https://doi.org/10.1369/0022155420955246 DOI: 10.1369/0022155420955246 Journal of Histochemistry & Cytochemistry 2020, Vol. 68(10) 679–690 © The Author(s) 2020 Article reuse guidelines: [sagepub.com/journals-permissions](https://us.sagepub.com/en-us/journals-permissions)

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Expression of p11 and Heteromeric TASK Channels in Rat Carotid Body Glomus Cells and Nerve Growth Factor–differentiated PC12 Cells

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Summary

TWIK-related acid-sensitive K⁺ (TASK) homomeric channels, TASK1 and TASK3, are present in PC12 cells. The channels do not heteromerize due plausibly to a lack of p11 protein. Single-channel recording reveals that most of the rat carotid body (CB) glomus cells express heteromeric TASK1-TASK3 channels, but the presence of p11 in glomus cells has not yet been verified. TASK1, but not TASK3, binds to p11, which has a retention signal for the endoplasmic reticulum. We hypothesized that p11 could facilitate heteromeric TASK1-TASK3 formation in glomus cells. We investigated this hypothesis in isolated immunocytochemically identified rat CB glomus cells. The findings were that glomus cells expressed p11-like immunoreactive (IR) material, and TASK1- and TASK3-like IR material mainly coincided in the cytoplasm. The proximity ligation assay showed that TASK1 and TASK3 heteromerized. In separate experiments, supporting evidence for the major role of p11 for channel heteromerization was provided in PC12 cells stimulated by nerve growth factor. p11 production took place there via multiple signaling pathways comprising mitogen-activated protein kinase and phospholipase C, and heteromeric TASK1-TASK3 channels were formed. We conclude that p11 is expressed and TASK1 and TASK3 heteromerize in rat CB glomus cells. (J Histochem Cytochem 68: 679–690, 2020)

Keywords

carotid body glomus cell, PC12 cell, p11, TASK1, TASK3

Introduction

TWIK-related acid-sensitive K⁺ (TASK) channels contribute to the resting membrane potential in a variety of cells, such as carotid body (CB) glomus cells,¹ brain neurons, $2,3$ and endocrine cells. $4-8$ They belong to the twopore domain K $^+(K_{2P})$ channel family, 9,10 which comprises 15 subunits in humans. Having two pore domains in each subunit, K_{2P} channels exist as a homodimer or heterodimer of subunits. The TASK subfamily consists of TASK1, TASK3, and TASK5: TASK1 and TASK3 function as homomeric or heteromeric channels, ^{11,12} whereas TASK5 is devoid of channel activity in exogenous expression systems.¹³ The molecular structure of TASK channels

constitutively expressed in some cells^{7,12,14,15} or exogenously expressed in oocytes¹⁶ or cell lines^{3,11} has been investigated with biochemical or electrophysiological approaches. Single-channel analysis in mouse¹² and rat CB glomus cells¹⁴ revealed that most of the TASK channels are heteromeric TASK1-TASK3 structures. As

Received for publication May 2, 2020; accepted August 14, 2020.

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homomeric TASK1 or TASK3 channel activity has been recorded in glomus cells of TASK3 or TASK1 knockout mice,¹² respectively, it has been surmised that heteromeric TASK1-TASK3 channels are more readily assembled and/or trafficked to the plasma membrane than either homodimer. In contradistinction, endogenous homomeric TASK1 and TASK3 channels are found in the plasma membrane of PC12 cells¹⁷ which originate from rat adrenal medullary chromaffin cells.¹⁸ Interestingly, when green fluorescent protein (GFP)-tagged TASK1 or TASK3 proteins are exogenously expressed in PC12 cells,17 they do not heteromerize with endogenous TASK subunits in the plasma membrane. We have recently proposed that a lack of heteromeric TASK1-TASK3 channels in PC12 cells is due to a lack of $p11$, 7 a cytosolic protein which contains a retention signal in the C-terminus.^{19,20} As p11 is not expressed in PC12 cells,²¹ TASK1 protein translated at the rough endoplasmic reticulum (ER) may be promptly folded in a mature form without encountering TASK3 and trafficked to the plasma membrane through the Golgi complex. If this notion is true, the cells where heteromeric TASK1-TASK3 channels are formed will be expected to produce p11. The present immunocytochemical study addresses this issue in rat CB glomus cells where heteromeric TASK1-TASK3 channels have been functionally identified in the plasma membrane,¹⁴ but the presence of p11 has not yet been substantiated. In addition, to enhance the understanding of how p11 expression is regulated, we explored the signal transduction pathways for the nerve growth factor (NGF)-induced p11 production in PC12 cells,⁷²¹ which have the same catecholamine synthesizing potency¹⁸ as the glomus cells do.22

