**Chronic stress impairs autoinhibition in neurons of**

**the locus coeruleus to increase asparagine endopeptidase activity**

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## Abstract

Impairments of locus coeruleus (LC) are implicated in anxiety/depression and Alzheimer's disease (AD). Increases in cytosolic noradrenaline (NA) concentration and MAO-A activity initiate the LC impairment through production of NA metabolite, 3,4-dihydroxyphenyl-glycolaldehyde (DOPEGAL), by MAO-A. However, how NA accumulates in soma/dendritic cytosol of LC neurons has never been addressed despite the fact that NA is virtually absent in cytosol while NA is produced exclusively in cytoplasmic vesicles from dopamine by dopamine-β-hydroxylase. Since re-uptake of autocrine-released NA following spike activity is the major source of NA accumulation, we investigated whether and how chronic stress can increase the spike activity accompanied by NA autocrine. Overexcitation of LC neurons is normally prevented by the autoinhibition mediated by activation of α2A-adrenergic receptor (AR)-coupled inwardly rectifying potassium-current (GIRK-I) with autocrine-released NA. Patch-clamp study revealed that NA-induced GIRK-I in LC neurons was decreased in chronic restraint stress (RS) mice while a similar decrease was gradually caused by repeated excitation. Chronic RS caused internalization of α2A-ARs expressed in cell membrane in LC neurons and decreased protein/mRNA levels of α2A-ARs/GIRKs in membrane fraction. Subsequently, chronic RS increased the protein levels of MAO-A, DOPEGAL-induced asparagine endopeptidase (AEP) and tau N368. These results suggest that chronic RS-induced overexcitation due to the internalization of α2A-ARs/GIRK is accompanied by $[Ca^{2+}]_i$ increases, subsequently increasing $Ca^{2+}$-dependent MAO-A activity and NA-autocrine. Thus, it is likely that internalization of α2A-AR increased cytosolic NA, as reflected in AEP increases, by facilitating re-uptake of autocrine-released NA. The suppression of α2A-AR internalization has a strong translational potential for AD.

**Keywords:** locus coeruleus; α2A adrenergic receptor; GIRK; autoinhibition; MAO-A; AEP

## Background

Locus coeruleus (LC) is implicated in anxiety and depression (*Itoi and Sugimoto, 2010*), and is often the first place where Alzheimer's disease (AD)-related pathology appears (*Mather and Harley, 2016*). Noradrenaline (NA) is known to inhibit amyloid-induced oxidative stress, and caspase activation in cortex/hippocampus (*Counts and Mufson, 2010*). Thus, the impairment of LC-noradrenergic system is potentially involved in the AD pathogenesis (*Ross et al., 2015; Chen et al., 2022*).

It was reported that cytosolic NA in the soma/dendrites of LC neurons is degraded by monoamine oxidase A (MAO-A) into 3,4-dihydroxyphenyl-glycolaldehyde (DOPEGAL) (*Burke et al., 1999*), which generates apoptotic free radicals (*Burke et al., 1998*) and activates asparagine endopeptidase (AEP) to produce Aβ and hypephosphorylated tau, consequently leading to AD (*Burke et al., 1999; Kang et al., 2020*). However, there appears normally no NA except dopamine in the cytosol of LC neurons because NA is produced exclusively in cytoplasmic vesicles by dopamine-β-hydroxylase (DBH) (*Potter and Axelrod, 1963*) from dopamine which is taken up into cytoplasmic vesicles by vMAT2.

LC neurons release NA following action potential not only from the axon terminals but also from their cell bodies as autocrine (*Huang et al., 2007*), and autocrine-released NA activates α2 adrenergic receptor (α2-AR) expressed on the soma/dendrites (*Aoki et al., 1994*), resulting in activation of α2-AR-coupled inwardly rectifying $K^+$ (GIRK) channel (*Arima et al., 1998*) to cause autoinhibition (spike-afterhyperpolarization) (*Williams et al., 1985*). Thus, overexcitation of LC neurons is normally prevented by the autoinhibition.

As autocrine-released NA is taken up by NA-transporter (NAT) (*Torres et al., 2003*), uptaken-NA may be the only source of free NA to be metabolized by MAO-A in the cytosol of LC neurons. However, the amount of free NA to be metabolized by MAO-A in the cytosol is thought to be very small due to a much lower affinity of NA to MAO-A compared to vMAT2 (*Costa, 2012*). Therefore, it is not clear how free NA to be metabolized by MAO-A is increased in the soma/dendites of LC neurons in AD pathogenesis. Because the passive NA-leakage occurs from cytoplasmic vesicles in a manner that achieves a dynamic equilibrium with active NA-storage into cytoplasmic vesicles by vMAT2 (*Eisenhofer et al., 2004*), increased NA-uptake by NAT in response to increased NA-autocrine following $[Ca^{2+}]_i$ increases or overexcitation may increase free NA by facilitating the passive NA-leakage from cytoplasmic vesicles. Furthermore, the affinity of MAO-A for NA may be modifiable as MAO-A activity is $Ca^{2+}$-dependent (*Cao et al., 2007*). Then, a question arises whether or not impairment of

autoinhibition that causes overexcitation and $[Ca^{2+}]_i$ increases can consequently increase AEP
activity through the production of DOPEGAL.

Neuronal activity in LC neurons is transiently increased by stress (*Valentino and Foote,*
*1988*) through activation of corticotropin releasing factor (CRF) receptor (*Valentino et al.,*
*1983; McCall et al., 2015*) together with glutamate receptors (*Valentino et al., 2001*). However,
even if firing activity is increased by synaptic activation (*Barcomb et al., 2022*), autoinhibition
would occur to suppress overexcitation in LC neurons. Nevertheless, it is known that long-
term overexcitation due to chronic stress ultimately causes degeneration of LC neurons
(*Kitayama et al., 2004*). Then, this raises an important question whether and how stress impairs
autoinhibition to cause sustained overexcitation. The whole mechanisms for the impairment
of LC would be revealed by solving these questions. Here we examined whether the
autoinhibition mediated by α2A-AR-coupled GIRK-I is impaired by overexcitation in LC
neurons or chronic stress, to clarify the mechanisms for NA accumulation and/or MAO-A
activity increase.

## 109  Results

**110  Spike frequency adaptation in LC neurons is mediated by α2-AR-coupled GIRK channel**
LC neurons invariably display spike-frequency adaptation (**Figure 1A, B**). We have
hypothesized that the autoinhibition would occur following respective spike firing and
contribute to the generation of the spike-frequency adaptation and the afterhyperpolarization
seen at the offset of current pulses.

When the cumulative number of spikes was plotted against the elapsed time from the
onset of the current pulse every 50 ms, the cumulative spike number increased non-linearly in
a saturation manner (**Figure 1C**). This non-linear relationship was fitted with a saturation
curve defined by Monod equation (see Extended Methods and Materials) in order to quantify
the spike-frequency adaptation. Because there was no significant difference in *b* between the
two groups of spike trains, the profile of the normalized cumulative spike number vs the elapsed
time remains the same regardless of firing frequency, suggesting that spike-frequency
adaptation occurs in the same way.

The spike-frequency adaptation observed was almost completely abolished by a blocker
of α2-AR, atipamezole (**Figure 1D, E**), which subsequently largely decreased pulse-
afterhyperpolarization (pulse-AHP) (**Figure 1D, E**, blue and red stars). The cumulative spike
number increased non-linearly in a saturation manner in the control condition while it increased

almost in a linear manner after atipamezole application (**Figure 1F**). In the relationship
between *a* and *b* examined in LC neurons, atipamezole invariably and significantly shifted both
*a* and *b* in a larger direction, and there was a significant difference in the relationship between
*a* and *b* (**Figure 1G**).

