

Research Article: Confirmation | Neuronal Excitability

# Refining the Identity and Role of Kv4 Channels in Mouse Substantia Nigra Dopaminergic Neurons

https://doi.org/10.1523/ENEURO.0207-21.2021

Cite as: eNeuro 2021; 10.1523/ENEURO.0207-21.2021

Received: 10 May 2021 Accepted: 17 May 2021

This Early Release article has been peer-reviewed and accepted, but has not been through the composition and copyediting processes. The final version may differ slightly in style or formatting and will contain links to any extended data.

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- 1 REFINING THE IDENTITY AND ROLE OF KV4 CHANNELS IN MOUSE
- 2 SUBSTANTIA NIGRA DOPAMINERGIC NEURONS
- 3 **Abbreviated Title**: Kv4 channels in substantia nigra dopaminergic neurons
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- 17 M.A. and J-M.G. wrote the manuscript.
- 18 **Conflict of interest**: the authors declare no competing financial interests.
- 19 Acknowledgements: this work was supported by the European Research Council (Consolidator grant
- 20 616827 CanaloHmics to J-M.G., supporting A.H-H. and M.T.), the Fondation de France (grant 00076344
- 21 to J-M.G. and M.A., supporting A.H-H.), and the Agence Nationale pour la Recherche (ANR Logik ANR-
- 22 17-CE16-0022, supporting J.R.F.). We thank O. Toutendji for technical assistance.

## ABSTRACT

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Substantia nigra pars compacta (SNc) dopaminergic (DA) neurons display a peculiar electrical phenotype characterized in vitro by a spontaneous tonic regular activity (pacemaking activity), a broad action potential and a biphasic post-inhibitory response. The transient A-type current (I<sub>A</sub>) is known to play a crucial role in this electrical phenotype, and so far this current was considered to be carried exclusively by Kv4.3 potassium channels. Using Kv4.3<sup>-/-</sup> transgenic mice, we demonstrate that the constitutive loss of this channel is associated with increased exploratory behavior and impaired motor learning at the behavioral level. Consistently it is also associated with a lack of compensatory changes in other ion currents at the cellular level. Using antigen retrieval immunohistochemistry, we then demonstrate that Kv4.2 potassium channels are also expressed in SNc DA neurons, even though their contribution to I<sub>A</sub> appears significant only in a minority of neurons (~5-10%). Using correlative analysis on recorded electrophysiological parameters and multi-compartment modeling, we then demonstrate that, rather than its conductance level, IA gating kinetics (inactivation time constant) appear as the main biophysical property defining post-inhibitory rebound delay and pacemaking frequency. Moreover, we show that the hyperpolarization-activated current (I<sub>H</sub>) has an opposing and complementary influence on the same firing features.

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## SIGNIFICANCE STATEMENT

Substantia nigra pars compacta (SNc) dopaminergic (DA) neurons are characterized by pacemaking activity, a broad action potential and biphasic post-inhibitory response. The A-type transient potassium current (I<sub>A</sub>) plays a central role in this electrical phenotype. While it was thought so far that Kv4.3 ion channels were fully responsible for I<sub>A</sub>, using a Kv4.3<sup>-/-</sup> transgenic mouse and antigen retrieval immunohistochemistry we demonstrate that Kv4.2 channels are also expressed in SNc DA neurons, although their contribution is significant in a minority of neurons only. Using electrophysiological recordings and computational modeling, we then demonstrate that I<sub>A</sub> gating kinetics and its functional complementarity with the hyperpolarization-activated current are major determinants of both pacemaking activity and post-inhibitory response in SNc DA neurons.

#### INTRODUCTION

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While the expression of only two types of voltage-gated ion channels in the squid giant axon allowed Hodgkin and Huxley to dissect the biophysical processes underlying action potential (AP) genesis and conduction (Hodgkin and Huxley, 1952), most neuronal types express a multitude of ion channel subtypes underlying their electrical activity (Cembrowski et al., 2016; Fuzik et al., 2016; Tapia et al., 2018; Northcutt et al., 2019). In spontaneously active neurons, a variety of voltage- and calcium-gated ion channels are not only responsible for the AP, but also govern the subthreshold oscillations leading to AP firing, determine firing frequency and control its regularity (Atherton and Bevan, 2005; Swensen and Bean, 2005; Bean, 2007; Gantz et al., 2018). Substantia nigra pars compacta (SNc) dopaminergic (DA) neurons spontaneously generate a regular tonic pattern of activity, also known as "pacemaking" activity (Grace and Onn, 1989; Gantz et al., 2018). Over the past 40 years, many studies contributed to the identification of the specific ion channels involved in shaping pacemaking activity (Nedergaard and Greenfield, 1992; Liss et al., 2001; Seutin et al., 2001; Wolfart et al., 2001; Neuhoff et al., 2002; Liss et al., 2005; Chan et al., 2007; Puopolo et al., 2007; Guzman et al., 2009; Putzier et al., 2009; Ji et al., 2012; Gantz et al., 2018). In particular, several studies have suggested that the transient A-type potassium current (I<sub>A</sub>) plays an essential role in controlling pacemaking rate and post-inhibitory firing delay in these neurons (Liss et al., 2001; Putzier et al., 2008; Amendola et al., 2012; Tarfa et al., 2017). In addition, single-cell PCR, in situ hybridization and immunohistochemistry experiments suggested that the A-type current is carried exclusively by Kv4.3 ion channels (Serodio and Rudy, 1998; Liss et al., 2001; Ding et al., 2011; Dufour et al., 2014a; Tapia et al., 2018). Interestingly, several studies also suggested that the H-type current (I<sub>H</sub>, carried by HCN channels) displays strong functional interactions with IA, having for instance an opposite

influence on post-inhibitory rebound delay (Amendola et al., 2012; Tarfa et al., 2017). The gating properties of these two currents were also shown to be co-regulated in rat SNc DA neurons (Amendola et al., 2012).

In the current study, we used in particular electrophysiological recordings from wild-type (WT) and Kv4.3<sup>-/-</sup> mice to refine the identity and role of Kv4 channels in the firing of SNc DA neurons. Using this mouse model, we show that the constitutive loss of Kv4.3 is associated with increased exploratory activity and impaired motor learning. Consistently, it is also associated with a lack of compensatory changes in other ion currents at the cellular level. We then demonstrate that the Kv4.2 subunit is expressed in SNc DA neurons, although its functional contribution is minor in most SNc DA neurons. Finally, we also demonstrate that pacemaking frequency and post-inhibitory rebound delay are mainly determined by I<sub>A</sub> time constant of inactivation and I<sub>H</sub> amplitude.

#### 92 MATERIAL AND METHODS

93	Animals
94	Female and male P15-P80 WT (n=68 animals) and Kv4.3 <sup>-/-</sup> (n=40, Deltagen) mice from
95	C57BL6/J genetic background were housed with free access to food and water in a temperature-
96	controlled room (24°C) on a 12:12 h dark-light cycle (lights on at 07:00 h). All efforts were
97	made to minimize the number of animals used and to maintain them in good general health
98	according to the European (Council Directive 86/609/EEC) and institutional guidelines for the
99	care and use of laboratory animals (French National Research Council).
100	Behavioral Experiments
100 101	Behavioral Experiments  A group of female and male WT (n=11) and Kv4.3 <sup>-/-</sup> (n=13) mice aged P56-P63 at the start of the
101	A group of female and male WT (n=11) and Kv4.3 <sup>-/-</sup> (n=13) mice aged P56-P63 at the start of the
101 102	A group of female and male WT (n=11) and Kv4.3 <sup>-/-</sup> (n=13) mice aged P56-P63 at the start of the behavioral testing were used to evaluate changes in motor function, using in particular locomotor

Actimetry was monitored in individual activity chambers (20 cm × 11.2 cm × 20.7 cm) housed within a sound-attenuating cubicle and under homogeneous illumination (Imetronic, Pessac, France). Each chamber was equipped with two pairs of infrared photobeams located 1.5 and 7.5cm above the floor level of the chamber. The number of back-and-forth movements (animals breaking the lower photobeams) as well as the number of vertical movements (animals breaking the upper photobeams) were recorded in 5-min bins over 90 min. Numbers of back-and-forth movements (locomotion) and vertical movements (rearing) are shown as mean ± SEM for each

time bin over the whole period of recording time. Locomotion and rearing activities over the whole 90-min period were normalized and displayed as a mean percentage of control littermates.

Motor learning

Motor learning was evaluated on the accelerating rotarod (10cm diameter rod) test at a speed of 5 to 40 rotations per min (RPM) for 5 min. On the first day, mice were allowed to freely explore the non-rotating apparatus for 60s and subsequently trained to hold on the rotating rod (5 RPM) for at least two 60s trials, each trial being separated by a 10min break. Mice were allowed to recover for one hour before the first test. The testing phase consisted in 10 consecutive trials on the accelerating rod separated by 15-min breaks that allowed consolidation of performance. Results are shown as the average latency to fall off the rod (mean  $\pm$  SEM) at each trial. A performance index was calculated for each individual and consisted in the average latency of the

last 3 trials divided by the average latency of the first 3 trials multiplied by 100

 $\left(\frac{average\ latency\ 8-10}{average\ latency\ 1-3} \times 100\right)$ .

#### Electrophysiology

- 126 119 neurons from 39 WT mice and 109 neurons from 24 Kv4.3<sup>-/-</sup> mice were recorded (current-
- 127 clamp and voltage-clamp).

