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Effects of aging and caloric restriction on dentate gyrus synapses and glutamate receptor subunits

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Abstract

Caloric restriction (CR) attenuates aging-related degenerative processes throughout the body. It is less clear, however, whether CR has a similar effect in the brain, particularly in the hippocampus, an area important for learning and memory processes that often are compromised in aging. In order to evaluate the effect of CR on synapses across lifespan, we quantified synapses stereologically in the middle molecular layer of the dentate gyrus (DG) of young, middle aged, and old Fischer 344 X Brown Norway rats fed *ad libitum* (AL) or a CR diet from 4 months of age. The results indicate that synapses are maintained across lifespan in both AL and CR rats. In light of this stability, we addressed whether aging and CR influence neurotransmitter receptor levels by measuring subunits of NMDA (NR1, NR2A, and NR2B) and AMPA (GluR1, GluR2) receptors in the DG of a second cohort of AL and CR rats across lifespan. The results reveal that the NR1 and GluR1 subunits decline with age in AL, but not CR rats. The absence of an aging-related decline in these subunits in CR rats, however, does not arise from increased levels in old CR rats throughout lifespan, but that are reached in AL rats only in old age.

1. Introduction

Caloric restriction (CR) is regarded as a method of decelerating senescence because of its ability to extend lifespan and to protect against many aging-related degenerative processes and diseases throughout the body [44,55]. Although its effects on the brain are less clear, moderate CR has been shown to attenuate aging-related deficits in performance on hippocampus-dependent learning and memory tasks in some rodent strains, such as Fischer 344 x Brown

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Norway (F344xBN) rats [25,26,43,60,61,73]. This attenuation suggests that CR modulates neurobiological processes that subserve cognition and that a more complete understanding of the impact of CR on the neural basis of learning and memory during aging could lead to therapies for human memory impairment.

In view of recent stereological evidence that the number of principal neurons in the hippocampus does not decrease with age [63,65], attention has shifted to synapses as potential substrates for the aging-related changes that lead to deficits in learning and memory [17,19, 67]. This possibility is supported by the positive correlation between learning and memory ability and the density of synaptic boutons and their spinous targets in the hippocampus [34, 49,52,70]. Similarly, enlargement of the postsynaptic density (PSD) has been associated with enhanced synaptic efficacy. The PSD expands through an NMDA receptor-mediated insertion of AMPA receptors [14,24,38,42,62]. It has been proposed that the enlarged PSD then perforates and the postsynaptic spine ultimately divides to create a multiple synapse bouton (MSB) complex consisting of several synapses on a single presynaptic terminal contacting more than one postsynaptic element [24,38,57]. Because perforated and MSB synapses increase after long-term potentiation (LTP), learning, and environmental enrichment, they are believed to represent morphological correlates of enhanced synaptic efficacy [6,18,21,29,30, 57,76,77].

In the hippocampus, NMDA and AMPA ionotropic glutamate receptors (NMDAR and AMPAR, respectively) are the chief mediators of excitatory synaptic transmission. While NMDARs mediate a major form of LTP, a functional correlate of learning and memory, AMPAR activation accounts for the fast postsynaptic responses to glutamate. Importantly, synaptic plasticity may rely upon AMPARs, since the absence of AMPARs at NMDARpositive synapses renders them functionally silent at resting potential, and NMDAR-mediated LTP at these synapses involves the rapid insertion of AMPARs [3,35,37]. Moreover, genetic and pharmacological studies have demonstrated the critical involvement of these receptors in LTP and hippocampus-dependent learning and memory [66]. In fact, spatial memory ability varies directly with levels of these glutamate receptors and their constituent subunits [1,10, 39,74]. In light of these data, evidence that NMDARs and AMPARs and their subunits decrease with age [9,15,40,41,71,81] has led to suggestions that this decline may contribute to the cognitive impairments observed in old animals. Evidence that CR not only prevents agingrelated deficits in LTP but also prevents aging-related changes in the obligatory NR1 subunit of the NMDAR [6,18,21,29,30,57,76,77], suggests that CR also might prevent the agingrelated decline in other glutamate receptor subunits. This question is important since maintenance of glutamate receptor subunits over lifespan represents a potential mechanism whereby CR could preserve learning and memory after middle age [43].

