

IRBIT regulates CaMKII α activity and contributes to catecholamine homeostasis through tyrosine hydroxylase phosphorylation

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Inositol 1,4,5-trisphosphate receptor (IP₃R) binding protein released with IP₃ (IRBIT) contributes to various physiological events (electrolyte transport and fluid secretion, mRNA polyadenylation, and the maintenance of genomic integrity) through its interaction with multiple targets. However, little is known about the physiological role of IRBIT in the brain. Here we identified calcium calmodulin-dependent kinase II alpha (CaMKII α) as an IRBIT-interacting molecule in the central nervous system. IRBIT binds to and suppresses CaMKII α kinase activity by inhibiting the binding of calmodulin to CaMKII α . In addition, we show that mice lacking IRBIT present with elevated catecholamine levels, increased locomotor activity, and social abnormalities. The level of tyrosine hydroxylase (TH) phosphorylation by CaMKII α , which affects TH activity, was significantly increased in the ventral tegmental area of IRBIT-deficient mice. We concluded that IRBIT suppresses CaMKII α activity and contributes to catecholamine homeostasis through TH phosphorylation.

IRBIT | CaMKII α | catecholamine | hyperactivity

Inositol 1,4,5-trisphosphate receptor (IP₃R) binding protein released with IP₃ (IRBIT) was originally identified as a molecule that interacts with the intracellular calcium channel, IP₃R. IRBIT binds to and suppresses IP₃R activity in the resting state by blocking IP₃ access to IP₃R (1, 2). Our group and others have reported that IRBIT contributes to electrolyte transport, mRNA processing, and the maintenance of genomic integrity (3–9) through its interaction with multiple targets. However, little is known about the physiological role of IRBIT in the brain, where it is most highly expressed (1).

Calcium calmodulin (CaM) dependent kinase II alpha (CaMKII α) is a Ser/Thr kinase that is abundant in the central nervous system and is activated by the binding of Ca²⁺-CaM. CaMKII α phosphorylates various target proteins and is involved in the regulation of synaptic transmission and plasticity (10, 11). CaMKII α is expressed in the hippocampus, neocortex, thalamus, hypothalamus, olfactory bulb, cerebellum, and basal ganglia (12, 13). Many studies involving mutant mice and also pharmacological studies have indicated that CaMKII α activity is essential for the acquisition of memory and learning (14, 15). In addition, the appropriate regulation of CaMKII α is required for cognitive function and mood control (16–18). Thus, aberrant CaMKII α activity is associated with several neuronal disorders such as schizophrenia, autism spectrum disorder, attention-deficit hyperactivity disorder (ADHD), and drug addiction, in which hyperactivity and social abnormalities are frequently observed (19–23). However, the precise mechanism linking CaMKII α dysregulation and mental disorders is poorly understood.

Recent behavioral studies using knockout (KO) mouse models or pharmacological approaches have revealed that the dysregulation of dopamine (DA) systems is correlated with a hyperactive phenotype and social abnormalities (24–26). The catecholamines, DA

and norepinephrine (NE) are biosynthesized from the amino acids phenylalanine and tyrosine. The sequence of steps starts with the enzyme, tyrosine hydroxylase (TH). Thus, TH is the rate-limiting enzyme for both DA and NE synthesis. The appropriate regulation of TH activity is important for the maintenance of normal brain function and mental state (27).

In this study, we identified CaMKII α as an IRBIT-interacting molecule in the central nervous system. IRBIT binds to and suppresses the kinase activity of CaMKII α by inhibiting the binding of CaM to CaMKII α . In addition, we found that mice deficient in IRBIT present with hyperactivity and social abnormalities. In IRBIT KO mice, we observed increased catecholamine levels and hyperphosphorylation of Ser19 on TH, which is known to enhance TH activity and increase the biosynthesis of DA and NE (27, 28). Thus, we have concluded that IRBIT regulates CaMKII α activity and contributes to catecholamine homeostasis through TH phosphorylation.

Results

IRBIT Interacts with CaMKII α and Is Dissociated by an Excess of Ca²⁺-CaM. Because IRBIT is highly expressed in dendrites and spine-like structures and colocalizes with Homer 1, which is a postsynaptic marker in hippocampal neurons (29) (Fig. 1A), we speculate that IRBIT interacts with synaptic molecules and contributes to neuronal function. To identify the proteins that interact with IRBIT, we immunoprecipitated (IP) IRBIT from a crude membrane fraction

Significance

Appropriate homeostatic regulation of catecholamines (dopamine, norepinephrine) is important for the maintenance of normal brain function and mental state. The dysregulation of dopamine systems has been correlated with a hyperactive phenotype and social abnormalities, which are frequently observed in patients with psychiatric disorders. In this report, we found that IRBIT regulates catecholamine homeostasis by binding to calcium calmodulin-dependent kinase II alpha and subsequently controlling tyrosine hydroxylase phosphorylation. In addition, mice lacking IRBIT present with increased locomotor activity and social abnormalities. Our finding provides new insight into the homeostatic regulation of catecholamines.

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The authors declare no conflict of interest.

