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Increased MCP-1 and Microglia in Various Regions of the Human Alcoholic Brain

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Abstract

Cytokines and microglia have been implicated in anxiety, depression, neurodegeneration as well as the regulation of alcohol drinking and other consumatory behaviors, all of which are associated with alcoholism. Studies using animal models of alcoholism suggest that microglia and proinflammatory cytokines contribute to alcoholic pathologies (Crews et al., 2006). In the current study, human postmortem brains from moderate drinking controls and alcoholics obtained from the New South Wales Tissue Resource Center were used to study the cytokine, monocyte chemoattractant protein 1 (MCP-1, CCL2) and microglia markers in various brain regions. Since MCP-1 is a key proinflammatory cytokine induced by chronic alcohol treatment of mice, and known to regulate drinking behavior in mice, MCP-1 protein levels from human brain homogenate were measured using ELISA, and indicated increased MCP-1 concentration in ventral tegmental area (VTA), substantia nigra (SN), hippocampus and amygdala of alcoholic brains as compared with controls. Immunohistochemistry was further performed to visualize human microglia using ionized calcium binding adaptor protein-1 (Iba-1), and Glucose transporter-5 (GluT₅). Alcoholics were found to have brain region-specific increases in microglial markers. In cingulate cortex, both Iba-1 and GluT₅ were increased in alcoholic brains relative to controls. Alternatively, no detectable change was found in amygdala nuclei. In VTA and midbrain, only GluT₅, but not Iba-1 was increased in alcoholic brains. These data suggest that the enhanced expression of MCP-1 and microglia activities in alcoholic brains could contribute to ethanol-induced pathogenesis.

Keywords

Alcohol; MCP-1; microglia; Iba-1; GluT₅; human

Introduction

Alcohol (ethanol) is a common dietary constituent that modulates the immune system. Although moderate alcohol consumption has a protective effect on heart diseases and appears to have health benefits, heavy drinking increases mortality by escalating the risk of many diseases, especially disorders of the central nervous system (Nelson and Kolls, 2002). In humans, chronic alcohol consumption is associated with increases in serum

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proinflammatory cytokines including TNF α and IL-1 β (McClain and Cohen, 1989; McClain et al., 1999). Monocytes isolated from the blood of alcoholics produce greater amounts of TNF α , a proinflammatory cytokine, spontaneously and in response to endotoxin challenge (McClain et al., 2004). The current study is designed to further investigate the roles of microglial recruitment and activation in alcohol induced inflammatory neurodegeneration (Lee et al., 2004; Crews et al., 2006; Pascual et al., 2007).

Microglia, as the resident monocyte/macrophage in CNS parenchyma (Gehrmann et al., 1995), play an important role in the brain having both a beneficial healing and a toxic inflammatory role. Under resting conditions, microglia exhibit a quiescent phenotype as indicated by a ramified morphology, and constitutive expression of macrophage antigens such as ionized calcium binding adaptor protein –1 (Iba-1) (Ladeby et al., 2005). In response to a multitude of CNS pathological conditions and systemic infectious processes, microglia transform from a resting to an active state characterized by cellular enlargement, amoeboid morphology, increased membrane ruffles and projections and upregulation of several cell surface molecules and proteins (Block et al., 2007). Iba-1 is a microglia/macrophage-specific protein that participates in the ruffling and phagocytosis of activated microglia in human, rats and mice (Ohsawa et al., 2004). Glucose transporter type 5 (GluT₅) is another microglia specific protein (Horikoshi et al., 2003; Sasaki et al., 2004), whose expression contributes to the kinetics of cerebral metabolism (Vannucci et al., 1997). Thus, studying the expression of Iba-1 and GluT₅ can provide critical information on microglia recruitment, morphological transformation and functional activation after life-long alcohol exposure in human alcoholic brains.

Furthermore, monocyte chemoattractant protein-1 (MCP-1, also known as CCL₂) is a key cytokine mediating CNS inflammation and alcohol drinking behavior (Blednov et al., 2005). As a predominant chemoattractant, MCP-1 causes the migration and activation of microglia (McManus et al., 2000), whose secretions of proinflammatory cytokines can be neurotoxic (Kaul et al., 2001; Little et al., 2002; Mahad and Ransohoff, 2003; Persidsky and Gendelman, 2003). As a potential neurotoxin, enhanced expression of MCP-1 increases the volume of an infarct after middle cerebral artery (MCA) occlusion (Chen et al., 2003), whereas MCP-1 knockout mice have smaller infarcts and less neuronal loss compared to their wild-type controls (Hughes et al., 2002). The production of the proinflammatory cytokines such as IL-1 β (interleukin-1 beta) and TNF α (tumor necrosis factor alpha) is significantly reduced in MCP-1 knockout mice challenged with LPS (lipopolysaccharide) (Rankine et al., 2006). These knockout mice also exhibit a substantial reduction in alcohol consumption and preference (Blednov et al., 2005). Therefore, the level of MCP-1 expression in human alcoholic brain is critical for further understanding the mechanisms of alcohol-induced neuroinflammation.

The current study was designed to investigate the inflammatory status in postmortem alcoholic brains by determining microglial activities using Iba-1 and GluT₅ labeling, and the key inflammatory cytokine, MCP-1, expression. Results indicate increased MCP-1 protein levels in ventral tegmental area (VTA), substantia nigra (SN), hippocampus and amygdala of alcoholic brains compared to controls. Furthermore, increased microglia expressions of Iba-1 and/or GluT₅ in specific regions of alcoholic brain were found.

Methods

Tissue Source

Human postmortem brain tissue was obtained from the New South Wales Tissue Resource Center in Australia [ethics committee approval number: HREC2002/2/3.14 (1441) and X03-0117]. Both paraffin sections for microglial staining and fresh frozen brain tissue for

ELISA were used in this study. The detailed patients' medical history is presented in tables 1 and 2.