In the present study, we quantified synapses as well as relative protein levels of NMDAR and AMPAR subunits in the dentate gyrus (DG) of young, middle-aged and old rats fed *ad libitum* (AL) or a CR diet. The DG is the first subregion in the hippocampal trisynaptic pathway and has been shown in both lesion and pharmacological studies to play a critical role in hippocampus-dependent learning and memory [23,27,83]. Furthermore, stereologically quantified synapses in the DG have been reported to decrease across the lifespan [19]. Using unbiased stereological techniques, we determined the synapse per neuron ratio in the DG across life span in AL fed and CR rats to test the hypothesis that there are quantitative, aging-related changes in synapses that are attenuated by CR. In the second part of the study, we used Western blot analysis to address the possibility that CR affects aging-related synaptic changes at the receptor level. We specifically tested the hypotheses that NMDAR and AMPAR subunits decline with age in the DG of AL rats and that CR attenuates this decline. Our findings indicate that neither aging nor CR affects the number of DG synapses per neuron. In contrast with this stability, the NR1 and GluR1 glutamate receptor subunits in the DG demonstrate a decline

across lifespan in AL rats. CR, however, decreases receptor subunits within 6 months to levels that are maintained across lifespan.

2. Methods

2.1. Animal model

The F344xBN F1 hybrid is a widely-used model for studies of aging and CR [36,46,69,82] and has been reported by Markowska and colleagues to demonstrate CR-mediated protection of learning and memory after middle age [43]. The central focus of the present study was to analyze synaptic structural and biochemical changes as a function of aging and CR. In order to avoid potential direct structural changes due to the environmental enrichment of the testing procedure itself, behavioral testing was not included in the present study. Nevertheless, the experimental design was modeled closely after the Markowska study to permit analysis of synaptic changes in a model of CR-induced amelioration of aging-related cognitive decline. For this study, adult male rats were obtained from the NIA Caloric Restriction Colony (Harlan Industries) and were maintained in our facility on a 12-hour light-dark cycle for two months prior to sacrifice. At sacrifice, groups consisted of young (10 months), middle age (18 months), and old (29 months) AL and CR rats with 6 rats per group for electron microscopy studies and 8 rats per group for Western blot analysis. Rats were housed individually in order to monitor daily food intake. The animal facility at the Wake Forest University School of Medicine is fully accredited by the American Association for Accreditation of Laboratory Animal Care and complies with all Public Health Service, National Institutes of Health, and institutional policies and standards for laboratory animal care. All procedures described herein were approved by the institutional Animal Care and Use Committee. The following notations (age/ diet) are used in the Results section to refer to young, middle-aged, or old rats in AL or CR groups: 10AL, 18AL, 29AL, 10CR, 18CR, and 29CR.

2.2. Caloric Restriction

At the NIA facility, all rats were fed AL (NIH-31 diet) until 14 weeks of age, at which time the CR regimen was initiated by incremental reduction of 10% per week over four weeks, reaching full 60% restriction by 17 weeks of age. The vitamin-fortified NIH-31 diet fed to CR rats provided 60% of the calories and 100% of the vitamins consumed by AL rats (NIH-31). All rats were fed daily, shortly before dark cycle onset, and had free access to water. Daily monitoring of food consumption verified the caloric consumption by each group. Rats were weighed weekly, immediately prior to feeding.

2.3. Electron Microscopy

For electron microscopy studies, rats were anesthetized with ketamine/xylazine (80 mg/kg/12 mg/kg, i.m.) and then perfused transcardially with 1.3 M cacodylate buffer (pH 7.4, 20 ml), followed by fixative (2% paraformaledyde/2% glutaraldehyde), at a flow rate of 40 ml/min for a minimum of 30 minutes using a peristaltic pump. The brains were removed and post-fixed overnight at 4°C. The next day, the brain was sectioned coronally on a vibratome at 200 µm. A similar number of sections was derived through the hippocampus of all animals, indicating no gross evidence of a difference in hippocampal size across ages and diets. Sections were viewed with a dissecting microscope and a lightfield-darkfield base and trapezoids containing the DG were cut from sections containing the dorsomedial hippocampus (Bregma -2.8 through -4.0, [59]). Analysis was restricted to the dorsomedial hippocampus because, in contrast with the ventrolateral portion, this area is crucial for spatial memory acquisition and retrieval in the Morris water maze [31,50,51]. The six resulting blocks were washed thoroughly in 0.15M cacodylate buffer, postfixed in 2% osmium tetroxide, dehydrated, embedded in Araldite 502 plastic (Ted Pella). Two of the six blocks were selected randomly, resulting in an equal probability of sampling along the rostrocaudal axis and were trimmed so that the middle