Materials and Methods

Animals

Eleven male Wistar rats weighing 150 to 300 g were used. Rats were housed in standard cages with free access to food and water. Animals were kept under a light/dark cycle of 12/12 hr under the normoxic condition. All procedures for the care and treatment of animals were carried out according to the Japanese Act on the Welfare and Management of Animals and the Guidelines for the Proper Conduct of Animal Experiments issued by the Science Council of Japan. The experiments were approved by the Institutional Animal Care and Use Committee of the University of Occupational and Environmental Health in Kitakyushu (permit AE19-014). The procedures complied with the Animal Research: Reporting of In Vivo Experiments (ARRIVE) guidelines. All efforts were made to minimize potential animal distress.

Immunocytochemistry

The rats were anesthetized with a mixture of medetomidine, midazolam, and butorphanol, i.p., according to a published protocol.²³ After tracheostomy, the trachea-esophagus bundle was cut through and retracted rostrally to get bilateral access to the carotid artery bifurcation area. The superior sympathetic cervical ganglions covering the CBs were removed. The CB was carefully sliced off from the artery wall on either side under an operating microscope and immediately immersed in ice-cold Ca^{2+} -deficient saline solution, where 1.8 mM CaCl₂ was omitted. The standard saline contained 137 mM NaCl, 5.4 mM KCl, 1.8 mM CaCl₂, 0.5 mM MgCl₂, 0.53 mM NaH₂PO₄, 5 mM p-glucose, and 5 mM Hepes with pH adjusted to 7.4 with 4 mM NaOH. The preparations were incubated in a 0.5% collagenase-containing Ca^{2+} -free solution at 36C for 30 min. Then, one CB was placed in a 35-mm glassbottom dish (P35GCOL-1.0-14-C; MatTek, Ashland, MA), and cells were dissociated using fine needles under a microscope. Dissociated CB cells, allowed to attach to the glass for 30 min, were fixed in 4% paraformaldehyde in phosphate-buffered saline (PBS; pH 7.2) for 1 hr and preincubated in PBS with 5% fetal bovine serum (FBS; cat. no. 172012; Sigma-Aldrich, Tokyo, Japan) and 0.3% Triton-X for 30 min. For indirect immunofluorescence studies, cells were treated overnight with a combination of mouse anti–tyrosine hydroxylase (TH) (cat. no. MAB318; Millipore, Temecula, CA) (RRID: AB_2201528) and goat anti-p11 antibodies (cat. no. AF2377; R&D Systems, Minneapolis, MN) (RRID: AB_2183469), rabbit anti-TASK1 (cat. no. sc-28635; Santa Cruz Biotechnology, Dallas, TX) (RRID: AB_661017) and goat anti-TASK3 antibodies (cat. no. sc-11317; Santa Cruz Biotechnology) (RRID: AB_2131233), or anti-TASK1 and anti-p11 antibodies. The specific immunoreactivities of anti-p11, anti-TASK1, and anti-TASK3 antibodies used have been immunocytochemically confirmed in our previous experiments where the proteins tagged with GFP or myc were exogenously expressed in PC12 cells, $6,17$ and the Kcnk3 or Kcnk9 gene was deleted in the mouse germ line.⁷ The selectivity of anti-TH antibody has been well documented.²⁴ After incubation, cells were washed $3\times$ with PBS and treated with a respective secondary antibody conjugated with Alexa 488, 546, or 633 (Molecular Probes; Eugene, OR). Fluorescence was observed under a laser confocal microscope (LSM5 Pascal; Carl Zeiss, Tokyo, Japan). The objective lens was an oil immersion with a magnification of $63\times$ and a numerical aperture of 1.4. For Alexa 488, a 488-nm laser was used and 510 to 560 nm emission was observed (FITC-like fluorescence); for Alexa 546, a 543-nm laser was used and emission above 560 nm was observed (rhodamine-like fluorescence). For GFP, a 514-nm laser was used and 530 to 600 nm emission was observed. The cell periphery was defined as an area of 1 µm width at the cell boundary, which was determined in a differential interference contrast (DIC) image. The immunofluorescence of the whole cell and at the cell periphery was measured with ImageJ software (NIH; Bethesda, MD). The coincidence of immunoreactivities to two different antibodies was assessed with MetaMorph software (Universal Imaging; Downingtown, PA). The fluorescence intensity in the nucleus was set as a threshold for a specific immunoreaction.