Bath application of NA induced a GIRK-I in response to the ramp pulse (**Figure 1H**, red
trace) whereas addition of atipamezole to NA almost completely abolished the NA-induced
GIRK-I (**Figure 1H**, blue trace), showing that GIRK-I reversed at –97 mV. The amplitudes of
the inward component of NA-induced GIRK-I were invariably larger than those of the outward
component, consistent with the inwardly rectifying property of GIRK-I. Atipamezole
significantly decreased the amplitudes of the inward and outward components (**Figure 1I**).
Taken together, NA autocrine is likely to be caused following respective spikes in the LC
neuron and NA is accumulated around the LC neuron to activate α2-AR-coupled GIRK
channels, to cause spike-frequency adaptation by enhancing AHP.

**$Ca^{2+}$-dependent rundown of NA-induced GIRK-I in LC neurons**
In PC12 cells, heterologously expressed α2A-ARs undergo internalization following agonist
application (*Taraviras et al., 2002*). Such internalization of α2A-ARs would result in an
abolishment of GIRK-I or desensitization given the functional coupling between α2A-AR and
GIRK as in LC noradrenergic neurons. However, in LC neurons, NA-induced GIRK-I showed
almost no rundown/desensitization after repeating the ramp pulse 20 times in the persistent
presence of NA (**Figure supplement S1A-D**). Then, we next examined whether GIRK-I
shows rundown under the condition of "increased excitability" or increased $[Ca^{2+}]_i$, because
acute/chronic stress increases the excitability of LC neurons (*Borodovitsyna et al., 2018;*
*Campos-Lira et al., 2018*) and various channels/receptors are known to be internalized in a
$Ca^{2+}$-dependent manner (*Beattie et al., 2000; Aziz et al., 2012; Zaccor et al., 2020*).

To examine the $Ca^{2+}$-dependency of the rundown of GIRK-I, the respective ramp
pulses to evoke GIRK-I were preceded by various pulse trains which would induce $Ca^{2+}$
currents differentially (compare **Figure supplement 1F** and **Figure 2B**). After examining
various pulse protocols (**Figures supplement 1 and 2**), the best protocol to induce a prominent
rundown of GIRK-I was found as shown in **Figure 2B**. First, bath application of NA induced
GIRK-I (**Figure 2A** red trace; 2) that showed the inwardly rectifying profile as revealed by
subtraction of the control response from that obtained after NA application (**Figure 2D**, red
trace; 2−1). Subsequently, the combined command pulse (**Figure 2B**, upper trace) was applied

every minute. The positive pulse trains evoked large inward $Ca^{2+}$ currents in the presence of

extracellular 30 mM TEA (**Figure 2B**, lower traces) whereas similar trains of the same positive

pulse in the absence of TEA hardly induced $Ca^{2+}$ currents (**Figure supplement 1F**, lower trace).

With repetition of the combined command pulse 20 times, the NA-induced GIRK-I (**Figure**

**2C**, red trace; 2) was decreased to almost the same current as the control response (**Figure 2C**,

blue trace; 3). Indeed, no GIRK-I remained after 20 times repetition of the combined command

pulse as revealed by the subtraction between trace 1 and trace 3 (**Figure 2D**, blue trace; 3−1).

Both the amplitudes of the inward and outward components of the NA-induced GIRK-I

promptly decreased to zero along with repetition of the combined command pulse (**Figure 2E**,

blue and red circles; **Figure 2F**). Thus, it is likely that NA-induced GIRK-I displays a

prominent rundown in a $Ca^{2+}$-dependent manner. Indeed, we further confirmed that these

rundowns were completely dependent on the number and application timing of the command

pulse in the presence or absence of TEA (see **Figure 2** and **Figures supplement 1 and 2**),

which would differentially increase $Ca^{2+}$ transients in amplitude and time course (**Figure**

**supplement 3**).

To investigate the effects of the rundown of GIRK-I on the spike-frequency adaptation,

spike trains were examined before and after the voltage-clamp experiment (**Figure 2A-F**). The

rundown of GIRK-I invariably resulted in an abolishment of spike-frequency adaptation

(**Figure supplement 4**), similar to the effects of atipamezole on spike-frequency adaptation

(**Figure 1**). Thus, $Ca^{2+}$ increases can cause the rundown of NA-induced GIRK-I and almost

abolished the spike-frequency adaptation, leading to an increase in the excitability of LC

neurons. These observations suggest that $[Ca^{2+}]_i$ increases cause  rundown of NA-induced

GIRK-I to suppress autoinhibition, leading to excitability increases that cause further increases

in $[Ca^{2+}]_i$ to further cause a rundown of the GIRK-I.

The agonist-dependency of the rundown of NA-induced GIRK-I was also confirmed

with α2A-AR blocker and GIRK blocker; α2-AR blockade but not GIRK blockade prevented

the rundown of NA-induced GIRK-I caused by excitability increases (see **Figures supplement**

**5** and **6**).

**β-Arrestin is involved in the rundown of NA-induced GIRK-I**

We further examined whether the internalization of α2-AR is involved in the rundown of NA-

induced GIRK-I or not. As β-arrestin/AP2 complex is known to be involved in the agonist-

dependent internalization of GPCR (*DeGraff et al., 1999; Cottingham et al., 2011*), we

examined the effects of barbadin, which prevents AP2 from binding with β-arrestin by forming

barbadin/AP2 complex, on the rundown of NA-induced GIRK-I. With the patch pipettes filled

with barbadin-including internal solution, it was examined whether NA-induced GIRK-I

displays rundown in response to the same combined command pulse (**Figure 2H**, upper trace)

as that in the control experiment (**Figure 2B**, upper trace). The NA-induced GIRK-I (**Figure**

**2G-I**, red traces) was hardly decreased (**Figure 2H, I**, blue traces) after repeating the combined

pulse 20 times, in contrast to the control rundown (**Figure 2C, D**). Barbadin largely suppressed

the rundown of the both components of GIRK-I (**Figure 2J, K**). These results suggest that the

rundown of NA-induced GIRK-I is caused by the internalization of α2-AR-coupled GIRK

channels.

**Restraint stress alters α2A-AR expression in LC neurons**

We next investigated whether the expression of α2A-AR is altered in LC neurons by applying

restraint stress (RS) to mice. After mice were subjected to daily 12-h RS for 3 days followed

by non-stress condition for 2 days, the immunoreactivity for α2A-ARs (see **Figure supplement**

**7**) seen along the membrane of TH-positive LC neurons was apparently lower compared to the

control (**Figure 3A, B**). Consistent with this observation, Western blot analysis revealed

significantly lower protein levels of α2A-ARs in LC neurons obtained after 3-day RS compared

to the control (**Figure 3C**). Especially when α2A-ARs expression was examined in the

membrane fraction identified using $Na^+/K^+$-ATPase, the immunoreactivity for α2A-ARs

expression was apparently lower in LC neurons of RS mice than in those of the control mice

(**Figure 3D, E**; *, compare the rightmost lower panels). Consistent with this observation,

Western blot analysis revealed a significantly lower protein level of α2A-ARs in the membrane

fraction of LC neurons of RS mice compared to the control whereas there was no significant

difference in the cytosol between the control and RS mice (**Figure 3F, G**). Thus, the

internalization of α2A-ARs is not necessarily accompanied by increases in cytosolic levels of

α2A-ARs because the internalized receptors are either recycled or degraded (*Koenig and Ikeda,*

*1996; Takei et al., 1996*).

