutes [45, 46], and a large fraction of the new
533 mRNA is transported into dendrites where it accumulates in
534 region of activated synapses, presumably to be translated lo-
535 cally [47]. Because Arc mRNA is subject to translation-
536 dependent degradation, it needs to be translationally repressed
537 in order to reach synapses on distal dendrites [48–50]. The
538 precise dendritic spatiotemporal mechanisms regulating Arc
539 mRNA translation and repression of translation are yet to be
540 elucidated. However, in the CA1 region of the hippocampus,
541 application of the metabotropic glutamate receptor agonist
542 (S)-3,5-dihydroxyphenylglycine (DHPG) induces a
543 transcription-independent LTD that requires translation of
544 dendritic Arc and expression of protein within 10 min of treat-
545 ment [51]. In isolated synaptoneurosome preparations,
546 in vitro stimulation with BDNF induces Arc protein syn-
547 thesis in a time range of 10–30 min [52, 53]. As men-
548 tioned, Arc translation is necessary for LTP consolida-
549 tion in vivo. The process depends on the provision of
550 newly transcribed Arc, and Panja et al. [26] showed that
551 Arc mRNA shifts from the monosome/mRNP fraction to
552 polysomes during LTP consolidation. Recent imaging
553 studies from hippocampal neuronal cell cultures also
554 suggest that, at the basal state, Arc mRNA in dendrites
555 is held quiescent in stalled polysomes. In response to
556 glutamate stimulation, Arc is translated in less than
557 1 min [54]. Given the profound effect of anti-miR-34a
558 at the basal state, it is tempting to consider that en-
559 hanced expression of Arc and other miR34a targets is
560 due to activation of mRNA on stalled polysomes. Both
561 Arc mRNA and miR-34a are found in the excitatory
562 synaptic compartment of the dentate gyrus at the
563 unstimulated basal state [5].

564 The selective modulation of basal transmission by anti-miR-
565 34a, with no effect on established LTP, is extraordinary and
566 the underlying mechanism is unknown. The lack of effect
567 during the LTP state might be due to (1) inability of
568 anti-miR-34a to access miR-34a or (2) degradation of
569 the relevant miR-34a pool. Previously we reported that
570 LTP induction in the dentate gyrus of urethane-
571 anesthetized rats is associated with increased, NMDAR-
572 dependent association of miR-34a with the Ago2/RISC [9].
573 Perhaps this process of miR-34a loading onto Ago2 inhibits
574 anti-miR binding. Assuming miR-34a regulates a preexisting
575 pool of Arc mRNA on stalled polysomes, it is possible that
576 miR-34a is degraded or sequestered in P-bodies following
577 HFS-induced translation. In this case, the basis for anti-miR-
578 34a induced synaptic depression will be lost. A third possibil-
579 ity, so far unexplored, is that newly synthesized Arc mRNA
580 [53] is immediately translated, thus bypassing the stage of

miR-34a binding and repression. In terms of the action of
the Arc protein itself, to enhance or depress synaptic trans-
mission, this is known to be dependent on the cell sig-
naling context [25].

miR-34a Regulation of Novel Predicted Targets: Ctip2, TCF4, PKI1- α and Ube2g1

The bioinformatic mRNA target prediction and the in vivo
evidence allows us to suggest four novel miR-34a targets.
However, they are not formally identified as targets by muta-
tion of miRNA binding sites. Here, we show that acute inhi-
bition of miR-34a initiates acute upregulation of Arc along
with four additional bioinformatically predicted targets of
miR-34a: Ctip2, PKI-1 α , TCF4 and Ube2g.