#### 128 Acute midbrain slice preparation

Acute slices were prepared from P15-P25 animals of either sex. Mice were anesthetized with isoflurane (CSP) in an oxygenated chamber (TEM SEGA) before decapitation. After decapitation the brain was immersed briefly in oxygenated ice-cold low-calcium aCSF containing the following (in mM): 125 NaCl, 25 NaHCO3, 2.5 KCl, 1.25 NaH2PO4, 0.5 CaCl2, 4 MgCl2, and

133	25 D-glucose, pH 7.4, oxygenated with 95% O2/5% CO2 gas. The cortices were removed and
134	then coronal midbrain slices (250 $\mu m$ ) were cut in ice-cold oxygenated low-calcium aCSF on a
135	vibratome (Leica VT1200S and vibrating microtome 7000smz, Camden Instruments, UK).
136	Following 20-30 min incubation in oxygenated low-calcium aCSF at 33°C, the acute slices were
137	then incubated for a minimum of 30 min in oxygenated aCSF (containing in mM: 125 NaCl, 25
138	NaHCO3, 2.5 KCl, 1.25 NaH2PO4, 2 CaCl2, 2 MgCl2, and 25 glucose, pH 7.4, oxygenated with
139	95% O2/5% CO2 gas) at room temperature before electrophysiological recordings.
140	<u>Drugs</u>
141	Kynurenate (2 mM, Sigma-Aldrich) and picrotoxin (100 μM, Sigma-Aldrich) were used to block
142	excitatory and inhibitory synaptic activity, respectively. AmmTX3 (1µM, Alomone) was used to
143	block the transient potassium current (IA) carried by Kv4 channels. Drugs were bath applied via
144	continuous perfusion in aCSF.
145	Electrophysiology recordings and analysis
146	All recordings (228 neurons from 63 mice) were performed on midbrain slices continuously
147	superfused with oxygenated aCSF at 30-32°C. Picrotoxin and kynurenate were systematically
148	added to the aCSF for all recordings to prevent contamination of the intrinsically generated
149	activity by glutamatergic and GABAergic spontaneous synaptic activity. Patch pipettes (1.9-2.7
150	MOhm) were pulled from borosilicate glass (GC150TF-10, Harvard Apparatus) on a DMZ-
151	Universal Puller (Zeitz Instruments) and filled with a patch solution containing the following (in
152	mM): 20 KCl, 10 HEPES, 0.5 EGTA, 2 MgCl2, 0.4 Na-GTP, 2 Na <sub>2</sub> -ATP, 4 Mg-ATP,0.3 CaCl2,
153	SUPERase RNase inhibitor (0.1 $U/\mu l$ ) and 115 K-gluconate, pH 7.4, 290–300 mOsm. For
154	AmmTX3 experiments, patch pipettes (3.2-4.0 MOhm) were filled with a patch solution

containing the following (in mM): 20 KCl, 10 HEPES, 0.5 EGTA, 2 MgCl2, 2 Na-ATP, and 120 K-gluconate, pH 7.4, 290–300 mOsm. Whole-cell recordings were made from SNc DA neurons visualized using infrared differential interference contrast videomicroscopy (QImaging Retiga camera; Olympus BX51WI microscope), and were identified based on their location, large soma size (>25μm), and electrophysiological profile (regular slow pacemaking activity, large spike half-width, large sag in response to hyperpolarizing current steps). For voltage-clamp experiments, only whole-cell recordings with an uncompensated series resistance <7 MOhm (compensated 85–90%) were included in the analysis. For current-clamp pharmacology experiments, higher series resistances were tolerated as long as the bridge compensation was properly adjusted to 100%. Liquid junction potential (-13.2 mV) and capacitive currents were compensated on-line. Recordings were acquired at 50kHz and were filtered with a low-pass filter (Bessel characteristic 2.8kHz cutoff). For current-clamp recordings, 1s hyperpolarizing current steps were injected to elicit a hyperpolarization-induced sag (due to I<sub>H</sub> activation).

#### Current-clamp recordings and protocols

The spontaneous firing frequency was calculated from a minimum of 30 s of stable recording in cell-attached mode and from current-clamp recording (with no injected current) within the first 5 min after obtaining the whole-cell configuration. The coefficient of variation of the interspike interval (CV<sub>ISI</sub>) was extracted from the same recording. Action potentials (APs) generated during this period of spontaneous activity were averaged and several parameters were extracted: AP threshold, AP amplitude, AP duration at half of its maximal height (AP half-width), AHP trough voltage, AHP latency. Hyperpolarizing current steps and depolarizing current steps were used to characterize the post-inhibitory rebound and the excitability properties. The gain start, gain end

and spike frequency adaptation (SFA) index used to define excitability were calculated as described before (Dufour et al., 2014b).

#### Voltage-clamp recordings

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The I<sub>A</sub> current was elicited by a protocol consisting in a 500 ms prestep at -100 mV (to fully deinactivate I<sub>A</sub>) followed by a 500 ms voltage step to -40 mV (to activate I<sub>A</sub> without eliciting delayed rectifier potassium currents). The current generated by the same protocol using a prestep at -40mV (to fully inactivate IA) was subtracted to isolate IA. IA properties (peak amplitude and total charge) were measured after subtracting the baseline at -40mV. Total charge was calculated by integrating the current over the whole duration (500ms) of the voltage step. The peak of the current elicited at -40mV was then plotted against the voltage of each corresponding prestep, and was fitted with a Boltzmann function to obtain IA half-inactivation voltage (V50 IA) (see (Amendola et al., 2012)). The inactivation time constant (IA tau) was extracted from a monoexponential fit of the decay of the current. A two-step voltage-clamp protocol was used to determine the voltage-dependence of activation of I<sub>H</sub> (V<sub>50</sub> I<sub>H</sub>) and obtain the maximum I<sub>H</sub> amplitude (see (Amendola et al., 2012) for details). For voltage-clamp recordings of delayed rectifier current (I<sub>KDR</sub>), tetrodotoxin (1 µM, Alomone), nickel (200 µM, Sigma-Aldrich) and cadmium (400 µM, Sigma-Aldrich) were also added to the aCSF. I<sub>KDR</sub> was elicited by using a protocol consisting in a prestep at -40mV (to fully inactivate I<sub>A</sub>) followed by incremental depolarizing voltage steps up to +40mV.

#### Data acquisition

Data were acquired using an EPC 10 USB patch-clamp amplifier (HEKA) and the Patchmaster software acquisition interface (HEKA). Analysis was performed using FitMaster v2x73 (Heka).

#### *Immunohistochemistry*

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Adult (P21-P28) C57BL/6 WT mice (n=2) or Kv4.3<sup>-/-</sup> littermates (n=2) of either sex were euthanized with ketamine-xylazine mix (100mg/Kg ketamine, 10mg/Kg xylazine), and transcardially perfused with PBS and ice-cold 4% paraformaldehyde in PBS. Brains were removed and post-fixed overnight (o/n) at 4°C in the same fixative solution. Coronal brain slices of 50 µm were obtained using a vibratome (vibrating microtome 7000smz, Camden Instruments, UK) and collected as floating sections. When indicated, antigen retrieval was performed by incubating the slices in sodium citrate (10mM, Sigma-Aldrich) during 30 min at 80°C (Jiao et al., 1999). Subsequently, slices were blocked for 1 h 30 min at room temperature (RT) in a solution containing 0.3% Triton X-100 (SIGMA) and 5% Normal Goat Serum (NGS, Vector Laboratories) in PBS. After blocking, sections were incubated with primary antibodies in a solution containing 0.3% Triton X-100 and 1% NGS in PBS (o/n; 4°C). The following primary antibodies were used in this study: Chicken anti-TH (1:1000; Abcam, ab76442, RRID:AB 1524535), Rabbit anti Kv4.3 (1:500, Alomone Labs, APC-017, RRID:AB 2040178), Mouse anti Kv4.2 (1:200, Neuromab, 75-361, clone L28/4, RRID:AB 23155873). After three washes (15 min/each) in PBS containing 0.3% Triton X-100, the floating sections were incubated with the following secondary antibodies: Alexa 488-Goat anti-mouse (1:200, Jackson ImmunoResearch), Alexa 488-Goat anti-rabbit (1:200, Jackson ImmunoResearch) and Alexa 594-Goat anti-chicken (1:200, Jackson ImmunoResearch) in a PBS solution containing 0.3% Triton X-100 and 1% NGS for 2 h at RT. Finally, sections were washed three times (15min/each), incubated with DAPI (1.5 µg/mL; Sigma-Aldrich) for 10 min, and mounted in Vectashield (Vector Laboratories). Sections were stored at 4°C, and images were acquired on a Zeiss LSM-780 confocal scanning microscope. All experiments involving WT and Kv4.3<sup>-/-</sup>

comparisons were performed in parallel applying the same acquisition settings to both genotypes. Images were processed and analyzed with ImageJ (NIH). Kv4.2-positive cells were visually identified in both genotypes on the basis of a perimembranous-like Kv4.2 staining and expressed as a percentage of the total number of TH+ cells. In order to compare the labeling pattern of Kv4.2 and Kv4.3, the line selection tool was used to trace 3µm-length lines perpendicular to the cell perimeter in individual optical sections. In each cell, 3 regions were analyzed, and 5 cells were used to calculate the average profile in each condition. Raw intensity values were collected, normalized (0-1 range) to the maximal value, and plotted as a function of distance (0 corresponding to Kv4 peak fluorescence signal, negative distances to extracellular space and positive distances to intracellular space; see Figure 5B). All the images shown are one single optical slice. For the sake of representation and to overcome differences regarding levels of TH expression in different DA neurons, different minimum and maximum display settings were applied in Figure 5C exclusively for the TH channel.

#### Modeling

Simulations were performed using NEURON 7.5 software (Hines and Carnevale, 2001) as previously described (Moubarak et al., 2019). Realistic morphologies of 22 rat SNc DA neurons obtained previously were used to build multicompartment models (Moubarak et al., 2019). For each compartment, membrane voltage was obtained as the time integral of a first-order differential equation:

$$\frac{dV}{dt} = -\frac{1}{C_m} * \sum [g_i * (V_m - E_{rev})] - I_{axial}$$

where  $V_m$  is the membrane potential,  $C_m$  the membrane capacitance,  $g_i$  are ionic conductances and  $E_{rev}$  their respective reversal potentials. The axial flow of current ( $I_{axial}$ ) between adjacent

compartments is calculated by the NEURON simulation package (Hines and Carnevale, 2001).
Cytoplasmic resistivity, specific membrane capacitance and specific membrane resistance were
set to 150 Ohm.cm, 0.75 $\mu F/cm^2$ , and 100,000 Ohm.cm <sup>2</sup> , respectively, with $E_{rev}$ for the leak
conductance set at -50 mV. Six active conductances were included in the model: fast sodium
$(I_{Na}), \ delayed \ rectifier \ potassium \ (I_{KDR}), \ transient \ potassium \ (I_{A}), \ L\text{-type \ calcium \ } (I_{CaL}),$
hyperpolarization-activated ( $I_{\text{H}}$ ) and small conductance calcium-activated potassium ( $I_{\text{SK}}$ )
currents (Moubarak et al., 2019). Active conductances followed activation-inactivation Hodgkin-
Huxley kinetics (Table 2). Parameters for $I_A$ , $I_{CaL}$ , $I_{SK}$ , $I_{Na}$ , $I_{KDR}$ and $I_H$ were based on our
previous model and published values for SNc DA neurons (Gentet and Williams, 2007; Seutin
and Engel, 2010; Amendola et al., 2012; Philippart et al., 2016; Moubarak et al., 2019).
Intracellular calcium uptake was modeled as a simple decaying model according to Destexhe
(Destexhe et al., 1993). Conductance values were set according to our own measurements or
published values (see Table 2). Consistent with the literature (Kole and Stuart, 2008; Hu et al.,
2009), $g_{Na}$ and $g_{KDR}$ were set to higher values in the axon initial segment (AIS) than in the rest of
the neuron so that the action potential always initiated in the AIS. For sake of simplicity,
activation and inactivation kinetics of IA were voltage-independent but coupled to each other,
such that activation rate was 50 times faster than inactivation rate. In addition the inactivation and
activation $V_{50}s$ were also coupled (50 mV shift). As $I_A$ and $I_H$ voltage-dependences have been
shown to be positively correlated in rat SNc DA neurons (Amendola et al., 2012), both values
were forced to co-vary in the model according to the equation $V_{50}$ inact. $I_A = 0.814*(V_{50} \ act.$
$I_{H}$ )+3.36.
Initializing potential was set at -70 mV and pacemaking frequency was let to stabilize (4 spikes)
before further analysis. Each simulation run had a duration of 8000 ms with a dt of 0.02 ms.
before further analysis. Each simulation run had a duration of ovol his with a dt 01 0.02 his.

