Properties of $\omega$ conotoxin MVIIC receptors associated with $\alpha$1A calcium channel subunits in rat brain

Nicole Martin-Moutot*a*, Christian Levequea, Kazuki Satob, Rika Katoa, Masami Takahashib, Michael Seagarb

*INSERM U374, Institut Jean Roche, Faculté de Médecine Secteur Nord, Boulevard Pierre Dramard, 13916 Marseille Cedex 20, France

bMitsubishi Kasei Institute of Life Science, Machida, Tokyo 194, Japan

Received 22 March 1995; revised version received 20 April 1995

Abstract Solubilized $^{125}$I-$\omega$ conotoxin MVIIC receptors from rat cerebellum were immunoprecipitated by antibodies directed against the calcium channel $\alpha$1A subunit. Anti-$\omega$A antibodies recognized a 240–220, 180 and 160 kDa proteins in immunoblots of cerebellar membranes. Disuccinimidyl suberate cross-linked $^{125}$I-$\omega$ conotoxin MVIIC to an $\alpha$2δ-like 200–180 kDa subunit, which migrated at 150–140 kDa after disulfide reduction. These observations are consistent with a heteromeric structure in which high affinity $\omega$ conotoxin MVIIC binding sites formed by $\alpha$1A subunits are located in close proximity to peripheral $\alpha$2 subunits.

Key words: Calcium channel; $\omega$ conotoxin MVIIC receptor; $\alpha$1A subunit; $\alpha$2 subunit

1. Introduction

Voltage-gated calcium channels in neurons have been classified as L, N, P, Q and T type, on the basis of their electrophysiological and pharmacological properties [1–3]. Calcium channels contain a pore-forming $\alpha$1 subunit that carries binding sites for channel blockers, and is generally associated with peripheral $\alpha$2δ and $\beta$ subunits [4–6]. A family of cDNAs encoding homologous $\alpha$1 (A, B, C, D, and E) subunits have been isolated from neuronal libraries [7,8]. Important aims at present are to establish a correlation between calcium channel antagonist binding sites expressed in the brain and a given class of $\alpha$1 subunit and to examine the polypeptide composition of the different calcium channel classes expressed in the brain.

Specific neurotoxins are powerful tools in this field. For example, irreversible block by $\omega$ conotoxin GVIA ($\omega$GVIA) from the marine mollusc Conus geographus [9], is a property unique to N-type channels and $\omega$GVIA receptor sites have been subsequently shown to be associated with the $\alpha$1B subunit in the brain [5,6,10] and in heterologous expression systems [11]. A structurally related peptide from Conus magus, $\omega$ conotoxin MVIIC ($\omega$MVIIC), blocks a population of $\omega$GVIA-insensitive calcium channels that are expressed in both the soma and synaptic terminals of central neurones [3,12,13]. Binding studies with rat brain membranes indicate that $\omega$MVIIC and $\omega$GVIA have distinct receptor sites [12,14]. Expression of the $\alpha$1A subunit in Xenopus oocytes supports calcium currents that display sensitivity to sub-micromolar concentrations of $\omega$MVIIC [15], however little is known about the polypeptide composition of the native calcium channels in the mammalian brain that constitute high affinity $\omega$MVIIC receptors. In the present study we have used affinity labeling methods and sequence directed anti-$\alpha$1 antibodies to address this question.

2. Experimental

2.1. Materials

Synthetic $\omega$MVIIC was obtained from Bachem (Bubendorf), $\omega$GVIA from the Peptide Institute (Osaka), and disuccinimidyl suberate (DSS) from Pierce (Interchim, France). Protein A Sepharose CL4B was from Sigma (St. Louis MO) and N-glycosidase F from Boehringer (Mannheim).

