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Investigating the role of striatal dopamine receptor 2 in motor coordination and balance: insights into the pathogenesis of DYT1 dystonia

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Abstract

DYT1 or DYT-TOR1A dystonia is early-onset, generalized dystonia. Most DYT1 dystonia patients have a heterozygous trinucleotide *GAG* deletion in *DYT1* or *TOR1A* gene, with a loss of a glutamic acid residue of the protein torsinA. DYT1 dystonia patients show reduced striatal dopamine D2 receptor (D2R) binding activity. We previously reported reduced striatal D2R proteins and impaired corticostriatal plasticity in *Dyt1 GAG* heterozygous knock-in (*Dyt1* KI) mice. It remains unclear how the D2R reduction contributes to the pathogenesis of DYT1 dystonia. Recent knockout studies indicate that D2R on cholinergic interneurons (ChIs) has a significant role in corticostriatal plasticity, while D2R on medium spiny neurons (MSNs) plays a minor role. To determine how reduced D2Rs on ChIs and MSNs affect motor performance, we generated ChI- or MSN-specific D2R conditional knockout mice (*Drd2* ChKO or *Drd2* sKO). The striatal ChIs in the *Drd2* ChKO mice showed an increased firing frequency and impaired quinpirole-induced inhibition, suggesting a reduced D2R function on the ChIs. *Drd2* ChKO mice had an age-dependent deficient performance on the beam-walking test similar to the *Dyt1* KI mice. The *Drd2* sKO mice, conversely, had a deficit on the rotarod but not the beam-walking test. Our findings suggest that D2Rs on ChIs and MSNs have critical roles in motor control and balance. The similarity of the beam-walking deficit between the *Drd2* ChKO and *Dyt1* KI mice supports our earlier notion that D2R reduction on striatal ChIs contributes to the pathophysiology and the motor symptoms of DYT1 dystonia.

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Keywords

DYT1/DYT-TOR1A dystonia; dopamine D2 receptor; cholinergic interneurons; medium spiny neurons; corticostriatal plasticity

1. Introduction

Dystonia is a movement disorder characterized by sustained or intermittent muscle contractions causing abnormal, often repetitive movements, postures, or both (Albanese et al., 2013). It can be caused by genetic mutations, brain injury, or the side effects of drugs. DYT1 or DYT-TOR1A dystonia (OMIM: #128100) is the most common type of early-onset generalized dystonia (Bressman et al., 2000) with symptoms onset from 5 to 28 years old. The majority of the individuals affected by DYT1 dystonia share a trinucleotide deletion (*GAG*) in the exon 5 of the *DYT1* or *TOR1A* gene, leading to a loss of a glutamate amino acid residue for torsinA (torsinA^E) (Klein and Ozelius, 2002). Other mutations in the gene were also reported in rare cases (Leung et al., 2001; Zirn *et al.*, 2008; Ritz et al., 2009; Calakos et al., 2010; Cheng et al., 2014; Harata, 2014; Vulinovic et al., 2014; Dobricic et al., 2015). Moreover, recent reports suggest that homozygous mutation in *DYT1* causes developmental delay and motor deficits (Kariminejad et al., 2017; Reichert et al., 2017; Isik et al., 2019). As a member of the AAA+ (ATPases associated with diverse cellular activities) superfamily (Ozelius et al., 1997), torsinA is expressed at a higher level during early development of the brain (Xiao et al., 2004; Siegert et al., 2005; Vasudevan et al., 2006), especially for the neurons located in the basal ganglia (Augood et al., 1999; Shashidharan et al., 2000). In DYT1 dystonia patients, mutant torsinA might exert a dominant-negative effect and decrease wildtype torsinA activity due to a homo-oligomeric feature of AAA+ proteins (Konakova and Pulst, 2005; Pham et al., 2006; Zhao et al., 2013; Demircioglu et al., 2016; Chase et al., 2017).

It is generally believed that dystonia is not a muscle disease but is caused by brain deficits. Deep brain stimulation (DBS) targeting the globus pallidus internus (GPi) or subthalamic nucleus (STN) ameliorates the dystonia symptom (Sun et al., 2007; Panov et al., 2013; Guzzi et al., 2016; Cury et al., 2018; Wu et al., 2019), suggesting a functional alteration of the basal ganglia circuits. The striatum functions in controlling movement based on the balanced modulation of striatal circuits between cholinergic interneurons (ChIs) and the dopaminergic terminals from the substantia nigra (Stoof et al., 1992; Pisani et al., 2007; Aosaki et al., 2010). Growing evidence suggests that an alteration of the cholinergic systems plays a critical role in the pathophysiology of DYT1 dystonia (Eskow Jaunarajs et al., 2015). Anticholinergic drugs such as trihexyphenidyl (THP) are clinically effective in treating DYT1 dystonia, indicating an abnormal cholinergic tone in DYT1 dystonia. Decreased dopamine D2 receptor (D2R) binding activity and expression level in the striatum have been identified in DYT1 dystonia patients (Augood et al., 2002; Asanuma et al., 2005) and in multiple lines of *Dyt1* rodent models (Napolitano et al., 2010; Yokoi et al., 2011; Dang et al., 2012; Bonsi et al., 2019; Yokoi et al., 2020b), whereas high levels of D2Rs are present on the ChI in wild type mice (Le Moine et al., 1990). Activation of D2R slows the spontaneous firing of ChIs and acetylcholine release (Maurice et al., 2004; Chuhma et al.,

2014). Previous studies showed a paradoxical excitation in response to quinpirole, a D2R agonist, in multiple DYT1 dystonia genetic models (Sciamanna et al., 2011; Grundmann et al., 2012; Sciamanna et al., 2012; Martella et al., 2014; Eskow Jaunarajs et al., 2015; Scarduzio et al., 2017), suggesting D2R alterations contribute to the DYT1 pathophysiology. However, whether the reduction of striatal D2R is the result or cause of DYT1 dystonia is not known.