Enzyme-Linked Immunosorbent Assay (ELISA)

Frozen brain samples were homogenized in the buffer containing: 20mM Tris, 0.25mM sucrose, 2mM EDTA, 10mM EGTA, and 1% Triton X-100. One protease inhibitor cocktail tablet (Roche Diagnostics, Germany) was added into each 10ml of the homogenization buffer. The homogenate was then centrifuged at 100,000g (Beckman Ultracentrifuge) for 45 min at 4°C and supernatants were collected and total protein content was determined by using BCA protein Assay Reagent Kit (Pierce, Rockford, IL) to ensure that an equal amount of protein from each sample was used for the assay. The MCP-1 ELISA was conducted using Human MCP-1 immunoassay kit (R&D systems, Minneapolis, MN) and the manufacture's protocol was followed. The optical density of each sample was determined using Spectra Max microplate reader (Molecular Devices, Sunnyvale, CA) and the concentration of MCP-1 was calculated based on standards and expressed in pg/mg of total protein content.

Immunohistochemistry

Paraffin sections were deparaffinized in Xylene for 30 min and rehydrated in a series of ethanol from 100% to 50%. The slices were then washed in Tris-buffered saline (TBS) buffer for 15min. Antigen retrieval was done by incubating the section in Citra solution (BioGenex, San Ramon, CA) in a steamer for 30min. The antibodies against rabbit Iba-1 (Wako Pure Chemical Industries, Japan) or rabbit-GluT₅ (IBL, Japan) were used at a dilution of 1:400 and 1:20 respectively with overnight incubation at 4°C. Sections were then rinsed TBS, and incubated with biotinylated goat anti-rabbit secondary antibody (Vector Laboratories, Burlingame, CA, USA) at a dilution of 1:200 at room temperature for 2hr. Subsequently, avidin-biotin-peroxidase complex (ABC Elite Kit, Vector Laboratories) was applied for 1hr at room temperature. Finally, the positive cells were visualized using nickel-enhanced diaminobenzidine (DAB) as a chromagen.

Microscopic Quantification

Both Iba-1 and GluT₅ (Glucose transporter 5) positive cells were quantified using image analysis software (Bioquant Nova Advanced Image Analysis, R&M Biometric, Nashville, TN). Images were captured on an Olympus BX50 microscope and Sony DCX-390 video camera at 40X. Light levels were normalized to preset levels and the microscope, camera, and software were background corrected to ensure reliability of image acquisition (Crews et al., 2004). In each region (cingulate cortex, midbrain, VTA, and amygdala), three random images from each brain sample were captured within a standard ROL (Region of Interest), and staining density was measured in pixels within this area (pixels/mm²). Subsequently, the average of the three measurements was used to represent the immunoreactivity of each sample.

Statistics

All values were reported as mean ± SEM, and analyzed by ANOVA, and Pearson correlation test using SPSS. Differences were considered significant if the p value was smaller than 0.05.

Results

The demographic characteristics of control and alcoholic subjects are given in Table 1 and 2. Mean ages in MCP-1 study using frozen brain homogenates were not significantly different

being 60.8 ± 3.6 years old in controls and 63.8 ± 3.1 years old in alcoholics. The mean postmortem interval (PMI) was 29.0 ± 3.0 in controls and 36.6 ± 6.0 in alcoholics. One-way ANOVA analyses indicated no significant differences in age ($F_{(1,13)} = 0.38, p=0.55$), or PMI (post-mortem interval) ($F_{(1,13)} = 0.96, p=0.35$) between alcoholics and controls. For the study using cingulate cortex paraffin sections, the mean ages of the subjects were 47 ± 3.8 yr in the control group and 49 ± 4.4 yr in the alcoholic group. The average PMIs in these populations were 24.5 ± 1.8 hr in controls and 24.5 ± 4.4 hr in alcoholics. ANOVA indicated no significant differences in age ($F_{(1,7)} = 0.89, p=0.77$) or PMI ($F_{(1,7)} = 0.00, p=1.0$) between control and alcoholic groups. The mean ages of the subjects providing paraffin sections for VTA, midbrain, and amygdala were 64.1 ± 3.7 yr in control group and 63.4 ± 3.5 yr in alcoholic group. The average PMIs for these subjects were 25.4 ± 3.6 hr in controls and 31.4 ± 7.9 hr in alcoholics. No significant differences were found in age ($F_{(1,14)} = 0.02, p=0.86$) or PMI ($F_{(1,14)} = 0.48, p=0.50$) between control and alcoholic groups. In addition, we looked at smoking history and gender as factors. However, we were not able to analyze these variables separately due to the small sample size. These results suggest that neither age nor PMI were different between control and alcoholic groups.

Increased MCP-1 Protein Concentration in Various Regions of the Alcoholic Brains

The MCP-1 protein concentration (pg/mg of total protein) determined by ELISA was compared in each brain region between alcoholics and control brains (Fig. 1). In VTA, the average MCP-1 concentrations were 212.6 ± 43.5 pg/mg in alcoholic brains ($n=7$), and 90.2 ± 20.9 pg/mg in controls ($n=6$), and a significant two-fold increase in MCP-1 in alcoholic brains was detected by ANOVA ($F_{(1,11)} = 5.8, p=0.035$). In the substantia nigra (SN), the MCP-1 concentration was 314.04 ± 59.6 pg/mg in alcoholics ($n=6$), which is significantly higher than controls (121.0 ± 14.3 pg/mg in controls, $n=5$) [ANOVA ($F_{(1,9)} = 8.3, p=0.018$)]. In the hippocampus, MCP-1 levels were 243.14 ± 40.0 pg/mg in alcoholics ($n=8$) and 106.46 ± 24.1 pg/mg in controls ($n=6$) [ANOVA ($F_{(1,13)} = 6.6, p=0.023$)]. Similarly, in the amygdala nuclei, the MCP-1 concentration in alcoholics was 322.55 ± 70.5 pg/mg ($n=8$), which is almost three-fold higher than that of the controls (113.26 ± 25.6 pg/mg, $n=6$) [ANOVA ($F_{(1,12)} = 6.0, p=0.03$)]. In addition, Pearson tests indicated no significant correlations were found between MCP-1 concentrations and age or PMI in all regions studied (p values > 0.05). These data indicate that alcoholic brains have significantly higher MCP-1 levels than those of controls across multiple limbic brain regions.