2.2. Membrane preparations

Cerebella or cerebra from one-month-old Wistar rats were dissected and homogenized in a glass/tetlon Thomas apparatus in a 0.32 M sucrose, 10 mM HEPES buffer adjusted to pH 7.4 with NaOH, in the presence of a mixture of protease inhibitors: 0.2 mM phenylmethylsulfonyl fluoride, 1 mM iodoacetamide, 2 mM pepstatin A, 1 mM EDTA, 2 µM leupeptin, 0.15 µM aprotinin, and 2 mM benzamidine. After a 10 min centrifugation at 800 x g, the supernatant was recovered and spun at 27,000 x g for 40 min. Protein was assayed by the BCA method (Pierce) using a BSA standard.

2.3. Conotoxin iodination and binding experiments

Preparation and binding of mono-$^{125}$I-$\omega$MVIIC were performed as previously described [16,17]. Mono-$^{125}$I-$\omega$MVIIC was prepared by the iodogen method using NaI-$^{125}$I (New England Nuclear, 2,000 Ci/mmol) and purified by HPLC on an analytical C18 column (Beckmann) [12]. The binding of $^{125}$I-$\omega$MVIIC to synaptosomal membranes was measured at 30°C in 100 µl Tris-buffered saline: 150 mM NaCl, 25 mM Tris base adjusted to pH 7.4 (TBS), containing 0.1% BSA. Binding was initiated by addition of 20 µg of membrane protein, stopped by rapid filtration on glass fiber filters (GFC-Whatman) pre-treated with 0.3% polyethyleneimine and washed three times with 2 ml of ice-cold TBS containing 0.1% BSA, 1.5 mM CaCl$_2$, 0.05% Tween-20.

2.4. Affinity labeling

Synaptosomes (300 µg) were incubated for 2 h at 4°C with 1 nM $^{125}$I-$\omega$MVIIC in the presence or absence of 1 µM unlabeled $\omega$MVIIC in 300 µl of 20 mM HEPES, 0.15 M NaCl, adjusted to pH 8.3 with NaOH. DSS solubilized in dimethylsulfoxide was added to a final concentration of 30 µM, and the mixture was incubated for 20 min at 20°C. The synaptosomes were sedimented, washed once with 20 mM HEPES, 0.15 M NaCl, 10 mM CaCl$_2$, adjusted to pH 8.3 with NaOH, and once with the same buffer from which CaCl$_2$ was omitted. Synaptosomes were again incubated with 1 nM $^{125}$I-$\omega$MVIIC and the cycle repeated as described above. The labelled membranes were processed for electrophoresis as described below.

2.5. Preparation of antibodies

Polyclonal antisera were produced in rabbits using synthetic peptides

Abbreviations: $\omega$GVIA, $\omega$ conotoxin GVIA; $\omega$MVIIC, $\omega$ conotoxin MVIIC; $\omega$AgalIVA, $\omega$ agatoxin IVA; DSS, disuccinimidyl suberate; BSA, bovine serum albumin; TBS, Tris-buffered saline; CHAPS, (3-[C(3-Cholamidopropyl)-(dimethylammonio)]-1-propane-sulfonate); DHP, 1,4-dihydropyridine.

*Corresponding author. Fax: (33) 91090506.
selected from calcium channel z1 sequences, coupled to keyhole limpet hemocyanin. RbA-1 antibodies were raised against the peptide (C)PSERAPGREGPYGRE, and rbA-2 antibodies against the peptide LRQTARPRESARDPDA corresponding to residues 865-881 and 845-860, respectively from the rat brain z1A sequence [18]. Antisera were raised against peptides from the two alternative C-terminal regions of the rabbit brain sequence BI [19]. BI-1 antibodies against the peptide (C)RDQRWSRSPSEGROHTTHHRQ and BI-2 antibodies against the peptide (C)GGRRWTAASAGKGGGPRA SA, corresponding to BI-1 residues 2,254-2,273 and BI-2 residues 2,402-2,421 respectively. IgG were purified by affinity chromatography on protein A-Sepharose CL4B.