Ctip2 is a C₂H₂ zinc-finger transcription factor, highly
expressed in dentate granule cells. Selective ablation of
Ctip2 in the adult dentate gyrus results in morphological
changes leading to functional impairment [39]. Experiments
in cultured striatal cell lines indicate that Ctip2 regulates a
multitude of genes, and enhances brain-derived neurotrophic
factor (BDNF) signalling [55] which activates cascades such
as PLC/PKC, PI3K/Akt, Ras/Erk, AMPK/ACC and NF κ B
pathways [56]. BDNF plays a critical role in plasticity at glu-
tamatergic and GABAergic synapses by both pre- and post-
synaptic mechanisms [57] and contributes to both acute and
homeostatic alterations in hippocampal synaptic function
[58]. BDNF is also known to enhance Arc expression. Thus,
Ctip2 regulation could potentially impact synaptic transmis-
sion indirectly. However, the rapid decrease in synaptic trans-
mission is less likely to involve transcription and regulation of
BDNF secretion and signalling.

TCF4 is a transcription factor known to regulate synaptic
plasticity and memory function [38]. TCF4 haploinsufficient
mice has been studied as an animal model of autism spectrum
disorder. These TCF4-deficient mice have impaired Arc ex-
pression and enhanced LTP induction in hippocampal region
CA1. In our investigation of anti-miR-34a effect, expression of
TCF4 and Arc protein were both enhanced while basal syn-
aptic transmission was depressed. Though the mechanisms are
unknown, the opposite effects on synaptic transmission are
consistent with a homeostatic role of Arc. Interestingly, abla-
tion of TCF4 results in downregulation Ube2g1 gene tran-
scription, another predicted target of miR-34a [59]. Our west-
ern blots on total homogenates showed that infusion of
anti-miR-34a enhances Ube2g1 expression 1.5-fold during
basal conditions. However, Ube2g1 is not a dendritically lo-
cated transcript [60]. This indicates that our observed increase
in Ube2g1 expression may be secondary to enhanced TCF4
expression, not a result of direct miR-34a-mediated mRNA
derepression. Mechanisms involving somatic translation of
Ube2g1 do not present a likely explanation for the observed
acute modulation of synaptic efficacy.

632 PKI-1 α is a member of the cAMP-dependent protein ki- 683
 633 nase (PKA) inhibitor family. PKA enhances excitatory synap- 684
 634 tic transmission in the dentate gyrus [61], whereas PKI nega- 685
 635 tively modulates synaptic activity and regulates gene expres- 686
 636 sion induced by PKA [61]. In the dentate gyrus, PKI1 α 687
 637 mRNA and protein is down regulated by strong depolarization 688
 638 [62]. However, chronic infusion of antisense oligonucleotides 689
 639 against PKI α into the rat brain results in a dramatic reduction 690
 640 of excitability and ability to exhibit LTP and LTD [62]. Here, 691
 641 we demonstrate that infusion of antimiR-34a during baseline 692
 642 conditions results in a 2.2-fold increase in PKI1 α protein ex- 693
 643 pression. However, if PKI1 α is a direct mRNA target regulat- 694
 644 ed by miR-34a, we would expect an effect of antimiR-34a 695
 645 infusion also during LTP. 696

646 Gano1 (G protein subunit alpha 1) constitutes up to 0.5% of 697
 647 cerebral membrane protein [63], and mutations in Gnao1 are 698
 648 linked to epileptic encephalopathy [41], indicating an impor- 699
 649 tant role in brain function. Here, western blots on total homog- 700
 650 enates show that loss of miR-34a action does not initiate syn- 701
 651 thesis of Gano1, indicating that Gano1 is not implicated in 702
 652 acute regulation of synaptic transmission during baseline con- 703
 653 ditions in the dentate gyrus. 704

654 **miR-34a Regulation of Synaptic Transmission**
 655 **in Relation to Brain Disorders**

