Astrocytic gap junctional communication is reduced in amyloid-\(\beta\)-treated cultured astrocytes, but not in Alzheimer’s disease transgenic mice

Nancy F Cruz, Kelly K Ball and Gerald A Dienel1
Department of Neurology, University of Arkansas for Medical Sciences, Little Rock, AR 72205, U.S.A.

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ABSTRACT

Alzheimer’s disease is characterized by accumulation of amyloid deposits in brain, progressive cognitive deficits and reduced glucose utilization. Many consequences of the disease are attributed to neuronal dysfunction, but roles of astrocytes in its pathogenesis are not well understood. Astrocytes are extensively coupled via gap junctions, and abnormal trafficking of metabolites and signalling molecules within astrocytic syncytia could alter functional interactions among cells comprising the neurovascular unit. To evaluate the influence of amyloid-\(\beta\) on astrocyte gap junctional communication, cultured astrocytes were treated with monomerized amyloid-\(\beta\)1–40 (1 \(\mu\)mol/l) for intervals ranging from 2 h to 5 days, and the areas labelled by test compounds were determined by impaling a single astrocyte with a micropipette and diffusion of material into coupled cells. Amyloid-\(\beta\)-treated astrocytes had rapid, sustained 50–70% reductions in the area labelled by Lucifer Yellow, anionic Alexa Fluor \(H\) dyes and energy-related compounds, 6-NBDG (a fluorescent glucose analogue), NADH and NADPH. Amyloid-\(\beta\) treatment also caused a transient increase in oxidative stress. In striking contrast with these results, spreading of Lucifer Yellow within astrocytic networks in brain slices from three regions of 8.5–14-month-old control and transgenic Alzheimer’s model mice was variable, labelling 10–2000 cells; there were no statistically significant differences in the number of dye-labelled cells among the groups or with age. Thus amyloid-induced dysfunction of gap junctional communication in cultured astrocytes does not reflect the maintenance of dye transfer through astrocytic syncytial networks in transgenic mice; the pathophysiology of Alzheimer’s disease is not appropriately represented by the cell culture system.

Key words: amyloid protein, astrocyte, connexin, dye transfer, gap junction, metabolite trafficking.

INTRODUCTION

Astrocytes are extensively linked to each other via gap junctional channels formed by Cx (connexin) proteins, Cx43, Cx30 and Cx26 (Nagy et al., 2004). These channels enable transcellular movement of fuel, metabolites, signalling molecules and electrolytes, with preferential transfer of material that is governed by molecular size, charge and connexin composition (Harris, 2007). The physiological functions of cell-to-cell transfer of energy-related metabolites within the astrocytic syncytium in brain are not well understood, but astrocytes have important roles in brain energetics (Hertz et al., 2007). Sharing among astrocytes of fuel and redox compounds and rapid elimination of metabolic by-products may be essential astrocytic responses to brain activation. For example, the astrocytes that remove glutamate and potassium from an active synaptic cleft incur an increased demand for energy that would be reduced if the neurotransmitters and electrolytes quickly diffuse into neighbouring gap junction-coupled astrocytes. Also, astrocytic uptake and

1 To whom correspondence should be addressed (email gadienel@uams.edu).

Abbreviations: A350, Alexa Fluor \(R\) 350; A568, Alexa Fluor \(R\) 568; aCSF, artificial cerebral spinal fluid; APP, amyloid-\(\beta\)-precursor protein; Cx, connexin; dBcAMP, dibutyryl cAMP; DCF, dichlorofluorescein; DMEM, Dulbecco’s modified Eagle’s medium; FBS, fetal bovine serum; GFAP, glial fibrillary acidic protein; H2DCF-DA, carboxy-
dihydrodichlorofluorescein diacetate; L-LME, L-leucine methyl ester hydrochloride; 6-NBDG, 6-[N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl)amino]-6-deoxyglucose; ROS, reactive oxygen species; SR101, sulforhodamine 101; STZ, streptozotocin.
dispersed of lactate among gap junction-coupled cells (Gandhi et al., 2009b) in conjuction with its discharge from perivascular ependet that are linked to extensive gap junctional networks (Ball et al., 2007) could serve to help up-regulate blood flow in a larger volume of tissue compared with that of focally activated cells, by raising the level of lactate in the interstitial fluid near the endothelium (Gordon et al., 2008). Furthermore, release from activated astrocytes of molecules used as signals for brain imaging and spectroscopic assays (e.g. NADH, NADPH and ¹³C- or ¹²C-labelled metabolites of glucose or acetate) could cause underestimation of cellular redox status and metabolic rates calculated from accumulation of labelled products unless appropriately taken into account.