**RS reduces NA-induced GIRK-I**

As 3-day RS caused the internalization or downregulation of α2A-ARs, we next examined if

the daily 12-h RS for up to 5 days can differentially cause reductions of NA-induced GIRK-I

in LC neurons. NA-induced GIRK-I decreased with an increase in the RS period (**Figure 4A-**

**E**).  The amplitudes of inward and outward components of GIRK-I decreased with the increase

in the period of RS (**Figure 4F, G**), in a way that is described by a saturation function (red

interrupted lines).  The amplitudes of inward and outward components of GIRK-I in 1-day

mice were significantly smaller than those of the control mice, and those in 3-day RS mice

were significantly smaller than those in 1-day RS mice while no significant difference in those

between 3-day and 5-day RS mice (**Figure 4F, G**).  Thus, the downregulation of GIRK-I was

increased as RS period is increased, but in a saturation manner.

Quantitative PCR analysis revealed that LC neurons expressed much higher mRNA

level of α2A-AR than that of α2C-AR (**Figure 4H**) and also expressed higher mRNA level of

GIRK1 compared to GIRK2 while GIRK3 was hardly expressed (**Figure 4I**).  Consistent with

the significant decreases in NA-induced GIRK-I in the mice subjected to 3-day RS, mRNA

levels of α2A-AR, GIRK1 and GIRK2 in mice subjected to 3-day RS were significantly smaller

than those of the control mice (**Figure 4J-L**).     Thus, RS downregulated the

expression/production of α2A-AR-coupled GIRK1 prominently in LC neurons.

**Histological and biochemical changes in LC neurons after 5-day RS**

Next, immunohistochemical experiments demonstrated that 5-day RS appeared to slightly

increase the expressions of TH and MAO-A in LC neurons (**Figure 5A**).  Furthermore, AEP

expression also appeared to increase in cytosol after 5-day RS compared to the control (**Figure

5B**).  Because TH catalyzes the rate limiting step in the synthesis of catecholamines, if the RS

enhances TH activity, free NA concentration in LC neurons would be directly and greatly

increased.  Therefore, we first carefully examined whether TH is really upregulated in RS mice

or not by performing Western blot analysis.  No significant increase in the protein level of TH

was observed in 5-day RS mice compared to the control (**Figure 5C,** upper panel; **Figure 5D**).

This result clearly indicates that TH activity was not significantly upregulated in 5-day RS

mice, in spite of the TH immunohistochemical experiment.  We further examined whether

protein levels of MAO-A and AEP are increased or not by 5-day RS.  Western blot analysis

revealed significant increases in the protein levels of MAO-A (**Figure 5C**, lower panel; **Figure

5E**) and of pro-AEP and active AEP (**Figure 5F-H**) in LC neurons after 5-day RS compared

to the control.  Western blot analysis also revealed a significant increase in the expression of

tau and tau N368 fragment after 5-day RS (**Figure 5I-K**).  Thus, an increased activity of active

AEP caused a production of tau N368 fragments as observed at 37 and 35 kDa (**Figure 5I**),

which is consistent with a previous report (*Zhang et al., 2014*).  Relative protein levels of the

tau N368 fragments (**Figure 5K**) were the mean values obtained from the two band intensities,

which may reflect the two N-terminal fragments (upper and lower bands) generated by AEP

cleavage at Asn-368 of the 4 and 3 repeat-tau isoforms, respectively (*Goedert et al., 1989*).

Finally, Y-maze test revealed that 5-day RS decreased spatial memory (**Figure supplement 8**).

## Discussion

We have found that LC neurons display spike-frequency adaptation which is caused by the

autoinhibition mediated by α2A-AR-coupled GIRK-I following respective spikes (**Figure 1**)

and that α2A-AR-coupled GIRK-I showed $Ca^{2+}$-dependent rundown due to the internalization

of α2A-AR (**Figure 2**, **Figure supplement 5A-D** and **Figure supplement 6E-H**), leading to

the abolishment of spike-frequency adaptation and the excitability increase (**Figure**

**supplement 4**). Immunohistochemical examination and Western blot analysis revealed that

α2A-AR expression in cell membrane was decreased following RS (**Figures 3 and 4J**).

Consistently, NA-induced GIRK-I was decreased in LC neurons with an increase in the period

of RS (**Figure 4A-G**). Following chronic RS, protein levels of MAO-A, pro-/active-AEP, and

tau and tau N368 fragment were significantly increased (**Figure 5A-C, E-K**).

Based on these results, we propose a novel mechanism for how chronic RS impairs

autoinhibition and increases cytosolic free NA to be metabolized by enhanced $Ca^{2+}$-dependent

MAO-A activity, to consequently increase AEP activity and produce tau N368 protein (**Figure**

**6**). In non-stress mice, autocrine released NA is slowly taken up by NAT after binding to α2-

ARs to cause autoinhibition (**Figure 6A**), while in chronic RS mice, autocrine-released NA

may be directly and rapidly taken up by NAT due to the absence (internalization) of α2-ARs

(**Figure 6B**). Such increased rate of re-uptake of NA by NAT would increase free NA

concentration by facilitating the leakage of NA from cytoplasmic vesicles in the cell body,

which would be metabolized by upregulated MAO-A activity to DOPEGAL, as revealed by

the increased activity of AEP in chronic RS mice (**Figure 6B**).

**Spike frequency adaptation**

LC neurons display spike-frequency adaptation. The mechanism for the spike-frequency

adaptation may be different among neurons while M currents are usually responsible for the

adaptation in hippocampal pyramidal neurons (*Aiken et al., 1995*). In LC neurons, the spike-

frequency adaptation was mediated by α2A-AR-coupled GIRK-I as the spike-frequency

adaptation was abolished by atipamezole (**Figure 1D-G**) and by causing the rundown of GIRK-

I (**Figure supplement 4**). These results suggest that autocrine NA release following respective

spikes activated α2A-ARs expressed in the soma/dendrites of LC neurons, and subsequently

activated α2A-AR-coupled GIRK-I to cause autoinhibition as spike-frequency adaptation.

Thus, spike-frequency adaptation clearly reflects NA autocrine.

**Rundown of α2A-AR-coupled GIRK-I**

α2A-AR-coupled GIRK-I displayed $Ca^{2+}$-dependent rundown in the presence of agonist. The

degree and time course of rundown of α2A-AR-coupled GIRK-I were dependent on the number

and timing of positive pulses applied under voltage-clamp condition to cause $Ca^{2+}$ currents

(**Figure 2A-F** and **Figures supplement 1E-J, 2, 3**). Given the $Ca^{2+}$-dependency of the

rundown of autoinhibition, the excitability of LC neurons would increase as LC neurons were

excited more such as under RS condition, resulting in a vicious cycle between excitability

increase accompanied by $[Ca^{2+}]_i$ increase and rundown of autoinhibition/$Ca^{2+}$-dependent

internalization of α2A-ARs.

**Internalization of α2A-ARs**

Activation of GPCR by agonist binding usually causes the formation of GPCR/β-arrestin

complexes. Subsequently, the GPCR/β-arrestin complex is assembled through the interaction

between β-arrestin and AP2 in clathrin-coated pits (CCPs), which is formed by the association

of AP2 with phosphatidylinositol 4,5-bisphosphate (PIP2) at the plasma membrane, leading to

endocytosis of GPCRs (*Pierce and Lefkowitz, 2001*). PIP2 is locally produced from the

membrane phosphatidylinositol (PI) by PI4/PIP5 kinases in a $Ca^{2+}$-dependent manner

(*Sarmento et al., 2014*). Barbadin prevented the stable formation of sufficient GPCR/β-

arrestin/AP2 complexes in CCPs as barbadin prevents AP2 from binding with β-arrestin by

forming barbadin/AP2 complex, thereby hampering GPCR internalization (*Beautrait et al.,*

*2017*). In the present study, intracellular barbadin effectively suppressed the rundown of

GIRK-I (**Figure 2G-K**), indicating that the rundown of GIRK-I was caused by the

internalization of α2A-ARs in LC neurons.