Cell Culture and Transfection

PC12 cells (RRID: CVCL_0481) originating from male rat adrenal medullary cells¹⁸ and PC12nnr5 cells lacking TrkA (RRID: CVCL_128)²⁵ were cultured in Dulbecco's modified Eagle medium (DMEM; Gibco/ Life Technologies, Tokyo, Japan) supplemented with 10% FBS, as previously decribed.17 The Lipofectamine 2000 reagent (Invitrogen/Life Technologies) was used to transfect the cells with expression vectors, according to the manufacturer's instructions. Briefly, plasmid (1.6 µg) and lipofectamine 2000 (4.0 µl) were added to an individual Ependorf tube that was filled with 50 µl of DMEM, and then left for 5 min at a room temperature before mixing. After 20 min, the mixture was added to 2 ml of DMEM in a 35-mm glass-bottom dish where PC12 cells had been cultured. The transfected cells were further cultured for 24 hr. The cells were fixed with 4% paraformaldehyde in PBS for 30 min at room temperature. After washing 3× with PBS, the cells were incubated in PBS with 0.1% Triton X-100 for 30 min and then with PBS containing 1% FBS for 1 hr at room temperature. The cells were treated with primary and secondary antibodies. Coverslips were mounted in 50% glycerol containing 1 mg/ml 4-diaminobenzene. To differentiate PC12 cells into neuron-like cells, the cells were treated with NGF (50 ng/ml) in the absence of FBS.

Plasmids

The following mammalian expression vectors were used: GFP-tagged TASK1 (GFP-TASK1) and GFPtagged TASK3 (GFP-TASK3)³ where rat TASK1 and TASK3 cDNAs were each subcloned into pEGFP with the consequent ligation of GFP to the N-terminus of TASK; myc-DDK-tagged p11 (myc-p11) where mouse p11 was into pCMV6-Entry with the consequent ligation of myc-DDK to the C-terminus of p11; hemagglutinin

(HA)-tagged TrkA26 where rat TrkA cDNA was into pCMX with the insertion of HA tag at the N-terminus of TrkA; HA-tagged TrkA Y794F and HA-tagged TrkA 499F26,27 where Y at amino acid position 794 or 499 was replaced with F using PCR mutagenesis; and Flagtagged TrkA Y760F²⁸ where Y at 760 in rat TrkA subcloned into pcDNA3 was replaced with F using PCR mutagenesis.