Alzheimer's disease is a devastating neurodegenerative disease associated with cognitive dysfunction, increased levels of amyloid-$\beta$ deposits and neurofibrillary tangles in brain, oxidative stress, neuroinflammation and neuronal dysfunction (Craft, 2009; Fotuhi et al., 2009; Querfurth and LaFerla, 2010), and one of the earliest and most reliable markers for Alzheimer's disease and its progression is reduced glucose utilization in brain (Alexander et al., 2002). Astrocytic pathology is associated with Alzheimer's disease, including hypertrophy, oxidative stress, DNA damage and reduced expression of the EAAT2 glutamate transporter (Wharton et al., 2010). Furthermore, release from activated astrocytes of molecules used as signals for brain imaging and spectroscopic assays (e.g. NADH, NADPH and ¹³C- or ¹²C-labelled metabolites of glucose or acetate) could cause underestimation of cellular redox status and metabolic rates calculated from accumulation of labelled products unless appropriately taken into account.

Astrocyte culture
Cultured astrocytes were prepared essentially as described by Hertz et al. (1998) after harvest of cells from cerebral cortex of 1-day-old albino Wistar–Hanover rats (Taconic Farms, Germantown, NY). Astrocytes were grown in T75 culture flasks containing DMEM (Dulbecco’s modified Eagle’s medium; catalogue number 12320-032), penicillin, streptomycin, Amphotericin B and trypsin obtained from Invitrogen and FBS (fetal bovine serum) was from Jackson Laboratories. The mutant mouse lines originated from Invitrogen (Molecular Probes). Human amyloid-$\beta_{1-40}$ (product no. A1075, formula mass 4330 Da; Sigma) was treated with hexafluoro-2-propanol to monomerize the amyloid-$\beta$ (Klein, 2002); freeze-dried samples were stored at ~80°C and dissolved in PBS immediately prior to use.

Brain slices
Breeder pairs of transgenic mice (PDGF-APPSwInd, stock #004661) that contain the human APP with Swedish (K670N/M671L) and Indiana (V717F) mutations (Mucke et al., 2000). We report that treatment of cultured astrocytes with amyloid-$\beta$ markedly reduced gap junctional transfer of six test compounds, whereas dye-transfer among astrocytes in brain slices from three regions of transgenic Alzheimer’s mice is similar to that of aged littermate controls, contrasting the results of previous brain slice studies.
deposition occurs in 50–60% of the mice at 5–7 months of age and 100% at 8–10 months of age (Mucke et al., 2000). All animal use procedures were in strict accordance with the NIH Guide for Care and Use of Laboratory Animals and were approved by the local Animal Care and Use Committee.

Mixed-gender heterozygous transgenic Alzheimer’s mice and control littermates (n=6/group) ranging in age from 8.5 to 14 months were anaesthetized with isoflurane, decapitated, their brains were quickly removed and chilled by immersion in oxygenated, ice-cold aCSF (artificial cerebral spinal fluid) solution containing 26 mmol/l NaHCO₃ (pH 7.3) and 248 mmol/l sucrose, and 250 μm-thick slices were prepared as described by Moyer and Brown (1998). Coronal sections of brain were cut using a Leica VT 1000S tissue slicer, slices were incubated in oxygenated aCSF containing sucrose for 30 min at 35˚C, then for ≥1 h at 22˚C, and transferred to an open bath perfusion chamber (Warner Instruments).

**Gap junctional communication assays**

Gap junctional communication among astrocytes was assayed by impaling a single cell with a micropipette containing a test compound, allowing the material to diffuse within the astrocytic network for a fixed time (2 or 5 min in cultured cells or brain slices respectively), then measuring the labelled astrocytic network for a fixed time (2 or 5 min in cultured cells or brain slices respectively). The areas labelled by the probes were determined with MetaVue software. Counts of the number of dye-labelled cells in brain slices were carried out as follows. The 250 μm-thick brain slices that were previously injected with Lucifer Yellow were removed from the microscope stage and immediately fixed by immersion in 4% paraformaldehyde in PBS (pH 7.4), quickly followed by mixing for 30 s, then gentle shaking for 30 min at room temperature (approx. 20˚C) before being stored at 4˚C. Prior to sectioning, the slices were rinsed in PBS, cryoprotected in 20% sucrose/PBS for 5–7 days, frozen and cut into 7 μm-thick serial sections at −20˚C. Sections were mounted on SuperfrostPlus glass slides, air dried for 24 h and then stored in slide boxes at 4˚C. After the sections were coverslipped (Gelmount medium and glass coverslips), the Lucifer-Yellow-containing astrocytes were counted in each 7 μm-thick serial section using a Zeiss Axioscope2 with Zeiss Plan Neofluar objectives (×10/0.30 numerical aperture or ×20/0.50), an MTI CD72 camera and an MCID™ Imaging System (InterFocus Imaging Limited), then summed to obtain the total number of dye-labelled cells for each 250 μm-thick brain slice.