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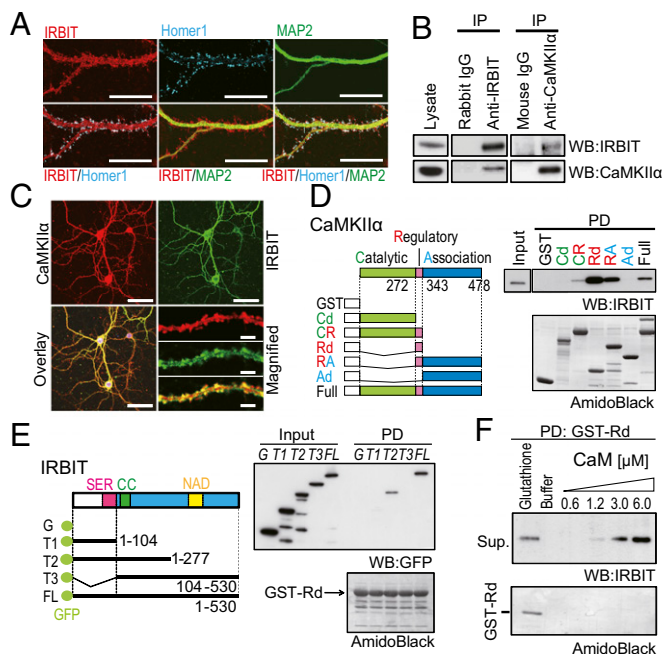


Fig. 1. IRBIT interacted with CaMKII α . (A) Cultured hippocampal neurons were stained with antibodies against IRBIT (red), Homer 1 (blue), and MAP2 (green). (Scale bar, 20 μ m.) (B) Co-IP of IRBIT and CaMKII α from the hippocampus. (C) The expression of CaMKII α (red) and IRBIT (green) antibodies. (Scale bar, 50 μ m.) The magnified images of the dendrite are shown. (Scale bar, 5 μ m.) (D) Identification of CaMKII α binding region to IRBIT in vitro. (Left) Schematic diagram of GST–CaMKII α truncated mutants. CR, catalytic domain plus regulatory domain; Rd, regulatory domain; RA, regulatory domain plus associated domain; Cd, catalytic domain; Ad, associated domain. (Right) Pull-down assay using various GST–CaMKII α mutants and purified IRBIT (from Sf9 cells). (E) Identification of the binding region of IRBIT to CaMKII α . (Left) Schematic diagram of GFP–IRBIT truncated mutants. (Right) Pull-down assay using GST–Rd and GFP–IRBIT truncation mutants. (F) Ca²⁺–CaM and IRBIT competitively bind to CaMKII α . Purified IRBIT was pulled down with GST–Rd and eluted with various concentration of Ca²⁺–CaM. Elution with glutathione was used as a positive control.

of the hippocampus, and IRBIT-associated proteins were identified using mass-spectrometry analysis. We found several peptides that matched the sequence of CaMKII α (SI Materials and Methods and Fig. S14). Physical interactions between IRBIT and CaMKII α have also been reported in proximal tubule cells (7). To confirm the interaction between IRBIT and CaMKII α in the brain, we performed a co-IP assay of samples from the hippocampus using anti-IRBIT and anti-CaMKII α antibodies. IRBIT was co-IP with CaMKII α using anti-CaMKII α antibody, and vice versa (Fig. 1B). Moreover, the immunostaining of primary-cultured hippocampal neurons showed that IRBIT and CaMKII α were extensively colocalized in spines and dendrites (Fig. 1C).

To further verify the direct interaction between IRBIT and CaMKII α , we performed a pull-down (PD) assay with the purified CaMKII α deletion mutants and purified IRBIT (from *Spodoptera frugiperda* 9 (Sf9) cells) and determined their binding domains. IRBIT bound to full-length CaMKII α protein in vitro (Fig. 1D). We also found that IRBIT binds to CaMKII α fragments that contain the regulatory domain (CR, Rd, and RA), but not to fragments containing the catalytic domain (Cd) or associated domain (Ad). As a control, no binding of IRBIT was observed to GST alone. To determine the region of IRBIT that is responsible for the interaction with CaMKII α , we performed a PD assay using various GFP–IRBIT deletion mutants and the regulatory domain of CaMKII α . CaMKII α bound to the T2 fragment and full-length (FL) IRBIT, but not to the T1 and T3

fragments, or GFP alone (Fig. 1E). These results indicate that IRBIT directly binds to CaMKII α in vitro. The amino acid residues 272–343 of CaMKII α (Rd) are sufficient for binding to IRBIT, and the two segments including the amino acid residues 1–104 and 105–277 of IRBIT are both necessary for binding to CaMKII α . Because the Ca²⁺–CaM complex also binds to the regulatory domain of CaMKII α , we next tested whether IRBIT and the Ca²⁺–CaM complex bind competitively to the regulatory domain. An excess of Ca²⁺–CaM dissociated IRBIT from the regulatory domain of CaMKII α (Fig. 1F).