Animal models are essential for investigating the pathophysiology of human disease and contribute to the development of effective treatments. DYT1 animal models do not have overt dystonia, but they do show impairments in motor coordination and balance with the beam-walking and rotarod tests (Dang et al., 2005; Sharma et al., 2005; Dang et al., 2006a; Grundmann et al., 2007; Yokoi *et al.*, 2008; Zhao et al., 2008; Page et al., 2010; Sciamanna et al., 2012; Song et al., 2012; Yokoi et al., 2015; Sciamanna et al., 2020; Yokoi et al., 2020b). *Dyt1/Tor1a* GAG heterozygous knock-in (*Dyt1* KI) mice have the corresponding in-frame trinucleotide GAG deletion in the endogenous *Dyt1* and model DYT1 patients (Dang et al., 2005; Goodchild et al., 2005). *Dyt1* KI mice exhibit the reduced level of striatal D2R protein (Dang et al., 2012; Bonsi et al., 2019), elevated extracellular acetylcholine (Scarduzio et al., 2017; Downs et al., 2019), impaired long-term depression (LTD) in the corticostriatal pathway (Dang et al., 2012; Maltese et al., 2014; Martella et al., 2014), abnormal sustained co-contractions of agonist and antagonist muscles (DeAndrade et al., 2016), and motor deficits in the beam-walking test (Dang et al., 2005; Song et al., 2012). The deficits in the LTD, beam-walking, and muscle co-contractions can be ameliorated by THP (Dang et al., 2012; Maltese et al., 2014; DeAndrade et al., 2016). Since D2R on ChIs have been identified as a key regulator of the striatal bidirectional synaptic plasticity (Wang et al., 2006; Augustin et al., 2018) and the dopamine/acetylcholine balance (Kharkwal et al., 2016), we proposed a model that links reduced D2R signaling on ChIs, the elevation of acetylcholine level, impairment of LTD, and motor abnormality in DYT1 dystonia (Dang et al., 2012). To further test the role of D2R on the ChIs in the pathogenesis of DYT1 dystonia directly, we used ChI-specific *Drd2* conditional knockout mice (*Drd2* ChKO). We characterized their biochemical, electrophysiological, and behavioral phenotypes. We also analyzed MSN-specific *Drd2* conditional knockout mice (*Drd2* sKO) to explore the role of D2R on MSNs in motor performance. Finally, we characterized an individual line of *Dyt1* KI mice for their motor performance to compare with the *Drd2* ChKO and sKO mice.

2. Materials and methods

All experiments were carried out by investigators blind to the genotypes and in compliance with the ARRIVE guidelines and the National Institutes of Health guide for the care and use of laboratory animals (NIH Publications No. 8023, revised 1978). All animal experiments were approved by the Institution Animal Care and Use Committee at the University of Florida. The mice were housed with *ad libitum* access to food and water with 12 hours-light and 12 hours-dark.

2.1 Animals

Chat-cre mice were obtained by crossing ChAT-IRES-Cre::*flrt-neo-flrt* mice (Jackson Lab. Stock No: 006410) (Rossi et al., 2011) with flippase (FLP) deleter strain (Jackson Lab. Stock No: 003946; ROSA26::FLPe knock-in) (Farley et al., 2000). The recombinase FLP recognizes flippase recognition target (*flrt*) sites and removed *neomycin* cassette in ChAT-IRES-Cre::*flrt-neo-flrt* mice by FLP-mediated recombination (Sauer and Henderson, 1988). *Chat-cre* mice can be used for deleting the floxed gene in ChIs, without disrupting endogenous *Chat* expression (Rossi et al., 2011). *Drd2 loxP*^{-/-} mice (Bello et al., 2011), which have floxed exon 2 of the *Drd2* gene, were imported (Jackson Lab. Stock No: 020631) and bred with *Chat-cre*^{+/-} mice. Recombination between these two *loxP* sites leads to the loss of D2R activity (Bello et al., 2011). *Chat-cre*^{+/-}*Drd2 loxP*^{+/-} (double heterozygous, ChDHet) mice were generated by crossing the *Chat-cre*^{+/-} mice and *Drd2 loxP*^{-/-} mice. *Drd2* ChKO mice, ChDHet mice, and their control (CT) littermates (*Drd2 loxP*^{-/-} or *Drd2 loxP*^{+/-}) were generated by crossing the ChDHet mice with *Drd2 loxP*^{-/-} mice. The MSN-specific *Drd2* KO mice were generated by breeding *Drd2 loxP* mice with *Rgs9-cre* mice (Dang et al., 2006b), to produce *Rgs9-cre*^{+/-}*Drd2 loxP*^{+/-} (sDHet) which were used for breeding with homozygous *Drd2 loxP* mice to generate the sKO, sDHet, and CT littermates. *Dyt1* KI mice (*Tor1a^{tm2Wtd}/J*; Jackson Lab. stock no. 025637) were bred as previously described (Goodchild et al., 2005).