**Congo Red staining of amyloid plaques**

An alkaline Congo Red staining method (Puchtler et al., 1962; Wilcock et al., 2006) was used to identify amyloid plaques in the mouse brain tissue. After the Lucifer Yellow-labelled cells were counted, the coverslips were removed from the slides by soaking in water overnight, then rinsed several times to remove any remaining Gelmount. Next, the tissue sections

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were soaked in deionized water for 30 s, transferred to Gill’s Haematoxylin #3 (Polysciences Incorporated) for 30 s, and rinsed in tap water until the water was clear. A critical aspect of the Congo Red staining procedure was that each of the two alkaline solutions was freshly prepared and used within 15 min. The pretreatment solution was prepared by adding 2.5 ml of 1% (w/v) aqueous NaOH to 250 ml of previously prepared NaCl-saturated 80% ethanol; then the solution was mixed and filtered. The staining solution was similarly prepared by adding the NaOH to a 0.2% Congo Red (Chroma #10460) solution in 80% ethanol saturated with NaCl. The slides were first incubated for 20 min in the pretreatment solution, placed in the staining solution for 20 min, then quickly passed through three changes of absolute ethanol followed by three changes of xylene and mounted with Permount medium (Fisher Chemical). Brightfield and fluorescent images of the Congo Red-stained tissue (Puchtler and Sweat, 1965) were obtained and composite images (100 and ×200 magnification) were produced by overlaying the brightfield images with their corresponding colorized fluorescent images. The presence of amyloid plaques was confirmed by microscopic examination under cross-polarized light and demonstration of red–green birefringence (Jin et al., 2003 and references cited therein).

Assay for ROS (reactive oxygen species)/RNS (reactive nitrogen species)
H2DCF-DA is cell-membrane-permeable, cleaved by intracellular esterases and, after oxidation mediated mainly by H2O2 and peroxyxinitrite, becomes fluorescent DCF (dichlorofluorescein) (Cruthirds et al., 2005). H2DCF-DA (10 μmol/l) was added to the culture medium, cells returned to the CO2 incubator for 30 min at 37°C, cells washed with perfusion solution and fluorescence intensity determined with Nikon E600 microscope and MetaVue software.

Statistics
All statistical analyses were performed with GraphPad Prism® software, version 5.02 (GraphPad Software). Comparisons between two groups of independent samples were made with two-tailed, unpaired t tests. P<0.05 is considered to be statistically significant. Reported values are means and vertical error bars in Figures represent 1 S.D.

RESULTS

Amyloid–β reduces Lucifer Yellow transfer through gap junctions in cultured astrocytes
Previous studies of glutamate uptake, signalling pathways and oxidative stress (Matos et al., 2008) and calcium waves (Haughey and Mattson, 2003; Chow et al., 2010) exposed cultured astrocytes to 5 μmol/l amyloid-β1–40 or amyloid-β1–42 for various time intervals; this dose is considered to be the most likely level of astrocytic exposure in Alzheimer’s brain (Haughey and Mattson, 2003). A higher level of amyloid-β1–35 (20 μmol/l) has been used to evaluate effects on gap junctional inhibition by cytokines released by microglia (Mème et al., 2006). The 5 μmol/l dose of amyloid-β1–42 was stated to be non-toxic to astrocytes within the 24 h (Haughey and Mattson, 2003), but even much lower doses (10 or 100 nmol/l and 1 μmol/l) of amyloid-β1–40 enhanced the generation of ROS and advanced glycation end-products in vascular endothelial cells grown in high (25 mmol/l) compared with low (5 mmol/l) glucose (Burdo et al., 2009). ‘Aged’ solutions of amyloid-β1–40 and amyloid-β1–42 in which oligomer and fibrillar aggregates of the peptide are formed are also non-toxic in the range 1–10 μmol/l, but do stimulate oxidative stress in astrocytes grown in high-glucose DMEM (Matos et al., 2008); aged amyloid also reduces free intracellular calcium levels in astrocytes, alters their morphology and increases expression of immunoreactive S100β and GFAP (Meske et al., 1998). In our initial studies, dye transfer was tested with 1, 2, 5 or 10 μmol/l of monomerized amyloid-β1–40, and the lowest dose was chosen for use to minimize formation of vacuoles that were visible in cells given the highest doses. The effects of amyloid-β1–40 were first assessed in cultured astrocytes grown in low (5.5 mmol/l) or high (25 mmol/l) glucose because we recently found that experimental diabetes, a risk factor for Alzheimer’s disease, reduces dye transfer through gap junctions in cultured astrocytes and STZ (streptozotocin)-diabetic rats (Gandi et al., 2010), and STZ-induced diabetes also enhances cognitive deficits and expression of oxidative stress markers in presymptomatic Alzheimer’s transgenic mice (Burdo et al., 2009).