**Chronic RS increases the activities of MAO-A and pro-/active-AEP in the cell body**

Chronic cold stress reduced α2-AR mediated inhibition of LC neurons (*Jedema et al., 2008*),

and chronic social stress downregulated α2-ARs in the LC (*Flugge, 1996*). Our present results

were consistent with those previous findings: Following RS, α2A-ARs expression was

decreased due to the internalization of α2A-ARs (**Figure 3**), together with the decreases in NA-
induced GIRK-I (**Figure 4A-G**) which mediates the autoinhibition.  As long as RS persists, the
impairment of autoinhibition leads to the persistent overexcitation in LC neurons. This would
further increase NA autocrine, because respective spikes were accompanied by NA autocrine
as reflected in spike-frequency adaptation or autoinhibition.

As extracellular NA is mostly taken up into cytosol by NAT (*Torres et al., 2003*), it is
likely that under chronic RS the increased autocrine-released NA is directly and more rapidly
taken up by NAT due to the absence of α2A-ARs (**Figure 6B**). Such an increase in the rate of
NA uptake by NAT is similar to the effects of uptake inhibitor of transmitter ligand on the
activity of receptors. After re-uptake of NA by NAT, a large portion of NA in the cytosol is
restored in cytoplasmic vesicles by vMAT2 while only a small portion of NA can escape from
vMAT2 to be metabolized by MAO-A (*Eisenhofer et al., 1992*), because vMAT2 has a much
higher affinity for NA than MAO-A (*Bloom, 2006; Costa, 2012*).  However, there is an
additional component of NA to be metabolized by MAO-A, as NA can leak from cytoplasmic
vesicles into cytosol (*Eisenhofer et al., 2004*).  When re-uptake of NA by NAT is facilitated
due to the internalization of α2-ARs in response to NA autocrine, active NA storage into
cytoplasmic vesicles by vMAT2 would increase.  Then, the rate of NA leakage from
cytoplasmic vesicles would also increase because passive NA leakage from cytoplasmic
vesicles and active NA storage into cytoplasmic vesicles by vMAT2 are in a dynamic
equilibrium in cytoplasmic vesicles (*Eisenhofer et al., 2004*).  Subsequently, such increases in
the rate of NA leakage would result in increases in a MAO-A metabolite, DOPEGAL
(*Goldstein, 2021*) (**Figure 6B**).  Indeed, it has previously been demonstrated that a partial
inhibition of NAT in human subjects reduced the original level of a derivative of DOPEGAL
(3,4-dihydroxyphenylglycol; DHPG) in CSF by more than 50% (*Kielbasa et al., 2015*).  Thus,
free NA to be metabolized by MAO-A would be increased when autocrine-released NA is
taken up by NAT.  Therefore, under the chronic RS condition, more NA leaked from
cytoplasmic vesicles is likely to be metabolized by MAO-A, which activity is enhanced $Ca^{2+}$-
dependently and by chronic RS (*Cao et al., 2007*) (**Figures 5C, E and 6B**), to DOPEGAL
compared to the control condition, as revealed by the increased protein levels of pro-AEP and
active-AEP (**Figure 5B, F-H**).  Thus, re-uptaken NA by NAT may largely be degraded to
DOPEGAL by the increased activity of MAO-A.  Subsequently, DOPEGAL converts pro-AEP
into active-AEP to produce tau N368 protein (**Figure 5I-K**).

It has never been addressed how DOPEGAL production was caused in the cell body

of LC neurons while the DOPEGAL production in the cell bodies of LC neurons is well

established in AD patients (*Burke et al., 1999; Kang et al., 2020*) and in transgenic AD model

mice (*Kang et al., 2020*).  We demonstrated for the first time that 5-day RS can impair the

autoinhibition by causing the internalization of α2A-AR and consequently induces active-AEP

to produce tau N368 fragment.  Indeed, spatial memory impairment was detected in 5-day RS

mice (**Figure supplement 8**).  It is likely that, in addition to NAT inhibitor, the suppression of

the internalization of α2A-AR has a strong translational potential for developing drugs that

prevent the degeneration of LC.

**Limitations of this study**

One may argue against the usage of 3-week-old juvenile mice to study the neurodegenerative

disorder which is normally found in aged ones.  However, a stringent autoinhibition together

with potent NAT activity were previously demonstrated in adult rats using intracellular

recordings (*Williams et al., 1985*). Thus, regardless of the age of mice, LC neurons showed the

autoinhibition that is mediated by the activity of α2A-AR-coupled GIRK channels. We also

confirmed that spike-frequency adaptation reflecting autoinhibition can be induced in adult

mice as prominently as in juvenile mice regardless of sexes (**Figure supplement 9**), although

we did not investigate the sex differences in RS effects (*Leistner and Menke, 2020*). The

downregulation of autoinhibition by the internalization of α2A-AR is agonist- and $Ca^{2+}$-

dependent regardless of the age of mice.  However, it is not known if there is any difference in

the susceptibility of α2A-AR internalization to RS between juvenile and aged mice.  Thus, a

further study may be necessary to examine if autoinhibition in LC neurons is more easily

downregulated in aged mice compared to juvenile ones.

It has also been reported that re-uptake of NA by NAT in LC neurons may decrease

with age (*Sanders et al., 2005; Mitchell et al., 2017*).  However, NAT expression in LC in 3-

week-old juvenile mice was only slightly larger (by 4%) than in adult mice although the

difference was significant (*Mitchell et al., 2017*).  Potent NAT activity stringently affecting

autoinhibition was reported in adult rats (*Williams et al., 1985*).  Furthermore, partial inhibition

of NAT in human subjects largely reduced the original level of a derivative of DOPEGAL in

CSF as described above.  Thus, based on the present study, it is likely that increased re-uptake

of NA by NAT following excessive NA autocrine in the absence of α2A-AR, induced in

response to chronic RS, resulted in the increase in active-AEP and subsequent production of

tau N368.  However, a further experiment to directly assess NAT activity would be necessary

to firmly justify our hypothesis.

**Conclusion**

Our study unveiled the cellular mechanism for the degeneration of LC neurons under chronic

RS as follows: Chronic RS impaired the autoinhibition in LC neurons by inducing $Ca^{2+}$- and

agonist-dependent internalization of soma/dendritic α2A-AR coupled with GIRK channels.

This subsequently caused overexcitation/$[Ca^{2+}]_i$ increases and increased $Ca^{2+}$-dependent

MAO-A activity and active AEP activity, leading to a production of tau N368 fragment

together with an impairment of spatial memory.  These results clearly indicate that chronic RS

increases cytosolic free NA to be metabolized by MAO-A, presumably by facilitating the re-

uptake of autocrine-released NA due to the absence/internalization of α2A-AR.  Thus, in

addition to NAT inhibitor, the suppression of α2A-AR internalization may have a strong

translational potential for developing drugs that prevent the degeneration of LC.