IRBIT Inhibits Kinase Activity of CaMKII α in Vitro. Although a previous report showed that IRBIT is a substrate of CaMKII α (7), the effect of IRBIT on CaMKII α activity is unknown. To explore the functional effect of IRBIT binding on CaMKII α kinase activity, we performed an in vitro kinase assay using purified IRBIT (from Sf9 cells). IRBIT significantly inhibited CaMKII α kinase activity in a concentration-dependent manner (Fig. 2A). When we added a relatively lower concentration of IRBIT (0.1 or 0.5 μ M), the double-reciprocal analysis of these data yielded straight lines intersecting on the y axis, indicating simple competitive inhibition of CaMKII α by IRBIT with respect to the Ca²⁺–CaM complex (Fig. 2B). Interestingly, the double-reciprocal plot for 1.0 μ M IRBIT showed complex inhibition. Boiled IRBIT did not affect the CaMKII α kinase activity (Fig. S1B). In addition, a large excess of CaM (50 μ M) completely blocked the inhibition of CaMKII α by 1.0 μ M IRBIT (Fig. S1C). In contrast, the inhibition by IRBIT was noncompetitive with respect to ATP (Fig. S1D and Fig. 2C) and was not affected by an excess of substrate

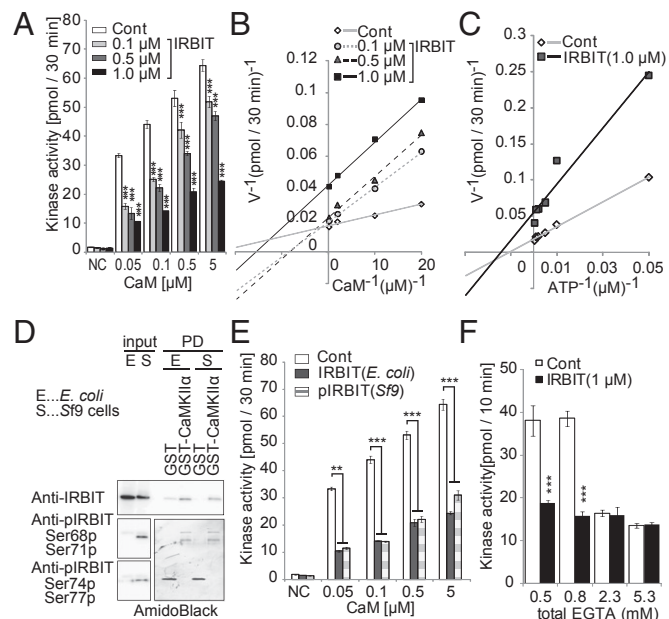


Fig. 2. IRBIT inhibits the CaMKII α kinase activity in vitro. (A) Effect of IRBIT on the Ca²⁺–CaM dependency of CaMKII α activity. Synthetic peptide (syntide-2) was phosphorylated by purified CaMKII α (4 nM) with or without IRBIT. CaM and IRBIT concentrations are indicated. Negative control (NC): kinase assay without CaM. ATP: 200 μ M. Syntide-2: 10 μ M. (B) The double-reciprocal plot analysis of Fig. 2A. (C) Effect of IRBIT on the ATP dependency of CaMKII α activity. CaM: 0.1 μ M. Syntide-2: 10 μ M. (D) Effect of IRBIT phosphorylation on the IRBIT–CaMKII α interaction. Purified IRBIT from *E. coli* or Sf9 cells were pulled down with GST–CaMKII α . (E) Effect of nonphospho-IRBIT (*E. coli*) and phospho-IRBIT (Sf9 cells) on the CaMKII α activity. ATP: 200 μ M. Syntide-2: 10 μ M. (F) Effect of IRBIT on the Ca²⁺–CaM-independent CaMKII α activity. In vitro kinase assay was performed with various concentrations of EGTA. $n = 3$. ATP: 200 μ M. Syntide-2: 10 μ M.

(Fig. S1 E and F). Therefore, we concluded that IRBIT inhibits CaMKII α kinase activity by competing with the Ca²⁺-CaM complex.

Because IRBIT interacts with various binding partners in a phosphorylation-dependent manner (2–4), we next examined the phosphorylation dependency of the IRBIT–CaMKII α interaction using nonphospho-IRBIT and phosphorylated IRBIT, purified from *Escherichia coli* and Sf9 insect cells, respectively. Both nonphospho-IRBIT and phospho-IRBIT interacted similarly with GST–CaMKII α (Fig. 2D). Phosphorylation of IRBIT was confirmed using antibodies specific for phospho-IRBIT (S68p/S71p and S74p/S77p). In addition, nonphospho-IRBIT inhibited CaMKII α kinase activity as effectively as phospho-IRBIT (Fig. 2E). These results indicated that the phosphorylation of IRBIT (at S68/S71 and S74/S77), which is essential for other known interactions, is not necessary for the regulation of CaMKII α .