Genotyping for *Drd2* ChKO, *Drd2* sKO, and *Dyt1* KI mice was performed by PCR using a set of *Drd2loxP*F (5'- TCTCCCTCATCTCTGGACTCA -3') and *Drd2loxP*R (5'- TGGGAAAGGGCTACAGCA -3') primers for *Drd2 loxP*, a pair of *creA* (5'- ATCTCCGGTATTGAAACTCCAGCGC-3') and *cre6* (5'- CACTCATGGAAAATAGCGATC- 3') primers for *cre* (Campos et al., 2004), and a set of 22117 (5'-GTGCATCAGAGTGGAGATGC- 3') and 22118 (5'- CAGGAGCTTCAGGTCCATGT- 3') primers for *Dyt1* KI (Goodchild et al., 2005). Tail DNA was used as the template.

A minimum number of animals was used following the principles of the 3Rs (Replacement, Reduction, and Refinement). We further used systematic heterogenization of study samples and conditions recommended by others (Voelkl et al., 2020). The mice were first used for behavioral tests, followed by Western blot analysis and electrophysiology, which were delayed due to the pandemic.

2.2 Behavioral analysis

Motor behaviors were assessed by semi-quantitative assessments, accelerated rotarod tests, and beam-walking, in this order. Mice were allowed to rest for at least 4 days in between the behavioral tests. A group of 8 *Drd2* ChKO, 6 ChDHet, and 11 CT littermates of either sex was tested two times at 3 and 7 months of age (Fig. 4A). Another male cohort of 12 *Drd2* sKO, 6 sDHet, and 11 CT littermates was tested for motor performance at the average age of 6 months (P121-P270). The last group of 21 male *Dyt1* KI and 19 WT littermates was also tested for motor performance at the average age of 4.5 months (P108-P150). All motor behaviors were tested within the last 8 h of the light period after acclimation to a sound-attenuated testing room for 1 h. Body form assessment of motor disorders was performed as

described earlier (Fernagut et al., 2002; Dang et al., 2005). Mice were placed individually on the table. Truncal distortion and balance adjustments to a postural challenge were examined. Truncal distortion was assessed as the flexed posture. The postural challenge was observed by flipping the mouse onto its back, and its capability of righting was noted.

The accelerated rotarod test assesses mice's ability to maintain balance and coordination on an accelerating rotating rod (Ugo Basile) as previously described (Sciamanna et al., 2012). The apparatus started at an initial speed of 4 rpm, and then each mouse was put on the same slot one by one. The rod speed was gradually increased at a rate of 0.2 rpm/s. The latency to fall was measured with a cutoff time of 3 min at a final rate of 40 rpm. Mice were tested for three trials at about 1 h intervals on each day for 2 days.

The beam-walking test was performed as described earlier (Dang et al., 2005; Dang et al., 2012). The mice were trained to transverse a medium square beam (14 mm wide) in three consecutive trials each day for 2 days, and they were tested twice each on the medium square beam and a medium round beam (17 mm diameter) on the third day. The mice were then tested twice, each on a small round beam (10 mm diameter) and a small square beam (7 mm wide) on the fourth day. The number of hind paw slips on each side was counted by investigators blind to the genotypes. All 4 beams were 100 cm long, and the slips traversing the middle 80 cm were counted.

2.3 Western blot analysis

Two groups of mice were used for Western blot analysis. 5 *Drd2* ChKO mice, 3 ChDHet mice, and 5 controls of either sex at an average of 10 months of age were used in the ChKO group. Moreover, 4 *Drd2* sKO male mice with its 4 sDHet, and 5 control littermates at an average of 10 months of age were used in the sKO group. The striata were dissected and were homogenized in 200 μ l of ice-cold lysis buffer, and the proteins were extracted in 1% Triton X-100 buffer and quantified as previously described (Yokoi et al., 2020a; Yokoi et al., 2020b). Protein concentration was measured by Bradford assay (Bradford, 1976). Protein samples were separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to Millipore Immobilon-FL transfer polyvinylidene difluoride (PVDF) membranes. The membranes were processed as previously described (Yokoi et al., 2015). The PVDF membranes were blocked with LI-COR Odyssey blocking buffer and incubated at 4°C overnight with goat anti-choline acetyltransferase (ChAT) antibody (EMD Millipore, AB144P; 1:1,000 dilution), rabbit anti-acetylcholinesterase (AChE) antibody (Santa Cruz, sc-11409; 1:1,000 dilution), mouse Anti-D2R antibody (Santa Cruz, sc-5303; 1:500 dilution), or rabbit anti-glyceraldehyde-3-phosphate dehydrogenase (GAPDH) antibody (Santa Cruz, sc-25778; 1:1,000 dilution). LI-COR IRDye 800CW donkey anti-goat IgG (H+L) or LI-COR IRDye 680RD donkey anti-rabbit IgG (H+L) were used as appropriate at a dilution of 1:15,556. The signals were captured with an LI-COR Odyssey imaging system and analyzed with Image Studio Lite Ver. 5.2. The density of the corresponding band signals was normalized to those of GAPDH. GAPDH is a reasonable western blot loading control in neuronal tissues (Bauer et al., 2009; Bangaru et al., 2012). GAPDH was chosen in the current study because of the proper molecular weight and stability between ages (Yu et al., 2011). Western blot analysis was performed in duplicate.