**Methods**

**Restraint stress**

Three-weeks-old male C57BL/6 mice (JAPAN HAMAMATSU SLC, Hamamatsu, Japan)

were used and housed at $24 \pm 2°C$.  Food and water were available ad libitum.  The mice in the

stressed groups were restrained in a modified 50 ml clear polystyrene conical centrifuge with

multiple air holes for ventilation.  The mice were subjected to 12-h restraint stress (RS; 20:00

p.m. to 8:00 a.m. of the next day) for 1, 2, 3 and 5 days.  In the previous study, the duration of

RS widely ranged from 0.5-h (*Gray et al., 2010*) to 24-h (*Chu et al., 2016*) while the RS with

the duration less than 12-h has been applied daily for a period (6-h for 2 weeks (*Sakaguchi and*

*Nakamura, 1990*); 8-h for up to 16 weeks (*Sugama et al., 2016*); 12-h for 2 days (*Zhang et al.,*

*2008*)).  As the prolonged stress is known to be a potential risk factor for AD (*Bisht et al.,*

*2018*), the protocol of daily 12-h RS is relevant for the study of functional and cytological

disorders of LC neurons.  Once the restraint ended, the mice were returned to their home cages

with access to food and water ad libitum.  The control group remained in their home cages

without the restraint procedure.

**Slice preparation and electrophysiological recordings**

Slices were prepared from 3-weeks-old male C57BL/6 mice to investigate the effects of RS for varying periods on GIRK-I, and from C57BL/6 mice of either sex at 16–23 postnatal days (PD) to investigate the effects of $[Ca^{2+}]_i$ increases on GIRK-I, similar to our previous study (*Toyoda et al., 2022*). Using an Axopatch 200B (MDS Analytical Technologies, Sunnyvale, CA), whole-cell recordings were made from LC neurons that were viewed under Nomarski optics (BX51WI; Olympus, Tokyo, Japan). The internal solution of the patch pipettes had the following ionic composition (in mM): 123 K-gluconate, 18 KCl, 10 NaCl, 2 $MgCl_2$, 2 ATP-$Na_2$, 0.3 GTP-$Na_3$, 10 HEPES, and 0.2 EGTA; pH 7.3. The membrane potential values given in the text were corrected for the junction potential (10 mV) between the internal solution for the whole-cell recording (negative) and the standard extracellular solution. The pipette resistances were 4–6 MΩ. The series resistance was < 10 MΩ. All recordings were made at room temperature (RT). Records of currents and voltages were low-pass filtered at 5 kHz (3-pole Bessel filter), digitized at a sampling rate of 10 kHz (Digidata 1322A, MDS Analytical Technologies) and stored on a computer hard disk. Spike-frequency adaptation was quantitatively estimated by fitting with a saturation curve defined by Monod equation (y = (***a*** * x) / (***b*** + x)) with a saturation level (***a***) and a half saturation constant (***b***). The amplitudes of inward and outward components of NA-induced GIRK-I were measured at –130 mV and –60 mV.

**Procedure for immunohistochemical examination**

C57BL6/J male mice (4-5 weeks) were perfused transcardially with 5 ml of PBS, followed by 10 ml of a freshly prepared 4% paraformaldehyde in PBS. The whole brain including LC (–9 to –10 mm from bregma) was dissected out and immersed in 4% paraformaldehyde solution for 4 h at 4°C fallowed by cryoprotection with 30% (w/w) sucrose in PBS.

Brainstem including LC regions was cut into 8-20 μm-thickness coronal sections on a freezing microtome (Leica Microsystems, Wetzlar, Germany). To examine the distribution of α2A-AR in LC neurons, mouse anti-TH (1:500; sc-25269, Santa Cruz Biotechnology, Santa Cruz, CA, USA), rabbit anti-sodium potassium ATPase (1:1000; ab76020, Abcam, Cambridge, MA, USA) and goat anti-α2A-AR (1:200; ab45871, Abcam) or rabbit anti-α2A-AR (1:500; RA14110, Neuromics, Northfield, MN, USA) were used as primary antibodies, while cy5 donkey anti-mouse IgG (1:500; cy-2500, Vector Laboratories, Burlingame, CA, USA), FITC

donkey anti-rabbit IgG (1:500; 711-095-152, Jackson ImmunoResearch, West Grove, PA, USA) and Cy3 donkey anti-goat IgG (1:500; 706-165-147, Jackson ImmunoResearch) were used as secondary antibodies. We used the affinity purified IgG antibody for α2A-AR immunohistochemistry, in which antibody specificity was ensured as examined previously (*Stone et al., 1998; Tan et al., 2009*). To examine the expression in LC neurons, mouse anti-TH (1:500) and rabbit anti-MAO-A (1:2000; ab126751, Abcam) were used as primary antibodies, while FITC donkey anti-mouse IgG (1:500; 715-545-150; Jackson ImmunoResearch) for TH and Cy3 donkey anti-rabbit IgG (1:2000; 711-165-152; Jackson ImmunoResearch) were used as secondary antibodies. To examine the distribution of legumain in LC neurons, mouse anti-legumain (1:200; sc-133234, Santa Cruz) and rabbit anti-DBH (1:500; 22806, ImmunoStar, Hudson, WI, USA) were used as primary antibodies, while FITC donkey anti-mouse IgG (1:500; 715-545-150, Jackson ImmunoResearch) and Cy3 donkey anti-rabbit IgG (1:500; 711-165-152, Jackson ImmunoResearch) were used as secondary antibodies. The slices were coverslipped with mounting media (H1000, Vector Laboratories), and sections were visualized with a confocal microscope (LSM980, Carl Zeiss, Oberkochen, Germany) at 10x and 63x magnification.

**Western blot study**

Brain slices from the control or RS-treated mice were sectioned with vibratome (VT1000P, (Leica Microsystems) to obtain 500 μm-thickness coronal sections containing LC region (bregma –5.34 mm to –5.80 mm). LC dissection was done via micro punch using 1.5 mm biopsy punch (BP-15F, Kai medical, Gifu, Japan) bilaterally under light microscope. Dissected LC were quickly frozen in liquid nitrogen and stored at −80°C until processed. Because LC region was too small to collect sufficient tissue from one mouse for the analysis, the punched LC tissues were collected from two to three mice to obtain a protein lysate sample for each western blotting analysis. LC tissues were homogenized by bead homogenizer in T-PER™ Tissue Protein Extraction Reagent (78510, Thermo Fisher) containing protease and phosphatase inhibitor cocktails (MilliporeSigma, St. Louis, MO, USA). The membrane and cytosol proteins from LC lysate were isolated using Mem-PER Plus Kit (78840, Thermo Fisher) in accordance with the manufacture's protocol. The protein concentration of the lysate was determined using a Lowry protein assay (Bio-Rad, Hercules, CA, USA). Protein samples (20 μg) were heated to 95°C for 5 min, separated by 12% sodium dodecyl sulfate-polyacrylamide gel electrophoresis, and transferred to a polyvinylidene fluoride membrane

using a Transblot SD apparatus (Bio-Rad). After the blots had been washed with tris-buffered saline containing Tween-20 (TBST, 10 mM Tris-HCl, 150 mM NaCl, 0.05% Tween-20), the membranes were blocked with 2% BSA and 4% skim milk/TBST at RT for 1 h and incubated at 4°C overnight with primary antibodies, followed by an anti-rabbit IgG horseradish peroxidase (HRP)-conjugated secondary antibody (1:5,000, 7074; Cell Signaling Technology, Danvers, MA, USA) at RT for 1 h. The western blot membrane was stripped the primary and secondary antibodies and reprobed with the anti-β-actin antibody followed by the anti-rabbit IgG HRP-conjugated secondary antibody. The following primary antibodies were used for the blots: rabbit anti-α2A-AR (1:500; AAR020, alomone), rabbit anti-GAPDH (1:1000; LF-PA0018, Abfrontier, Seoul, Korea), mouse anti-beta actin (1:2000, A5441, MilliporeSigma) and rabbit anti-sodium potassium ATPase (1:1000; ab76020, Abchem) (**Figure 3C, F and G**); rabbit anti-TH (1:1600; GTX113016, GeneTex, Irvine, CA, USA), rabbit anti-MAO-A (1:6400; ab126751, Abcam), mouse anti-legumain (AEP) (1:400; sc-133234, Santa Cruz), mouse anti-TAU-5 (1:1500, Santa Cruz Biotechnology, SC-58860) and mouse anti-beta actin (1:12800, 66009-1-Ig, Proteintec, Rosemont, IL, USA) (**Figure 5C, F and I**); rabbit anti-tau AEP-cleaved (N368) (1:5000, ABN1703, MilliporeSigma) and mouse anti-beta actin (1:1000, A5441, MilliporeSigma) (**Figure 5I, K**). Western blot analysis was performed by measuring band intensity with ImageJ 1.50i software (National Institutes of Health, Bethesda, MD, USA).