CaMKII α has Ca²⁺-CaM-independent activity in its T286 autophosphorylated form (30). We therefore investigated the effect of IRBIT on the Ca²⁺-CaM-independent activity of CaMKII α . Purified CaMKII α was preincubated with Ca²⁺-CaM and ATP to achieve autophosphorylation. Phosphorylated CaMKII α was mixed with substrate and IRBIT in the presence of various concentrations of EGTA (details in Fig. S1G). As shown in Fig. 2F, IRBIT inhibited the CaMKII α kinase activity at low concentrations of EGTA (0.3, 0.8 μ M). However, IRBIT did not affect kinase activity at higher concentrations of EGTA (2.3, 5.3 μ M). Thus, IRBIT did not regulate Ca²⁺-CaM-independent activity of CaMKII α . In addition, we confirmed the effect of IRBIT on CaMKII α kinase activity using recombinant Homer 3 protein, which is a substrate for CaMKII α (31). IRBIT significantly inhibited the phosphorylation of Homer 3 in a concentration-dependent manner (Fig. S1 H and I). Therefore, we concluded that IRBIT negatively regulates CaMKII α kinase activity in vitro.

IRBIT Regulates Kinase Activity of CaMKII α in Living Cells. To further explore the functional effect of IRBIT on CaMKII α activity in living cells, we examined the effect of IRBIT overexpression on the Ca²⁺-ionophore (4-Bromo-A23187, BrA) induced phosphorylation of Homer 3 in HEK-293 cells that stably express GFP–CaMKII α (GFP–CaMKII α cells). Because IRBIT has been reported as a CaMKII α substrate (7), we used an IRBIT T1 fragment (indicated in Fig. 1E), with disrupted binding to CaMKII α , but containing the consensus sequence for the CaMKII α substrate motif as a negative control. The magnitude of the BrA-induced increase in intracellular calcium was not affected by the overexpression of IRBIT or the IRBIT T1 fragment compared with control cells (Fig. S2 A and B). However, overexpression of IRBIT significantly reduced the BrA-induced phosphorylation of Homer 3 compared with the control (Fig. 3 A and B), whereas overexpression of the IRBIT T1 fragment did not. We then investigated the effect of IRBIT knockdown on CaMKII α kinase activity within cells. Transfection with siRNAs targeting IRBIT clearly abolished the IRBIT signal in GFP–CaMKII α cells (Fig. S2C), but did not affect the calcium increase induced by BrA (Fig. S2 D and E). Then we expressed HA–Homer 3 with control siRNAs or IRBIT siRNAs in GFP–CaMKII α cells and stimulated the cells using 2.5 μ M BrA. The knockdown of IRBIT significantly increased the BrA-induced phosphorylation of Homer 3 compared with the control siRNA (Fig. 3 C and D).

To further confirm the effect of IRBIT on CaMKII α activity, we expressed a fluorescence resonance energy transfer (FRET)-based CaMKII α activity probe (Camuix) (32) in mouse embryonic fibroblast (MEF) cells from IRBIT KO and wild-type (WT) mice. The morphology of IRBIT KO MEF cells was similar to WT MEF cells (Fig. S3A). Both WT and KO MEF cells showed roughly normal growth rates and expressed other Ca²⁺-associated proteins at a similar level (Fig. S3B). Interactions between IRBIT and Camuix were confirmed by co-IP (Fig. S3C). We performed simultaneous imaging of the Ca²⁺ change and Camuix FRET. We found that the overexpression of IRBIT in WT MEF

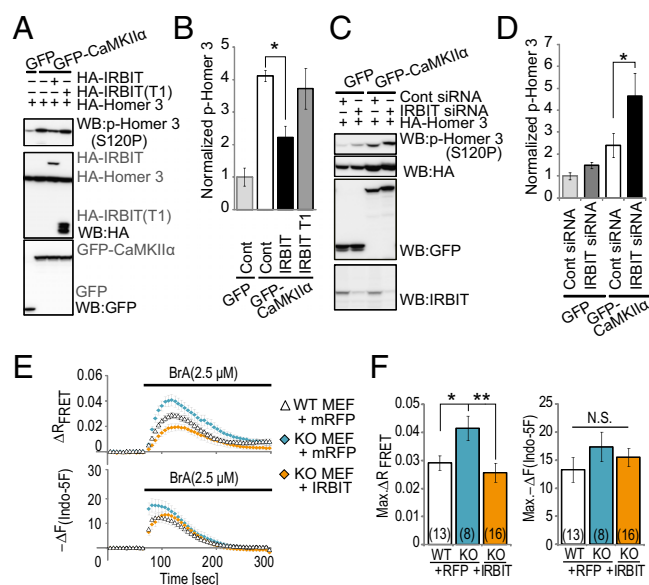


Fig. 3. IRBIT suppresses the CaMKII α kinase activity in living cells. (A) IRBIT overexpression inhibited phosphorylation of Homer 3. GFP or GFP–CaMKII α cells were transfected with HA–Homer 3 and HA–IRBIT or HA–IRBIT (T1, 1–104). After 24 h, cells were stimulated with 2.5 μ M BrA for 2 min. (B) Quantitative analysis of phospho-Homer 3 in Fig. 3A. $n = 3$. (C) Phosphorylation of Homer 3 was enhanced in IRBIT knock-down cells. GFP–CaMKII α cells were transfected with HA–Homer 3 and control siRNA or IRBIT siRNA. After 48 h, cells were stimulated with 2.5 μ M BrA for 2 min. (D) Quantitative analysis of phospho-Homer 3 in Fig. 3C. $n = 4$. (E) Simultaneous imaging of FRET and Ca²⁺ change in IRBIT WT or KO MEF cells transfected with mRFP or IRBIT/mRFP. (Upper) Representative FRET changes are shown. (Lower) Representative Ca²⁺ responses (Indo-5F) are shown. (F) Quantitation of FRET and Ca²⁺ peak amplitude. Peak amplitude FRET (Max. Δ R FRET) and Ca²⁺ responses [Max. $-\Delta$ F (Indo-5F)] that were expressed as the averaged amplitude of 0–50 s are equal to zero. Results show three independent experiments. The total cell numbers are indicated in each graph.