of the MML of the supragranular blade of the dentate gyrus was aligned along one edge of the trapezoid. Thin sections (700 Å) were cut on an LKB IV ultramicrotome, collected onto formvar-coated copper slot grids, stained with uranyl acetate and lead citrate, and viewed with a Zeiss 10-C electron microscope. Areas to be sampled were chosen in a systematically random fashion and photographed at 8,000X. Specifically, following random placement of the first photomicrograph site along the edge of the section that was aligned with the middle of the MML, subsequent photomicrographs were taken at regular intervals along the full extent of that edge. To obtain the serial pairs, comparable areas were photographed in the adjacent section.

Neurons were quantified stereologically in 1 μ m sections at the light microscopic level (Nikon Labophot, 20X objective) using the physical disector method [72] (disector height = 4 μ m). Counting frames of known area were superimposed on a field within the granule cell layer of the DG region. The counting units were neuronal nuclei and were included in the total if they appeared in only one of the counting frames in the physical disector pair and did not contact the two exclusion lines of the counting frame. Six section pairs were quantified per block, and two blocks were analyzed per animal.

Synapses were quantified in sections from the same Araldite-embedded block used for neuron quantification to ensure that any changes in tissue volume during processing would affect neuron and synapse numbers equally. Synapses were quantified stereologically in 24 disector pairs per animal (12 disector pairs per block) comprising 48 electron micrographs from serial pairs of 700 Å thin sections. The thickness of the thin sections was determined by the Small fold method [80]. The number of blocks and disector pairs were determined a priori based on power analysis calculations using variance from preliminary studies to achieve a power of 80% to detect a 20% change with a p value less than 0.05. The electron micrographs were taken in the MML, midway between the two boundaries of the molecular layer, i] the hippocampal fissure and ii] the granule cell layer. Since the MML occupies the middle 40% of the molecular layer [64], all electron micrographs were taken well within the limits of the MML. Negatives were digitized for subsequent stereological quantification on a computer. Synapses were defined as having a PSD associated with a presynaptic element containing at least three vesicles (Figure 1, s), with the PSD considered as the counting object. MSB synapses were defined as those synapses that were part of multiple synapse bouton complexes; specifically, MSB synapses shared a single presynaptic terminal, but contacted different postsynaptic elements (Figure 1, asterisks). The numerical densities of all synapses and of MSB synapses in the MML of DG were determined in serial sections using the physical disector method and StereoInvestigator software (MicroBrightfield). Digitized images of the serial sections were viewed at a final magnification of 56,000X on the computer screen and superimposed with a counting box having 2 inclusion and 2 exclusion lines. First, all synapses within the box, but without PSDs (the counting objects) lying on the exclusion lines, were indicated with a marker on the reference section. Those markers then were temporarily removed by the software and the counting box was placed in the same position on the look-up section using fiduciary marks. Next, all synapses were indicated with a different marker on the look-up section. Finally, both sets of markers were revealed, and, only those synapses marked in either the look-up or reference sections, but not both, were counted. Evidence suggests that this method of counting MSB synapses underestimates their number, but generates relative differences between groups that are comparable to those generated by the serial section bouton reconstruction method [28,30]. This method of quantification was suitable for the present study, since the aim was not to determine actual number of MSB synapses, but instead, to examine relative changes in MSB synapses.

The numerical densities of neurons, synapses and MSB synapses were calculated for each animal using the equation $Nv = \Sigma Q - \Sigma a \cdot h$, where Nv is the numerical density, Q- is the number

of unique counting objects in a section, a is the counting frame area, and h is the height of the physical disector. In order to control for the possible confound of differential changes in tissue volume during processing, the numerical densities of synapses and MSB synapses were normalized by dividing by the numerical density of neurons in the same Araldite-embedded block of tissue, thus deriving the number of synapses per neuron. This normalized quantification of synapses is not intended to represent true number of synapses, but rather to

provide a measure of the relative changes in synapses as a function of aging and caloric restriction. Finally, all quantification was performed blindly in order to eliminate investigator bias.