**Quantitative real-time PCR**

cDNA was synthesized with Super Script II Reverse Transcriptase (18064, Invitrogen, Carlsbad, CA, USA) with 1 mg of LC neuron RNA. Using a 7300 Gene Amp PCR system (Applied Biosystems, Carlsbad), quantitative real-time PCR was run in Power SYBR Green PCR master mix (4367656; Life Technologies, Carlsbad) under the following thermal cycling conditions; an initial step of 94°C/5 min, followed by 40 cycles of a series of steps of 94°C/45 s, 61°C/45 s, 72°C/45 s and ended with a series of steps of 95°C/60 s, 60°C/1 min, 95°C/15 s, 60°C/15 s. The following primers were used: α2A (forward GGAATCATGGCTGTGGAGAT, reverse CAGAAATCCCTTCCCTGTCA); α2C (forward TCATCGTTTTCACCGTGGTA, reverse GCTCATTGGCCAGAGAAAAG); GIRK1 (forward ACCCTGGTGGATCTCAAGTG, reverse GGCCACACAGGGAGTGTAGT); GRIK2 (forward CAACCTCAACGGGTTTGTCT, reverse ATCCTACCATGAAGGCGTTG); GIRK3 (forward CGTCTCACCTCTCGTCATCA, reverse

CATCCACCAGGTACGAGCTT); GAPDH (forward CCAGAACATCATCCCTGCAT,
reverse GCATCGAAGGTGGAAGAGTG).

**Behavioral test**

The control mice as non-stress group and mice exposed to RS for 1 day, 3 days and 5 days as
1-day, 3-day and 5-day RS groups were tested on the Y maze apparatus, consisted of three
identical arms in a "Y" shape made of white acryl (35 cm in length, 5 cm in width, and 20 cm
in height) to assess working spatial memory. All tests were performed between 9 AM and
noon. Investigators were blinded to the group when performing experiments. Each mouse was
placed at the extremity of the "start arm" and left for a training session of 15 minutes, with one
of the 2 other arms closed with a white acrylic board (the closed arm was randomly chosen for
each mouse). Then, the mouse was brought back to its cage for 5 minutes, the maze was
cleaned, the closed arm was opened, and the mouse placed at the extremity of the start arm for
a 5-minute test session, under video tracking. The time spent and distance traveled in each arm
(start, familiar, and novel arms) were measured and analyzed with Smart 3.0 video tracking
software (Panlab, Harvard Apparatus, Holliston, MA, USA). Given their natural curiosity,
mice are expected to spend more time exploring the new arm.

**Drug application**

Atipamezole as a blocker of $\alpha$2-AR and tertiapin-Q as a GIRK blocker were bath-applied at 5-
10 μM and 200 nM, respectively. Barbadin as an arrestin/AP2 blocker was added to the internal
solution at a final concentration of 100 μM (*Beautrait et al., 2017*). All chemicals except
tertiapin-Q were purchased from MilliporeSigma unless specified otherwise. Tertiapin-Q was
purchased from Tocris Bioscience (Bristol, UK).

**Statistical Analysis**

Statistical analysis was performed using STATISTICA10J (StatSoft Japan, Tokyo, Japan).
Numerical data were expressed as the mean ± SD. The statistical significance was assessed
using Student's *t*-test, one-way ANOVA followed by post hoc Fisher's PLSD and Wilk's
lambda. The significance level was set at 5% or less ($p < 0.05$). Statistical results are given as

a $p$ value, unless $p < 0.001$.  All the statistical data contained in this study are presented in
Table S1.

## Acknowledgements

This research was funded by Japan Society for the Promotion of Science (26290006 and
21K06441 to Y.K.: 20K06926 and 23K06346  to H.T.).  Y.K. was supported by Brain Pool
Program (NRF) funded by the Korean government (Ministry of Science and ICT).  H.T. was
supported by Naito Foundation.  This research was partly supported by the Basic Science
Research Program through the National Research Foundation of Korea grants (NRF-
2018R1A5A2024418 and NRF-2021R1A2C3003334 to S.B.O) funded by the Korean
government (Ministry of Science and ICT).

## Author Contributions

561 YK, HT, and SBO conceived and designed the study.  HT, DK, BGK, and TS performed
562 experiments.  HT, DK, BGK, TS, TK, and YK analyzed data.  YK, HT and SBO wrote the
563 manuscript.

## Ethics

566 This study was conducted following the Helsinki Declaration, and carried out according to the
567 ARRIVE guidelines.  All experimental procedures were approved and performed in accordance
568 with the relevant guidelines of the Ethical Guidelines Committee at Osaka University, Seoul
569 National University and Kyushu University.

## Data Availability

572 All data and/or analyses generated during the current study are available from the
573 corresponding author upon request.

## Competing Interests

576 The authors declare no competing financial or non-financial interests.

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 **Figure legends**

 **Figure 1. Atipamezole abolishes spike-frequency adaptation and blocks NA-induced**

 **GIRK-I in LC neurons.**

 (**A, B**) Sample traces of spike trains induced in an LC neuron by injection of current pulses at

 75 and 100 pA. Note the spike-frequency adaptation and pulse-afterhyperpolarization (pulse-

 AHP).

 (**C**) Plotting of the cumulative spike numbers vs the elapsed time from the onset of the current

 pulses every 50 ms at 75 pA (blue circles) or 100 pA (red circles) ($n = 7$). Two-way RM

 ANOVA, *$p < 0.001$. The saturation level (***a***) and the half saturation constant (***b***) were

 measured by fitting the saturation function defined as y = (***a*** * x) / (***b*** + x) to the data points.

 The values of ***a*** at 75 and 100 pA current pulses were $44.6 \pm 16.2$ and $52.5 \pm 12.3$, respectively

 (paired *t*-test, $p = 0.010$). The values of ***b*** at 75 and 100 pA current pulses were $2.1 \pm 0.7$ and

 $2.1 \pm 0.4$, respectively (paired *t*-test, $p = 0.823$).

 (**D, E**) Sample traces of spike trains induced in an LC neuron evoked by current pulses at 100

 pA under control condition and after application of 10 μM atipamezole. Note that atipamezole

 suppresses spike-frequency adaptation and pulse-AHP (blue and red stars).

 (**F**) Plotting of the cumulative spike numbers vs the elapsed time during the current pulses

 every 50 ms before and after application of atipamezole (blue and red circles, respectively) (*n*

 = 10). Two-way RM ANOVA, *$p < 0.001$.