In agreement with our previous results, the spread of Lucifer Yellow through gap junctions was much greater in cells grown in low (Figures 1Aa and 1Ab) compared with high (Figures 1Bd and 1Be) glucose. Treatment of both culture groups with 1 μmol/l monomerized amyloid-β for 2 h markedly reduced dye transfer compared with their respective controls (Figures 1Ac and 1Bf). This decrement averaged approx. 50–75% in astrocytes grown in low (Figure 2A) or high (Figure 2B) glucose compared with paired control cultures. Pretreatment of low-glucose control astrocytes with octanol reduced the Lucifer Yellow-labelled area by approx. 75% (P<0.001), from 11220±4566 μm² (mean±S.D.) to 2913±950 μm², a magnitude similar to that caused by amyloid treatment. Maximal inhibition of dye transfer by amyloid-β occurred within 2 h, and increased duration of exposure to the single dose of monomerized amyloid from 2 h to 5 days did not alter the decrement (Figure 2); no attempt was made to determine the proportion of amyloid monomers, oligomers and fibrils over time. Because high glucose reduced dye transfer in control cultures and blunted the percentage change after amyloid-β treatment (Figure 2), all subsequent assays used astrocytes grown in low glucose.
Amyloid-\(\beta\) causes modest, transient oxidative stress in astrocytes grown in low glucose

We previously identified oxidative stress as a probable causative factor in disruption of trafficking of small molecules through the astrocytic network in experimental diabetes (Gandhi et al., 2010), and amyloid-\(\beta\) exacerbates the generation of ROS in brain vascular endothelial cells (Burdo et al., 2009). DCF fluorescence was, therefore, assayed to evaluate the extent of amyloid-\(\beta\)-induced oxidative stress in the low-glucose cultures. DCF fluorescence approximately doubled after 2 h exposure to amyloid-\(\beta\), but the increase did not persist at 24 h, and a slight decrement was evident at 5 days (Figure 3). The rapid onset of reduced dye transfer (Figures 1 and 2) and low, transient increase in oxidative stress with amyloid treatment (Figure 3) sharply contrast the responses of hyperglycaemic astrocytes. These cells had a rapid, progressive rise in DCF fluorescence, but there was a 3–5 day delay before the onset of inhibition of dye transfer, suggesting that other factors mediate the effects of amyloid-\(\beta\) on dye transfer.

Amyloid-\(\beta\) impairs gap junctional transfer of compounds with different size or charge

Cell-to-cell transfer of anionic dyes that are either larger (A568, 730 Da; Figures 4a and 4b) or smaller (A350, 311 Da when ionized; Figures 4c and 4d) than Lucifer Yellow VS (536 Da when ionized; Figures 1 and 2) is also markedly lowered by amyloid-\(\beta\) treatment. The area labelled by A568 was reduced within 2 h by amyloid-\(\beta\) and it persisted at 1 and 2 days (Figure 5A), whereas that labelled by A350 was statistically significantly only at day one due to lower values for the control group in the 2 day treatment group (Figure 5B). Octanol treatment of control astrocytes reduced the A568-labelled area by approx. 80%, from 7379 ± 3080 \(\mu\)m\(^2\) to 1434 ± 658 \(\mu\)m\(^2\) (\(P<0.01\)). Diffusion within the astrocytic network of molecules related to glucose metabolism was also inhibited by amyloid-\(\beta\). The area labelled by 6-NBDG (342 Da), a non-phosphorylatable uncharged fluorescent analogue of glucose (Aller et al. 1997; Speizer et al.1985) that is used to assay transport, was reduced by 40–70% after 1–2 days treatment of astrocytes with amyloid-\(\beta\) (Figure 6A). NADH and NADPH are phosphorylated anions involved in redox reactions in the glycolytic and pentose phosphate shunt pathways respectively. Movement of NADH (663 Da) within the astrocytic syncytium was depressed after exposure to amyloid-\(\beta\) for 2 h to 2 days (Figure 6B), and spreading of NADPH (723 Da) was also reduced by 24 h treatment with amyloid-\(\beta\) (data not shown). Octanol pretreatment of control astrocytes reduced NBDG labelling from 12816 ± 4305 \(\mu\)m\(^2\) to 1153 ± 492 \(\mu\)m\(^2\) (\(P<0.001\)) and diminished NADH labelling from 6109 ± 2873 \(\mu\)m\(^2\) to 1323 ± 519 \(\mu\)m\(^2\) (\(P<0.01\)).