2.4. Western Blot

For Western blot analyses, rats were anesthetized with sodium pentobarbital (200 mg/kg, i.p.) and decapitated in accordance with the recommendations of the Panel on Euthanasia of the American Veterinary Medical Association. The brains were removed immediately and cooled for 2 minutes in chilled artificial cerebrospinal fluid (140 mM NaCl, 5 mM KCl, 1 mM MgCl₂, 1 mM CaCl₂, 24 mM dextrose, 10 mM HEPES, pH 7.4). The hippocampi then were dissected on ice. With the aid of a dissecting microscope, each hippocampus was bisected and the dorsomedial half was divided into four slabs perpendicular to the long axis of the hippocampus. As in the electron microscopy experiments, Western blot analysis was confined to the dorsomedial hippocampus due to its importance for Morris water maze learning and memory tasks [31,50,51]. Using anatomical landmarks, each slab was dissected further to isolate the DG; CA3 was separated from the DG with a cut connecting the ends of the supragranular and infragranular blades of the DG, and then the DG and CA1 were separated along the hippocampal fissure. Samples were weighed, frozen on dry ice, and stored at -80°C. Tissue from the right hippocampus was prepared for the present study and tissue from the left hippocampus was prepared for a separate study [54]. Samples were homogenized using a ground glass dounce with 50 volumes of Laemmli homogenization buffer (60 mM TrisCl, pH 6.8, 10% glycerol, 2% SDS, 200mM EDTA) per gram of tissue, with protease inhibitor cocktail (P8340, 1:250, Sigma-Aldrich, Inc., St. Louis, MI) and phosphatase-inhibitors (P5726, 0.5%, Sigma-Aldrich, Inc., St. Louis, MI). Use of this homogenization buffer has been reported to permit 100% solubilization of the NMDAR in each sample [4]. The soluble supernatant was stored at ^{-80°}C. The protein concentration of each supernatant was determined using a BCA protein assay (Pierce Technology, Rockford, IL).

For NMDAR and AMPAR subunit analysis, 7.5µg of each homogenized sample were loaded into wells on the left half of a 15-well, 7.5% BioRad gel in a cohort of six (one sample per condition, 2 diets X 3 ages; Figure 2) that was duplicated on the right half of the gel. Samples were separated under reducing conditions using SDS-PAGE and then electrophorectically transferred onto Immobilon membranes using a BioRad Mini Transfer Cell. Rabbit polyclonal antibodies to the NR1 (0.6 µg/ml, Chemicon International, Temencula, CA), NR2A (0.07 µg/ ml, Chemicon), NR2B (0.07 µg/ml, Chemicon), GluR1 (0.01 µg/ml, Upstate, Lake Placid, NY) and GluR2 (0.25 μ g/ml, Chemicon) subunits were used for immunodetection. All of the antibodies were affinity purified and tested for specificity; no antibody cross-reacted with other receptor subunits in the family. A mouse monoclonal antibody to actin (0.002 µg/ml, Chemicon) was used to verify equal protein loading. Peroxidase-conjugated donkey anti-rabbit IgG (10 ng/ml, Jackson ImmunoResearch Laboratories, Inc., West Grove, PA) and peroxidaseconjugated donkey anti-mouse IgG (10 ng/ml, Jackson) secondary antibodies were used with the SuperSignal West Pico Chemiluminescent Substrate (Pierce Technology, Rockford, IL) for detection. Blots were exposed on Kodak Biomax film and quantitated densitometrically with BioRad VersaDoc and Quantity One analysis software (BioRad Laboratories, Hercules, CA) in order to determine the relative amounts of each protein. The Gaussian trace optical density of each band was normalized to the average density of all of the matching bands within

each cohort to permit comparison between blots and experiments. A repeated measurement was excluded if it was greater than two standard deviations away from the mean. Each blot was repeated at least once, thus generating a minimum of four acceptable measurements per antibody per sample.

2.5. Statistical Analysis

Statistical analyses were performed using GraphPad Prism (version 4.00 for Windows; GraphPad Software, San Diego, CA) and SigmaStat (Systat Software, Inc., Richmond, CA). The main effects and interaction of the two independent variables, age and diet, were evaluated using two-way univariate analysis of variance with Bonferroni's post test. A two-tailed significance level for each of the main effects was held at 0.05; means are reported +/- standard deviation. A Bonferroni adjustment was used for multiple comparisons to control for a type I error. In the absence of a main effect of diet, AL and CR groups were combined to increase the power to detect change with age. In these cases, the effect of age on the dependent variables was determined using a one-way univariate analysis of variance with Tukey's post test. Hartley's F-max test was used to test the homogeneity of variances among combined AL and CR groups for the number of overall synapses and MSB synapses per neuron.