2.4 Electrophysiology

Electrophysiological recordings of striatal ChIs were obtained from 3 *Drd2* ChKO, 4 ChDHet, and 2 CT littermates of either sex at an average of 12 months of age, as described previously (Lyu et al., 2019). The animals were anesthetized by the inhalation of isoflurane, decapitated, and the brains were rapidly removed and briefly chilled in an ice-cold cutting solution containing (in mM) 190 Sucrose, 2.5 KCl, 1.25 NaH₂PO₄, 25 NaHCO₃, 1 CaCl₂, 10 MgCl₂, and 10 D-glucose and was oxygenated with 95% O₂-5% CO₂ (pH 7.35~7.45). Coronal brain slices 300 μm-thick were cut with a Vibratome (LEICA VT 1000S, Leica Microsystems, Wetzlar, Germany) in the same ice-cold cutting solution. Slices were first incubated on a brain slice keeper (AutoMate Scientific, Inc. Berkeley, CA) and covered by a thin layer of artificial cerebrospinal fluid (ACSF) containing (in mM) 127 NaCl, 2.5 KCl, 1.25 NaH₂PO₄, 25 NaHCO₃, 2 CaCl₂, 1 MgCl₂, and 10 D-glucose and were constantly oxygenated with 95% O₂ and 5% CO₂ (pH 7.35~7.45) at 35°C for 60 min, followed by incubation at room temperature. After incubation of at least 60 min, a slice was then transferred to a submerged recording chamber with the continuous flow (2 ml/min) of ACSF. All experiments were carried out at 32°C under visual guidance using an inverted microscope equipped with infrared differential interference contrast (IR-DIC) videomicroscopy (Axioskop-FS; Carl Zeiss, Jena, Germany) and a 40× water-immersion lens.

Cholinergic interneurons were identified based on morphology and size, as they are irregularly polygonal with large cell soma (>20 μm), and were further confirmed by characteristic electrophysiological properties observed in whole-cell current-clamp recordings (Oswald et al., 2009). The patch electrodes had a resistance of 5-10 MΩ when filled with a K-gluconate-based solution containing the following intracellular solution (in mM): 112.5 K-gluconate, 4 NaCl, 17.5 KCl, 0.5 CaCl₂, 5 MgATP, 1 NaGTP, 5 EGTA, 10 HEPES, 0.1% Biocytin; with pH 7.2 (osmolality 270-280 mOsm/l). While approaching the cell, positive pressure was applied to the patch electrode. The seal (> 5 GΩ) between the recording pipette and the cell membrane was obtained by applying suction to the electrode. The spontaneous action potentials were recorded in the voltage-clamp mode that maintained an average of 0 pA holding current with an Axopatch 1D amplifier (Axon Instruments, Foster City, CA). After recording the baseline with cell-attached mode, subsets of the neurons were investigated further with the bath application of D2R agonist quinpirole (10 μM, 90 seconds) followed by a washout for 5 min with ACSF. ChIs were recovered with an average recovery time of 2 min.

The recording data was acquired using pClamp 10 software and further analyzed by the Mini Analysis Program (Synaptosoft). The electrophysiological recordings and analysis were performed by investigators who were blind to the genotypes. The spontaneous firing rates of 30 seconds before quinpirole treatment and the last 30 seconds during the application of quinpirole were compared to determine the quinpirole effect. Signals were filtered at 5 kHz and digitized at 10 kHz with a DigiData 1440 (Molecular Devices).

2.5 Statistics

All data were tested for normality using the JMP Pro software. Electrophysiological recording data were analyzed using the student's t-test. The Western blot signals and the latency to fall in the accelerated rotarod test were analyzed by either the mixed model ANOVA (SAS) for normally distributed data or GENMOD procedure (SAS) with a log link for gamma distribution for not normally distributed data (Yokoi et al., 2020b). Slip numbers of the medium round, small round, and small square beams in the beam-walking test were analyzed by using the GENMOD procedure (SAS) with a negative binomial distribution (Dang et al., 2005). Sex, age, body-color, and body weight were used as covariates in all behavioral analyses. Two-way interactions between genotypes and other covariates were explored before final analysis. Significance was assigned at $p = 0.05$.

3. Results

3.1 Generation of *Drd2* ChKO mice

To investigate whether decreased D2R on ChIs causes motor deficits similar to *Dyt1* KI mice (Dang et al., 2005; Dang et al., 2012), *Drd2* ChKO was generated as described (Figure 1). The *Drd2* ChKO mice were obtained with Mendel's ratio ($p = 0.999$, Chi-Square test), suggesting that *Drd2* ChKO mice are neither embryonic nor neonatal lethal. *Drd2* ChKO mice grew up to adulthood without noticeable developmental delay.

3.2 Impaired quinpirole mediated inhibition in *Drd2* ChKO mice

To assess the knockout of D2R, we measured the spontaneous firing properties of the ChIs in *Drd2* ChKO and its modulation by D2R agonist quinpirole. The firing frequency in both *Drd2* ChKO and ChDHet mice was significantly increased compared with control mice (Fig. 2B, Mean \pm S.E.M.: CT: 2.31 ± 0.45 Hz, $n = 11$ cells/2 mice; ChKO: 4.84 ± 0.69 Hz, $n = 12$ cells/3 mice; ChDHet: 4.11 ± 0.70 Hz, $n = 9$ cells/4 mice; CT and ChKO: $p = 0.006$; CT and ChDHet: $p = 0.0497$; ChKO with ChDHet: $p = 0.47$, student's t-test). After quinpirole treatment, the firing frequency was significantly decreased in all genotypes (Fig. 2C, After quinpirole application: Mean \pm S.E.M.: CT: 1.01 ± 0.36 Hz, $p = 0.005$; ChKO: 3.49 ± 0.60 Hz, $p = 0.001$; ChDHet: 3.03 ± 0.58 Hz, $p = 0.013$, student's t-test). However, the ratio of the firing frequency (after/prior treatment) showed a significant increase in *Drd2* ChKO and ChDHet mice compared with CT mice (Fig. 2D, Mean \pm S.E.M.: CT: 0.38 ± 0.11 ; ChKO: 0.68 ± 0.06 ; ChDHet: 0.74 ± 0.08 ; CT and ChKO: $p = 0.03$; CT and ChDHet: $p = 0.02$; ChKO with ChDHet: $p = 0.55$, student's t-test). There was no change of resting membrane potential of ChIs in *Drd2* ChKO and ChDHet mice compared to the control mice (Supplemental Figure 1). These results indicate the successful knockout of D2Rs on ChIs.