Spatial discretization followed the "d\_lambda rule" (Hines and Carnevale, 2001). All dendritic compartments and the axon-start compartment contained all currents whereas AIS and axon only contained fast sodium ( $I_{Na}$ ) and delayed rectifier potassium ( $I_{KDR}$ ) currents. To measure post-inhibitory rebound delay, current injection was performed by inserting a virtual electrode in the soma. A 1s pulse of current was injected into the model. Negative current amplitude was adjusted to achieve a peak hyperpolarization around -120 mV in each neuron and condition. Firing frequency, rebound delay and action potential property analyses were computed online by handmade routines directly written in NEURON hoc language (Hines and Carnevale, 2001). This model is derived from a previous model available at model DB database under the number 245427.

#### **Statistics**

All statistical analyses were conducted under the R environment with appropriate packages. For behavioral experiments, normality was assessed by visual inspection of quantile-quantile (Q-Q) plots for the different scores per animal (ggpubr package). Nearly all the data points did not depart for normality estimated within a 95% coefficient interval. For electrophysiology experiments, normality was checked using the Shapiro-Wilk normality test (stat package).

#### Behavior

To assess locomotor and exploratory behavior, number of horizontal (locomotion) and vertical (rearing) photobeam breaks was measured per 5-min bin over 90 min and compared between genotypes. As no significant difference was found between males and females in any of the behavioral tests, both sexes were pooled and analyzed as a single sample. Data are represented as line and scatter plots for the number of horizontal and vertical photobeam breaks per 5-min bin.

The number of total movements was then normalized to the level of WT spontaneous locomotion. To assess the locomotor phenotype of Kv4.3<sup>-/-</sup> mice, two-way repeated measures ANOVA tests with groups (WT/Kv4.3<sup>-/-</sup>) as the independent between-factor and time as the within-factor (training sessions or time-bins for locomotion) were performed (stat package). When the ANOVA was significant, multiple comparisons (false-discovery rate (fdr) adjustment, multicomp and emmeans packages) were used to evaluate differences between groups at different time points (Benjamini and Hochberg, 1995). P-values <0.05 were considered as statistically significant for all analyses. For motor learning, the average latency to fall off the accelerating rotarod was measured for each trial. Statistical difference in motor learning was assessed by comparing the *performance index*. Data are represented as line and scatter plots for the average latency to fall off the rod. Data are represented as mean ± SEM.

#### Electrophysiology and immunohistochemistry

The univariate statistical analysis of electrophysiological data, performed according to the distribution properties of the data using a Shapiro-Wilk normality test, included paired t-test or Wilcoxon signed rank test; t-test or Mann-Whitney Wilcoxon test with p < 0.05 considered to be statistically significant (stat package). In most figures, data are represented as scatter plots or box and whisker plots, with all individual points appearing on the graphs and dotted lines indicating the distribution of data (violin plots). For pharmacological experiments, data are represented as mean  $\pm$  SEM (scatter or bar plots). Correlation, linear regression and multiple linear regression analysis (**Figure 1**, **9**) were performed in R. For every pair of variables, correlation parameters, rho (Spearman correlation factor) or r (Pearson correlation factor), were selected after performing a Shapiro-Wilk normality test on the linear regression residuals and p values were corrected for multiple comparisons by an fdr adjustment (stat package). For multiple linear regression,

variables (extracellular ISI, rebound delay, I<sub>A</sub> tau, I<sub>H</sub> amplitude and I<sub>A</sub> amplitude) were first log transformed, and then dependent variables were standardized (subtracting the mean and dividing by the SD). A selection of the best subsets of dependent variables for each model size (1 to 4 for the model and 1 to 5 for real data) was first performed (leaps package) according to several criteria (adjusted R², AIC, BIC). The best model was then selected by comparing the prediction error of each model after performing a repeated (20 times) 10-fold cross-validation on test data (caret package). The best linear model, corresponding to the minimum cross-validation error (i.e. a model with the best predictive power) was then obtained. Multicolinearity was assessed by computing a score called the variance inflation factor (VIF package) and VIF was < 1.5 for all variables retained in the different models. For the immunohistochemistry experiments, a Fisher exact test was used to compare the proportions of Kv4.2-positive cells among TH-positive cells in WT and Kv4.3<sup>-/-</sup> mice. Figures were prepared using R, SigmaPlot 11.0, GraphPad Prism 6 and Adobe Illustrator CS5/CS6.

#### RESULTS

#### Motor learning impairment of the Kv4.3<sup>-/-</sup> transgenic mice

In order to define the precise contribution of Kv4.3 channels to SNc DA neuron electrophysiological phenotype, we used Kv4.3 constitutive KO mice (Niwa et al., 2008; Carrasquillo et al., 2012; Granados-Fuentes et al., 2012). These mice have been used previously to study Kv4.3 function in cardiac ventricles (Niwa et al., 2008), in neurons of the suprachiasmatic nucleus (Granados-Fuentes et al., 2012) and in neocortical pyramidal neurons (Carrasquillo et al., 2012). No major defect in cardiac function or circadian locomotor behavior

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was reported in these studies. As Kv4.3 is also strongly expressed in the SNc and in the ventral tegmental area (VTA) (Serodio and Rudy, 1998), we first sought to determine whether Kv4.3 loss could affect SNc or VTA-related behaviors, such as locomotion and motor learning (Figure 1). As illustrated in Figure 1A, horizontal locomotor activity assessed in photocell activity chambers was similar between Kv4.3<sup>-/-</sup> mice and their WT littermates (F<sub>1,21</sub>=0.31, p=0.58, between factor effect), both strains displaying a significant decrease of locomotion over time  $(F_{17,357}=20.5, p=2 \cdot 10^{-16}, within factor effect)$ . The exploratory behavior was higher in Kv4.3<sup>-/-</sup> mice than WT littermates in particular during the first part of the session (Figure 1A), although the overall exploratory activity was not significantly different between the two strains ( $F_{1,21}=1.42$ , p=0.25, between factor effect). Again, both strains displayed a strong run-down of activity over the whole 90-min testing time as a function of habituation to the locomotor activity chambers (F<sub>17,357</sub>=13.97, p=2•10<sup>-16</sup>, within factor effect). Consistent with the increased exploratory activity at the start of the session for the Kv4.3<sup>-/-</sup> mice, the two-way repeated measures ANOVA revealed a significant interaction between strains and time (F<sub>17,357</sub>=1.87, p=0.019). Motor learning abilities were then assessed using the rotarod test for ten consecutive trials (Figure 1B). While WT mice improved their performance across trials, as shown by the increase in the latency-to-fall, Kv4.3<sup>-/-</sup> mice were not able to adjust their performance over time. While the latency averaged over trials was not different between the two strains (F<sub>1,22</sub>=1.26, p=0.28, between factor effect, two-way repeated measures ANOVA), performance changed significantly across trials (F<sub>9, 198</sub>=4.5, p=2•10<sup>-5</sup>, within factor effect), especially for WT mice. Consistent with the differences in latency for the late trials (Figure 1B), the difference in learning between the two strains was revealed by the strains\*trials interaction statistics (F<sub>9,198</sub>=2.95, p=0.0026) and significant differences in falloff latencies of Kv4.3<sup>-/-</sup> and WT mice was found for the late trials (7 to 9, Figure 1B). In order to better quantify the difference in learning between strains, we then calculated a performance index

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based on the difference in latency for the first 3 trials against the last 3 trials. Consistent with the ANOVA results, the performance index in this motor learning task was significantly lower for the Kv4.3<sup>-/-</sup> mice (WT, 176.9  $\pm$  21.4, n=11 vs Kv4.3<sup>-/-</sup>, 111.2  $\pm$  8.3, n=13, p=0.006, unpaired t-test, **Figure 1B**).

### Changes in electrophysiological phenotype of Kv4.3<sup>-/-</sup> SNc DA neurons

Following the approach already used in a previous study (Dufour et al., 2014b), we then performed an exhaustive current-clamp characterization of the firing properties of SNc DA neurons to determine changes in phenotype associated with Kv4.3 deletion (Table 1). Passive properties, spontaneous activity, post-inhibitory rebound, action potential shape and excitability were assessed by measuring 16 different electrophysiological parameters in 75-101 WT and 66-77 Kv4.3<sup>-/-</sup> neurons. The first obvious electrophysiological change observed was that spontaneous activity (extracellularly recorded in cell-attached mode) was dramatically modified in Kv4.3<sup>-/-</sup> SNc DA neurons (Figure 2A, B). Spontaneous firing rate was increased by ~2-fold in Kv4.3<sup>-/-</sup> mice, as demonstrated by the significant decrease in interspike interval (ISI, Figure 2A, B; Table 1). Pacemaking regularity, measured by the coefficient of variation of the ISI (CV<sub>ISI</sub>) was also significantly different in Kv4.3<sup>-/-</sup> mice, although CV<sub>ISI</sub> values were very low (<20%) in both WT and Kv4.3<sup>-/-</sup> mice, indicating a highly regular tonic activity. Post-inhibitory rebound delay was also dramatically decreased in Kv4.3<sup>-/-</sup> mice (Figure 2C, D; Table 1). However, the I<sub>H</sub>-mediated voltage sag observed during prolonged hyperpolarization was not modified (Figure 2C, D; Table 1). Interestingly, most action potential parameters were unchanged in Kv4.3<sup>-/-</sup> mice, except for action potential half-width, which was slightly larger (Figure 3A; Table 1). We also analyzed neuronal excitability by measuring the responses of the neurons to increasing depolarizing current steps (Figure 3B). Excitability was slightly increased in the DA neurons of Kv4.3<sup>-/-</sup> mice,

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although this change only affected the initial response of neurons (gain start) to current injection (**Figure 3B**; **Table 1**). Consistently, the frequency of the response of Kv4.3<sup>-/-</sup> neurons to a 100pA step was also found to be significantly higher (**Figure 3B**, **Table 1**).