Increased Microglial Expression of Iba-1 and GluT₅ in Different Brain Regions of the Alcoholic Brains

To investigate microglia in various brain regions, the immunoreactivity of two specific microglia markers: Iba-1 and GluT₅ were determined. Stages of microglia activation were assessed based on their morphological characteristics from ramified resting stage to bushy-looking early activation, and then to activated amoeboid-like brain macrophages (Ladeby et al., 2005). In both controls and alcoholic brains, microglia were found in all stages of activation (Fig.2). Within all brain regions examined, many of the microglia had a variable morphology but there were no clear differences between control and alcoholic brains in degree of microglial activation, as indicated by morphology. However, alcoholic brains had considerable upregulation in microglial marker immunoreactivity in several brain regions when measured with image analysis software (BioQuant). In cingulate cortex (Fig. 3, Table 3), the immunoreactive density (pixel/mm²) of alcoholics was significantly increased for $214 \pm 11\%$ in Iba-1 ($F_{(1,7)} = 11.7, p=0.01$) and $194 \pm 25\%$ in GluT₅ ($F_{(1,7)} = 9.9, p=0.016$). In VTA (Fig. 4, Table 3), GluT₅ immunoreactivity was increased significantly in alcoholics for $175 \pm 17\%$ ($F_{(1,14)} = 12.5, p=0.003$) compared to controls, but no statistical difference in Iba-1 was found ($F_{(1,14)} = 1.18, p=0.30$). Similarly, in the midbrain (Fig. 5, Table 3), alcoholics had increased GluT₅ immunoreactivity for $156 \pm 13\%$ ($F_{(1,9)} = 22.66, p=0.01$), but

not Iba-1 ($F_{(1,9)} = 0.48$, $p = 0.51$) as compared with controls. In amygdala nuclei (Fig. 6, Table 3), the immunoreactivity to neither Iba-1 ($F_{(1,14)} = 0.32$, $p = 0.58$) nor GluT₅ ($F_{(1,14)} = 0.58$, $p = 0.46$) was significantly different between alcohol and control brains. In addition, none of these immunoreactive measures was significantly correlated with either age or PMI by Pearson tests (p values > 0.05). Taken together, these data indicated that the expression of the specific microglial markers in alcoholics is increased in cingulate cortex, but not in amygdala, as compared to controls. In VTA and midbrain, enhanced GluT₅ staining was found in alcoholic brains.

Discussion

We report here, for the first time, significantly increased MCP-1(CCL2) across multiple alcoholic brain regions. MCP-1 is a member of the Beta chemokine subfamily that signals through a G protein coupled receptor, CCR2 (Banisadr et al., 2002). Within the brain, MCP-1 is produced mostly by activated astrocytes and microglia (Glabinski et al., 1996). The impact of MCP-1 on neuroinflammation involves multiple pathways. As a classic chemoattractant, MCP-1 plays an important role in driving monocyte recruitment to injured tissue (Lu et al., 1998), possibly by increasing the permeability of the blood-brain-barrier (BBB) (Yamamoto et al., 2005; Stamatovic et al., 2003; Stamatovic et al., 2005). The subsequent migration and activation of these monocytes (microglia in the brain) leads to excessive production of proinflammatory/neurotoxic cytokines such IL-1 β and TNF α (Kaul et al., 2001; Mahad and Ransohoff, 2003; Persidsky and Gendelman, 2003). Mice that overexpress MCP-1 show increased Iba-1 immunoreactivity and accelerated senescent neurodegeneration (Yamamoto et al., 2005). Moreover, the upregulation of MCP-1 in CNS tissue can exacerbate neuronal death and other pathologies occurs before the detectable monocyte recruitment (Rankine et al., 2006), indicating that the contributions of MCP-1 to neuroinflammation are far beyond its role as a chemoattractant (Hughes et al., 2002; Rankine et al., 2006). A recent study found that trimethyltin induced hippocampal degeneration involved marked MCP-1 induction, without TNF α , IL-1, IL-6, or other proinflammatory cytokines (Little et al., 2002). Endogenous MCP-1 is also known to directly induce neuronal apoptosis (Kalehua et al., 2004). Thus, increased MCP-1 in alcoholic brains could directly cause neuronal damage, and could be one of the mechanisms contributing to alcohol-related neuronal loss and brain atrophy (Brooks, 2000; Harper et al., 2003; Ikegami et al., 2003). Lastly, it has been proposed that MCP-1 can act as a “priming” stimulus for microglia (lowering their “threshold sensitivity”), enhancing their synthesis of proinflammatory cytokines in response to subsequent stimulation (Rankine et al., 2006). This priming of microglia can occur as a consequence of a wide range of CNS diseases and other stressors including alcohol exposure. Interestingly, transgenic mouse studies have indicated that downregulation of MCP-1 expression increases hypnotic response and decreases alcohol preference (Blednov et al., 2005). These data consistently indicate an association between the level of MCP-1 expression and alcoholic pathologies. Taken together, the current findings of increased MCP-1 in human alcoholic brain further support the critical involvement of neuroinflammation in both behavioral and neurodegenerative pathologies associated with alcoholism.

Since enhanced levels of MCP-1 expression can lead to activation or migration of microglia, we chose two specific microglial markers, Iba-1 and GluT₅ to identify and study microglia in alcoholic brains as compared to controls. Microglia undergo a series of characteristic morphological transformations that are thought to represent stages of increasing proinflammatory and phagocytic activity that coincide with induction of many proteins, including the constitutively expressed microglial markers, Iba-1 and GluT₅. Morphologically, we observed all forms of microglia: ramified (resting), activated, and amoeboid (phagocytic) microglia in both alcoholic brains and controls across multiple brain

regions. Microglia are known to have increasingly diverse morphologies, including the highly activated phagocytic phenotypes in normal senescent brains as well as increased numbers of activated monocytes in degenerative diseases. Our tissues used for histochemistry are from patients with ages ranging from 52–82 years at death. We could not detect differences between alcoholic and control brains in the spectrum of microglial morphologies. However, the advanced age of our subjects may have masked alcohol induced changes.