3.3 Normal striatal cholinergic enzyme levels in *Drd2* ChKO mice

To examine whether selective knockout of D2R on ChIs leads to an altered striatal cholinergic system, we determined the protein level of the striatal ChAT and AChE, which are two key metabolic enzymes of acetylcholine (Fig. 3A–B). There were no significant difference in either ChAT (Fig. 3A, Mean \pm S.E.M.: CT: 0.16 ± 0.012 ; ChKO: 0.14 ± 0.012 ; ChDHet: 0.12 ± 0.017 ; CT and ChKO: $p = 0.34$; CT and ChDHet: $p = 0.11$; ChKO with

ChDHet: $p = 0.40$, $F_{[2,11]} = 1.57$, $p = 0.25$, ANOVA) or AChE (Fig. 3B, Mean \pm S.E.M.: CT: 1.95 ± 0.096 ; ChKO: 2.14 ± 0.096 ; ChDHet: 1.81 ± 0.16 ; CT and ChKO: $p = 0.21$; CT and ChDHet: $p = 0.47$; ChKO with ChDHet: $p = 0.11$, $F_{[2,11]} = 1.80$, $p = 0.21$, ANOVA) between *Drd2* ChKO mice ($n = 5$), ChDHet ($n = 3$), and *Drd2 loxP* mice ($n = 5$). The results indicate that there was no ChI neurodegeneration in *Drd2* ChKO striatum, which is consistent with most of the DYT1 dystonia mouse models.

3.4 Reduced striatal D2R levels in *Drd2* sKO mice

To assess the knockout of D2R in MSNs, we measured the protein level of strial D2R in *Drd2* sKO mice ($n = 4$), which showed significantly reduced D2R expression level compared with their sDHet ($n = 4$) and control ($n = 5$) littermates (Fig. 3C, Mean \pm S.E.M.: CT: 0.20 ± 0.01 ; sKO: 0.14 ± 0.0096 ; sDHet: 0.24 ± 0.04 ; CT and sKO: $p < 0.001$; CT and sDHet: $p = 0.41$; sKO with sDHet: $p = 0.003$, GENMOD with a gamma distribution). The results indicate the successful knockout of D2Rs on MSNs.

3.5 Impaired motor coordination and balance in *Drd2* ChKO mice

We next determined the motor performance of the *Drd2* ChKO mice. *Drd2* ChKO mice and ChDHet mice exhibited strong righting reflexes when tipped on their side and did not show any overt dystonic behavior in the body form assessment. We further performed rotarod and beam-walking tests at 3 months of age. There were no differences between *Drd2* ChKO mice ($n = 8$), control mice ($n = 11$), and the ChDHet mice ($n = 6$) on the rotarod (Fig. 4B–C) and beam-walking tests (Fig. 4D). However, at 7 months of age, *Drd2* ChKO and ChDHet mice displayed excessive slips on the beam-walking test (Fig. 4G, Mean \pm S.E.M.: CT: 1.02 ± 0.26 slips; ChKO: 2.27 ± 0.51 slips; ChDHet: 2.27 ± 0.44 slips; CT and ChKO: $p = 0.007$; CT and ChDHet: $p = 0.001$; ChKO with ChDHet: $p = 0.988$, GENMOD with a negative binomial distribution), suggesting an age-dependent impairment in motor performance. There was no significant difference between *Drd2* ChKO mice, ChDHet mice, and control mice on the accelerated rotarod test across 6 trials (Fig. 4E–F, Mean \pm S.E.M.: CT: 145.25 ± 16.82 sec; ChKO: 129.82 ± 11.83 sec; ChDHet: 125.39 ± 23.25 sec; CT and ChKO: $p = 0.30$; CT and ChDHet: $p = 0.46$; ChKO with ChDHet: $p = 0.87$, GENMOD with a gamma distribution).

3.6 Impaired motor coordination and balance in the *Drd2* sKO mice

MSNs can regulate the activity of ChIs (Lim et al., 2014). D2Rs on MSNs in the indirect pathway also contribute to synaptic plasticity in the striatum (Wang et al., 2006; Shen et al., 2008; Augustin et al., 2018). To further study the role of striatal D2R in the DYT1 dystonia, we generated MSN-specific *Drd2* conditional knockout mice to test motor coordination and balance. *Drd2* sKO mice were bred with the same breeding strategy with *Drd2* ChKO (Fig. 1) by using *Rgs9-cre* mice that induce recombination restricted to MSNs (Dang et al., 2006b; Lyu et al., 2019). *Drd2* sKO mice displayed significantly reduced latency to fall on the rotarod test across 6 trials (Fig. 5A–B, Mean \pm S.E.M.: CT: 111.44 ± 7.94 sec; sKO: 82.00 ± 8.13 sec; sDHet: 112.18 ± 9.32 sec; CT and sKO: $p = 0.018$; CT and sDHet: $p = 0.95$; sKO and sDHet: $p = 0.021$; $F_{[2,29]} = 3.89$, $p = 0.032$, ANOVA). Conversely, there was no difference between *Drd2* sKO mice, the sDHet mice, and the control littermates on the beam-walking test (Fig. 5C, Mean \pm S.E.M.: CT: 6.44 ± 2.20 slips; sKO: 7.09 ± 1.81 slips;

sDHet: 3.88 ± 0.85 slips; CT and sKO: $p = 0.83$; CT and sDHet: $p = 0.30$; sKO with sDHet: $p = 0.09$, GENMOD with a negative binomial distribution).