## Voltage-clamp characterization of $I_A$ in Kv4.3 $^{-1}$ SNc DA neurons

We then directly investigated changes in the properties of I<sub>A</sub> by performing voltage-clamp recordings in WT and Kv4.3<sup>-/-</sup> SNc DA neurons (Figure 4). A dramatic decrease in I<sub>A</sub> amplitude was observed in Kv4.3<sup>-/-</sup> mice (Figure 4A, B; Table 1). However, a small transient residual current with I<sub>A</sub>-like properties (voltage-dependent inactivation) was still present in all Kv4.3<sup>-/-</sup> recordings (Figure 4A). Most interestingly, this residual current was completely blocked by the Kv4-specific toxin AmmTX3 (n=4, no measurable residual current after toxin application; Figure 4A), suggesting that a Kv4 subunit other than Kv4.3 is expressed at a low level in Kv4.3<sup>-/-</sup> SNc DA neurons. We then measured its time constant of inactivation (IA tau) and calculated the overall charge carried by the current (Figure 4C-E). Both parameters were strongly decreased in Kv4.3<sup>-/-</sup> SNc DA neurons (Figure 4C, D; Table 1), although a minority of cells (n=5/42) displayed values similar to the WT measurements for both of these parameters. Plotting I<sub>A</sub> charge versus I<sub>A</sub> tau revealed the clear separation of values between the Kv4.3<sup>-/-</sup> and WT measurements, except for the 5 cells identified before (Figure 4E). Based on these voltage-clamp data, it appears that, although the Kv4.3 subunit by far predominates in WT SNc DA neurons, another Kv4 subunit is also expressed, at least in the Kv4.3<sup>-/-</sup> neurons. Although in most cases, the expression level of this unidentified subunit is too low to compensate for the loss of Kv4.3, it generates an A-type current that provides a minority of Kv4.3<sup>-/-</sup> SNc DA neurons (5/42=12%) with a "wildtype" voltage-clamp phenotype.

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#### Using antigen retrieval to reveal the expression of Kv4.2 channels by SNc DA neurons

Several studies have investigated the expression of A-type Kv channels in SNc DA neurons, using in situ hybridization (Serodio et al., 1996; Serodio and Rudy, 1998), single-cell quantitative PCR (Liss et al., 2001; Ding et al., 2011; Tapia et al., 2018) and immunohistochemistry (Liss et al., 2001; Dufour et al., 2014a). A high level of expression for Kv4.3 (Liss et al., 2001; Dufour et al., 2014a; Tapia et al., 2018) and the absence of Kv4.1 (Serodio and Rudy, 1998; Liss et al., 2001; Ding et al., 2011) were consistently reported, while the presence of Kv4.2 is still debated. In particular, while Kv4.2 mRNA has been detected in several studies (Ding et al., 2011; Tapia et al., 2018), the protein was not detected by classical immunohistochemistry (Liss et al., 2001; Dufour et al., 2014a). Interestingly, it has been shown that, depending on the brain region and the subcellular location of the ion channel of interest, an antigen retrieval (AR) procedure may be required to uncover potassium channel antigen epitopes before performing immunolabeling (Lorincz and Nusser, 2008). We first confirmed that Kv4.3 was strongly expressed in SNc DA neurons in WT mice, with the expected membrane profile of immunostaining, and that it was absent from Kv4.3<sup>-/-</sup> SNc DA neurons (**Figure 5A**). We then performed Kv4.2 immunolabeling with or without AR on the neocortex and the CA1 region of the hippocampus where this ion channel is highly expressed (Serodio and Rudy, 1998). As can be seen in Figure 5-1, Kv4.2 immunostaining was greatly improved by AR, revealing a strong perisomatic and dendritic staining of pyramidal cells in both regions. Therefore we implemented AR before performing Kv4.2 immunostaining on WT midbrain slices (Figure 5B). Similar to what we observed in the hippocampus and in the cortex, Kv4.2 immunostaining in the SNc was greatly improved by AR, although only a minority of DA neurons (TH-positive) displayed a clear perisomatic Kv4.2 signal compatible with membrane expression of the channel. In fact, the AR Kv4.2 staining profile of Kv4.2-positive cells was very similar to the membrane staining profile observed for Kv4.3 (**Figure 5B**). This distinctive staining profile was then used to quantify the percentage of Kv4.2-positive SNc DA neurons in both WT and Kv4.3<sup>-/-</sup> mice (**Figure 5C**, **D**). The percentage of Kv4.2-positive cells was not significantly different between WT and Kv4.3<sup>-/-</sup> mice (WT 20/423=4.7% *vs* Kv4.3<sup>-/-</sup> 19/382=5%, p=1, Fisher exact test, **Figure 5D**), suggesting that Kv4.2 pattern of expression is not modified by the loss of Kv4.3. This percentage is not statistically different from the percentage of Kv4.3<sup>-/-</sup> SNc DA neurons presenting an atypically large and slow I<sub>A</sub> reported in **Figure 4C-E** (5/42=11.9% vs 19/382=4.6%, p=0.077, Fisher exact test). Moreover, the percentage of Kv4.2-positive DA neurons was very similar in the medial and lateral SNc of both WT (SNc medial 11/227=4.8% *vs* SNc lateral 9/196=4.6%, p=1, Fisher exact test) and Kv4.3<sup>-/-</sup> mice (SNc medial 13/218=6% *vs* SNc lateral 6/164=3.7%, p=0.35, Fisher exact test).

#### Lack of compensation in the face of Kv4.3 loss

Genetic deletion of Kv4.2 channels in cortical pyramidal neurons is associated with compensatory modifications in a delayed rectifier-like (I<sub>KDR</sub>-like) potassium current (Nerbonne et al., 2008). Moreover, I<sub>H</sub> and I<sub>A</sub> have complementary influences on post-inhibitory firing in SNc DA neurons (Amendola et al., 2012; Tarfa et al., 2017). In order to reveal putative homeostatic compensations of Kv4.3 deletion in SNc DA neurons, we first performed a series of current-clamp recordings on a subset of neurons (n=18 for WT, n=32 for Kv4.3<sup>-/-</sup>) to compare the effect of acutely blocking Kv4 channels using the scorpion toxin AmmTX3 (Vacher et al., 2002) to the changes observed in the Kv4.3<sup>-/-</sup> mouse. Consistent with previous reports (Amendola et al., 2012; Tarfa et al., 2017), AmmTX3 strongly increased pacemaking frequency and dramatically reduced

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post-inhibitory rebound delay (Figure 6). Most interestingly though, the magnitude of the effects of the toxin was very similar to that observed in the Kv4.3<sup>-/-</sup> neurons: firing frequency was increased by ~68% in both conditions (Figure 6B), while rebound delay was decreased by 87% after AmmTX3 and by 82% in Kv4.3<sup>-/-</sup> mice (Figure 6D). Consistent with the data presented earlier, these results strongly suggest that the Kv4-mediated A-type current is virtually completely abolished in Kv4.3<sup>-/-</sup> SNc DA neurons and that its loss is not compensated by changes in other Kv channels (and associated currents). We then used voltage-clamp recordings to directly assess whether a decrease in I<sub>A</sub> could be compensated by a parallel decrease in I<sub>H</sub> or a compensatory increase in I<sub>KDR</sub> (Nerbonne et al., 2008) (Figure 7). Unlike what has been described in cortical neurons following Kv4.2 deletion, I<sub>KDR</sub> was not modified in Kv4.3<sup>-/-</sup> SNc DA neurons (**Figure 7A**). I<sub>H</sub> was found to be slightly larger in Kv4.3<sup>-/-</sup> SNc DA neurons (Figure 7B, Table 1), but its voltage sensitivity was unchanged. Altogether, the voltage-clamp recordings of IA, IKDR and IH and the AR Kv4.2 immunostaining suggest that Kv4.3 loss is not compensated by changes in expression and/or function of functionally-overlapping channels. These data provide a biophysical explanation for the observation made earlier that the acute blockade of Kv4 channels produces an electrophysiological phenotype qualitatively and quantitatively virtually identical to the Kv4.3 genetic deletion (Figure 6). Bridging the gap between biophysical changes in  $I_A$  and  $I_H$  and variation in electrophysiological phenotype We then decided to investigate whether the cell-to-cell variations in I<sub>A</sub> biophysical properties in WT and Kv4.3<sup>-/-</sup> SNc DA neurons were predictive of variations in electrophysiological

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phenotype (Figure 8). We first looked at potential correlations between firing parameters. As already presented in Figure 2, the most significant alterations in firing observed in Kv4.3<sup>-/-</sup> SNc DA neurons are a strong increase in spontaneous firing frequency (strong decrease in the extracellularly-measured ISI) and a strong decrease in post-inhibitory rebound delay. We found that extracellular ISI and rebound delay (log transformed) were strongly positively correlated with each other in both WT and Kv4.3<sup>-/-</sup> neurons (Figure 8B), although the slope of this relationship seemed slightly different between the two genotypes. We therefore tried to determine whether specific IA biophysical properties were better predictors of these variations in ISI or rebound delay (Figure 8C). We also analyzed the relationship between I<sub>H</sub> properties and ISI or rebound delay. Out of the 5 biophysical parameters analyzed ( $I_A$  tau,  $I_H$  amplitude,  $I_A$  amplitude,  $I_A$  inactivation  $V_{50}$ ,  $I_H$  activation  $V_{50}$ ), only two parameters were significantly correlated with ISI or rebound delay: IA tau and IH amplitude (log transformed) were positively and negatively correlated, respectively, with both ISI and rebound delay in both WT and Kv4.3-/- neurons (Figure 8C). Surprisingly, neither I<sub>A</sub> amplitude (measured at -40mV) nor its voltage-dependence (inactivation V<sub>50</sub>) were predictive of variations in ISI or rebound delay (**Figure 8C**). I<sub>H</sub> activation  $V_{50}$  was also unable to predict variations in these firing parameters (data not shown). In addition, reminiscent of the observation made in rat neurons (Amendola et al., 2012), I<sub>H</sub> activation and I<sub>A</sub> inactivation V<sub>50</sub>s were found to be positively correlated (Figure 8D). I<sub>A</sub> tau and I<sub>H</sub> amplitude were also found to be negatively correlated (Figure 8D). Based on these observations, we then tested whether combining several of these 5 biophysical parameters could improve the prediction of ISI or rebound delay using multiple linear regression (Figure 8E). We first standardized these parameters (subtracting the mean and dividing by the SD), and then looked for the best subset of variables predictive of ISI or rebound delay. While ISI was best predicted by a multiple linear regression involving only IA tau and IH amplitude, rebound delay was better predicted when IA

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tau,  $I_H$  amplitude and  $I_A$  inactivation  $V_{50}$  were included in the linear regression (**Figure 8E**). Rebound delay prediction was much more accurate than ISI prediction ( $r^2$ =0.771 compared to  $r^2$ =0.421). Based on the scaling factors given by the multiple linear regression, it is important to note that both ISI and rebound delay in real neurons seem to be most sensitive to variations in  $I_A$  tau.