 (**G**) Plotting of ***a*** against ***b***, which were measured by fitting the saturation function to the data

 points in **F**. Both the values of ***a*** and ***b*** obtained by curve fitting to the data points after

 application of atipamezole (open and filled red circles) were significantly shifted to larger

 values than those obtained under control condition (open and filled blue circles) (paired *t*-test,

 ***a*** and ***b***, [‡]$p = 0.003$ and [‡]$p = 0.002$, respectively), and there was a significant difference in the

 relationship between ***a*** and ***b*** (Wilk's lambda, [§]$p = 0.002$) ($n = 10$).

 (**H**) Top, Voltage command pulse. Lower panel, Superimposed current traces obtained under

 control condition (1), after application of 100 μM NA (2), and after application of 10 μM

 atipamezole (Atip) in addition to NA (3). The amplitudes of inward and outward components

 in NA-induced GIRK-I were $-70 \pm 22$ and $36 \pm 11$ pA, respectively (paired *t*-test, [‡]$p = 0.005$,

 $n = 6$).

 (**I**) A graph showing the effects of atipamezole on amplitudes of the inward and outward

 components of NA-induced GIRK-I at $-130$ mV and $-60$ mV, obtained by subtraction of

 currents recorded under control condition from those recorded after application of NA (2–1)

and from those recorded after application of NA and atipamezole (3–1). Inward component,

paired *t*-test, ‡$p < 0.001$; outward component, paired *t*-test, ‡$p < 0.001$ ($n = 6$).

**Figure 2. Ca²⁺-dependent rundown of NA-induced GIRK-I and its suppression by**

**barbadin.**

(**A**) Upper panel, Ramp command pulse. Lower panel, Superimposed current traces obtained

under control condition (1) and after application of 100 μM NA for 5 min (2).

(**B**) Upper panel, A combined command pulse applied every minute; five trains of 20 positive

pulses (5 ms duration to 0 mV at 20 Hz) at an interval of 2 s in the presence of extracellular 30

mM TEA and intracellular 0.2 mM EGTA, which were followed by the ramp pulse after an

interval of 25 s. Lower panel, representative Ca²⁺ currents in response to positive pulses in the

presence of NA and 30 mM TEA.

(**C**) Upper panel, Ramp command pulse. Lower panel, Superimposed current traces obtained

after application of NA for 5 min (2) and in response to application of the 20th combined pulse

in the presence of NA and TEA (3).

(**D**) The I-V relationship of NA-induced GIRK-I obtained by subtraction of currents recorded

under control condition from that recorded after application of NA for 5 min (red trace, 2–1)

and that obtained by subtraction of the control current from that recorded in response to

application of the 20th combined pulse in the presence of NA and TEA (blue trace, 3–1).

(**E**) Plotting of amplitudes of inward components at –130 mV (blue circles) and outward

components at –60 mV (red circles) against time.

(**F**) Plotting of normalized amplitudes of inward components at –130 mV and outward

components at –60 mV before and 20 min after application of positive pulse trains in the

presence of NA and TEA. The amplitudes of inward components at –130 mV and those of

outward components at –60 mV were normalized by those recorded in the presence of NA

before applying positive pulse trains. Inward component, paired *t*-test, ‡$p < 0.001$; outward

component, paired *t*-test, ‡$p < 0.001$ ($n = 8$).

(**G**) Upper panel, Ramp command pulse. Lower panel, Superimposed current traces obtained

under control condition (1) and after application of 100 μM NA for 5 min (2) in LC neurons

dialyzed with 100 μM barbadin (β-arrestin/AP2 blocker).

(**H**) Upper panel, A combined command pulse applied every minute; five trains of 20 positive

pulses (5 ms duration to 0 mV at 20 Hz) at an inter-train interval of 2 s in the presence of

extracellular 30 mM TEA and intracellular 0.2 mM EGTA, which were followed by the ramp

pulse after an interval of 25 s.  Lower panel, Superimposed current traces obtained after

application of NA for 5 min (2) and in response to application of the 20th combined pulse in

the presence of NA and TEA (3) in LC neurons dialyzed with barbadin.

(**I**) The I-V relationship of NA-induced GIRK-I obtained by subtraction of currents recorded

under control condition from that recorded after application of NA for 5 min (red trace, 2–1)

and that obtained by subtraction of the control current from that recorded in response to

application of the 20th combined pulse in the presence of NA and TEA (blue trace, 3–1).

(**J**) Plotting of normalized amplitudes (left vertical axis) of inward components at –130 mV

(filled blue circles) and outward components at –60 mV (filled red circles) against time ($n = 8$).

Amplitudes of GIRK-I were normalized to the values obtained before applying pulse trains.

Right vertical axis refers to the original amplitudes.  Open blue and red circles represent the

normalized amplitudes of GIRK-I shown in **E**.

(**K**) Plotting of normalized amplitudes of inward components at –130 mV and outward

components at –60 mV before and 20 min after application of positive pulse trains in the

presence of NA and TEA ($n = 8$).  The amplitudes of inward components at –130 mV and those

of outward components at –60 mV were normalized to those recorded in the presence of NA

before applying positive pulse trains.  Open light pink circles represent results obtained under

the control condition in **F**.  Inward component, unpaired $t$-test, \*$p < 0.001$; outward component,

unpaired $t$-test, \*$p < 0.001$.

**Figure 3. RS reduces expression of α2A-ARs in LC Neurons.**

(**A, B**) Confocal images of LC neurons showing immunoreactivities for TH and α2A-ARs,

together with a merged one in non-stress and 3-day RS mice. Arrowheads indicate the

membrane regions of TH-positive neurons, along which α2A-ARs were differentially

expressed between the control and the RS mice. Asterisks indicate LC neurons.

(**C**) Western blotting analyses showing the expression of α2A-ARs in non-stress and 3-day RS

mice ($n = 6$ and 7, respectively).  Unpaired $t$-test, [†]$p < 0.001$.

(**D, E**) Confocal images of LC neurons in non-stress and RS mice.  Upper panels from left to

right showing the respective immunoreactivity for TH, α2A-ARs (with rabbit anti-α2A-AR)

and $Na^+$-$K^+$-pump and an enlarged image of the region enclosed with a rectangle (arrowheads)

in its immediate left panel.  Lower panels from left to right showing the merged image of TH

and α2A-ARs, that of TH, α2A-ARs and $Na^+$-$K^+$-pump, that of α2A-ARs and $Na^+$-$K^+$-pump

and an enlarged image of the region enclosed with a rectangle in its immediate left panel.

(**F, G**) Western blotting analyses showing expressions of α2A-ARs and $Na^+$-$K^+$-pump in

membrane fraction and those of α2A-ARs and β-actin in cytosol fraction in non-stress and 5-

day RS mice ($n = 3$ and 3 samples, respectively). Each sample represents the analysis result in

the LC tissues obtained from 2-3 mice. Membrane; unpaired $t$-test, $^†p = 0.016$. Cytosol;

unpaired $t$-test, $^†p = 0.083$.

**Figure 4. RS reduces NA-induced GIRK-I in LC neurons.**

(**A-E**) Representative traces of NA-induced GIRK-I obtained from LC neurons in non-stress

and 1-day, 2-day, 3-day, and 5-day RS mice.