Proximal Ligation Assay (PLA)

To detect heteromeric TASK1-TASK3 channel formation in CB cells and PC12 cells, an in situ PLA,²⁹ which visualizes subcellular localization and protein–protein interaction, was performed using the Duolink in situ PLA kit (Olink Bioscience, Uppsala, Sweden), according to the manufacturer's instructions. The fixed cells were treated overnight with a combination of rabbit anti-TASK1 and goat anti-TASK3 antibodies or mouse anti-GFP (cat. no. sc-9996; Santa Cruz Biotechnology) (RRID: AB_627695) and rabbit anti-myc antibodies (cat. no. sc-789: Santa Cruz Biotechnology) (RRID: AB_631274). After treatment with primary antibodies, cells were incubated with secondary antibodies conjugated with oligonucleotides (PLA probe MINUS and PLA probe PLUS). The oligonucleotides of two PLA probes were hybridized by ligase if they were in close proximity (<40 nm). A rolling-circle amplification was carried out with polymerase and fluorescently labeled oligonucleotides, resulting in amplification of the fluorescent signal. The fluorescence emanating from PLA products was observed with the LSM 5 Pascal Microscope System.

Sources of Reagents

The plasmid for myc-DDK-tagged p11 (cat. no. MR226807) was purchased from OriGene Technologies (Rockville, MD), NGF (cat. no. N6009) was from Sigma-Aldrich, mouse anti-HA antibody (cat. no. sc-7392) (RRID: AB_1537399) was from Santa Cruz Biotechnology, and mouse anti-Flag antibody (cat. no. 200-301-383) (RRID: AB_627809) was from Rockland Immunochemicals (Limerick, PA).

Statistics

Data are expressed as mean \pm SEM. The significance of differences between experimental and control results was assessed with unpaired Student's *t*-test for data with normal distribution (Shapiro–Wilk test). Otherwise, Mann–Whitney rank-sum test or Kruskal– Wallis analysis of variance on ranks was used. A value of *p*<0.05 defined a statistically significant difference.

Statistical analysis was performed with Sigma Plot v13.0 software (Systat Software; San Jose, CA).

Results

Expression of p11 in Tyrosine Hydroxylase– positive Cells

Carotid bodies are composed of excitable glomus or type I cells that detect hypoxia, and glia-like sustentacular or type II cells that surround clusters of type I cells. Type II cells are non-excitable and do not contain voltage-dependent channels.³⁰ TH is a rate-limiting enzyme for catecholamine biosynthesis in both CB glomus and PC12 cells.

First, immunoreactivities to anti-HT and anti-p11 antibodies were examined in NGF-stimulated and nonstimulated PC12 cells. As is evident in Fig. 1A to C, TH-like IR material was detected in both stimulated and non-stimulated PC12 cells, whereas p11-like IR material predominantly occurred in the stimulated cells, consistent with references.^{7,21} These results warranted double immunostaining with these antibodies.

TH expression was immunocytochemically examined in isolated CB glomus cells to verify their identity.³¹ Mechanical dispersion of collagenase-treated CBs (Fig. 1D) yielded glomus cell clusters (Fig. 1E) and single cells (*n*=10; Fig. 1F). All of them showed both TH- and p11-like IR material (Fig. 1B and C). Thus, clustered cells comprised only glomus cells with no contribution by type II cells.

Expression of TASK1 and TASK3

Because p11-immunopositive cells turned out to be glomus cells, isolated cells were subjected to double immunostaining for p11 and TASK1. All the seven p11-immunopositive cells showed TASK1-like IR material as exemplified in Fig. 2A. The expression of TASK1 and TASK3 was simultaneously explored with rabbit anti-TASK1 and goat anti-TASK3 antibodies. Most of TASK1- and TASK3-like IR material was present in the cytoplasm, and both immunoreactivities were mainly colocalized (Fig. 2B). The fraction of TASK1-like IR material colocalized with TASK3-like IR material was 60.4 \pm 5.8%, whereas that of TASK3-like IR material with TASK1 was $58.1 \pm 5.2\%$ (Fig. 2C).