2.6. Immunoprecipitation
Membranes were incubated with 0.5 nM $^{125}$I-oMVIIIC overnight at 4°C in binding buffer, pelleted at 10,000 × g for 10 min, and solubilized in 1% CHAPS in buffer A (0.32 M sucrose, 10 mM HEPES adjusted to pH 7.4). Protein A-Sepharose CL4B was swollen in TBS containing 0.5% BSA and incubated with antibodies for 1 h at 4°C. Following centrifugation, the pellet was washed with 0.4% CHAPS in buffer A and solubilized membrane proteins containing 3 fmol of oMVIIIC receptor were added in a final volume of 100 μl and incubated for 3 h at 4°C. The pellet was washed once with TBS, 0.5% BSA, and the immunoprecipitated radioactivity was counted.

2.7. Electrophoresis and Western blotting
Proteins were denatured at room temperature for 30 min in 3% SDS, 6% sucrose, in 70 mM Tris-HCl at pH 8.9, in the presence or absence of 25 mM diithiothreitol. Iodoacetamide was then added to a final concentration of 80 mM. SDS-PAGE was performed on 5-12% gradient gels. Proteins were transferred to nitrocellulose membranes (Hybond, Amersham) at 100 mA overnight. The membrane was blocked for 2 h at room temperature with 10% dried milk in TBS and then incubated with antibodies (50 μg/ml) overnight at 4°C. After extensive washing in 0.05% Tween-20 TBS, detection was performed with anti-IgG peroxidase and the ECL system (Amersham).

2.8. Deglycosylation
Affinity-labeled protein bands were electroeluted from SDS gels in 50 mM Tris, 0.1% SDS, 0.38 M glycine adjusted to pH 8.8 with HCl, diluted five-fold in 0.1 M NaPO$_4$, 0.1% SDS, 0.5% CHAPS, 10 mM EDTA, 1% β-mercaptoethanol, pH 8, and concentrated on Ultrafree TTK24 (Millipore) membranes. N-glycosidase F (2 units) was added and incubated for 24 h at 37°C. Proteins were denatured and separated by SDS PAGE as described.

3. Results and discussion

3.1. $^{125}$I-ω conotoxin MVIIIC binding to cerebellar membranes
Specific binding of 0.5 nM $^{125}$I-ωMVIIIC to cerebellar membranes reached equilibrium in about 30 min at 30°C and pH 7.4 (Fig. 1A). In these conditions non-specific binding determined in the presence of 0.1 μM unlabeled ωMVIIIC, accounted for approximately 50% of the total bound radioactivity. The addition at equilibrium of a large excess of unlabeled ωMVIIIC initiated dissociation with 50% release occurring at about 10 min (Fig. 1B). Semi-logarithmic plots of the association and dissociation kinetics (insets to Fig. 1A, B) allowed measurement of the apparent association rate constant ($k_{app}$) = 7.7 × 10$^{-4}$ s$^{-1}$ and the dissociation rate constant ($k_{diss}$) = 5.5 × 10$^{-4}$ s$^{-1}$, and subsequent calculation of the association rate constant $k_1 = 0.5 × 10^8$ s$^{-1}$ M$^{-1}$. The equilibrium dissociation constant was therefore ($K_D = k_{diss}/k_{app}$) = 1.8 nM, which was in good agreement with results obtained from displacement of $^{125}$I-ωMVIIIC by unlabeled ωMVIIIC ($K_D = 1$ nM) at equilibrium (not shown). Direct receptor saturation experiments were not performed. However from experiments carried out with a ligand concentration = $K_D$, a total receptor capacity of approximately 0.9 pmol/mg protein in cerebellar synaptosomes and 0.4 pmol/mg protein in cerebral synaptosomes was estimated. $^{125}$I-ωMVIIIC binding was not displaced by unlabeled ωGVIA, or ω agatoxin IVA (ωAgatIVA) but was inhibited by Ca$^{2+}$ with a $K_i < 1$ mM (Table 1).