(**F, G**) Amplitudes of inward and outward components of NA-induced GIRK-I obtained from

LC neurons in non-stress and 1-day, 2-day, 3-day, and 5-day RS mice ($n = 8, 7, 5, 6$ and 8,

respectively) decreased with the increase in the period of RS, in a way that can be described by

a saturation function (red interrupted lines). The saturation level ($\boldsymbol{a} + \boldsymbol{b}$) and the half saturation

constant ($\boldsymbol{c}$) were determined by fitting the saturation function, defined as y = $\boldsymbol{a}$ + ($\boldsymbol{b}$* x) / ($\boldsymbol{c}$ +

x), to the data points. The values of $\boldsymbol{a}, \boldsymbol{b}$ and $\boldsymbol{c}$ for the inward component of GIRK-I were −151.1,

108, and 0.9, respectively, and those for the outward component of GIRK-I were 81.1, –67,

and 1.6, respectively. Inward component: one-way ANOVA, $p < 0.001$, post hoc fisher's

PLSD, 1-day; $^#p < 0.001$ vs Non-stress and $^*p < 0.05$ vs 3-day and 5-day, 2-day; $^#p < 0.001$ vs

Non-stress, 3-day; $^#p < 0.001$ vs Non-stress, 5-day; $^#p < 0.001$ vs Non-stress. Outward

component: one-way ANOVA, $p < 0.001$, post hoc fisher's PLSD, 1-day; $p = 0.004$ vs Non-

stress and $^*p < 0.05$ vs 3-day and 5-day, 2-day; $p = 0.015$ vs Non-stress, 3-day; $p < 0.001$ vs

Non-stress, 5-day; $p < 0.001$ vs Non-stress.

(**H**) Relative expressions of α2A and α2C mRNAs, normalized to GAPDH in LC neurons ($n =$

5). Paired $t$-test, $^‡p = 0.014$.

(**I**) Relative expressions of GIRK1, GIRK2, and GIRK3 mRNAs, normalized to GAPDH in

LC neurons ($n = 5$). One-way RM ANOVA, $p < 0.001$, post hoc fisher's PLSD; $^*p = 0.008$

for GIRK1 vs GIRK2, $^*p < 0.001$ for GIRK1 vs GIRK3, $^*p = 0.008$ for GIRK2 vs GIRK3.

(**J, K**) Relative expressions of α2A and GIRK1 mRNAs, respectively, normalized to GAPDH

in LC neurons in non-stress mice ($n = 8$), and 3-day RS mice ($n = 6$). α2A-AR: unpaired $t$-test,

$^†p = 0.020$; GIRK1: unpaired $t$-test, $^†p = 0.013$.

(**L**) Normalized relative expressions of GIRK2 mRNA in LC neurons in non-stress mice ($n =$

8) and 3-day RS mice ($n = 6$), normalized to the ratio of the mean value of the relative

expressions of GIRK1 mRNA to that of GIRK2 mRNA in LC neurons in non-stress mice (**I**).

Unpaired *t*-test, $^{\dagger}p = 0.037$.

**Figure 5. Expressions of TH, MAO-A, DBH, AEP and tau proteins in LC neurons.**

(**A**) Confocal images of neurons in LC at a low magnification showing the immunoreactivities

for TH and MAO-A, together with a merged one.  RS appeared to cause slight increases in the

expressions of TH and MAO-A.  Brightness of background in sample image obtained from the

control was corrected to be equal to that obtained from the RS.

(**B**) Confocal images of LC neurons at a high magnification showing the immunoreactivities

for DBH and AEP, together with a merged one.  RS caused increases in the expression of AEP.

(**C**) Western blotting of TH and MAO-A in LC neurons obtained from non-stress and 5-day

RS mice.

(**D, E**) Western blot analysis revealing no significant change in TH expression ($n = 10$ samples)

but a significant increase in MAO-A expression ($n = 6$ samples) in 5-day RS group compared

to non-stress group.  Unpaired *t*-test, $^{\dagger}p = 0.029$.

(**F**) Western blotting of pro-AEP and active AEP in LC neurons obtained from non-stress and

5-day RS mice.

(**G, H**) Western blot analysis revealing significant increases in pro-AEP and active AEP ($n =$

6 and $n = 6$ samples, respectively) in 5-day RS group compared to non-stress group.  Unpaired

*t*-test, $^{\dagger}p = 0.034$ (**G**) and $^{\dagger}p = 0.047$ (**H**).

(**I**) Western blotting of tau and tau N368 fragment in LC neurons obtained from non-stress and

5-day RS mice.  Unpaired *t*-test, $^{\dagger}p = 0.025$.

(**J, K**) Western blot analyses revealing a significant increase in tau and tau N368 fragment in

LC neurons ($n = 8$, and $n = 6$, respectively) in 5-day RS group compared to non-stress group.

Each sample represents the analysis result in the LC tissues obtained from 2-3 mice.  Unpaired

*t*-test, $^{\dagger}p < 0.05$.

**Figure 6. Presumed cellular mechanisms for the degeneration of LC by stress.**

(**A, B**) Differential free concentrations of NA to be metabolized by MAO-A into DOPEGAL

between control and stress conditions: Under the control condition (**A**), NA in the cytosol is

mostly taken up into the cytoplasmic vesicles by vMAT2 (thick arrow 3), rather than being

directly metabolized by MAO-A into DOPEGAL (thin interrupted arrow).  Vesicular NA is

released from cell bodies as autocrine following $[Ca^{2+}]_i$ increases caused by action potentials,

and the released NA activates α2A-AR-coupled GIRK channels, causing autoinhibition. Subsequently, the autocrine released NA is slowly taken up into the cytosol of LC neurons by NAT after dissociation from α2A-ARs. Under the stress condition (**B**), an activation of CRF receptors in LC neurons by stress inhibits leak $K^+$ channels and increases firing activities in LC neurons, subsequently causing a larger $[Ca^{2+}]_i$ increase together with $Ca^{2+}$-induced $Ca^{2+}$ release (CICR). Impairment of autoinhibiton due to $Ca^{2+}$ dependent internalization of α2A-ARs-coupled GIRK channels leads to the persistent excitation in LC neurons, which enhances autocrine release of NA (thick arrow 1). Subsequently, the excessively autocrine released NA is taken up directly and rapidly by NAT into the cytosol without binding to α2A-ARs (thick arrow 2). Such a facilitation of re-uptake of NA by NAT would increase active NA storage into cytoplasmic vesicles by vMAT2 (thick arrow 3), while the rate of NA leakage from cytoplasmic vesicles would also increase (arrow 4) due to a dynamic equilibrium in cytoplasmic vesicles between active NA storage into cytoplasmic vesicles and passive NA leakage from cytoplasmic vesicles. Subsequently, such an increase in the rate of NA leakage would result in an increase in a MAO-A metabolite, DOPEGAL and AEP, leading to a production of cleaved tau N368 fragment and an impairment of learning/memory.

# Figure 1

# Figure 2

# Figure 3

# Figure 4

# Figure 5

# Figure 6

## A Control

Slow taken-up of NA after dissociation from α2A-AR

Activation of α2A-AR coupled GIRK (Autoinhibition)

NAT

α2A-AR

K+

GIRK

β-arrestin

Gβγ

Gα GTP

MAO-A

vMAT2

DOPEGAL

DA — DBH → NA

vMAT2

Autocrine release of NA

Inhibition

⊖ Ca2+ channel

⊕

Ca2+

## B Stress

Direct and rapid taken-up of NA

NAT

NA

α2A-AR

GIRK

Ca2+-dependent internalization

β-arrestin

Gα Gβγ

⊕

MAO-A

vMAT2

DA — DBH → NA

vMAT2

Excessive autocrine release of NA due to impaired autoinhibition

DOPEGAL

⊕

Dis-inhibition

⊕

AEP

ER

CICR

Ca2+ channel

Tau N368

Leak K+ channel

⊕

Ca2+

Impaired learning/memory

CRF

⊖

⊕

Stress