Heteromeric Channel Formation

The coincidence of TASK1- with TASK3-like IR material strongly suggests that both form a heteromeric channel. This notion was further explored using PLA with rabbit anti-TASK1 and goat anti-TASK3 antibodies. The PLA products between TASK1 and TASK3 developed only in p11-expressing PC12 cells and were mainly present in the cytoplasm (Fig. 3A), although all the cells endogenously exhibited TASK1- and TASK3-like IR material (Fig. 4E), which is consistent with a previous report.7 Similarly, five CB cells exhibited the PLA reaction in the cytoplasm (Fig. 3B and C), indicating that TASK1 and TASK3 heteromerized there.

p11 Binding to Heteromeric TASK1-TASK3 Channels

The results outlined above indicate that the expression of p11 results in the formation of heteromeric TASK1- TASK3 channels. Because the binding site for p11, called i20, is in the C-terminus of TASK1, but not TASK3,²⁰ the heteromeric TASK1-TASK3 channels are expected to complex with p11. This possibility was examined with PLA in PC12 cells where myc-p11 and either GFP-TASK1 or GFP-TASK3 were exogenously expressed. In this constellation, GFP-TASK and PLA products between GFP and myc were colocalized in the cytoplasm and at the cell periphery (Fig. 4A), irrespective of whether the TASK subtype was TASK1 or TASK3. This result suggests that GFP-TASK3 forms a heteromeric channel with endogenous TASK1 and p11 suppresses its ER exit by binding to the TASK1 subunit of the heteromeric channel.

Detailed inspection of the PLA reaction reveals that the amount of PLA products between GFP-TASK1 and myc-p11 was significantly more than that of PLA products between GFP-TASK3 and myc-p11 (Fig. 4B). This difference might be ascribed to a difference in the number of TASK dimer channels capable of binding to p11. Exogenous GFP-TASK1 will make a heteromeric channel with endogenous TASK3 and a homomeric channel with endogenous TASK1 or exogenous GFP-TASK1. On the contrary, exogenous GFP-TASK3 will make a heteromeric channel with endogenous TASK1 and a homomeric channel with endogenous TASK3 or exogenous GFP-TASK3 that is incapable of binding to p11. If endogenous TASK1 and TASK3 and exogenous GFP-TASK1 and GFP-TASK3 are expressed to a similar level, PLA products between GFP and myc in GFP-TASK1-expressing cells reflect the number of both heteromeric and homomeric channels, whereas those in GFP-TASK3-expressing cells reflect the number of heteromeric channels alone. Thus, it would be a reasonable assumption that the amount of PLA products in GFP-TASK1-expressing cells was larger than that in GFP-TASK3-expressing cells.

To verify the validity of our reasoning, the amounts of TASK1 and TASK3 endogenously expressed in PC12

Figure 1. Immunocytochemistry for TH and p11 in rat CB glomus cells and PC12 cells. (A) Confocal images of TH- and p11-like immunofluorescence in PC12 cells stimulated and non-stimulated (−) with NGF for 3 days (3 d). First and second columns represent THand p11-like immunofluorescence images, respectively; third, DIC images; fourth, merge of second and third images. Unless otherwise noted, green and red represent FITC- and rhodamine-like fluorescence (this and following figures), respectively. (B and C) Summaries of TH- and p11-like fluorescence present in stimulated (PC NGF) and non-stimulated (PC) PC12 cells and CB glomus cells. The amounts

Figure 1. *(continued)*

of fluorescence are expressed in arbitrary unit (au). Data are mean ± SEM (PC: *n*=20 from two culture dishes; PC NGF: *n*=20 from two culture dishes; CB: *n*=10 from three CBs); Student's *t*-test. (D) Transmitted image of a CB treated with collagenase. The CB was obtained from a rat weighing 150 g. (E) Confocal images of TH- and p11-like immunofluorescence and DIC image in clustered CB cells. (F) Confocal images of TH- and p11-like immunofluorescence in a CB cell. Bars in A, E and F, D indicate 5 µm and 0.5 mm, respectively. Abbreviations: CB, carotid body; DIC, differential interference contrast; NGF, nerve growth factor; PC, pheochromocytoma; TH, tyrosine hydroxylase.