3.2. Immunoprecipitation of ω conotoxin MVIIIC receptors
RbA-1 antibodies were raised against a peptide from the cytoplasmic loop between domains II and III that is specific to the rat brain z1A subunit sequence [18]. The ability of these antibodies to recognize the ωMVIIIC receptor was tested. The addition of protein A-Sepharose CL4B beads coated with increasing amounts of rBA-1 antibodies to CHAPS solubilized
125I-oMVIIC receptors resulted in concentration-dependent immunoprecipitation (Fig. 2A). After subtraction of background precipitation determined with non-immune rabbit IgG, approximately 50% of oMVIIC receptors were specifically recognized at the plateau level. Immunoprecipitation was shown to be specific by two other criteria: block of recognition by the synthetic peptide used to raise rbA-1 antibodies, and lack of immunoprecipitation of solubilized 125I-oGVIA receptors (Fig. 2B). These results indicate that high affinity receptors for 125I-oMVIIC are associated with the z1A subunit.

BI-1 and BI-2 antibodies were designed to distinguish between two z1A splice variants expressed in rabbit brain that contain alternative C-terminal segments [19]. The peptide used to raise BI-1 displayed 85% identity with the rat brain z1A C-terminus [18]. Specific immunoprecipitation of about 15% of 125I-oMVIIC receptors was achieved with BI-1 antibody, but no reaction was detected with the BI-2 antibody (Fig. 2B). Some of the z1A subunits that constitute the oMVIIC sensitive calcium channel in the cerebellum thus contain the variant BI-1 C-terminus.

3.3. Detection of z1A polypeptides by immunoblotting

In order to detect the z1A subunits expressed in the brain, immunoblots were probed with rbA-2 antibodies. RbA-2 antibodies were raised against a second peptide specific to the loop linking domains II and III of z1A. Although this antibody specifically immunoprecipitated a smaller fraction (approximately 35%) of oMVIIC sensitive calcium channels than rbA-1, it was found to be more sensitive for staining of immunoblots. RbA-2 reacted with several bands in cerebellar synaptosomes (Fig. 3, lane 1). Specificity of recognition was ascertained by parallel experiments in which the cognate peptide was used to block antibody binding (Fig. 3, lane 2). A 240-220 kDa doublet that sometimes appeared as a single band, and two polypeptides of 180 and 160 kDa were identified, with the 180 kDa band producing the most intense reaction. Similar experiments with cerebral synaptosomes (Fig. 3, lanes 3 and 4), produced much lower levels of immunostaining, and only the 180 kDa band was consistently detected.

These results suggest that the 240-220 kDa doublet represents z1A subunits whereas the 180 and 160 kDa size forms may result from post-translational cleavage of the larger polypeptides. The reduction of disulfide bridges did not produce any change in electrophoretic mobility of these proteins (not shown).

3.4. Affinity labeling of 125I-o conotoxin MVIIC receptors

A homobifunctional reagent DSS was used to crosslink 125I-oMVIIC to membrane polypeptides in proximity to the binding site (Fig. 4). When electrophoresis was performed in non-reducing conditions, labeled proteins of 200 and 180 kDa were detected by autoradiography (Fig. 4A, lane 1). However after disulfide reduction with dithiothreitol (Fig. 4A, lane 2), these proteins migrated as a poorly resolved 150-140 kDa doublet. Labeling was blocked by the presence of 0.1 nM 125I-oMVIIC (Fig. 4A lanes 3 and 4), but not by 0.1 nM oGVIA (not shown).

Treatment with dithiothreitol, which had no effect on the mobility of the immunostained z1A polypeptides, thus apparently dissociated an unlabeled disulfide-linked polypeptide of 40–50 kDa from both the 200 and the 180 kDa proteins. The change in electrophoretic mobility of these proteins upon reduction is similar to that of the z2 subunit of the skeletal muscle L type calcium channel, which is disulfide linked to a δ polypeptide, and displays extensive N-glycosylation [4].