3. Results

3.1. CR rats weighed less than AL rats

All CR rats weighed less than age-matched AL rats (10CR, 275 g +/- 10 vs 10AL, 425 g +/- 26; 18CR, 300 g +/- 10 vs 18AL, 502 g +/- 25; 28CR, 311 g +/- 8 vs 28AL, 558 g +/- 20), but appeared healthy and had full coats of fur. The differences in mean weight rats varied significantly with both age ($F_{(2,75)} = 103.81$, p < 0.0001) and diet ($F_{(1,75)} = 1667.44$, p < 0.0001). For both AL and CR rats, the average weights at 18 and 29 months were higher than those at 10 months (p's<0.001). Only in AL rats, however, was the average weight at 29 months significantly higher than that at 18 months (p < 0.001). On average, the weight of CR rats was 40% lower than that of AL rats (35%, 40%, 44%, at 10, 18, and 29 months, respectively). At no point did the weight of any CR rat exceed that of any age-matched AL rat.

3.2. Effects of age and CR on DG synapses

The numerical densities of neurons and synapses were determined stereologically (Table 1) and the overall synapses per neuron and MSB synapses per neuron were derived from these values. There were no main effects of age or diet on the overall number of synapses per neuron (age, $F_{(2,30)}=0.08$, p=0.922; diet, $F_{(1,30)}=0.73$, p=0.401, Figure 3A) or the number of MSB synapses per neuron (age, $F_{(2,30)}=0.40$, p=0.676; diet, $F_{(1,30)}=1.05$, p=0.314, Figure 3B).

Because AL and CR groups did not differ statistically at any age, the data from those two groups were combined at each age to increase the power to detect potential changes across lifespan. Even with the resulting N of 12 per age, there was no significant main effect of age for either overall synapses per neuron ($F_{(1,30)}=0.087$, p=0.917, Figure 4A) or MSB synapses per neuron ($F_{(1,30)}=0.422$, p=0.659, Figure 4B). Although the mean number of synapses did not differ with age, Hartley's F-max test revealed significantly larger variances in the number of synapses per neuron (p<0.05) at 29 months compared to 18 months as well as in the number of MSB synapses per neuron at 10 and 29 months compared to 18 months (p<0.05).

3.4. Effect of aging and CR on relative levels of NMDAR and AMPAR subunits

In the Western Blot experiments, we compared relative protein levels of the NR1, NR2A, and NR2B subunits of the NMDAR and of the GluR1 and GluR2 subunits of the AMPAR in young, middle aged, and old AL and CR rats. For **NR1** (Figure 5A), there was a significant main effect

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of age ($F_{(2,38)}=6.359$, p<0.01) and diet ($F_{(1,38)}=6.233$, p<0.05) and a significant interaction between the two variables (F_(2,38)=7.581, p<0.01). Levels of NR1 in both 18AL and 29AL groups were significantly lower than in the 10 AL group (5.8% decrease in mean, p < 0.05; and 12.1% decrease, p < 0.001; respectively) and 29AL levels were lower than 18AL levels (7% decrease, p < 0.05). NR1 levels in the 10CR group also were significantly lower than in the 10AL group (9.1%, p<0.001), but did not differ between the two dietary conditions at 18 (p=0.059) or 29 months (p=0.139). NR1 levels in CR animals did not differ significantly among ages (all p's>0.05). For NR2A (Figure 5B), there was no main effect of age (F_(2.41)=0.927, p=0.404) or diet (F_(1,41)=0.831, p=0.367) nor an interaction between the two variables $(F_{(2,41)}=0.391, p=0.679)$. Statistical analysis of **NR2B** levels (Figure 5C) demonstrated a main effect of age ($F_{(2,39)}$ =5.333, p<0.01), but not diet ($F_{(1,39)}$ =2.136, p =0.152), and no interaction between the two variables ($F_{(2.39)}$ =0.912, p =0.912). Within CR animals, NR2B levels in the 29CR group were slightly, yet significantly, lower than in the 10CR group (3.6% decrease, p < 0.05). There were no other significant differences among age or diet groups for NR2B. There was a main effect of age ($F_{(2,40)}=6.080, p<0.01$), but not diet ($F_{(1,40)}=0.138, p=0.712$) on GluR1 levels (Figure 5D) as well as a significant interaction between the two variables $(F_{(2,40)}=4.331, p<0.05)$. Within the AL, but not CR groups, there were significant differences in GluR1 levels between individual age groups. Specifically, GluR1 levels were lower in both the 18AL and 29AL groups than in the 10AL group (8.9% decrease, p < 0.01; and 12.4% decrease, p < 0.001; respectively). Within the 10 month cohort, CR levels were significantly lower than the AL levels (7.2% decrease, p < 0.05), but there were no differences between diet groups at 18 (p=0.570) or 29 months (p=0.178). Finally, for **GluR2** (Figure 5E), there was a main effect of age ($F_{(2,39)}=6.612$, p<0.01), but not diet ($F_{(1,39)}=0.291$, p=0.593). No significant interaction between the variables was found ($F_{(2,39)}=1.315$, p=0.280). Within the CR group, GluR2 levels in the 29 month group were lower than at 10 months (4.7% decrease, p < 0.01) and 18 months (3.5% decrease, p < 0.05). GluR2 levels did not differ among age groups in AL animals. For actin, there were no main effects of age or diet, nor was there an interaction between the two variables (all *p*'s>0.05, Figure 5F).