We found increased microglial markers, both Iba-1 and GluT₅ immunoreactivity, in cingulate cortex of alcoholics. Cingulate cortex, along with prefrontal areas, contribute to the frontal attention system with cingulate cortex representing a cross road between the limbic system and frontal cortex (Fuster, 1997). It has been suggested that the disruption of cognitive function by alcohol exposure could be due to its effects on frontal cortical areas (Tu et al., 2007). Increases in microglial immunoreactivity indicate the enhanced microglia number or their activation in this region. Increases in microglial specific staining could be due to more microglia and/or larger activated microglia. Since we did not see significant changes in morphology, these findings likely indicated that the cingulate cortex of alcoholics has about 2-fold more microglia than moderate drinking controls. This could at least partially result from the increased MCP-1 expression, which in turn leads to the migration and/or proliferation of microglia in cingulate cortex.

In VTA and midbrain, we found a significant increase in GluT₅ staining, but not Iba-1. This discrepancy could be the result of the higher sensitivity with human-specific GluT₅ antibody as compared to Iba-1, which identifies microglia from human, mouse and rat (Ohsawa et al., 2004). Microglia are known to alter expression of multiple proteins as they progress through various states of activation. However, both Iba-1 and GluT₅ are microglia markers that are expressed at all stages of microglial activation, although their expression increases with activation. It is possible that alcohol-induced changes in VTA and midbrain GluT₅ are relatively subtle, which can be detected only by the more sensitive GluT₅ marker, but not by Iba-1. Regardless, the increase in VTA and midbrain GluT₅ staining is consistent with increased microglia activation. Although the precise role of VTA in drinking behavior is not known, studies have suggested the dopamine D₂ receptors in VTA neurons are responsible for drinking behaviors (Eiler and June, 2007). However, these receptors are also expressed on microglia in VTA (Farber et al., 2005), which imply the involvement of VTA microglia in alcohol-related pathogenesis. Thus, increased VTA microglia activation could one of the mechanisms contributing to alcoholism. In amygdala, no detectable differences were found in the immunoreactivity of either GluT₅ or Iba-1 between alcoholic and control groups. Although studies have shown that amygdala is a key region mediating anxiety-like behaviors associated with chronic alcohol exposure and withdrawal (Knapp et al., 2007; Lack et al., 2007), the functional roles of the microglia in this region are unknown.

Taken together, the current findings indicate a region specific increase in microglia activities. Although the exact mechanisms of this regional discrepancy are unclear, we speculate that the increase in microglia in the cingulate cortex reflects higher vulnerability of this region to alcohol-induced inflammatory changes. Chronic alcohol exposure to mice has been shown to cause cellular changes in cingulate cortex (Marrero-Gordillo et al., 1998) and to disrupt cognitive function (Tu et al., 2007). Furthermore, although the exact mechanism of increased microglial activation is not known, our data suggest that elevated MCP-1 levels could at least partially, contribute to increased microglia staining intensity. As the first responder to environmental insults, microglia are believed to play dual roles in both neurodegeneration and neuroprotection (Nimmerjahn et al., 2005). While local microglia activation at the injured site might be neuro-destructive by releasing cytotoxins (Meda et al., 1995; Brown et al., 1996; Barger and Harmon, 1997; McDonald et al., 1997), the delayed

accumulation and activation of microglia resulting from MCP-1 could contribute to elimination of cellular debris and may be involved in neuroprotection by producing neurotrophic factors (Hermann et al., 2001; Laurenzi et al., 2001; Hashimoto et al., 2005; Lu et al., 2005). Although the relationship between microglia activation and MCP-1 expression in humans requires further investigation, our animal studies have found that chronic ethanol induces brain MCP-1 in mice (Qin et al., 2007) and binge alcohol withdrawal increases microglia in rat brains (Nixon and Crews, 2004). Even though many studies have found volume reduction in alcoholic brain, the specific loss of neurons in humans remains controversial (Harper, 1998). One study using unbiased sampling found a selective loss of superior frontal cortical neurons, particularly large pyramidal neurons in alcoholics (Kril et al., 1997). Our findings of increased MCP-1 and microglial markers could either contribute to or result from alcoholic neurodegeneration indicated by brain shrinkage and neuronal loss in human alcoholics. Altogether, our current findings support the role of inflammatory neurodegeneration in alcohol-related neuropathology of humans.

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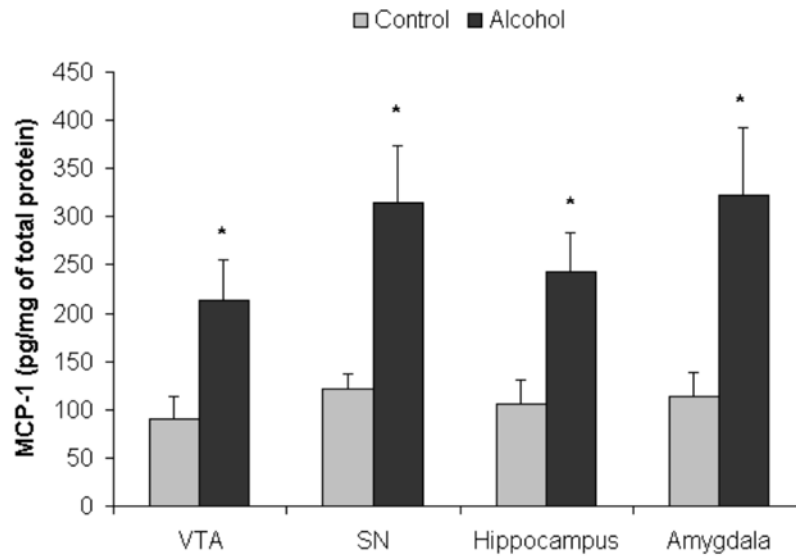


Fig. 1. Increased MCP-1 protein concentrations in alcoholic brains. MCP-1 protein concentrations (pg/mg of total protein) from brain homogenate were measured using ELISA represented as mean \pm SEM. Using ANOVA, significantly increased MCP-1 expression was detected in alcoholics as compared to control of in VTA (ventral tegmental area) of alcoholics as compared with controls (*, $p < 0.05$, $N=5$ controls, $N=7$ alcoholics), in substantia nigra (SN) (* $p < 0.05$, $N=5$ controls, $N=6$ alcoholics), in hippocampus (*, $p < 0.05$, $N=6$ controls, $N=8$ alcoholics), and in amygdala (*, $p < 0.05$, $N=6$ controls, $N=8$ alcoholics).