cells significantly inhibited the BrA-induced Camuix FRET change, but did not affect the Ca²⁺ change (Fig. S3 D and E). In addition, the BrA-induced Camuix FRET change was significantly increased in IRBIT KO MEF cells compared with WT MEF cells (Fig. 3 E and F). This enhancement of the FRET change was diminished by exogenous IRBIT expression in IRBIT KO MEF cells. On the other hand, we found that point mutation of CaMKII α regulatory domain (T286A and T286D) attenuated interaction between IRBIT and CaMKII α (Fig. S3F) and the overexpression of IRBIT did not affect BrA-induced FRET change of these Camuix mutants in WT MEF cells (Fig. S3 G and H). Thus, the effect of IRBIT on Camuix FRET change correlates with interaction between IRBIT and CaMKII α .

NMDA Receptor Activation Causes Prolonged CaMKII α Activity in IRBIT KO Hippocampal Neurons. To verify that IRBIT regulates CaMKII α activity in neurons, we transfected Camuix into cultured hippocampal neurons derived from WT or IRBIT KO mice and performed simultaneous imaging of FRET and Ca²⁺ change. Transfected neurons were transiently stimulated with 25 μ M NMDA with 1 μ M glycine. The transient NMDA-induced FRET change was significantly enhanced and more persistently observed in IRBIT KO neurons, whereas the Ca²⁺ change was not affected (Fig. 4 A and B). This enhancement of the FRET change was inhibited by exogenous IRBIT expression in IRBIT KO neurons. From these results, we concluded that IRBIT inhibits CaMKII α activity in neurons.

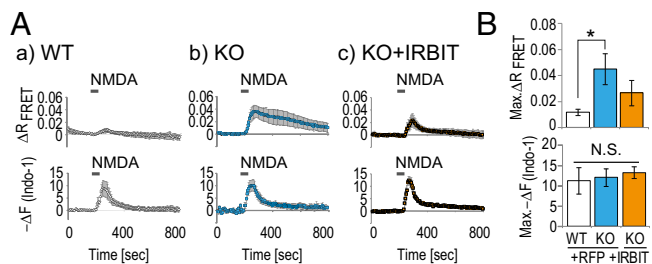


Fig. 4. CaMKII α activity is enhanced in hippocampal neurons from IRBIT KO mice. (A) Simultaneously imaging of FRET and Ca²⁺ change (Indo-1) after NMDA stimulation in WT or IRBIT KO cultured hippocampal neurons transfected with mRFP or IRBIT/mRFP. (B) Quantitation of FRET and Ca²⁺ peak amplitude [Max. $-\Delta F(\text{Indo-1})$] that were expressed as the averaged amplitude of 0–50 s are equal to zero.

IRBIT KO Mice Show Hyperactivity and Social Abnormalities. To examine the role of IRBIT on brain function, we performed behavioral testing on the IRBIT KO mouse (6). IRBIT KO mice showed slight dwarfism and low body weight, whereas their brain size and histological structures appeared normal (Fig. S4). We did not observe any significant differences between the WT and KO mice in tests of muscle strength, motor function, pain response, and fear responses (Fig. S5 A–I). However, interestingly, IRBIT KO mice showed increased locomotor activity compared with WT mice (Fig. 5 A–D). To evaluate the effect of environmental factors on the hyperactivity of IRBIT KO mice, we further tested their social interactions and monitored their activity in the home cage (Fig. S5 J and K). Consistent with the results of open field tests, IRBIT KO mice showed enhanced activity in the home cage (Fig. S5 J and K, Lower). In addition, IRBIT KO mouse showed an increment in contact between mice (Fig. S5 J and K, Upper). We also measured the social interaction in a novel environment. IRBIT KO mice also displayed enhanced social interaction in an open field (Fig. S5 L–P). From these results, we concluded that the mice lacking IRBIT present with hyperactivity and social abnormalities.

Dopamine and Norepinephrine Levels in IRBIT KO Mice. Recent behavioral studies using either KO mice or pharmacological approaches have revealed a correlation between dysregulated monoamine homeostasis and the hyperactive phenotype with social abnormalities (24–26, 33). Therefore, we examined monoamine levels in the brains of IRBIT KO mice. The sampling locations are illustrated in Fig. 5E. DA, NE, and their metabolites were significantly increased in the prefrontal cortex (PFC), hippocampus, striatum, and/or cerebellum of IRBIT KO mice (Fig. 5 F–K). By contrast, the level of serotonin and its metabolites showed no difference in the midbrain, including the raphe nucleus, which is the main source of serotonergic innervation. These results suggested that dysregulation of catecholamine homeostasis could be an explanation for the hyperactivity and the social abnormality of IRBIT KO mice.