4. Discussion

The results of the present study demonstrate that the number of synapses per neuron in the MML of the DG region of hippocampus do not change across lifespan and are unaffected by CR at young, middle, and old age. In contrast to the that stability of synapses across ages and diets, glutamate receptor subunits in the DG decline as a function of age and with CR. Aging-related decreases in NR1 and GluR1 occur in AL rats. In young rats, levels of these subunits are significantly lower in CR than in AL groups, but the levels in CR rats then remain stable across lifespan. These data demonstrate that CR eliminates the aging-related decline in NR1 and GluR1 subunits by inducing and then maintaining stable, lower levels of these subunits across lifespan.

4.1. Mean number of synapses per neuron in the MML of the DG does not change with diet or age

Although CR-induced changes in dendritic spines and LTP have been associated with alterations in synapse structure in previous experiments [15,48], CR produced no significant changes in the number of synapses per neuron at any age in the present study. Specifically, CR had no impact either on overall synapses or MSB synapses normalized to neuron number. For AL and CR groups combined, the results also indicated that overall synapses and MSB synapses per neuron did not change across the three ages examined. Other studies have reported a similar preservation of DG synapses over lifespan based on synaptophysin labeling of synaptic boutons [7,56,70] and a stereologic study examining CA1 synapses at 6 and 27 months also showed no aging-related change [20]. The only other stereological investigation of synapses in the rodent

DG reported a decline between the two ages examined in that study, 5 and 28 months [22]. The variation in findings between that study and the present one likely reflects the difference in animal ages. Although a 5 month old rat is sexually mature, it may well be that brain development continues beyond that age. Increases in dendritic extent observed in the supraoptic nucleus between 3 and 12 months of age have been interpreted to indicate the presence of ongoing maturational changes [16]. Moreover, synapses in CA3 have been demonstrated stereologically to decrease by 24% between 3 and 12 months of age [12]. Inclusion of intermediate ages would narrow the time period between 5 and 28 months during which synaptic loss occurs [11]. In light of the present results, we would predict a loss of synapses to occur between 5 and 10 months of age.

The present study used the physical disector to quantify synapses stereologically in the MML of the DG. Using this methodology, the disector height is determined based on the size of a specific counting object. In this study, the disector height was optimized for the quantification of simple synapse densities. Accordingly, the serial pairs of thin sections utilized here are not suitable for analysis of perforated synapses or entire MSB complexes. Such analysis requires serial reconstruction of these structures, and studies using this approach are currently underway in our laboratory. In the absence of a change in synapses per neuron, these studies may reveal subtle changes in synaptic structure.

4.2. Aging and caloric restriction modulate levels of glutamate receptor subunits

In light of the finding that neither aging nor CR affects the number of synapses per neuron in the MML of the DG, we used Western blot analysis to evaluate whether these variables affect synapses by influencing the levels of glutamate receptor subunits. The results indicated that subunit levels in AL rats declined (NR1 and GluR1) or did not change (NR2A, NR2B, GluR2) across lifespan; no subunit increased with age. These observations are consistent with the idea that a decline in one or more glutamate receptor subunits accompanies the aging-related learning and memory deficits. CR then might be expected to prevent aging-related declines in glutamate receptor subunits, since CR has been shown to prevent aging-related learning and memory deficits [25,26,43,60,61,73]. Interestingly, CR did not prevent the absolute declines in NR1 and GluR1 that occurred in AL rats. Instead, it resulted in significantly lower subunit levels by 10 months of age that then were maintained across lifespan. In comparison with NR1 and GluR1, levels of the other subunits were relatively stable across age groups were so small as to be of questionable biological significance.