656 As neuroscientists, a major aim in understanding the function 705
 657 of miR-34a and regulation of its target genes is to contribute 706
 658 to new knowledge that can aid the development of novel treat- 707
 659 ment strategies for people with psychiatric and neurodegener- 708
 660 ative brain disorders. Addressing neuropathology through 709
 661 antimiR strategies may be available in the near future, and 710
 662 the multiple benefits in using antimiR strategies have been 711
 663 described elsewhere [64, 65]. miR-34a targets genes that are 712
 664 linked to synaptic plasticity, energy metabolism and resting 713
 665 state network activity, and it plays a critical regulatory role 714
 666 in neurodegenerative diseases [15–18]. miR-34a’s role in 715
 667 AD is far from elucidated, but there is increasing evidence 716
 668 of miR-34a’s significance. miR-34a up- or downregulation, 717
 669 manipulation by overexpression or abolishment, all aid our 718
 670 work in understanding its memory-related mechanisms. A re- 719
 671 cent study by Sakar et al. showed that miR34-a overexpres- 720
 672 sion induces rapid cognitive impairment and AD-like pathol- 721
 673 ogy in mice [66]. Conversely, others have shown that rats 722
 674 overexpressing miR-34a in the brain have better learning abil- 723
 675 ities and reduced emotionality [21]. AD model mice injected 724
 676 with antimiR targeting the complete miR-34 family ‘rescues’ 725
 677 memory performance [67]. In miR-34a knockout/amyloid 726
 678 precursor protein/presenilin 1 mice (APP/PS1), it has been 727
 679 confirmed that miR-34a is involved in synaptic deficits in 728
 680 AD pathological development, partially due to inhibition of 729
 681 NMDA and AMPA receptor expression [68]. Although the 730
 682 latter experiments are carried out in transgenic animals, they

are consistent with our observed decrease in synaptic efficacy. 683
 In sporadic AD, up-regulated miR-34a in the neocortex ap- 684
 appears to down-regulate SHANK3, a postsynaptic scaffolding 685
 protein essential to post-synaptic structure and function [69, 686
 70]. Others have shown that stable hippocampal miR-34a in- 687
 hibition using adeno-associated virus-delivered miRNA 688
 sponges demonstrates transcriptome changes linked to neuro- 689
 active ligand-receptor transduction in cell communication, 690
 causing decreased capacity of reference memory in mice 691
 [27]. Supporting our findings, bioinformatical analyses de- 692
 scribed by Sarkar et al showed that miR-34a has the ability 693
 to affect molecular processes that are intrinsically linked to the 694
 regulation of pre- and post-synaptic neuronal excitability and 695
 resting state functional connectivity [15]. The same authors 696
 have also described that targets for miR-34a were profoundly 697
 reduced by miR-34a over-expression leading to cognitive de- 698
 cline and disease neuropathology [71]. Recently, it has been 699
 suggested that increasing Arc levels prior to the development 700
 of AD neuropathology could protect against cognitive impair- 701
 ment that accompany AD neuropathology [64]. Our results 702
 revealed that antimiR-34a infusion during basal conditions 703
 acutely depresses MMP-evoked fEPSPs due to sequence- 704
 specific mechanisms, a phenomenon that was absent during 705
 long-term potentiation. Multiple synaptic targets, including 706
 miR-34a target Arc, were upregulated during basal conditions, 707
 adding new knowledge to the complexity of miR-34a function 708
 in synaptic efficacy and plasticity. We realize that the temporal 709
 and spatial distribution of miR-34a and modulation of its tar- 710
 gets is highly heterogeneous and serve multiple synaptic 711
 mechanistic systems and cognitive trajectories. 712

713 **Conclusion**

714 Our results identify a set of miR-34a targets associated with 715
 716 regulation of basal synaptic transmission in the dentate gyrus. 717
 Further work is needed to elucidate the potential causal role of 718
 these specific miR34a targets. The results described in this 719
 study are novel in the field of microRNA regulation and syn- 720
 aptic transmission. Current knowledge on miRNA function in 721
 synaptic plasticity and transmission in vivo is predominately 722
 based on long-term manipulations. To our knowledge, 723
 no other study has demonstrated that miRNAs are in- 724
 volved in acute regulation of synaptic transmission 725
 in vivo. We conclude that miRNAs can generate rapid 726
 neuronal responses and, in this way, are ideally posi-
 tioned to modulate synaptic efficacy.

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- Q3. Figures 3 and 5 contains poor quality of text in image. Otherwise, please provide replacement figure file.
- Q4. References 65 and 72 based on original manuscript we received were identical. Hence, the latter was deleted and reference list and citations were adjusted. Please check if appropriate.
- Q5. Please provide complete bibliographic details of reference [13, 65].

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