Phosphorylation of TH Was Elevated in IRBIT KO Mice. To reveal the molecular mechanism by which IRBIT deletion leads to dysregulation of catecholamine homeostasis, we focused on TH, the rate-limiting enzyme for both DA and NE synthesis. Because CaMKII α regulates TH activity by phosphorylation (34), we first examined whether the expression of IRBIT affects TH phosphorylation via CaMKII α in GFP–CaMKII α cells. The over-expression of IRBIT significantly decreased the BrA-induced phosphorylation of TH compared with the control, whereas the IRBIT T1 fragment that lacked the ability to bind CaMKII α had no effect (Fig. 6 A and B). Next, we transfected GFP–IRBIT into rat pheochromocytoma-12 (PC-12) cells and evaluated the effect

of IRBIT on endogenous TH phosphorylation. In both the basal and stimulated states (50 mM KCl, 10 min), the phosphorylation level of endogenous TH was significantly decreased in PC-12 cells transfected with GFP–IRBIT compared with GFP alone (Fig. 6 C and D). We also examined TH phosphorylation levels in dopaminergic neurons in the ventral tegmental area (VTA) of IRBIT KO mice, where significant IRBIT expression was detected (Fig. 6E). We used anti-TH and antiphospho-TH (S19) specific antibodies to evaluate TH phosphorylation levels by immunohistochemistry. The phosphorylation level of TH was slightly but significantly increased in IRBIT KO dopaminergic neurons (Fig. 6 F–H), whereas the expression level of TH was not changed (Fig. 6I). These data suggested that the expression level of IRBIT determines the phosphorylation level of TH via CaMKII α , which might lead to increased TH activity and subsequently, to aberrant catecholamine homeostasis in IRBIT KO mice.

Discussion

We have investigated the physiological function of IRBIT in the brain. We showed that (i) IRBIT bound to CaMKII α and inhibited CaMKII α kinase activity in vitro and in living cells, (ii) IRBIT KO mice exhibited hyperactivity and social abnormalities, (iii) catecholamine levels were increased in IRBIT KO brain, (iv) TH phosphorylation in VTA neurons was increased in IRBIT KO mice, and (v) the expression level of IRBIT determined the phosphorylation of TH by CaMKII α . These data support a model in which IRBIT regulates CaMKII α activity and contributes to

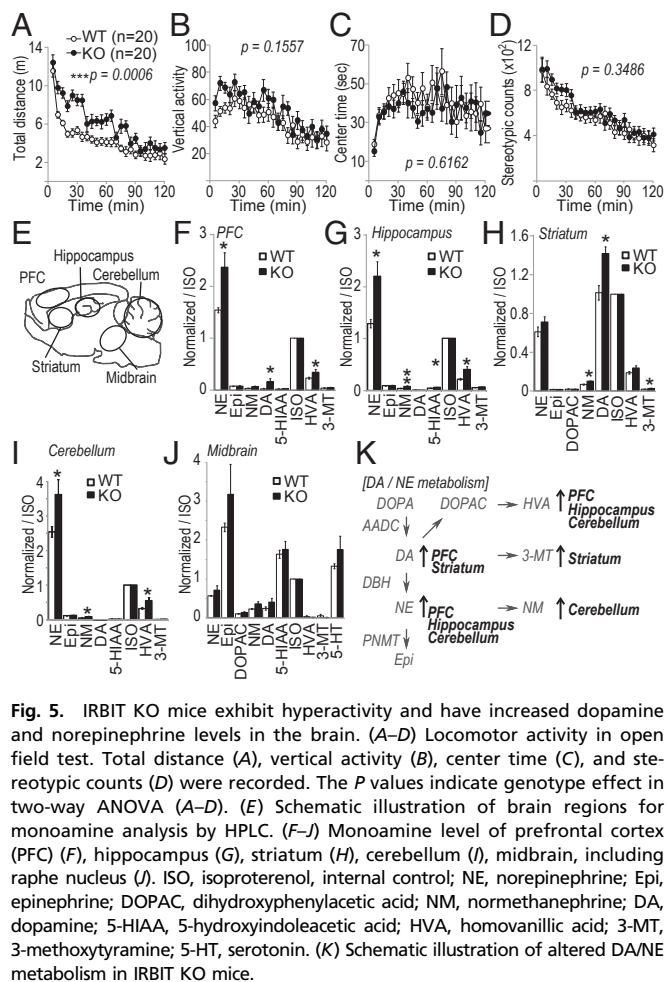


Fig. 5. IRBIT KO mice exhibit hyperactivity and have increased dopamine and norepinephrine levels in the brain. (A–D) Locomotor activity in open field test. Total distance (A), vertical activity (B), center time (C), and stereotypic counts (D) were recorded. The P values indicate genotype effect in two-way ANOVA (A–D). (E) Schematic illustration of brain regions for monoamine analysis by HPLC. (F–J) Monoamine level of prefrontal cortex (PFC) (F), hippocampus (G), striatum (H), cerebellum (I), midbrain, including raphe nucleus (J). ISO, isoproterenol, internal control; NE, norepinephrine; Epi, epinephrine; DOPAC, dihydroxyphenylacetic acid; NM, normethamphetamine; DA, dopamine; 5-HIAA, 5-hydroxyindoleacetic acid; HVA, homovanillic acid; 3-MT, 3-methoxytyramine; 5-HT, serotonin. (K) Schematic illustration of altered DA/NE metabolism in IRBIT KO mice.