Several studies reported in the literature have used Western blot analysis to examine protein levels of NMDAR and AMPAR subunits in rat hippocampus across lifespan [1,9,15,47,71] and following manipulation of food intake [15,47]. The results of those studies, however, cannot be compared directly to the present findings. In none of those studies were NMDAR or AMPAR subunit levels measured in separate hippocampal subregions, and subunit levels in CA1 and/or CA3 could mask a unique response in the DG. In addition, the published data were collected from different rat strains, and the effect of CR on hippocampus-dependent learning and memory performance is known to vary among strains [43]. The ages examined in some of those studies that assessed the effects of dietary manipulation, only one used the same 60% CR regimen as in the present study [43]; the other employed an every-other-day feeding schedule [47] recognized to have distinct biological effects [2].

Western blot analysis in the present study assessed the overall levels of receptor subunit protein in the DG, but did not distinguish between levels in the synaptic and cytosolic compartments. Modulation of receptor subunits may occur as alterations in receptor composition, trafficking, or distribution within the synapse. Determining whether receptor subunits in the synaptic

compartments specifically are affected by aging or CR requires ultrastructural localization of receptor subunits with postembedding immunogold electron microscopy. Although that technique could not be used in the present analysis as it requires tissue processing incompatible with the techniques used here, immunogold studies to localize receptor subunits that are underway in our laboratory will address this important issue.

In the present study, the only significant differences in relative protein levels of NMDAR and AMPAR subunits between CR and AL groups were in young animals; comparisons between middle aged and old animals revealed no differences between the two groups. In contrast, behavioral evaluation of spatial learning and memory in the same rat strain and at similar ages as in the present study revealed no difference between CR and AL groups at young or middle age [43]. In that study, only old CR animals demonstrated significantly better performance on learning and memory tasks as compared to old AL rats. These findings raise the critical question of whether CR-induced NMDAR and AMPAR subunit changes observed in young animals in the present study have effects restricted to early life or whether the early effects of CR may have important functional implications later in life.

4.3. Caloric restriction may promote biological stability

Considering the remarkable resistance of DG neurons to damage from acute insults such as ischemia or status epilepticus [33,58], it may not be surprising that aging and CR do not alter the number of synapses in the DG. Perhaps the mechanisms that exist to protect the DG from acute insults also act to buffer it against chronic, milder stressors such as aging and CR. CR is, in fact, a mild form of metabolic stress that induces some of the same changes that aging does, including increased glucocorticoids [53,68] and decreased levels of circulating insulinlike growth factor-I [5]. Results from our laboratory revealed a CR-induced increase in levels of brain-derived neurotrophic factor (BDNF) in CA1 similar to the aging-related BDNF increase that occurs in the DG and CA3 [54]. Given that BDNF is a neurotrophin that supports synapses and synaptic function [8,32,75,79], the aging-related increase in BDNF in the DG could contribute to the general stability of the number of synapses per neuron observed in the DG over lifespan. Thus, CR may not decelerate aging-related processes in the DG, but, instead, may stimulate natural protective mechanisms that occur normally during the aging process to help maintain homeostasis. This role for CR is consistent with the notion that CR is a form of "hormesis" [44,45,78], or a mild stressor that potentiates innate defenses that help to guard against subsequent, more egregious insults. If, in fact, the immediate benefits of CR are a consequence of stimulating adaptive responses normally triggered during the aging process, then the protection it affords during aging may reflect its ability to maintain these responses at stable levels over lifespan in order to preserve homeostasis. Such a scenario is consistent with the stability-longevity hypothesis [13] suggesting that the beneficial effects of CR are due to maintaining steady levels of metabolic byproducts. Although much investigation remains to be done to test the validity of this hypothesis, the data presented here are consistent with this concept of aging and the effects of CR.

4.4 Conclusion

In summary, the number of DG synapses per neuron does not change with aging or CR. Although some glutamate receptor subunit levels do decrease across lifespan in AL animals, CR does not prevent their decline. Instead, the levels of those subunits are reduced in CR animals early in life, by 10 months of age, after which they remain stable across lifespan. Further studies will be required to determine whether later introduction or more limited duration of CR will yield similar findings.

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Figure 1.

Electron micrograph illustrating synapses (s), two of which are involved in an MSB complex (asterisks).



Figure 2.