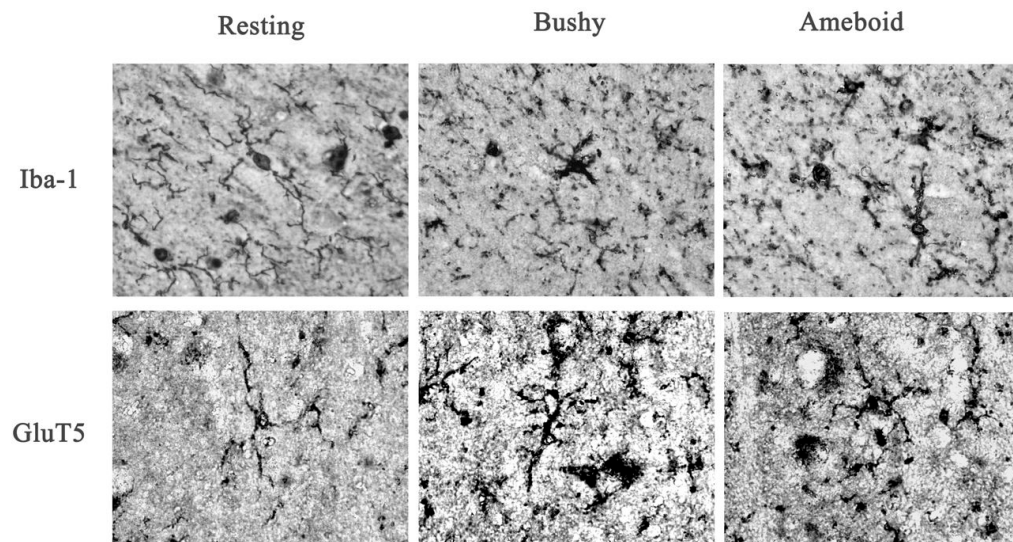


Fig.2. Representative pictures of different stages of microglia activation. Ramified microglial cells are believed to be in the resting stage; the bushy-looking microglia indicates early activation; ameboid microglia represent fully activated brain macrophages.

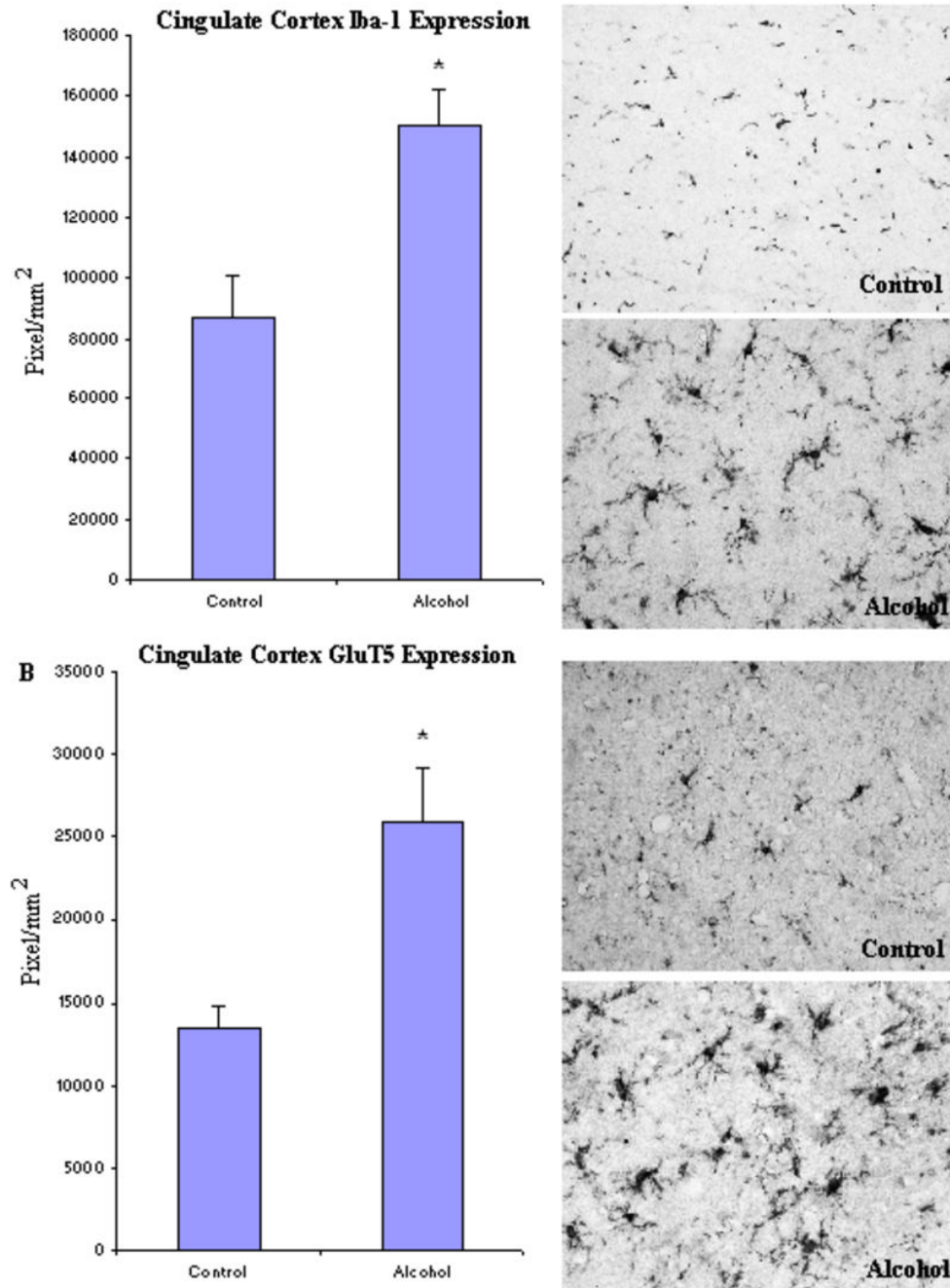


Fig. 3. In cingulate cortex, the expression of both Iba-1 and GluT₅ was significantly increased in the alcoholic brains (N=5) as compared to the controls (N=4). The level of immunoreactive density was quantified by BioQuant Nova analysis system as described in the Methods and presented as mean \pm SEM in pixel/mm². ANOVA indicated significant differences in immunoreactive density of both Iba-1 and GluT₅ between control and alcoholic groups (*, $p < 0.05$).