3.7 Impaired motor coordination and balance in the *Dyt1* KI mice

To further examine the altered motor coordination and balance we observed in *Drd2* ChKO, *Drd2* sKO, and our *Dyt1* KI mice (Dang et al., 2005), we performed rotarod and beam walking tests in another *Dyt1* knock-in model with the trinucleotide deletion commonly seen in DYT1 patients (Goodchild et al., 2005). In the accelerated rotarod test, the *Dyt1* KI mice showed no significant difference compared to WT littermates across all 6 trials (Fig. 6B, Mean \pm S.E.M.: WT: 115.08 ± 6.29 sec; *Dyt1* KI: 100.81 ± 7.06 sec; $p = 0.16$, GENMOD with a gamma distribution), but they had a significantly shorter time to fall in the 3 trials on the first day (Fig. 6A, $p = 0.048$). With the beam-walking test, the *Dyt1* KI mice displayed a 119% increase of slips over their WT littermates (Fig. 6C, Mean \pm S.E.M.: WT: 0.23 ± 0.07 slips; *Dyt1* KI: 0.51 ± 0.12 slips; $p = 0.016$, GENMOD with a negative binomial distribution). Taken together, the *Dyt1* KI mice displayed significant motor coordination and balance deficits.

4. Discussion

Both animal models and DYT1 dystonia patients show decreased striatal D2R binding or receptor protein levels. The D2R reduction appears to occur also in DYT6 and DYT11 dystonia (Beukers et al., 2009; Carbon et al., 2009; Zhang et al., 2012; Frederick et al., 2019). However, whether D2R reduction causes dystonia or is a consequence of dystonia is not clear. This is complicated further by the multiple locations of D2R in the striatum. D2Rs are present on at least five different types of neurons or axon terminals in the striatum: indirect-pathway MSNs, ChIs, a subset of GABA interneurons, and afferents to the striatum from midbrain dopaminergic neurons and prefrontal cortical neurons (Dobbs et al., 2017). Compared with the pharmacological methods, which may easily affect all these locations, genetic mouse models can precisely knockout D2Rs only on one subset of neurons. Here, we used genetic approaches to investigate the role of D2R reduction on ChIs and MSNs in the pathogenesis of DYT1 dystonia. We found that selective deletion or reduction of D2Rs from ChIs resulted in increased ChI firing rate, impaired quinpirole-mediated inhibition, and age-dependent motor deficits in beam-walking, which is a characteristic motor deficit found across multiple lines of DYT1 and other dystonia mouse models. Interestingly, knocking out D2Rs, but not reduction, from MSNs, produced a rotarod deficit, a motor deficit found in several DYT1 dystonia mouse models. Finally, rotarod and beam walking tests in another line of *Dyt1* KI mice showed similar motor deficits as the *Drd2* ChKO mice. Overall, these results indicate that knocking out or reduction of D2Rs in the ChI alone could lead to impaired coordination and balance, similar to *Dyt1* KI mice. Our results highlight the importance of D2R activity on ChIs to the pathogenesis of DYT1 dystonia.

Selective knockout or reduction of D2Rs on ChI led to age-dependent beam-walking deficits similar to *Dyt1* KI mice. Excessive slips were apparent only in 6-month-old *Dyt1* KI mice, not in 3-month-old mice (Dang et al., 2005). The age-dependent motor deficits have been shown in various DYT1 animal models (reviewed in Richter and Richter, 2014). Even

though it is challenging to compare mice's life phases with humans, a recent study suggested that 3-month-old mice are equivalent to 21-year-old humans while 7-month-old mice are equivalent to 30-year-old humans (Wang et al., 2020). Therefore the age-dependent beam-walking deficits suggest that motor deficits start to develop between 3 to 6 months of age. This is consistent with the human study that most DYT1 dystonia symptoms begin before the age of 26 years old (Opal et al., 2002). *Drd2* ChKO and ChDHet mice did not show a deficit on the rotarod, which is also consistent with the normal rotarod performance in the *Dyt1* KI mice (Dang et al., 2005) and the *Dyt1* KI line (Goodchild et al., 2005) used in this study. ChDHet mice displayed a beam-walking deficit suggesting that reduction of D2R expression on ChIs is sufficient to induce typical coordination and balance deficits in *Dyt1* KI mice. Overall, the similarity of the motor performance between *Drd2* ChKO, ChDHet, and both two *Dyt1* KI mouse lines supports what we proposed in an earlier study that reduced D2R on ChIs leads to impaired motor performance in DYT1 dystonia (Dang et al., 2012). There are limitations to this study. *Chat-cre* line can also affect cholinergic neurons outside of the striatum like pedunculo-pontine, laterodorsal tegmental nuclei, basal forebrain, and the spinal cord (Wang and Morales, 2009; Ballinger et al., 2016). However, D2R expression on those cholinergic neurons is low, and the contribution to the motor phenotypes remains to be investigated.