#### Influence of $I_A$ biophysical properties on SNc DA neuron firing

Many conductances other than I<sub>A</sub> and I<sub>H</sub> may vary in expression level and biophysical properties from neuron to neuron, potentially compensating or enhancing the effect of variations in the properties of these two currents on firing (Gentet and Williams, 2007; Puopolo et al., 2007; Seutin and Engel, 2010; Amendola et al., 2012; Philippart et al., 2016; Moubarak et al., 2019). In order to isolate the influence of specific biophysical properties of I<sub>A</sub> and I<sub>H</sub> on SNc DA neuron activity, we used a realistic multi-compartment Hodgkin-Huxley model of rat SNc DA neurons (Moubarak et al., 2019). Based on measurements obtained by different groups (Liss et al., 2001; Gentet and Williams, 2007; Amendola et al., 2012; Tarfa et al., 2017), each of the biophysical properties of I<sub>A</sub> (maximal conductance g<sub>A</sub>, voltage-dependence I<sub>A</sub> V<sub>50</sub>, inactivation rate I<sub>A</sub> tau) and I<sub>H</sub> maximal conductance (g<sub>H</sub>) were varied over a 10-fold range (20mV range for the voltagedependence) using 5 equi-distributed values (Figure 9A). IA and IH voltage-dependences were forced to co-vary in the model, based on our previous observations (Amendola et al., 2012). Using a sample of 22 realistic models and 5 independently-varying values for each biophysical property, a database of 13750 models (22\*5<sup>4</sup>) was generated (Figure 9). The average interspike interval (ISI) during spontaneous activity and the post-inhibitory firing delay in response to a hyperpolarizing pulse (rebound delay) were measured for each model and their average values were calculated for each combination of values of the 4 biophysical parameters (n=625).

Dimensional stacking (Taylor et al., 2006) was then used to represent the influence of the 4 biophysical parameters on these electrophysiological features in 2-dimensional heatmaps, allowing us to visually determine which parameters were most critical in controlling ISI and rebound delay (**Figure 9B**, **C**):  $g_A$  and  $I_A/I_H$   $V_{50}$  strongly modulated both ISI and rebound delay while  $I_A$  tau and  $g_H$  had a weaker influence on these firing properties. To quantify the contribution of each biophysical parameter to ISI and rebound delay variations, we then used the same strategy already presented in **Figure 8E** for experimental measurements: standardized parameters were used to run multiple linear regression against ISI or rebound delay (**Figure 9D**). Consistent with the visualization provided in **Figure 9B**, this sensitivity analysis revealed that the influence of  $g_A$  and  $I_A/I_H$   $V_{50}$  on ISI and rebound delay was 2-3 times stronger than that of  $g_H$  and  $I_A$  tau. While the results of this computational modeling are consistent with the general influence of  $I_A$  and  $I_H$  reported in these neurons (Liss et al., 2001; Seutin et al., 2001; Neuhoff et al., 2002; Puopolo et al., 2007; Amendola et al., 2012; Gantz et al., 2018), they reveal specific effects of  $I_A$  biophysical properties contrasting with the correlations identified in our experimental data.

#### DISCUSSION

The current study provides important elements regarding the identity of the ion channels underlying I<sub>A</sub> in SNc DA neurons and the relative influence of specific biophysical parameters of I<sub>A</sub> (voltage-dependence, gating kinetics, maximal conductance) on neuronal output. In particular, in contrast with previous studies (Serodio and Rudy, 1998; Liss et al., 2001; Dufour et al., 2014a), we show that the Kv4.2 subunit is expressed in SNc DA neurons, although its functional influence appears negligible in most neurons, due to a very low level of expression. In spite of

Kv4.2 presence, we show that the constitutive loss of Kv4.3 is not compensated by complementary intrinsic conductances (Kv4.2-mediated  $I_A$ ,  $I_{KDR}$ ,  $I_H$ ). In addition, while previous studies (Liss et al., 2001; Putzier et al., 2008) and the computational modeling performed here suggest a strong role of  $I_A$  conductance in controlling firing frequency, we demonstrate that  $I_A$  gating kinetics appear as the major determinant of both pacemaking frequency and post-inhibitory rebound delay. Our results also highlight the functional complementarity and correlation of biophysical properties of  $I_A$  and  $I_H$  in these neurons.

#### Kv4.2 is expressed in SNc DA neurons

One of the important results of the present work is the demonstration that Kv4.2 is expressed in mouse SNc DA neurons. So far, it was thought that only Kv4.3 was expressed and entirely responsible for the large A-type current observed in these neurons (Liss and Roeper, 2008; Gantz et al., 2018). Our results however unambiguously demonstrate that Kv4.2 is expressed by SNc DA neurons. First, in the absence of Kv4.1 (Liss et al., 2001; Ding et al., 2011), the presence of an AmmTX3-sensitive residual A-type current in the Kv4.3<sup>-/-</sup> neurons is only compatible with the expression of Kv4.2. While the Kv4.2-mediated residual I<sub>A</sub> is very small and fast in most neurons, in 12% of the voltage clamp-recorded neurons it is large enough to confer a "wild type" phenotype (see **Figure 5, 9**). Even though this residual I<sub>A</sub> current is much faster than its WT counterpart in most neurons, it still influences firing, particularly rebound delay, as suggested by the highly significant correlation between I<sub>A</sub> tau and rebound delay. Second, antigen retrieval immunohistochemistry confirmed that a Kv4.2 staining is observed in a minority (~5%) of SNc DA neurons in both Kv4.3<sup>-/-</sup> and WT mice. While the percentage of cells displaying a "high" expression of Kv4.2 is too small to allow a combined voltage-clamp/AR immunohistochemistry approach, the similarity in the proportion of AmmTX3-sensitive "large residual" I<sub>A</sub> (~12%) and

Kv4.2-positive neurons (~5%, not statistically different) strongly suggests that Kv4.2 is responsible for a large I<sub>A</sub> in a minority of SNc DA neurons. The presence of a small AmmTX3-sensitive residual I<sub>A</sub> in the rest of the Kv4.3<sup>-/-</sup> SNc DA neurons suggests that Kv4.2 is expressed and functional in all SNc DA neurons, although its influence might be minor in the presence of Kv4.3. The fact that the residual A-type current in Kv4.3<sup>-/-</sup> neurons inactivates faster than its WT counterpart is also consistent with the reported differences in inactivation kinetics of Kv4.2 and Kv4.3 (Serodio et al., 1994; Serodio et al., 1996). Altogether, these results suggest that, in contrast to the widely accepted view, a small subpopulation of SNc DA neurons (~5-10%) display an A-type current likely mediated by both Kv4.3 and Kv4.2 channels. Interestingly, these two subunits are very similar and co-immunoprecipitate from mouse brain lysates (Marionneau et al., 2009), suggesting that they could form heteromeric I<sub>A</sub> channels. At this point however, it is difficult to determine whether the Kv4.2-positive neurons in WT mice express Kv4.3/Kv4.2 heteromers or distinct Kv4.3 and Kv4.2 homomers.

## Lack of homeostatic compensation in Kv4.3<sup>-/-</sup> SNc DA neurons

Intriguingly, these results not only suggest that Kv4.2 expression is very low in most SNc DA neurons but also that it is not modified in the Kv4.3<sup>-/-</sup> SNc DA neurons. The structural and functional similarities between Kv4.3 and Kv4.2 means that in theory Kv4.2 channels should be able to compensate for the loss of Kv4.3. Despite this, the Kv4.2 pattern of expression is not modified in Kv4.3<sup>-/-</sup> SNc DA neurons. This is reminiscent of previous studies performed on the Kv4.2<sup>-/-</sup> mouse demonstrating that Kv4.3 expression pattern, assessed by western-blot or immunohistochemistry, is not modified following Kv4.2 loss (Menegola and Trimmer, 2006; Nerbonne et al., 2008). In addition, other currents (such as I<sub>KDR</sub> or I<sub>H</sub>) also appear to not be regulated in a compensatory direction in Kv4.3<sup>-/-</sup> SNc DA neurons. The current-clamp

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comparison between the acute blockade of Kv4 channels in WT neurons and the Kv4.3<sup>-/-</sup> neurons also supports this idea of a lack of compensatory modifications in functionally-overlapping currents. This is surprising in the light of the results obtained on neonatal cortical pyramidal cells (Nerbonne et al., 2008), where Kv4.2 genetic deletion is almost "fully" compensated by an increase in sustained potassium currents. To explain this difference, we may hypothesize that, although the alteration in electrophysiological phenotype observed in Kv4.3<sup>-/-</sup> neurons is striking (see Figure 3 in particular), the change in calcium dynamics associated with the elevated spontaneous activity may not be sufficient to trigger homeostatic regulatory mechanisms (O'Leary et al., 2014). Alternatively, modifications in the properties of incoming excitatory or inhibitory synaptic inputs (not analyzed in the current study) may compensate for the changes in intrinsic activity reported here, such that the overall in vivo activity of the SNc network is maintained in Kv4.3<sup>-/-</sup> animals. However, behavioral alterations suggest that the change in firing of SNc DA neurons is not totally compensated at the network level. Kv4.3<sup>-/-</sup> mice were more active than WT littermates in exploring the environment (increased rearing behavior) and could not adjust their motor control over sessions on the rotarod. This suggests that the changes in midbrain DA neuron tonic firing observed in the Kv4.3<sup>-/-</sup> SNc DA neurons might only be revealed when animals are exploring their environment or challenged in a motor learning task.