cells were immunocytochemically measured. To this end, the immunoactivity of anti-TASK1 antibody was compared with that of anti-TASK3 antibody in PC12 cells exogenously expressing GFP-TASK1 or GFP-TASK3 (Fig. 4C). The amount of TASK1- or TASK3-like immunofluorescence was plotted against that of GFP fluorescence. As shown in Fig. 4D, the slope for TASK1 like immunofluorescence was akin to that of TASK3. This indicates that both antibodies had similar immunoreactivity. When endogenous TASK1 and TASK3 were

dishes). PC12 cells were stimulated with NGF for 1 day (p11-expressing PC12 cells). Bars are 5 µm. Abbreviations: CB, carotid body; DIC, differential interference contrast; PC, pheochromocytoma; PLA, proximity ligation assay; NGF, nerve growth factor; TASK, TWIKrelated acid-sensitive K+.

stained with these antibodies (Fig. 4E), the amount of TASK1-like IR material was smaller than that of TASK3 (Fig. 4F), indicating that a larger amount of PLA products in GFP-TASK1-producing cells cannot be explained by a larger amount of endogenous TASK1.

Thus, the difference in the PLA reaction between GFP-TASK1- and GFP-TASK3-producing cells may be accounted for by the notion that PLA products represent both heteromeric TASK1-TASK3 channels and homomeric TASK1 channels in the former.

Figure 4. PLA for myc-p11 binding to GFP-TASK channels. (A) Confocal images of GFP-TASK1 or GFP-TASK3 and PLA reaction between GFP and myc in PC12 cells exogenously expressing myc-p11 and GFP-TASK1 (T1G) or GFP-TASK3 (T3G). (B) Summaries of PLA products in the total area and at the cell periphery of PC12 cells. Data are mean ± SEM; Mann–Whitney rank-sum test (T1G: *n*=10 from three culture dishes; T3G: *n*=10 from two culture dishes). Bars are 5 µm. (C) Exogenous expression of GFP-TASK1 or GFP-TASK3 in PC12 cells. First and second columns represent confocal images of GFP and TASK1- or TASK3-like IR material, respectively; third represents merge of first and second images. GFP-TASK and TASK-like IR material were visible as GFP and rhodamine-like fluorescence, respectively. (D) TASK1- and TASK3-like IR material are plotted against those of GFP fluorescence (TASK1: *n*=20 from five culture dishes; TASK3: *n*=19 from four culture dishes). Linear regression lines are *y* = −7911 + 2.811*x* (correlation coefficient *r* = 0.996) and *y* = −13330 + 2.662*x* (*r* = 0.994) for TASK1 and TASK3, respectively. (E) Immunocytochemical staining for TASK1 and TASK3 endogenously expressed in PC12 cells. First and second columns represent confocal images and merge of fluorescence and DIC images. (F) TASK1 and TASK3-like IR material in PC12 cells. Data are mean ± SEM (*n*=20 from two culture dishes for each of TASK1 and TASK3). Bars are 5 µm. Abbreviations: DIC, differential interference contrast; GFP, green fluorescent protein; IR, immunoreactive; PC, pheochromocytoma; PLA, proximity ligation assay; TASK, TWIK-related acid-sensitive K+.

Figure 5. Immunocytochemical analysis of signaling pathways involved in NGF-induced p11 production. (A) Confocal images of p11 and HA- or TrkA-like immunofluorescence in PC12nnr5 cells expressing wild-type (WT), Y499F-TrkA, Y760F-TrkA, or Y794F-TrkA. First and second columns represent p11- and HA- or Flag-like immunofluorescence images, respectively; third, merge of DIC and fluorescence images. HA-like fluorescence was observed in cells expressing WT, Y760F-TrkA, and Y794F-TrkA, whereas TrkA-like fluorescence was observed in cells expressing Y760F-TrkA. (B) p11-like fluorescence is plotted against those of HA- (WT, Y499F, and Y794F) or TrkA-like (Y760F) fluorescence observed in the same cells. Linear regression lines are as follows—WT: *y* = −0.634 + 0.810*x*, correlation coefficient *r* = 0.823; Y499F: *y* = −5.595 + 0.621*x, r* = 0.924; Y760F: *y* = −5.594 + 0.901*x, r* = 0.929; Y794F: *y* = −3.623 + 0.541*x, r* = 0.888. (C) Summary of p11-like IR material in PC12nnr5 cells expressing WT, Y499F-TrkA, Y760F-TrkA, or Y794F-TrkA. P11-like IR material was measured over the whole cell area including protrusions or neurites. Data are mean \pm SEM; Kruskal–Wallis one-way analysis of variance on ranks (WT: *n*=38 from three culture dishes; Y499F: *n*=33 from two; Y760F: *n*=38 from three; Y794F: *n*=38 from 3). Bars are 10 µm. Abbreviations: DIC, differential interference contrast; HA, hemagglutinin; IR, immunoreactive; NGF, nerve growth factor; PC, pheochromocytoma.