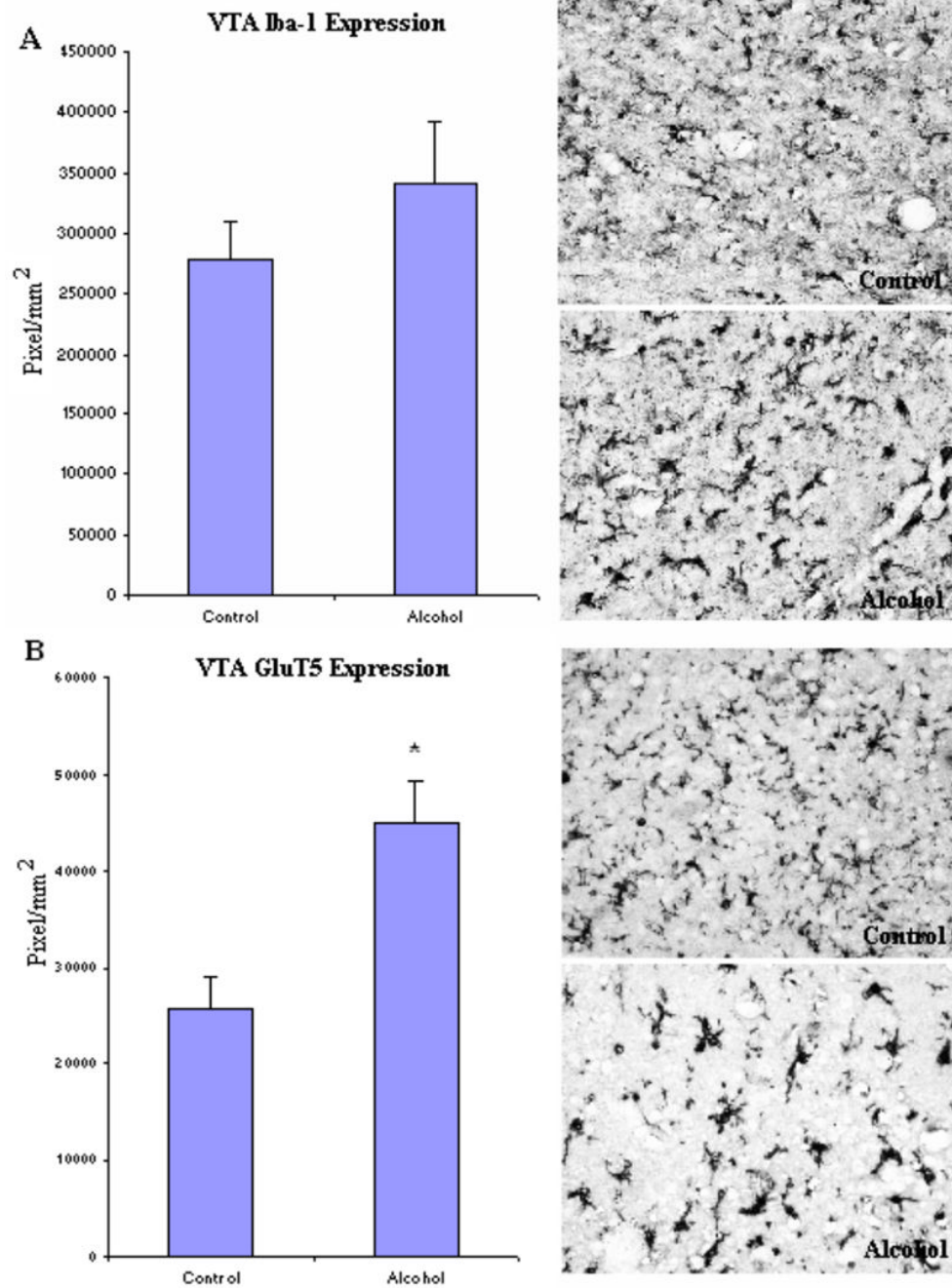


Fig. 4. In VTA, the expression of GluT₅, but not Iba-1 was significantly higher in the alcoholic brains (N=8) than controls (N=8). The immunoreactive density was measured by BioQuant Nova analysis system as described in the Methods and presented as mean \pm SEM in pixel/mm². ANOVA indicated a significant increase in GluT₅ immunoreactivity, but not Iba-1, in alcoholics as compared to controls (*, $p < 0.05$).