#### $I_A$ gating kinetics play a central role in SNc DA neuron output

The use of  $I_A$  voltage-clamp measurements and current-clamp recordings on a large number of neurons in WT and Kv4.3<sup>-/-</sup> mice allowed us to determine the impact of cell-to-cell variations in  $I_A$  biophysical properties on spontaneous activity and rebound delay. Interestingly, while realistic multi-compartment modeling suggested that  $I_A$  maximal conductance and voltage-dependence were the two factors most strongly influencing these electrophysiological features, our recordings

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revealed that IA inactivation rate was the dominant factor defining pacemaking frequency and rebound delay (Figure 9E). IH amplitude (proportional to IH maximal conductance in our measurements) was also found to play an important role in real neurons, while its influence was minor in the model. These differences may be explained by several factors. First, the database approach used for our simulations implies that all the tested biophysical properties are varied independently (except for the strict correlation applied to I<sub>A</sub> and I<sub>H</sub> V<sub>50</sub>s). While the independence between these biophysical parameters allowed us to precisely quantify the sensitivity of spontaneous activity and rebound delay to each parameter, it does not correspond to the observations made in real neurons: for instance I<sub>H</sub> amplitude and I<sub>A</sub> inactivation rate are negatively correlated, i.e. not independent from each other. On the other hand, reminiscent of a previous study performed on rat neurons (Amendola et al., 2012), we demonstrated that I<sub>A</sub> and I<sub>H</sub> V<sub>50</sub>s are also positively correlated in WT mouse neurons. However, this correlation is much lower than the one applied in the model (r=0.658,  $r^2$ =0.43 compared to  $r^2$ =1) and this difference may explain why this biophysical parameter appears as one of the most efficient in modulating firing in the model. Thus, the differences in independence of the biophysical properties may partly explain why the model and the experimental observations give different answers. Another factor may explain why IA maximal conductance has a strong effect on firing in the model, but not in real neurons. In order to isolate the effect of I<sub>A</sub> and I<sub>H</sub> biophysical properties on firing, all other conductances included in the model were held at fixed values. However, every ion current displays significant cell-to-cell variations in its properties (gating, conductance density) in a same neuronal population (Swensen and Bean, 2005; Schulz et al., 2006; Amendola et al., 2012; Moubarak et al., 2019). If happening at random, these variations in other currents would most likely dampen the effect of the variations in I<sub>A</sub> or I<sub>H</sub> specific properties on firing. In fact, we demonstrated in a previous study (Tapia et al., 2018) that the level of expression of Kv4.3 (at the

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mRNA level) in midbrain DA neurons co-varies with the expression levels of multiple somatodendritic ion channels, including Nav1.2, SK3 and GIRK2. If this co-variation is retained at the protein level, it would mean that cell-to-cell variations in IA maximal conductance occur in parallel with variations in density of other ion channels. Whether correlated or not, cell-to-cell variations in other conductances may thus explain why I<sub>A</sub> amplitude does not predict pacemaking and rebound delay in real neurons, and why IA inactivation rate appears as the main predictor of these electrophysiological features. In contrast with our findings, the results obtained by Liss and colleagues (2001) suggested that Kv4.3 expression level and channel density predicted pacemaking frequency in mouse neurons. Interestingly, it is noteworthy that I<sub>A</sub> inactivation rate showed restricted cell-to-cell variations in their recordings (2-fold range). On the other hand, I<sub>A</sub> charge density showed a 10-fold range of variation, suggesting that most of the variation in  $I_A$ function was due to variations in IA maximal conductance (Liss et al., 2001). In our recordings however, the levels of variability observed for I<sub>A</sub> (and I<sub>H</sub>) biophysical properties were rather similar, which led us to apply a 10-fold range to each parameter in our model. Thus, we postulate that the differences in our conclusions may be essentially related to differences in the cell-to-cell variability range of I<sub>A</sub> and I<sub>H</sub> biophysical parameters recorded in our samples.

#### Functional complementarity and co-regulation of $I_A$ and $I_H$ in SNc DA neurons

While our results confirm the well-established influence of I<sub>A</sub> on SNc DA neuron firing (Liss et al., 2001; Gentet and Williams, 2007; Putzier et al., 2008; Amendola et al., 2012; Tarfa et al., 2017), they also emphasize the functional complementarity between I<sub>A</sub> and I<sub>H</sub> in these neurons, and reinforce the idea that the channels underlying these currents are co-regulated (Amendola et al., 2012). Indeed, we confirm that I<sub>A</sub> and I<sub>H</sub> voltage-dependences are positively correlated and show that I<sub>H</sub> amplitude and I<sub>A</sub> inactivation rate are negatively correlated. While we do not have a

mechanistic explanation for this latter correlation, these results are reminiscent of the observations made by Tarfa and colleagues on nigrostriatal and mesoaccumbal DA neurons (2017). At the functional level, our results demonstrate that these two parameters are the main predictors of the cell-to-cell variations in pacemaking frequency and rebound delay, reinforcing the idea that  $I_A$  and  $I_H$  function as a complementary pair of currents tightly controlling post-inhibitory rebound delay in SNc DA neurons ( $r^2$ =0.77). The fact that pacemaking rate is not as accurately predicted by  $I_A$  and  $I_H$  properties ( $r^2$ =0.42) is consistent with the documented role of many other conductances and morphological parameters in defining this firing feature (Nedergaard and Greenfield, 1992; Wilson and Callaway, 2000; Wolfart et al., 2001; Liss et al., 2005; Puopolo et al., 2007; Putzier et al., 2009; Gantz et al., 2018; Moubarak et al., 2019).

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806	FIGURE LEGENDS								
807	Figure 1. Behavioral assessment of the Kv4.3-/- transgenic mouse. A, locomotion measured in								
808	actimetry chambers. Left, line and scatter plot showing the mean number of horizontal								
809	movements per 5-min bin (± SEM) in Kv4.3 <sup>-/-</sup> mice (red) compared to their WT littermates								
810	(black). Right, line and scatter plot showing the mean number of rearing events per 5-min bin in								
811	Kv4.3 <sup>-/-</sup> mice (red) compared to their WT littermates (black). Significant differences between								

strains for specific time bins (two-way repeated measures ANOVA) are indicated by an asterisk. **B**, changes in motor learning measured using a rotarod assay. The latency to falling off the rotating rod (with increasing rotating speed) was assessed over 10 consecutive trials (left). Significant differences between strains for specific trials (two-way repeated measures ANOVA) are indicated by an asterisk. The performance index ((average latency 8-10)/(average latency 1-3)\*100, right) was used to evaluate the learning ability of Kv4.3<sup>-/-</sup> mice (red) compared to their WT littermates (black). \* p<0.05, \*\* p<0.01.

Figure 2. Spontaneous activity and post-inhibitory rebound are profoundly altered in Kv4.3<sup>-/-</sup> SNc DA neurons. A, representative cell-attached recordings showing the spontaneous pattern of activity in WT (black trace, left) and Kv4.3<sup>-/-</sup> SNc DA neurons (red trace, right). B, box and whisker plots showing the distribution of values for extracellularly recorded spontaneous ISI (Extracellular ISI, left) and ISI coefficient of variation (CV<sub>extra. ISI</sub>, right) in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. C, representative current-clamp recordings showing the voltage response of SNc DA neurons to a current step (gray trace) hyperpolarizing membrane voltage to ~ -120mV in WT (black trace, left) and Kv4.3<sup>-/-</sup> mice (red trace, right). The recordings come from the same neurons as the cell-attached recordings presented in A. D, box and whisker plots showing the distribution of values for voltage sag amplitude (left) and post-inhibitory rebound delay (right) in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. \*\*\* p<0.001. Dotted lines in the box and whisker plots indicate the distribution of data (violin plots). Scale bars: A, horizontal 1s; C, top, horizontal 500ms, vertical 40mV; bottom, vertical 100pA; horizontal dotted lines indicate -60mV.

Figure 3. Action potential and excitability properties in Kv4.3<sup>-/-</sup> SNc DA neurons. A, left, current-clamp recordings showing the shape of the action potential in the WT (black traces) and Kv4.3<sup>-/-</sup> mice (red traces) on a slow (top) and fast time-scale (bottom). Right, box and whisker plots showing the distribution of values for action potential threshold (top left), amplitude (top right), half-width (AP half-width, bottom left) and afterhyperpolarization trough (AHP trough, bottom right) in WT and Kv4.3'. SNc DA neurons, B, left, current-clamp recordings showing the voltage response of SNc DA neurons to 100 and 200pA depolarizing current steps (gray traces) in WT (top, black traces) and Kv4.3<sup>-/-</sup> mice (bottom, red traces). Gray dotted rectangles indicate the interspike interval used to calculate the gain start and gain end on each train of action potentials. Right, box and whisker plots showing the distribution of values for gain start (top left), gain end (top right), spike frequency adaptation (SFA) index (bottom left) and start frequency at 100pA (bottom right) in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. \* p<0.05, \*\* p<0.01, \*\*\* p<0.001. Dotted lines in the box and whisker plots indicate the distribution of data (violin plots). Scale bars: A, top, horizontal 50ms, vertical 20mV; A, bottom, horizontal 2ms, vertical 20mV, horizontal dotted lines indicate -40mV; B, voltage, horizontal 500ms, vertical 40mV; current, vertical 100pA.

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Figure 4. Voltage-clamp analysis of I<sub>A</sub> in wild-type and Kv4.3<sup>-/-</sup> SNc DA neurons. A, voltage-clamp traces showing representative I<sub>A</sub> recordings obtained from a WT (black trace) and a Kv4.3<sup>-/-</sup> SNc DA neuron (red trace) in response to a voltage step to -40mV (gray trace). The small residual current present in the Kv4.3<sup>-/-</sup> mice is blocked by AmmTX3 (inset, orange trace). B, box and whisker plot showing the distribution of values for I<sub>A</sub> amplitude in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. C, box and whisker plot showing the distribution of values for I<sub>A</sub> time constant of

inactivation (I<sub>A</sub> tau) in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. The green dotted rectangle highlights 5 Kv4.3<sup>-/-</sup> outliers displaying unusually large values for I<sub>A</sub> tau. **D**, box and whisker plot showing the distribution of values for I<sub>A</sub> charge in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. The green dotted rectangle highlights 5 Kv4.3<sup>-/-</sup> outliers displaying unusually large values for I<sub>A</sub> charge (same cells as in **C**). **E**, left, scatter plot showing the relationship between I<sub>A</sub> tau and charge in WT (gray dots) and Kv4.3<sup>-/-</sup> SNc DA neurons (red dots). Please note that 5 of the Kv4.3<sup>-/-</sup> measurements lie in the WT region of space (green dotted ellipse). Right, voltage-clamp traces showing one example of the atypical I<sub>A</sub> recording (green trace, corresponding to the large green circle in the scatter plot) encountered in one of the 5 Kv4.3<sup>-/-</sup> outliers highlighted in panels **C** and **D**, compared to the typical recording obtained in Kv4.3<sup>-/-</sup> neurons (red trace, same as in panel **A**, corresponding to the large red circle in the scatter plot). \*\*\* p<0.001. Dotted lines in the box and whisker plots indicate the distribution of data (violin plots). Scale bars in **A** and **E**: horizontal 200ms, vertical InA.