Decreased D2R binding activity and expression level in the striatum has been found in DYT1 dystonia patients and animal models (Augood et al., 2002; Asanuma et al., 2005; Napolitano et al., 2010; Yokoi et al., 2011; Dang et al., 2012; Bonsi et al., 2019; Yokoi et al., 2020b). Besides, reduced D2R has been found in other dystonia patients and animal models, such as DYT6 (Carbon et al., 2009), DYT11 (Beukers et al., 2009; Zhang et al., 2012), cranial or hand dystonia (Perlmutter et al., 1997), and other related movement disorders like Parkinson's disease (Albin et al., 1989). Furthermore, the *DRD2* variants are related to myoclonus dystonia (Klein et al., 1999) and cervical dystonia (van der Weijden et al., 2020), highlighting that D2R is critical in motor control.

ChIs only account for 1 to 2% of striatal neurons, and most neurons in the striatum are MSNs. We further explored motor performance when D2Rs were selectively manipulated in the MSNs. *Drd2* sKO and sDHet mice did not display overt dystonia, consistent with other animals with striatal D2R reduction (Lemos et al., 2016; Shana et al., 2020). Furthermore, we found that *Drd2* sKO and sDHet mice did not have a beam-walking deficit. Conversely, *Drd2* sKO, but not sDHet mice, showed a rotarod deficit. Interestingly, other groups used *Adora2A-cre* transgenic mice instead of *Rgs9-cre* to direct *cre*-mediated recombination in indirect-pathway MSNs (Lemos et al., 2016; Shana et al., 2020), and the knockout mice showed impaired performance on the rotarod test. These results suggest that indirect pathway MSN-specific D2R knockout mice have a rotarod deficit, similar to a subset of *Dyt1* dystonia models (Sharma et al., 2005; Grundmann et al., 2007; Sciamanna et al., 2012; Sciamanna et al., 2020; Yokoi et al., 2020b). We conclude that D2R on MSNs also contributes to the coordination and balance deficits in *Dyt1* KI mice. It is interesting that MSNs also modulate nociception (Rooney and Sewell, 1989; Zarrindast and Moghaddampour, 1989; Verma and Kulkarni, 1993; Wall et al., 2013), and *Dyt1* KI mice show sensory deficit (Richter et al., 2017). Alteration of sensory perception will likely contribute to the motor performance in the rotarod and beam walking tests.

Rotarod and beam-walking tests are useful for determining motor coordination and balance deficits (Buccafusco, 2009). However, the challenge to animals is slightly different. Rotarod readout includes all 4 limbs, while the beam walking test is only focused on the hindlimbs. Furthermore, a previous study showed that only a 30% GABA-A receptor occupancy by benzodiazepine agonists was needed in order to observe motor deficits on the beam compared to over 70%-90% receptor occupancy for deficits on the rotarod (Stanley et al., 2005). Therefore, as motor coordination and balance measurements, the beam walking test is more sensitive than the rotarod test and is ideal for detecting subtle motor deficits in dystonia animal models. In the current study, knocking out D2R on ChI altered the firing of ChI, which in turn affected the modulatory effect of acetylcholine on both direct and indirect pathway MSNs. Knocking out D2R on MSNs, on the other hand, only affected the modulatory effect of DA on the indirect pathway MSNs. These differences may account for the different motor performance we observed in *Drd2* ChDHet and sDHet mice.

In our previous studies, we found impaired corticostriatal LTD and sustained co-contraction of hindlimb muscles in *Dyt1* KI mice. THP can restore LTD and beam-walking deficits and decrease muscle co-contraction (Dang et al., 2012; DeAndrade et al., 2016). These results suggest that impaired corticostriatal LTD could be correlated with the motor deficits in *Dyt1* KI mice. Interestingly, a recent study using *Drd2* conditional knockout mice replicated the LTD deficit found in *Dyt1* KI mice and showed that D2Rs on ChIs plays a major role in LTD, while the D2Rs on the indirect pathway MSNs only have a weaker modulatory effect on LTD induction (Augustin et al., 2018). Using similar *Drd2* conditional knockout mice, we recapitulated the motor coordination and balance deficits found in *Dyt1* KI mice.

Together, these results suggest that knocking out of striatal D2Rs alone in wild type mice could mimic the motor phenotypes of *Dyt1* KI mice, highlighting the importance of D2R in DYT1 pathophysiology. These results also further support that corticostriatal LTD deficits contribute to motor coordination and balance deficits in *Dyt1* KI mice (Dang et al., 2012). Correcting the corticostriatal LTD deficits might be able to treat the DYT1 dystonia patients (Maltese et al., 2014; Martella et al., 2014; Maltese et al., 2018). Interestingly, enhanced eIF2 α signaling can restore the impaired corticostriatal LTD in *Dyt1* KI mice and improve neonatal survival of *Dyt1* homozygous KI mice (Rittiner et al., 2016). The eIF2 α pathway integrates stress response. It is required for long-term synaptic plasticity (Trinh and Klann, 2013) by regulating the ATF4 and the CREB-dependent transcription and downstream synaptic plasticity-related genes. Therefore, it will be interesting to investigate whether correcting LTD deficit via either the eIF2 α signaling pathway or D2Rs on ChIs can rescue the motor deficits in *Dyt1* KI mice and provide symptom relief in DYT1 patients.