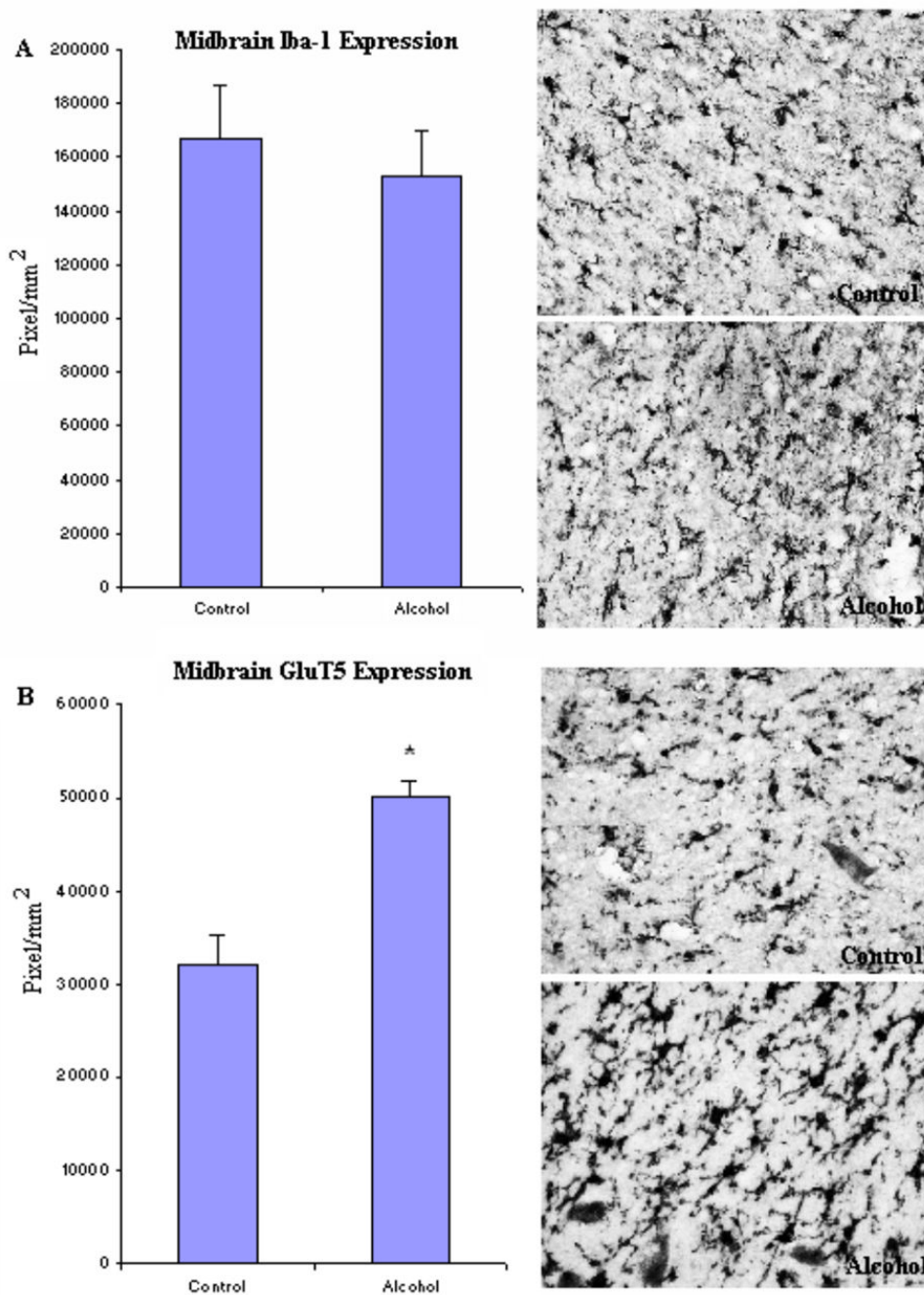


Fig. 5. In Midbrain, the expression of GluT₅, but not Iba-1 was significantly higher in the alcoholic brains (N=6) as compared to controls (N=5). The immunoreactive density was measured by BioQuant Nova analysis system as described in the Methods and presented as mean \pm SEM in pixel/mm². ANOVA indicated a significant increase in GluT₅ immunoreactivity, but not Iba-1, in alcoholics as compared to controls (*, $p < 0.05$).