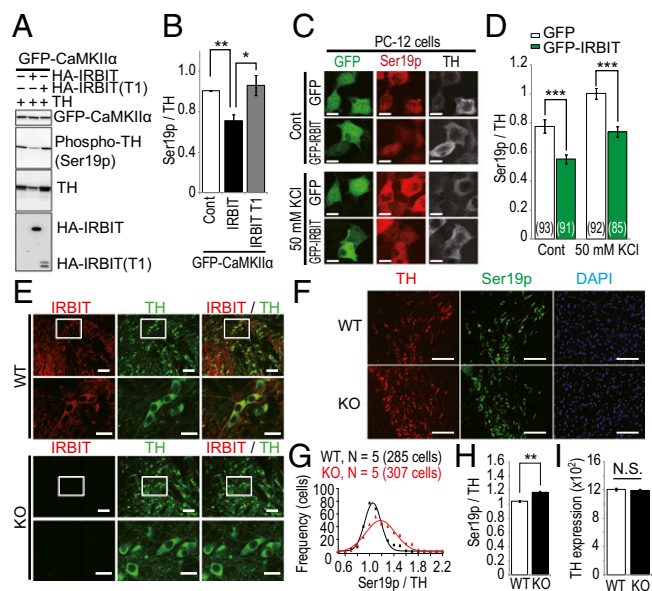


Fig. 6. IRBIT regulates the phosphorylation level of tyrosine hydroxylase (TH) in the dopaminergic neurons of the VTA. (A) GFP-CaMKII α cells were transfected with TH and HA-IRBIT or HA-IRBIT T1 fragment (1–104). After 24 h, cells were stimulated with 2.5 μ M BrA for 2 min. (B) Quantitative analysis of phospho-TH in A. (C) PC-12 cells were transfected with GFP or GFP-IRBIT. After 24 h, cells were stimulated with 50 mM KCl for 10 min. Nonstimulated and KCl-stimulated cells were fixed and stained with indicated antibodies. (D) Quantitative analysis of phospho-TH in C. Total number of cells is indicated in each bar. (E) Adult WT and IRBIT KO mice brain sections were stained with anti-IRBIT (red) and anti-TH (green) antibodies. (Scale bar, 50 μ m.) The boxed regions are shown at increased magnification. (Scale bar, 20 μ m.) (F) Adult WT and IRBIT KO mice brain sections were stained with anti-TH (red), anti-phospho-TH (Ser19p, green) antibodies, and DAPI (blue). (Scale bar, 100 μ m.) (G–I) Quantitative analysis of phospho-TH levels of dopaminergic neurons in the VTA of IRBIT KO mice. (G) Histogram of normalized TH phosphorylation. (H) Average of normalized TH phosphorylation. (I) Average of TH expression. $n = 5$. Total cell numbers of quantified TH positive neurons were 285 (WT) and 307 (KO).

catecholamine homeostasis through TH phosphorylation, and dysfunction of IRBIT causes hyperactivity and abnormal social phenotypes in mice.

Several reports have stated that CaMKII α activity is related to locomotor activity in mice. The transient overexpression of CaMKII α in the nucleus accumbens enhanced the hyperlocomotion response to amphetamine and AMPA (17, 35). The expression of CaMKII α was increased in the VTA after repeated cocaine injections, which induced locomotor hyperactivity (36). Moreover, the administration of the CaMKII α inhibitor into the VTA prevented the augmentation of cocaine-induced hyperactivity (37). In addition, CaMKII α activity was shown to be excessively elevated in the medial prefrontal cortex of ADHD model rats (38). On the other hand, CaMKII α heterozygous mice and autophosphorylation-deficient mice (T286A) also present with locomotor hyperactivity (18, 36). It has also been reported that overexpression of CaMKII α using the CaMKII α promoter leads to decreased locomotor activity in mice (39). Thus, both hyper- and hypoactivation of CaMKII α seems to cause hyperactive phenotypes in mice. Although the precise reasons for these contradictory observations are unknown, the contribution of the serotonergic pathway may reconcile the two observations. The serotonergic pathway plays a crucial role in the regulation of locomotor activity. For example, serotonin transporter KO mice show increased 5-HT release and hypolocomotion (40). It has been reported that CaMKII α

heterozygous and homozygous mice show a reduction in 5-HT release (41).