Astrocytic dye transfer in brain slices from aged control and Alzheimer’s mice

Because cultured astrocytes are not exposed to the same environment as cells in the aging and Alzheimer’s model mouse brain, Lucifer Yellow transfer was also assayed in 8.5–14-month-old transgenic mice that overexpress mutant human APP and in age-matched littermate controls. Mice with the mutant APP were identified by genotyping, and amyloid plaque formation was verified in all animals designated as ‘Alzheimer’s mice’ after the dye-labelled cells were counted. Congo Red-stained amyloid plaques were

Figure 1
Representative images of reduced Lucifer Yellow spread among astrocytes after brief exposure to amyloid-\(\beta\). Cultured astrocytes grown in (A) low (5.5 mmol/l) or (B) high (25 mmol/l) glucose-containing medium for 2–3 weeks. Cells were visualized and impaled with a micropipette (a and d), and Lucifer Yellow was diffused for 2 min into a single astrocyte in a culture treated with vehicle (b and e) or with 1 \(\mu\)mol/l of amyloid-\(\beta\)1–40 for 2 h prior to analysis (c and f). Scale bars indicated apply to all panels.
readily detected in parenchyma and perivascular space in hippocampus (Figure 7A) and cerebral cortex (Figure 7B) of transgenic mice. Some perivascular staining, but not Congo Red-stained plaques in parenchyma, was detected in the inferior colliculus of APP mice (Figure 7C). However, this does not rule out the presence of amyloid plaques in this structure, since a higher-sensitivity method to detect immunoreactive amyloid protein was not used. Because Congo Red can also bind to non-amyloid substances and some small red-stained areas (data not shown) were observed in white matter and in control brain, Congo Red-stained plaques were also examined by polarized microscopy; the plaques exhibited red–green birefringence (data not shown) which is characteristic of amyloid deposits (Jin et al., 2003).

Astrocytes in slices of hippocampus (Figure 8a), cerebral cortex and inferior colliculus from mixed-gender control and Alzheimer’s mice were identified as SR101-positive cells (Nimmerjahn et al., 2004; Ball et al., 2007), then a single astrocyte was impaled with a micropipette (Figure 8b) containing Lucifer Yellow VS (Figure 8c), and the dye was allowed to diffuse into the astrocytic syncytium for 5 min. Many dye-labelled astrocytes were readily detected in the 7 μm-thick serial sections of the 250 μm-thick slices from all three brain structures of Alzheimer’s and control mice. The labelling patterns differed in the three brain structures (Figures 8d–8f), although there was also some within-structure variation due to the exact location of the impaled astrocyte in each structure. In the hippocampus, labelling derived from impaled astrocytes located near the pyramidal neuronal layer was prominent and most dense in the astrocytes along the pyramidal neuronal layer, with labelled cells above and below the pyramidal layer (Figure 8d). In contrast, dye spread among astrocytes in the cerebral cortex caused labelling of cells in an approximate circular pattern around the impaled cell (Figure 8e). Astrocytic labelling in the
inferior colliculus in the thin sections tended to be scattered (Figures 8f), but most of the 250 μm-thick sections (data not shown) had a radial labelling pattern that extended from the injected cell towards the meningeal boundary, as we previously observed (Ball et al., 2007).

The total number of dye-labelled cells was quite variable in both control and APP mice, ranging from a low of approx. ten, up to several thousand coupled cells, with no statistically significant differences in the number of dye-labelled cells in control and amyloid-treated groups for any of the three brain regions (Figure 9). When data from all slices from all animals were plotted as a function of age, there were no obvious age-dependent trends over the relatively narrow range of ages of mice used in the present study; control and APP mice had variable and overlapping numbers of dye-labelled cells (Figure 10). Unfortunately, the number of APP mice available for this study was limited by breeding problems that persisted over approx. 2 years of the funding period; this issue reduced the number of ages and number of regional brain slices that were assessed, as well as the scope of planned work which included evaluation of gap junctional trafficking of endogenous metabolites and local rates of glucose utilization during brain activation.