**Figure 5. Kv4.2 channels are expressed by a minority of wild-type and Kv4.3**<sup>-/-</sup> **SNc DA neurons. A**, Top, low-magnification pictures of the SNc showing (from left to right) TH (red), Kv4.3 (green) and DAPI (blue) stainings in WT (top row) and Kv4.3<sup>-/-</sup> mice (bottom row). Bottom, high-magnification pictures of SNc DA neurons showing (from left to right) TH (red), Kv4.3 (green) and DAPI (blue) stainings in WT (top row) and Kv4.3<sup>-/-</sup> mice (bottom row). **B**, effect of the antigen retrieval procedure on Kv4.2 detection in the SNc. Top, low-magnification pictures showing (from left to right) TH (red), Kv4.2 (green) and DAPI staining (blue) without (top row) or with antigen retrieval (bottom row). Bottom, left, high-magnification pictures of the insets depicted in the merged low-magnification picture and corresponding to a Kv4.2-negative

(left) and a Kv4.2-positive SNc DA neuron (middle), for which inset pictures (right) illustrate the region selected to characterize the immunofluorescence profile of Kv4.2 and TH stainings (3μm yellow bar). Bottom, right, immunofluorescence profiles of Kv4.2 (green, top graph), Kv4.3 (green, bottom graph) and TH staining (red, both graphs), showing that both Kv4.2 and Kv4.3 profiles are very similar and strongly suggestive of specific plasma membrane expression. The average profiles (n=5) shown here were defined over 3μm selected regions of (yellow bar above the graph), such as the one shown on the inset pictures on the left. **C**, left, high-magnification pictures showing (from left to right) TH (red), Kv4.2 (green) and DAPI staining (blue) after antigen retrieval in WT (top row) and Kv4.3<sup>-/-</sup> mice (bottom row). Right, expanded view of the Kv4.2-positive cells (1, 2) highlighted in the merged pictures on the left. **D**, bar plot showing the counts of Kv4.2-negative (dark colors) and -positive (light colors) cells observed in the SNc of WT (black bar) and Kv4.3<sup>-/-</sup> mice (red bar). n.s. non-significant. Scale bars: **A**, top row 10μm, bottom row 10μm, inset 2μm; **C**, 10μm.

**Figure 5-1.** Validation of antigen retrieval Kv4.2 immunohistochemistry on neocortical and hippocampal neurons. **A**, low-magnification (top rows) and high-magnification pictures (bottom rows) showing the effect of the antigen retrieval procedure on Kv4.2 immunostaining (green) in the cortex. DAPI staining (blue) is also shown. **B**, low-magnification (top rows) and high-magnification pictures (bottom rows) showing the effect of the antigen retrieval procedure on Kv4.2 immunostaining (green) in the CA1 region of the hippocampus. DAPI staining (blue) is also shown. **A**, top row 100μm, bottom row 10μm; **B**, top row 100μm, bottom row 10μm.

Figure 6. Comparing the alterations in electrophysiological phenotype after acute blockade of Kv4 channels with the Kv4.3<sup>-/-</sup> mouse model. A, current-clamp recordings showing the spontaneous pattern of activity of a WT SNc DA neuron in control condition (black trace, left) and after AmmTX3 application (red trace, right). B, left, line and scatter plot showing the change in spontaneous firing frequency induced by AmmTX3 application in individual WT SNc DA neurons. Right, bar plot comparing the average change in spontaneous firing frequency after AmmTX3 application (left, light colors) or Kv4.3 channel deletion (right, dark colors). C, current-clamp recordings showing the voltage response of a WT SNc DA neuron to a hyperpolarizing current step (bottom gray traces) in control condition (left, black trace) and after AmmTX3 application (right, red trace). D, left, line and scatter plot showing the change in rebound delay induced by AmmTX3 application in individual WT SNc DA neurons. Right, bar plot showing the average change in rebound delay after AmmTX3 application (left, light colors) or Kv4.3 channel deletion (right, dark colors). \*\* p<0.01, \*\*\* p<0.001. Scale bars: A, horizontal 1s, vertical 20mV, horizontal gray dotted lines indicate -60mV; C, horizontal 500ms, vertical 20mV.

Figure 7. Absence of compensatory changes in delayed rectifier and  $I_H$  currents in the  $Kv4.3^{-/-}$  SNc DA neurons. A, properties of the delayed rectifier potassium current ( $I_{KDR}$ ) in SNc DA neurons in WT and  $Kv4.3^{-/-}$  mice. Left, voltage-clamp recordings of  $I_{KDR}$  obtained in a WT (black trace) and a  $Kv4.3^{-/-}$  mouse (red trace) in response to a voltage step to +20mV (gray trace). The peak and steady-state components of  $I_{KDR}$  are indicated by arrowheads. Right, line and

scatter plots representing the average current-voltage relationships of the peak (left) and steady-state I<sub>KDR</sub> (right) obtained from WT (gray dots) and Kv4.3<sup>-/-</sup> SNc DA neurons (red dots). **B**, properties of I<sub>H</sub> in SNc DA neurons in WT and Kv4.3<sup>-/-</sup> mice. Left, voltage-clamp recordings of I<sub>H</sub> obtained in a WT (black traces) and a Kv4.3<sup>-/-</sup> mouse (red traces) in response to increasingly hyperpolarized voltage steps. Right, box and whisker plot showing the distribution of values for I<sub>H</sub> amplitude (left) and voltage sensitivity (right) in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. \*\*\* p<0.001. Dotted lines in the box and whisker plots indicate the distribution of data (violin plots). Scale bars: **A**, horizontal 200ms, vertical 1nA; **B**, horizontal 2s, vertical 500pA.

Figure 8. Linking biophysical changes in I<sub>A</sub> to changes in electrophysiological phenotype in wild-type and Kv4.3<sup>-/-</sup> SNc DA neurons. A, recordings representative of the variation in spontaneous activity, post-inhibitory rebound and I<sub>A</sub> in WT and Kv4.3<sup>-/-</sup> SNc DA neurons. Left, cell-attached voltage-clamp recordings of spontaneous pacemaking activity in 2 WT (black traces) and 2 Kv4.3<sup>-/-</sup> neurons (red traces). The value of the average ISI is indicated above the trace. Middle, current-clamp recordings of the post-inhibitory rebound obtained in the same neurons. The value of rebound delay is indicated above the trace. Right, I<sub>A</sub> voltage-clamp recordings obtained at -40mV in the same neurons. The values of I<sub>A</sub> amplitude and tau are indicated above the trace. B, scatter plot showing the significant positive correlation between extracellularly recorded ISI and rebound delay observed in WT (gray dots) and Kv4.3<sup>-/-</sup> SNc DA neurons (red dots). The plain black and red lines correspond to the linear regression of the data (r, n and p values are shown on the graph), while the gray and pink dotted lines indicate the regression confidence intervals. The diamond and square symbols correspond to the recordings presented in panel A. C, scatter plots showing the relationships between biophysical variables

and neuronal output. Correlations between IA tau, IH amplitude, IA amplitude or IA V50 (from left to right) and rebound delay or extracellular ISI (from top to bottom) were tested in both WT (gray dots) and Kv4.3<sup>-/-</sup> neurons (red dots). Please note that only I<sub>A</sub> tau and I<sub>H</sub> amplitude were significantly correlated with both ISI and rebound delay in WT and Kv4.3<sup>-/-</sup> neurons. The plain black and red lines correspond to the linear regression of the data (r/rho, n and p values are shown on the graph), while the gray and pink dotted lines indicate the regression confidence intervals. Dashed black and red lines indicate non-significant correlations. The diamond and square symbols correspond to the recordings presented in panel A. D, scatter plots showing the significant correlations between I<sub>A</sub> and I<sub>H</sub> properties. Left, scatter plot showing the positive correlation between IA inactivation V50 and IH activation V50 in WT neurons. Right, scatter plot showing the negative correlation between IA tau and IH amplitude observed in both WT (gray dots) and Kv4.3<sup>-/-</sup> neurons (red dots). The plain black and red lines correspond to the linear regression of the data (r/rho, n and p values are shown on the graph), while the gray and pink dotted lines indicate the regression confidence intervals. E, left, scatter plot showing the multiple linear regression of extracellular ISI vs IA tau and IH amplitude (predicted ISI) in WT neurons. Right, scatter plot showing the multiple linear regression of rebound delay vs IA tau, IA inactivation V<sub>50</sub> and I<sub>H</sub> amplitude (predicted rebound delay) in WT neurons. The corresponding equations are indicated below the X axis of each graph. The plain black lines correspond to the linear regression of the data (r, n and p values are shown on the graph), while the gray dotted lines indicate the regression confidence intervals. Scale bars: A, left horizontal 1s; middle horizontal 500ms, vertical 50mV; right horizontal 250ms, vertical 2nA.

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Figure 9. Modeling the effect of the biophysical properties of  $I_{\rm A}$  and  $I_{\rm H}$  on spontaneous activity and post-inhibitory rebound. A, table presenting the 5 values tested for each biophysical parameter (gA, gH, IA/IH V50 and IA inactivation tau) in the multi-compartment model. Each property was varied independently, leading to 625 (54) versions of the model. B, dimensionally stacked heatmaps showing the variation in ISI (displayed as log(ISI), left) and in post-inhibitory rebound delay (displayed as log(rebound delay), right) as a function of g<sub>A</sub> (X axis, big scale), I<sub>A</sub>/I<sub>H</sub> V<sub>50</sub> (Y axis, big scale), I<sub>A</sub> tau (X axis, small scale) and g<sub>H</sub> (Y axis, small scale). C, example traces of spontaneous pacemaking (left) and post-inhibitory rebound (right) obtained for the minimal and maximal values of gA and IA tau (corresponding to the 4 squares surrounded in the panel **B** heatmaps). The gray horizontal bars above the traces help visualize the change in ISI (left) or rebound delay (right) as a function of changes in gA and IA tau. D, multiple linear regression reveals the relative contribution of each biophysical property to ISI (left) or rebound delay variation (right). The scatter plots show the relationship between the measured values of log(ISI) and log(rebound delay) and the values predicted using a linear combination of the 4 biophysical variables listed in panel A (the corresponding equations are shown above each graph). Scale bars: C, horizontal 500ms, vertical 50mV.