In IRBIT KO mice, DA, NE, and their metabolites were significantly increased in the PFC, hippocampus, striatum, and/or cerebellum. However, the levels of serotonin and its metabolites showed no difference in the midbrain, including the raphe nucleus (Fig. 5). In addition, we observed appreciable expression of IRBIT in the dopaminergic neurons of the VTA (Fig. 6A), but the immunosignal for IRBIT in 5HT-positive neurons of the raphe nucleus was very weak and below the limit of detection (Fig. S4G). Thus, it is possible that enhanced CaMKII α activity in the dopaminergic pathway, but not in the serotonergic pathway, contributes to the altered catecholamine homeostasis and increased locomotor activity of IRBIT KO mice.

The regulation of CaMKII α by IRBIT was competitive with respect to the Ca²⁺-CaM complex, because IRBIT bound to the regulatory domain of CaMKII α and was released by an excess of Ca²⁺-CaM (Fig. 1). Double-reciprocal analysis of the *in vitro* kinase assays indicated simple competitive inhibition of CaMKII α by IRBIT with respect to Ca²⁺-CaM at relatively lower concentrations of IRBIT (Fig. 2B, 0.1 or 0.5 μ M). However, the double-reciprocal plot at a higher concentration of IRBIT (1.0 μ M) apparently showed complex inhibition. The autophosphorylation of regulatory domain regulates CaMKII α activity (42–44). CaMKII α forms a dodecameric holoenzyme, and autophosphorylation of CaMKII α seems to occur *in trans*, between two subunits of the same holoenzyme (45). Because IRBIT is bound to the regulatory domain of CaMKII α (Fig. 1), it is possible that IRBIT affects the efficiency of autophosphorylation of CaMKII α and apparently leads to complex inhibition at a high concentration. We performed a binding assay using IRBIT and CaMKII α T286A or T286D mutants to separate the effects of Ca²⁺-CaM competition and the effects on autophosphorylation, but these mutations themselves affected the interaction between IRBIT and CaMKII α (Fig. S3F). This means that the interaction between IRBIT and CaMKII α is in close proximity to the T286 autophosphorylation site of CaMKII α . A detailed analysis of the kinetics using simulations is our next priority, to reveal the precise mechanism by which IRBIT regulates CaMKII α , because CaMKII α forms a dodecameric holoenzyme with multiple activity states brought about by phosphorylation, and IRBIT also seems to form a multimer (2).

In summary, we found that IRBIT regulates CaMKII α activity and contributes to catecholamine homeostasis through TH phosphorylation. IRBIT was originally identified as an IP₃R-interacting molecule and inhibited IP₃R-Ca²⁺ signaling. Deletion of IRBIT results in both enhancement of Ca²⁺ release from IP₃R and a lower threshold for CaMKII α activation. Thus, dysfunction of IRBIT would synergistically activate IP₃R-Ca²⁺-CaMKII α signaling. IRBIT has multiple targets in cells. Further studies using an IRBIT knock-in mutant mouse that specifically disrupts the interaction with each target molecule would help us to reveal the precise pathogenic mechanism underlying hyperactivity and social abnormalities. In addition, a multitude of mutant mouse studies and pharmacological studies has clearly demonstrated that CaMKII α activity is essential for memory and learning acquisition. It will be interesting to see how IRBIT contributes to the regulation of synaptic plasticity, learning, and memory in the brain.

Materials and Methods

A full description of the materials and methods can be found in *SI Materials and Methods*.

Animals. All animal experiments were performed in accordance with the RIKEN guidelines for animal experiments. Every effort was made to minimize the number of animals used.

In Vitro Kinase Assay. The kinase reaction was performed in kinase assay buffer [25 mM Hepes, pH 7.4, 1 mM MgCl₂, 0.3 mM EGTA, 0.33 mM CaCl₂, 1 mM 2-mercaptoethanol, 0.005% (wt/vol) Triton X-100, 0.87 μM BSA]. Reactions were performed on ice to avoid the autophosphorylation-dependent decrease in kinase activity. It is described in a previous report that performing experiments at 0 °C prevents autophosphorylation at the inhibitory sites on CaMKIIα (T305 and T306) (46). Recombinant Homer 3 or synthetic peptide, syntide-2 in kinase assay buffer, was incubated with 0.2 ng/μL recombinant CaMKIIα, various concentration of CaM, and with 200 μM ATP and [γ-³²P] ATP in the presence or absence of IRBIT. The reaction was stopped by addition of 83 mM EDTA. For phosphorylation of Homer 3, reaction mixture was separated by SDS/PAGE and phosphorylation level of Homer 3 was detected by Western blotting analysis with the antibody against phospho-Homer 3 (p-Ser120). For phosphorylation of syntide-2, reaction mixture was spotted onto the P81 phosphocellulose paper (GE Healthcare) and radioactivity was counted with a liquid scintillation counter.

Statistical Analysis. Statistical comparison between two independent groups of data were performed with the Student's *t* test. Other statistical analysis was conducted using IgorPro 4.0 or StatView (SAS Institute). Data were

analyzed by two-way ANOVA, or two-way repeated measures ANOVA, or one-way ANOVA followed by Bonferroni–Dunn test unless noted otherwise. Values in graphs were expressed as mean ± SEM **P* < 0.05, ***P* < 0.01, ****P* < 0.001.

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