**DISCUSSION**

Discordant results in tissue culture and brain slices: complications can arise from severe hyperglycaemia

The major finding of the present study is that exposure of cultured cerebral cortical astrocytes to low-dose monomerized amyloid-β markedly impaired transfer of material within the synctial network, but dye transfer was unaffected in aged transgenic mice overexpressing amyloid protein. In low-glucose cultures, amyloid-β treatment affected transcellular diffusion of six anionic and neutral compounds with...
molecular masses ranging from approx. 350 to 700 Daltons; it had a rapid onset (within 2 h), it was accompanied by modest, transient oxidative stress and the inhibition was sustained for at least 5 days after a single treatment (Figures 1–6). Because transcellular movement of NADH, NADPH and 6-NBDG were reduced by amyloid treatment, signals used to monitor redox status and glucose transport and utilization are affected and could influence metabolic studies, particularly in cultured astrocytes. Amyloid-β depressed gap junctional trafficking to a similar level in high- and low-glucose cultures, but those grown in high glucose had a smaller overall percentage decrease (Figures 1 and 2) owing to reduced dye transfer among the severely hyperglycaemic cells (Gandhi et al., 2010, Thi et al., 2010).

Diabetes is a risk factor that exacerbates pathophysiological changes in Alzheimer’s transgenic mice and cultured cells, with synergistic interactions that increase generation of advanced glycation products and ROS and diminish cognitive function (Burdo et al., 2009). STZ-diabetic rats have elevated levels of amyloid-β in brain (Li et al., 2007; Liu et al., 2008), but, to our knowledge, gap junctional trafficking has not been evaluated in a diabetic-Alzheimer’s rodent model. Interpretation of the inhibitory effects on glucose metabolic pathways, glutamate uptake, signalling pathways, gap junctional communication and other processes that have been ascribed to amyloid peptides in other studies are complicated by the use of astrocytes that were cultured in media containing 25–40 mmol/l glucose (Parpura-Gill et al., 1997; Même et al., 2006; Matos et al., 2008; Allaman et al., 2010) or by lack of information regarding medium glucose concentration (Meske et al., 1998; Haughy and Mattson, 2003; Chow et al., 2010). A glucose level of 25 mmol/l is approx. 10-fold greater than that in normal rat brain, and it exceeds that in diabetic rat brain by 3.6-fold; it is well known that prolonged hyperglycaemia causes oxidative stress, glycation reactions and sorbitol production (Gandhi et al., 2010 and references cited therein). In fact, astrocytes grown in high-glucose medium oxidize glucose and lactate at a rate half that of those grown in low glucose (Abe et al., 2006). Exposure to amyloid peptides and amyloid aggregates has deleterious consequences for cultured astrocytes and neurons, but the possibility of synergistic interactions of amyloid treatment with severely
diabetic conditions that potentiate damage needs to be examined and clarified.

Influence of inflammation, calcium level and age or brain region on dye coupling.

Rapid and sustained down-regulation of dye spread among cultured astrocytes by amyloid-β1-40 contrasts the results of Méme et al. (2006), who found no effect of amyloid-β25-35 on Lucifer Yellow diffusion through gap junctions in scrape-load assays; this smaller peptide did, however, enhance the inhibitory effects of cytokines released from LPS (lipopolysaccharide)-activated microglia that were grown in mixed cultures with astrocytes (38% microglia). Notably, non-activated mixed cultures had similar dye-labelled areas as astrocyte cultures (38% microglia). Notably, non-activated mixed cultures had similar dye-labelled areas as astrocyte cultures (38% microglia). Notably, non-activated mixed cultures had similar dye-labelled areas as astrocyte cultures (38% microglia). Notably, non-activated mixed cultures had similar dye-labelled areas as astrocyte cultures (38% microglia). Notably, non-activated mixed cultures had similar dye-labelled areas as astrocyte cultures (38% microglia).

High calcium levels can close gap junctions, but the physiological relevance of the level required is not clear (Rozental et al., 2001). Amyloid peptides are thought to form calcium channels in the astrocytic membrane, giving rise to intracellular calcium signals and oxidative stress due to activation of NADPH oxidase (Abramov et al., 2004). Exposure of astrocytes to amyloid-β1-42 increases intracellular calcium concentration, as well as the amplitude, velocity and travel distance of mechanically induced calcium waves (Haughey and Mattson, 2003). Acute dosing with amyloid-β1-40 causes

Figure 8 Representative images of Lucifer Yellow diffusion among astrocytes in slices from three brain regions of aged Alzheimer’s transgenic mice

Brain slices (250 μm thick) were prepared from hippocampus (a–b), cerebral cortex (e) and inferior colliculus (f) of aged Alzheimer’s transgenic mice. Astrocytes were identified as sulforhodamine-positive cells (a), then impaled with a micropipette (b) containing Lucifer Yellow that was allowed to diffuse into the astrocyte for 5 min (c). Then the sections were immediately fixed, cut into 7 μm-thick serial sections and dye-labelled cells counted in each section from each slice. Fluorescence images of typical sections are shown for each brain region to illustrate the different patterns of dye-labelled cells (bright spots) derived from impaling an astrocyte near the pyramidal neuronal layer in the hippocampus (d), or an astrocyte in cerebral cortex (e) and inferior colliculus (f). Arrowheads in (a), (e) and (d) denote dark spaces corresponding to unlabelled neuronal cell bodies. The scale bar in (a) also applies to (e).