Representative Western blots showing glutamate receptor subunits NR2A, NR1, and GluR1 in the blot on the left and NR2B and GluR2 in the blot on the right for AL and CR rats at young (10 months), middle (18 months) and old (29 months) ages. Actin was included on all gels as a control for equal protein loading. Samples were processed in cohorts of six, one sample per age/diet group, which was repeated on the right half of the gel. The bands directly above actin on the left are nonspecific bands associated with the NR1 antibody.

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Figure 3.

Scatter plot diagrams of the number of synapses per neuron (A) and the number of MSB synapses per neuron (B) for AL (open squares) and CR groups (filled triangles) at young (10 months), middle (18 months) and old (29 months) ages. The horizontal lines indicate the mean for each group. In no case did the means change as a function of age or diet.

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Figure 4.

Scatter plot diagrams of the number synapses per neuron (A) and the number of MSB synapses per neuron (B) for the AL and CR groups combined at young (10 months), middle (18 months) and old (29 months) ages. There are no significant differences in the means between any ages in either A or B; however, the variances at young and old ages in both are clearly greater than at middle age. The difference between variances was significant for synapses between middle and old age (A, p<0.05) and for MSB synapses between young and middle age and middle and old age (B, p<0.05 for each).

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Figure 5.

Relative levels of NMDAR subunits NR1 (A), NR2A (B), and NR2B (C), and AMPAR subunits GluR1 (D) and GluR2 (E), and actin (F) as a function of age (young, 10 months; middle aged, 18 months; and old, 29 months) and diet (AL, filled bars; CR, patterned bars). Asterisks above the CR bars at young age for NR1 (A) and GluR1 (D) indicate significant differences compared to AL animals (p<0.05). Asterisks above connector lines indicate significant aging-related decreases in receptor subunits (p<0.05). These data represent the mean \pm standard deviation of eight animals per experimental condition.

Table 1

Numerical Density of Neurons, Synapses, and MSB Synapses

		Neurons/ mm ³	Synapses/mm ³	MSB Synapses/mm ³
10AL N=6	mean	$3.10 imes 10^6$	$1.52 imes 10^9$	$4.17 imes 10^8$
	StDev	$0.09 imes 10^6$	$0.20 imes 10^9$	$1.10 imes 10^8$
	SEM	$0.04 imes 10^6$	$0.09 imes 10^9$	$0.49 imes 10^8$
	CE	0.01	0.06	0.12
	CV%	3	13	26
10CR N=6	mean	$3.04 imes 10^6$	$1.41 imes 10^9$	$3.69 imes 10^8$
	StDev	$0.18 imes 10^6$	$0.21 imes 10^9$	$0.90 imes 10^8$
	SEM	$0.81 imes 10^6$	$0.94 imes 10^9$	$0.40 imes 10^8$
	CE	0.03	0.07	0.11
	CV%	6	15	24
18AL N=6	mean	$3.20 imes 10^6$	$1.58 imes 10^9$	$4.50 imes 10^8$
	StDev	$0.11 imes 10^6$	$0.09 imes 10^9$	$0.64 imes 10^8$
	SEM	$0.05 imes 10^6$	$0.04 imes 10^9$	$0.29 imes 10^8$
	CE	0.02	0.03	0.06
	CV%	4	6	14
18CR N=6	mean	$3.17 imes 10^6$	$1.45 imes 10^9$	$4.07 imes 10^8$
	StDev	$0.32 imes 10^6$	$0.11 imes 10^9$	$0.46 imes 10^8$
	SEM	$0.14 imes 10^6$	$0.05 imes 10^9$	$0.21 imes 10^8$
	CE	0.05	0.03	0.05
	CV%	10	8	11
29AL N=6	mean	3.16×10^6	$1.48 imes 10^9$	$4.04 imes 10^8$
	StDev	$0.19 imes 10^6$	$0.34 imes 10^9$	$1.25 imes 10^8$
	SEM	$0.85 imes 10^6$	$0.15 imes 10^9$	$0.56 imes 10^6$
	CE	0.03	0.10	0.14
	CV%	6	23	31
29CR N=6	mean	$3.25 imes 10^6$	$1.46 imes 10^9$	$3.79 imes 10^8$
	Stdev	$0.29 imes 10^6$	0.24×10^{9}	0.64×10^{8}
	SEM	$0.13 imes 10^6$	$0.11 imes 10^9$	$0.29 imes 10^8$
	CE	0.04	0.07	0.08
	CV%	9	17	17