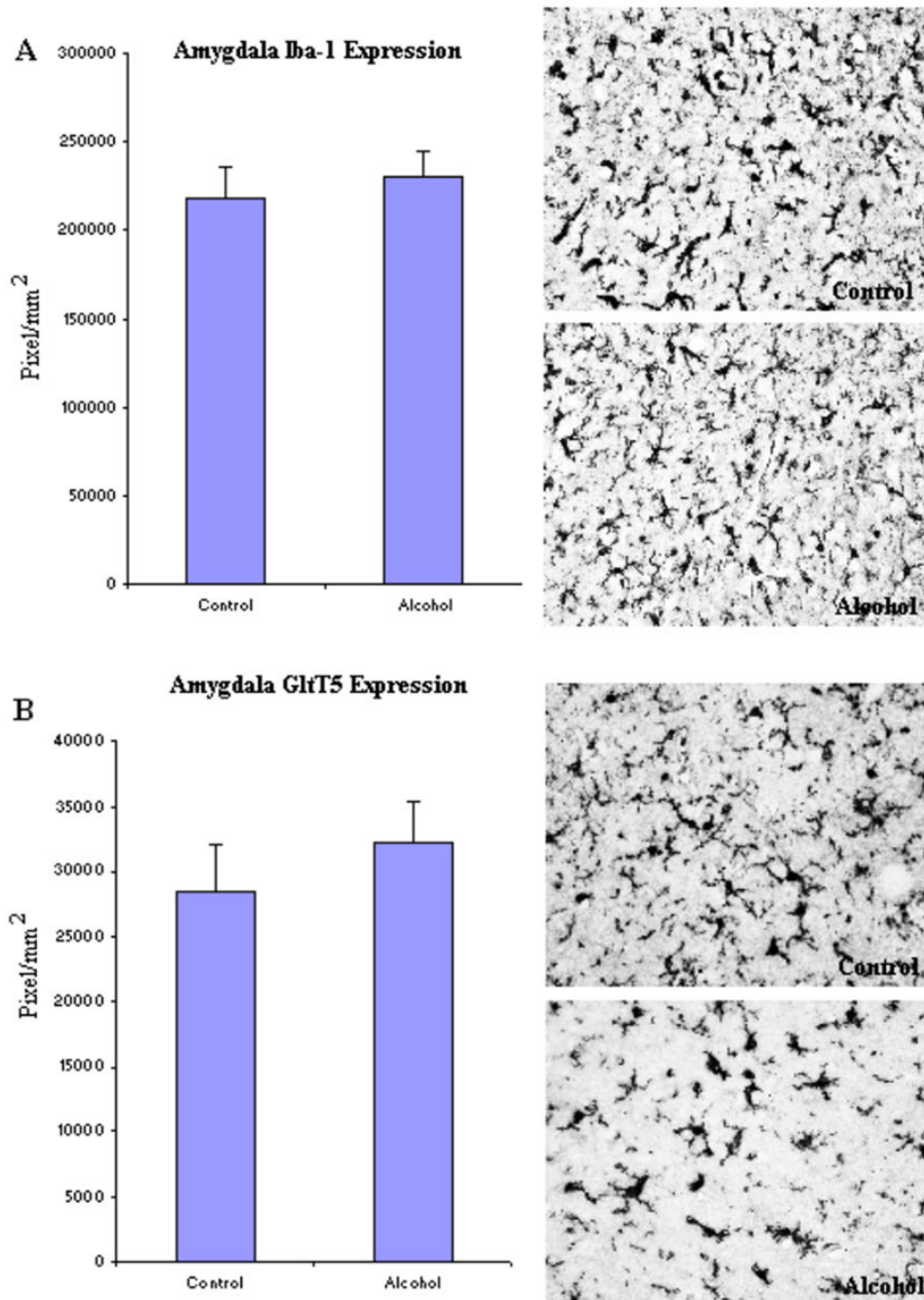


Fig. 6.

In Amygdala, no significant difference was detected in either Iba-1 or GluT₅ expression between alcoholic (N=8) and control brains (N=8). The immunoreactive density measured by BioQuant Nova systems as described in the Methods and presented as mean \pm SEM in pixel/mm². ANOVA indicated no significant differences in either marker between alcoholic and control groups.

Table 1

Case characteristics of subjects used for immunohistochemical analyses of the Cingulate Cortex.

Patient ID	Groups	Age	Sex	PMI	Clinical cause of death	Smoking	Daily intake/type
A094	Control	55	Male	20	Cardiac arrest	Ex smoker	Nil
A211	Control	50	Male	29	Ischaemic Heart Disease	Nil	Nil
A239	Control	37	Male	24	Electrocution	Unknown	Nil
A247	Control	46	Male	25	Mitral valve prolapse	Unknown	Nil
B180	Alcoholic	34	Male	8.5	Hanging	1 pkt/day	160 gms
B201	Alcoholic	46	Male	24	Alcohol toxicity	Unknown	150g beer
B210	Alcoholic	51	Male	27	Gastrintestinal Haemorrhage	1 pkt/day	150g beer/spirits
B221	Alcoholic	61	Male	28	Multiple Organ failure	Unknown	> 80g beer
B238	Alcoholic	52	Male	35	Alcohol toxicity	Nil	300g spirits

PMI: post-mortem interval

Case characteristics of subjects used for immunohistochemical analyses of Midbrain, VTA, Amygdala, and for ELISA analyses of VTA, SN, Amygdala and Hippocampus.

Table 2

Patient ID	Groups	Age	Sex	PMI	Clinical cause of death	Smoking	Daily intake/type
A312	Control	82	Male	23.5	Multiple organ failure	Unknown	<20g beer
A317	Control	78	Female	37	Acute myocardial infarction	Unknown	Nil
A339	Control	56	Male	37	Left ventricular scarring, hypertension, and cardiomegaly	15 cigs/day	10g
A357	Control	66	Female	6	Pneumonia and multiple CVA's	Nil	Nil
A367	Control	57	Male	18	Ischemic heart disease	Ex smoker	10g
A384	Control	60	Male	25	carcinomatosis, gastrointestinal stomach tumor	Nil	20g beer
A395	Control	60	Male	28	Ischemic heart disease	Nil	20g beer
A398	Control	54	Male	29	Coronary artery atheroma	Nil	<20g beer
B315	Alcoholic	52	Male	45.5	Lobar pneumonia and chronic Alcoholism	1 pkt/day	500g spirits
B221	Alcoholic	61	Male	28	Multiple Organ Failure	Unknown	>80g beer
B253	Alcoholic	56	Male	22	Coronary artery disease & upper gastro-intestinal hemorrhage	1 pkt/day	80g wine
B285	Alcoholic	67	Female	68	Hepatic & renal failure	Nil	80g beer
B298	Alcoholic	80	Male	28	Cardiopulmonary arrest	5 pkt/day	>80g beer
B322	Alcoholic	56	Male	29-31	Complications of liver disease	1 pkt/day	100 g beer
B401	Alcoholic	60	Male	51	Hepatic cirrhosis and its Consequences	Nil	>80g
B402	Alcoholic	75	Female	9	Hepatic encephalopathy and spontaneous bacterial peritonitis,	Nil	120 g beer

PMI: post-mortem interval; VTA: ventral tegmental area; SN: substantia nigra; ELISA: enzyme-linked immunosorbent assay.

Table 3Immunoreactivity of Iba-1 and GluT₅ in various brain regions

Regions	Control Group Mean \pm SEM; N	Alcoholic Group Mean \pm SEM; N	P value T Test
Iba-1 Immunoreactivity(pixel/mm² X10³)			
Cingulate	70.0 \pm 13.7 N = 4	150.1 \pm 12.3 N = 5	P = 0.01 *
VTA	277.9 \pm 31.4 N = 8	340.9 \pm 48.8 N = 8	P = 0.30
Midbrain	172.8 \pm 21.9 N = 6	152.7 \pm 17.4 N = 5	P = 0.50
Amygdala	217.7 \pm 18.5 N = 8	230.3 \pm 12.7 N = 8	P = 0.58
GluT₅ Immunoreactivity (pixel/mm² X10³)			
Cingulate	13.4 \pm 1.4 N = 4	26.0 \pm 3.3 N = 5	P = 0.02 *
VTA	25.7 \pm 3.3 N = 8	45.0 \pm 4.3 N = 8	P = 0.003 *
Midbrain	32.1 \pm 3.2 N = 6	50.1 \pm 1.6 N = 5	P = 0.01 *
Amygdala	28.4 \pm 3.6 N = 8	32.1 \pm 3.3 N = 8	P = 0.46

The immunoreactive density measured by BioQuant Nova systems as described in the Methods and presented as mean \pm SEM in pixel/mm². ANOVA was conducted in each region and p values are reported.

* : p<0.05 indicating statistical significance. Iba-1: ionized calcium binding adaptor protein-1; GluT₅: Glucose transporter-5; VTA: ventral tegmental area.