Figure 9 Dye transfer in brain regions in slices from aged control and Alzheimer’s mice

Total number of Lucifer-Yellow-labelled cells in three brain regions (A, B and C) from control and Alzheimer’s transgenic mice ranging in age from 8.5 to 14 months. Each point represent one brain slice; 1–3 slices were obtained from each region of six control and six transgenic mice. The number of slices for inferior colliculus, cerebral cortex and hippocampus respectively, were: controls, 10, 5, 6; transgenic, 10, 5, 5. Note that two of the control cortical slices had similar numbers of labelled cells and the values in the Figure overlap. There were no statistically significant differences between the control and experimental groups in each region.
time-delayed spontaneous calcium waves (Chow et al., 2010), whereas various ‘aged’ amyloid peptide preparations reduce intracellular calcium level in a dose- and time-dependent manner (Meske et al., 1998). In our gap junction assay procedure, impaling an astrocyte would initiate calcium waves, but dye spread does occur in control and amyloid-treated cells, and the contribution of calcium to differential channel closure in these two groups of cells, if any, is not clear. Other factors, such as amyloid-induced alterations in membranes that may modify Cx function or level, altered signalling cascades, and shifts in gene expression, may also be involved at different times after addition of monomerized amyloid-β1–40. The basis for the inhibitory bioactivity of amyloid-β on gap junctions in cultured astrocytes remains to be established in future studies.

Surprisingly, the robust blockade of dye transfer within networks of amyloid-β-treated cultured cerebral cortical astrocytes did not occur in cerebral cortex, hippocampus or inferior colliculus of APP-transgenic mice (Figures 8–10). These findings indicate that: (i) amyloid treatment alone is not sufficient to represent the more complex pathophysiology of astrocytic networks in the aged APP mouse brain, (ii) there may be compensatory responses that maintains syncytial connectivity in APP mouse brain, beyond the localized up-regulation of immunoreactive Cx43 observed in human amyloid plaques (Nagy et al., 1996a) and (iii) global compared with regional differences do not explain the different results in cultured cells and brain slices. Discordant results in tissue culture-brain slice assays were not, however, an issue in our recent study of experimental diabetes, in which both model systems exhibited reduced dye coupling and increased oxidative stress in spite of large differences in glucose level, time of exposure to high glucose and syncytial size (Gandhi et al., 2010). In the diabetic rat literature, plasma and brain glucose levels rise an average of approx. 3-fold to 27 and 7 mmol/l respectively (Gandhi et al., 2010), and the normal adult rat inferior colliculus has up to approx. 12000 dye-coupled cells compared with 10–20 cells in cultured cortical astrocytes (Ball et al., 2007 and references cited therein). In the present study, most slices from aged control and APP mice had > 100 coupled cells, with an average regional number of dye-coupled cells ranging from approx. 300 to 1100 (Figures 9 and 10), values that are much lower than in young adult rats. Peters et al. (2009) recently characterized properties of astrocytes in aged Alzheimer’s transgenic mice and observed modest syncytial labelling (≤ 114 cells) with biocytin [molecular mass 372 Da; diffused into cells for 20 min via patch clamp pipettes containing a 0.5% (13 mmol/l) solution] in young control mice and aged Alzheimer’s mice, with no apparent relationship to distance of the injected cell from an amyloid plaque. In contrast, their aged control mice (> 12 months) had no detectable networks in cerebral cortex and only three out of 20 cells in hippocampal slices were coupled. The reasons for lower coupling in aged APP mice and very poor dye transfer in their aged controls are not known. Conceivably, differences might arise from the genotype of the mice used in the studies by Peters et al. (2009) and by us, or from age-related differences in transfer of biocytin compared with Lucifer Yellow VS. Also, it is important to recognize that changes in transfer of a specific dye do not necessarily predict the passage of other compounds. For example, when diffused into single astrocytes in culture or in brain slices, hexose-6-phosphates are very poorly transferred through gap junctions compared with various fluorescent dyes that were used as internal permeant standards in the same assays (Gandhi et al., 2009a). Our dye-transfer assays used high concentrations of Lucifer Yellow VS to maximize sensitivity for detection of astrocytic coupling and to more easily count labelled cells in brain slices, since dye-labelled nuclei are more prominent with Lucifer Yellow VS compared with Lucifer Yellow CH. Dye leakage from impaled cells is unlikely to cause significant labelling due to washout via the perfusion solution and to negligible labelling by extracellular membrane-impermeant dye; labelling of brain slices caused by micropipette insertion or by damaging an impaled cell is very low. Also, in our previous studies, Lucifer Yellow labelling of cultured astrocytes located at some distance from the scrape-load site was negligible (Gandhi et al., 2010), and efflux of Lucifer Yellow (and, by inference, uptake from extracellular sources) from pre-loaded cells was not detectable unless cells were exposed to a divalent-cation-deficient medium to open ‘hemichannel’ or pannexin pores (Gandhi et al., 2009a). To summarize, astrocytic gap junctional communication in different model systems for Alzheimer’s disease varies considerably, and in vivo responses may be influenced by amyloid burden, age, local cytokine level, oxidative stress and other unidentified factors.