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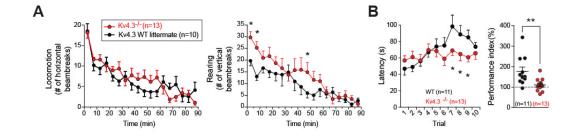
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**Table 1. Statistical analysis of electrophysiological parameters in wild-type and Kv4.3**<sup>-/-</sup> **SNc DA neurons.** The values for 16 electrophysiological parameters measured under current-clamp (corresponding to passive properties, spontaneous activity, post-inhibitory rebound, action potential and excitability) and 8 electrophysiological parameters measured under voltage-clamp (corresponding to I<sub>A</sub> and I<sub>H</sub> properties) are presented for WT and Kv4.3<sup>-/-</sup> SNc DA neurons. Mean and SD (black text) are reported for normally-distributed data, while Median and interquartile

992	range (IQR) are reported otherwise (gray text). Accordingly, statistical differences between WT
993	and Kv4.3 <sup>-/-</sup> neurons were tested using a t-test or a Mann Whitney test, depending on the
994	normality of the data. Asterisks indicate statistically significant differences (* p<0.05, ** p<0.01
995	*** p<0.001).
996	
997	Table 2. Equations governing the voltage dependence and kinetics of currents in the model
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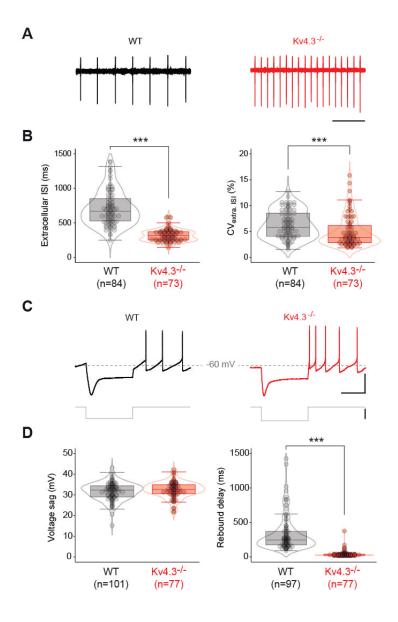


FIGURE 2

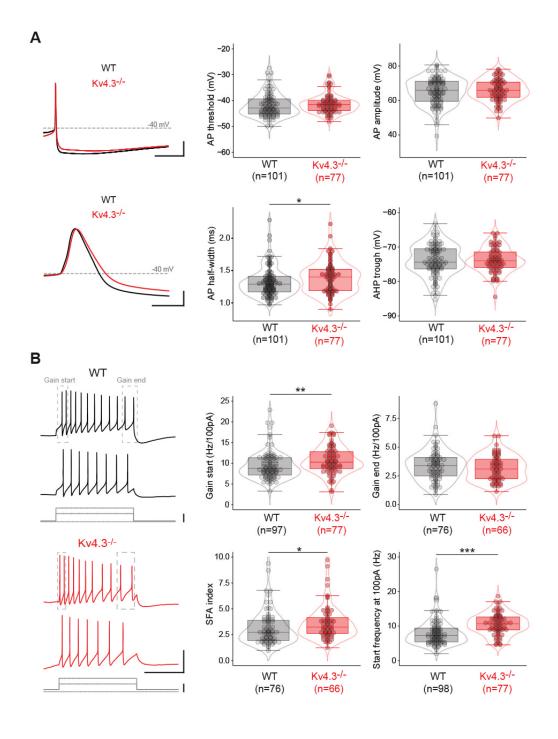


FIGURE 3

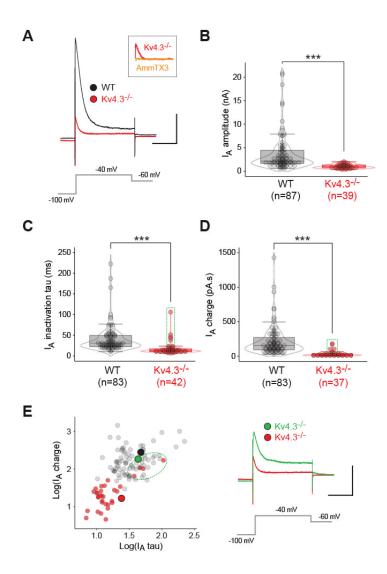


FIGURE 4

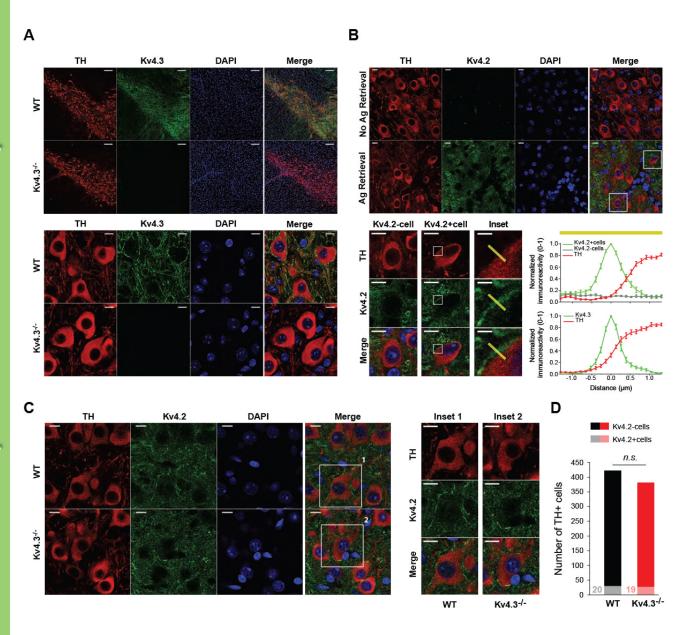


FIGURE 5

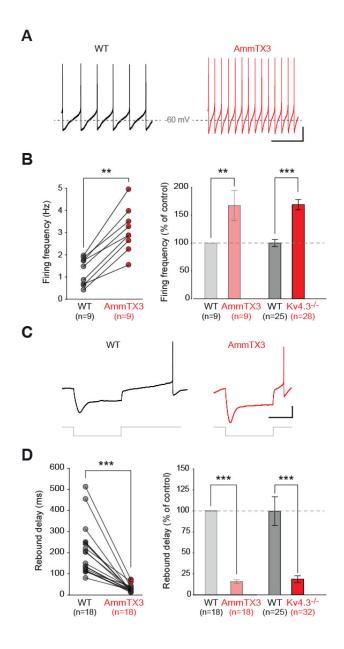


FIGURE 6

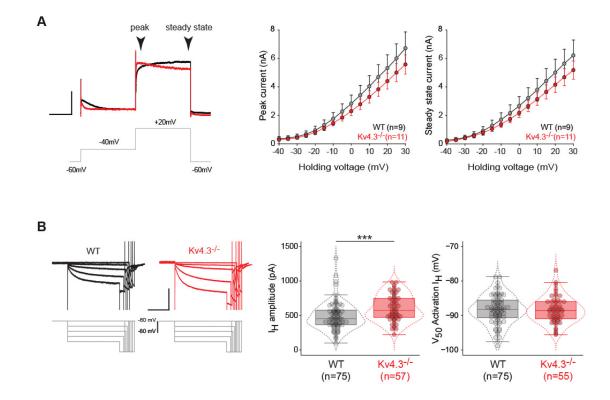


FIGURE 7

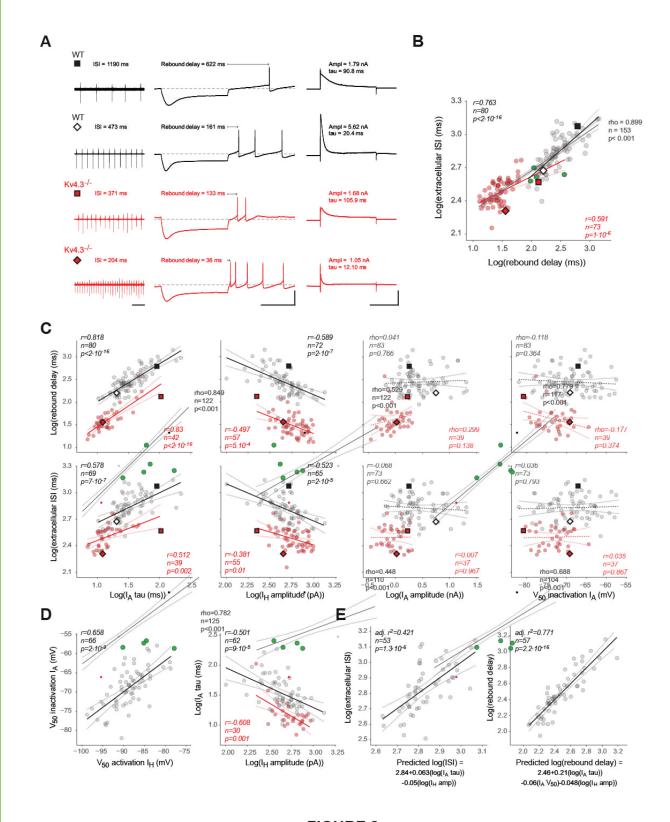


FIGURE 8

Α							
	Parameter	Values					Range
	g <sub>A</sub> (pS/µm <sup>2</sup> )	15	49	83	116	150	10-fold
	g <sub>H</sub> (pS/µm²)	0.25	0.8	1.35	1.9	2.5	10-fold
	I <sub>A</sub> /I <sub>H</sub> V <sub>50</sub> (mV)	-78/-100	-74/-95	-70/-90	-66/-85	-62/-80	20mV
	L. tau (me)	15	40	02	116	150	10 fold

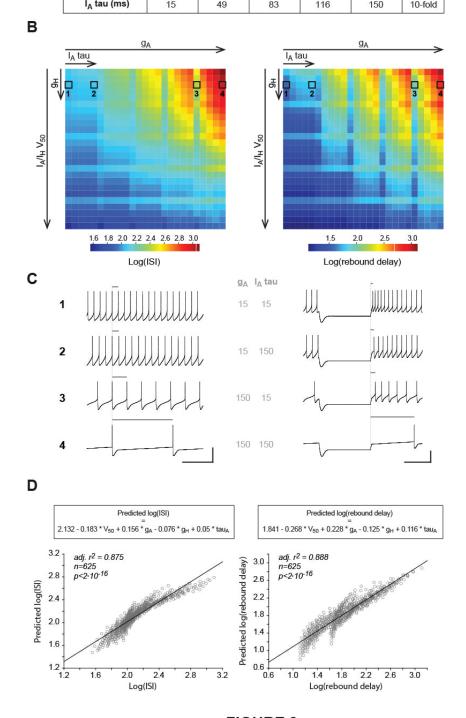


FIGURE 9