**Table 1**

<table>
<thead>
<tr>
<th>Assay conditions</th>
<th>Binding (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>100</td>
</tr>
<tr>
<td>0.3 μM ω conotoxin GVIA</td>
<td>103</td>
</tr>
<tr>
<td>0.3 μM ω agatoxin IVA</td>
<td>93</td>
</tr>
<tr>
<td>0.3 mM CaCl₂</td>
<td>64</td>
</tr>
<tr>
<td>1 mM CaCl₂</td>
<td>32</td>
</tr>
</tbody>
</table>

Assays were performed with 0.5 nM 125I-oMVIIC in the presence and absence of 0.1 μM unlabeled oMVIIC. Specific binding was calculated by subtraction and represented as a % of the control. Results are the mean of duplicate determinations from two independent experiments.
and N-glycosidase F treatment of the proteins cross-linked to 125I-\omega MVIIC confirmed the presence of N-linked carbohydrate and indicated a core polypeptide mass of about 100 kDa (not shown), which is in good agreement with the electrophoretic mobility of deglycosylated α2 subunits from skeletal muscle calcium channels [4]. Control incubations in the absence of N-glycosidase F did not affect migration. Alpha 2 and δ polypeptides are encoded by a single gene [20], of which a single splice variant α2b is expressed in the nervous system [21].

Two neurotoxins, ωAgIVA and ωMVIIC, are currently used to dissect αGVIA-insensitive, DHP-insensitive calcium channel currents in central neurons and heterologous expression systems [22]. Low nanomolar concentrations of ωAgIVA [23], but significantly higher concentrations of ωMVIIC [12], block P type channels, which are predominant in cerebellar Purkinje neurons. Preferential expression of class A transcripts in the cerebellum [18,19] has thus suggested that P type channels contain α1A subunits. However expression of α1A, α2 and β subunits in Xenopus oocytes induces currents with reversed sensitivity that are strongly blocked by ωMVIIC but only partially inhibited at high concentrations of ωAgIVA [15]. Subsequently a native calcium current with similar properties has been identified in cerebellar granule cells and designated as the Q channel [3]. Evidence therefore suggests that α1A subunits support both P and Q type channels, although little is known about the heteromeric composition of native channels.

Our data suggests that high affinity ωMVIIC binding sites in the cerebellum, which may correspond to Q type calcium channels, are associated with α1A subunits. Although we have not directly addressed the interaction of β-subunits with ωMVIIC receptors, heterologous expression of α1A has indicated that the generation of robust currents requires co-expression of a β subunit [15,19]. Co-expression of a β subunit with the cardiac α1C subunit has been shown to increase the binding affinity for dihydropyridines [24]. By analogy it is possible that β subunits may play a role in defining the sensitivity of α1A subunits to ωMVIIC and ωAgIVA.

Chemical cross-linking suggests proximity of an α2-like subunit to receptor-bound ωMVIIC, although conotoxin binding sites are thought to be formed by α1 subunits. This is not without precedent as cross-linking of αGVIA to an α2-like subunit in chick but not rat brain membranes has been reported [25], although the α1B subunit is now known to constitute the binding site. Subtle differences in the spatial relationships of homologous conotoxin binding sites to the α2 protein may therefore exist between different species and channel classes. Negative charges at the surface of these highly glycosylated subunits may conceivably affect the access of different toxins to their pharmacological sites. Consequently we may speculate that heterogeneity in the glycosylation of α2 subunits may also be relevant to differences in the potency of ωMVIIC and ωAgIVA in blocking P and Q type currents supported by the α1A subunit.

Acknowledgements: We thank Cécile Raymond and Farida Lassoued for technical assistance, and François Couraud for comments on the manuscript. Financial support was provided by the Association Française contre les Myopathies.
References