**Metabolite trafficking and brain imaging in Alzheimer’s disease**

The hippocampus and cerebral cortex are involved in memory and cognition, and these brain regions are, therefore,
commonly used as experimental systems for studies of Alzheimer’s disease and loss of higher brain function. However, other brain systems are also involved in the disease, and, for example, Alzheimer’s patients have hearing impairment (Palmer et al., 1999) and abnormal auditory evoked potentials and latency responses associated with midbrain–brainstem auditory processing (O’Mahony et al., 1994). The inferior colliculus, an auditory structure with the highest metabolic rate in brain (Sokoloff et al., 1977), has amyloid plaques (Sinha et al. 1993) and a lower level of cytochrome oxidase (Gonzalez-Lima et al. 1997), indicating a reduced capacity for oxygen utilization. Dye transfer was therefore examined in these three brain structures after testing a variety of fluorescent compounds in tissue culture. Assay of local rates of glucose utilization using \(^{18}\text{F}\)FDG-PET (\(^{18}\text{F}\)fluorodeoxyglucose–positron emission tomography) has been an important tool to evaluate the progression and treatment of Alzheimer’s disease (Alexander et al., 2002), and magnetic resonance spectroscopic studies using \(^{13}\text{C}\)glucose are increasingly used to study brain diseases because fluxes of labelled carbon into different pathways can be evaluated. Progressive mitochondrial dysfunction and impaired oxidative metabolism of glucose in brain of Alzheimer’s patients (Gibson et al., 2008) would increase lactate production and release from brain. Incomplete retention of labelled lactate and other diffusible compounds in regions of interest during metabolic assays would contribute to underestimation of glucose utilization rates determined with labelled glucose compared with fluorodeoxyglucose, the products of which are phosphorylated and trapped intracellularly. Also, increases in glycolytic flux in the early stages of the disease to compensate for energy deficits arising from mitochondrial damage could give the false impression that cellular function is normal, when, in fact, cellular damage is evolving, and by the time glucose utilization rates fall, the damage may be irreversible or reflect cell death. For these reasons, it is particularly important to use different approaches to evaluate functional metabolism in diseased brain. Our previous studies have focused on evaluating the cellular contributions to discordant images of auditory activation in rats derived from labelled glucose and deoxyglucose (Cruz et al., 2007). The results strongly support the conclusions that: (i) lactate release from activated brain accounts for most of the ‘missing’ glucose metabolites, (ii) lactate transporters and astrocytic gap junctional trafficking are involved in this process and (iii) release of some glucose, lactate, other metabolites and amyloid-\(\beta\) occurs by lymphatic drainage from the inferior colliculus through perivascular channels to the cervical lymph nodes (Dienel and Cruz, 2008; Gandhi et al., 2009a, 2009b; Ball et al., 2010). Amyloid deposits accumulate in the parenchyma and perivascular space in brain of animals with Alzheimer traits (e.g. Figure 7) and in Alzheimer’s patients, and partial blockade of perivascular fluid flow is postulated to contribute to the pathophysiology of Alzheimer’s disease (Carare et al., 2008; Weller et al., 2008). In diabetic patients, cerebrovascular disease and thickening of the extracellular matrix would exacerbate this condition by impairing vascular elasticity, diminishing arterial pulsations that power perivascular fluid flow and reducing the distribution within brain of fuel, signalling molecules, neurotransmitters and electrolytes, as well as clearance of metabolic by-products from activated regions. Further work is necessary to: (i) evaluate astrocytic networks in aged control brain compared with aged Alzheimer’s brain and (ii) determine whether changes in metabolite trafficking and functions of the neurovascular unit (i.e. interactions of astrocytes, neurons and endothelial cells involved in cerebrovascular and metabolic regulation) influence the outcome and interpretation of brain imaging and spectroscopic studies in Alzheimer’s patients.

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