Signal Transduction for p11 Production

PC12nnr5 cells, where p75NTR, but not TrkA, was expressed,²⁵ were used to explore the signaling mechanism for NGF-induced p11 production. The treatment with NGF of PC12nnr5 cells did not result in p11 production (not shown), indicating that TrkA is essential for p11 production. There are mainly three signaling pathways downstream of TrkA: mitogen-activated protein (MAP) kinase, phosphoinositide 3 (PI3)-kinase, and phospholipase C (PLC). 32 Thus, TrkA mutants were used to elucidate which signaling pathways were involved in p11 production. In PC12nnr5 cells expressing either wild-type or mutant TrkA (Fig. 5A), the amount of p11-like IR fluorescence depended on the

extent of TrkA expression. Thus, the amount of p11-like IR fluorescence was plotted against that of HA or Flaglike IR. The extent of p11 expression increased with that of exogenous TrkA, irrespective of its type (Fig. 5B). However, detailed inspection of the plots reveals that p11 expression differed among the TrkA mutants, which were expressed to similar extents. In TrkA Y499F-expressing cells where the MAP kinase pathway is mainly disrupted,²⁷ NGF stimulation induced short growth-cone-like protrusions, but not long neurites, and the amount of p11 elicited by NGF was significantly less than that in wild-type TrkA-expressing cells (Fig. 5C). A similar decrease in p11 production occurred in PC12nnr5 cells expressing Y794F-TrkA lacking in PLC signaling^{28,33} (Fig. 5C). In this TrkA mutant–expressing cells, however, a few long neurites were extended in response to NGF (Fig. 5A). On the contrary, cells expressing Y760F-TrkA, where the PI3 kinase pathway is mainly disrupted, $28,34$ produced p11 to the same extent as the control cells, but did not extend neurites in response to NGF (Fig. 5A and C). Thus, this mutation in TrkA impairs neurite extension and has no effect on p11 production in response to NGF.

Discussion

Heteromeric TASK Channels in CB Glomus and PC12 Cells

The main findings of this study were that (1) p11, TASK1, and TASK3 were expressed in rat CB glomus cells; (2) TASK1- and TASK3-like IR material mainly coincided in the cytoplasm; and (3) PLA products between TASK1 and TASK3 were predominantly located in the cytoplasm. These results suggest that TASK1 and TASK3 are present as a heteromeric channel mainly in the cytoplasm of glomus cells.

We have previously proposed that the cytoplasmic localization of heteromeric channels is due to retention of channels in the ER through binding to $p11^7$ which has a lysine-based retention/retrieval motif in the C-terminus.20 This notion is supported by the following evidence: (1) both TASK1 and TASK3, endogenously or exogenously expressed, are in the plasma membrane and do not heteromerize in PC12 cells in which p11 is not endogenously expressed^{7,21}; (2) exogenous expression of p11 in PC12 cells results in the formation of heteromeric TASK1-TASK3 channels, which are mainly located in the cytoplasm (present results)⁷; and (3) in CB glomus cells endogenously expressing p11, TASK1 and TASK3 proteins are mainly present in the cytoplasm (present results). The presence of TASK3 in the cytoplasm may be explained by the notion that endogenously expressed p11 suppresses the ER exit of TASK1 through binding to its terminus and subsequently facilitates the heteromeric channel formation of TASK1 and TASK3. As a result, heteromeric TASK channels are retained in the cytoplasm. Consistent with this notion, in PC12 cells expressing GFP-TASK3 and myc-p11, GFP-TASK3 proteins are predominantly located in the cytoplasm and PLA reaction develops between GFP-TASK3 and myc-p11 probably through endogenous TASK1.

Signal Transduction for p11 Production

The content of brain-derived neurotrophic factor (BDNF), which is known to produce p11 via TrkB, 35 decreases in humans with depression, and treatment

with antidepressants results in BDNF restoration in the cerebrospinal fluid.36 In in vitro experiments, p11 enhances plasma membrane expression of the serotonin receptors $5HT_{1B}R^{37}$ and $5HT_{4}R^{38}$ The biological plausibility that p11 plays an important role in the pathophysiology of depression arises. Thus, elucidation of signaling mechanisms for p11 production could enlighten the pathophysiology of depression and the development of new remedies.

Aside from brain neurons, 29 p11 is expressed in various kinds of cells, and its production is facilitated by growth factors and biologically active substances, such as NGF 7,21 and NO.³⁹ p11, which functions as a homodimer or heteromer with annexin II,⁴⁰ is involved in trafficking of membrane proteins,²⁰ such as receptors 37 and ion channels.⁴¹ Therefore, a change in p11 expression is assumed to induce a change in plasma membrane expression of receptors and ion channels, shaping cellular functions. Brain-derived neurotrophic factor is involved in neurite extension, synapse formation, plasticity in synaptic transmission, and cellular excitability.32 Several of these BDNF actions are mediated by p11.40 A pharmacological study shows that BDNF induces p11 formation in cultured neurons via the MAP kinase pathway.³⁵ This study also shows that Y499A mutation in TrkA, which mainly disrupts MAP kinase signaling,²⁷ resulted in a significant decrease in p11 expression in response to NGF. However, a similar extent of p11 diminution occurred in cells expressing the Y794A mutant lacking the PLC signaling.42,43 The result with the Y794F mutant might be due to a deficit in MAP kinase signaling. Protein kinase C has been suggested to activate MAP kinase signaling via Ras.³³ This possibility, however, is not feasible because Y794A mutation did not affect neurite extension in response to NGF. The MAP kinase pathway is known to be primarily involved in neurite extension. Thus, the fact that Y794F mutation did not affect neurite extension indicates that MAP kinase signaling is not disturbed in cells expressing Y794F-TrkA. The present findings that p11 production was diminished in TrkA mutated at Y499 or Y794, but not Y760, suggest that both MAP kinase and PLC pathways were involved in NGF-induced p11 production. It should be noted that p11 is expressed in glomus cells, but not adrenal medullary cells, which are developmentally closely related to glomus cells.⁴⁴ The molecular mechanisms elucidated for NGF-induced p11 expression in PC12 cells would pave the way to explore how p11 is expressed in glomus cells but not in adrenal medullary cells.

In conclusion, we elucidated the expression of p11 and heteromeric TASK1-TASK3 channels in rat CB glomus cells. In mouse hypoglossal motoneurons³ and cerebellar granule cells,² where heteromeric TASK1-TASK3 channels are putatively formed, p11 expression is confirmed at the mRNA level.⁴⁵ Those and the present findings suggest that p11 plays a major role in the generation of heteromeric TASK1-TASK3 channels. A further study, such as downregulation of p11 expression using a small interfering RNA approach, would be needed to elucidate whether p11 is essential for heteromeric channel formation in glomus cells or not.

Competing Interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

Author Contributions

HM, MP, RY, and MI contributed to the conception and design of the work; MP and MI made the preparation; HM acquired the data; HM and KH analyzed the data; and MI and MP wrote the manuscript. All authors have read and approved the final manuscript.

Funding

The author(s) disclosed receipt of the following financial support for the research, authorship, and/or publication of this article: This study was in part supported by grants JSPS KAKENHI (17K08555 to M. I. and 18K06865 to H. M.).

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