5. Conclusion

Reduced striatal D2R is found in both DYT1 patients and animal models. Selective knocking out or reduction of D2R on ChI alone led to the same age-dependent beam walking deficit observed in *Dyt1* KI mice. In contrast, the removal of D2R, but not the reduction of D2R, on MSNs led to the rotarod deficit found in a small subset of DYT1 animal models. The results suggest that D2R on ChIs is critical in the pathophysiology of DYT1 dystonia.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

AAA+	ATPases associated with diverse cellular activities
AChE	acetylcholinesterase
ACSF	artificial cerebrospinal fluid
ChAT	choline acetyltransferase
ChDHet	<i>Chat-cre Drd2 loxP</i> double heterozygote mice
ChI	cholinergic interneuron
CT	control
D2R	dopamine D2 receptor
DBS	deep brain stimulation
<i>Drd2</i> ChKO	<i>Drd2</i> ChI-specific KO mice
<i>Drd2</i> sKO	<i>Drd2</i> MSN-specific KO mice
<i>Dyt1</i> KI mice	<i>Dyt1</i> GAG heterozygous knock-in mice
GAPDH	glyceraldehyde-3-phosphate dehydrogenase
Gpi	globus pallidus internus
KO	knockout
LTD	long-term depression
MSN	medium spiny neuron
PVDF	polyvinylidene difluoride
sDHet	<i>Rgs9-cre Drd2 loxP</i> double heterozygote mice
SDS-PAGE	sodium dodecyl sulfate-polyacrylamide gel electrophoresis

STN	subthalamic nucleus
THP	trihexyphenidyl
WT	wild type

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Highlights

1. Lack or reduction of D2R on ChIs led to an age-dependent beam-walking deficit.
2. Lack or reduction of D2R on ChIs mimicked the motor deficit of *Dyt1* KI mice.
3. Lack or reduction of D2R on ChIs led to the increased spontaneous firing of ChIs.
4. Lack or reduction of D2R on ChIs in WT mice attenuated quinpirole response.
5. Loss of D2R on medium spiny neurons in WT mice led to a rotarod deficit.

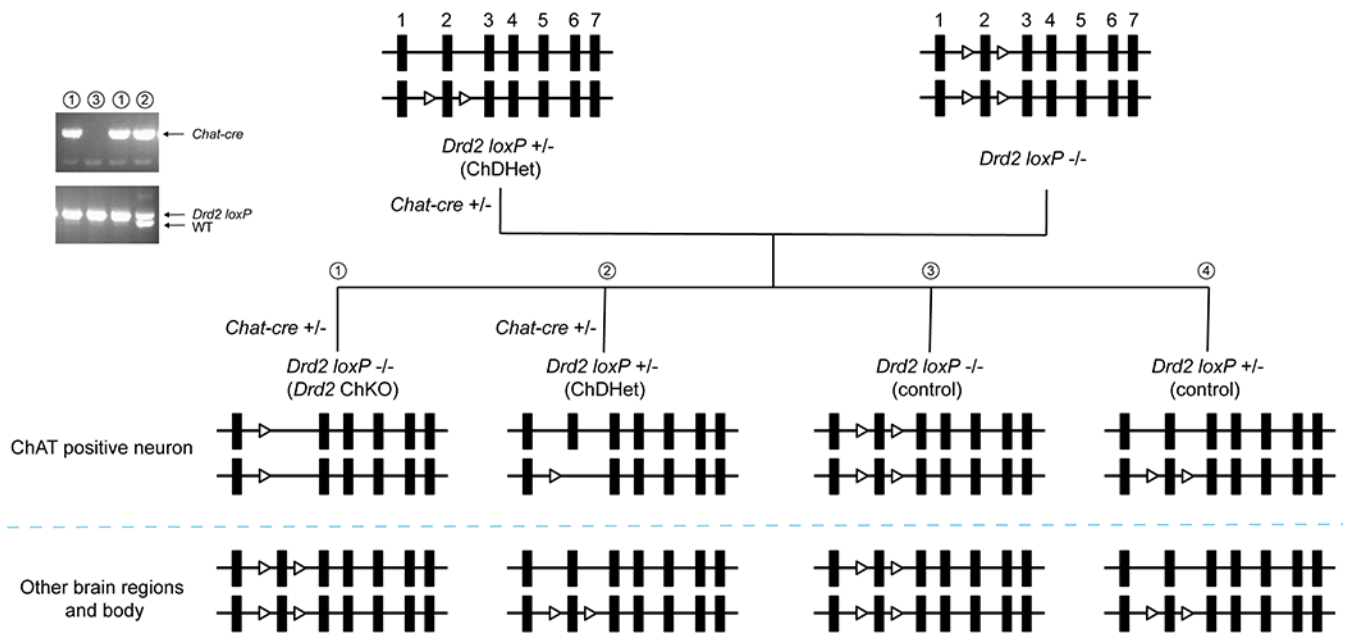


Figure 1. Schematic diagram of the generation of *Drd2* ChKO mice. *Chat-cre*^{+/-}*Drd2 loxP*^{+/-} (ChDHet) mice were crossed with *Drd2 loxP*^{-/-} mice to generate ① *Chat-cre*^{+/-}*Drd2 loxP*^{-/-} (*Drd2* ChKO), ② ChDHet, ③ *Drd2 loxP*^{-/-}, and ④ *Drd2 loxP*^{+/-} mice. PCR-based tail DNA genotyping of *Chat-cre* and *Drd2 loxP* loci (top left).

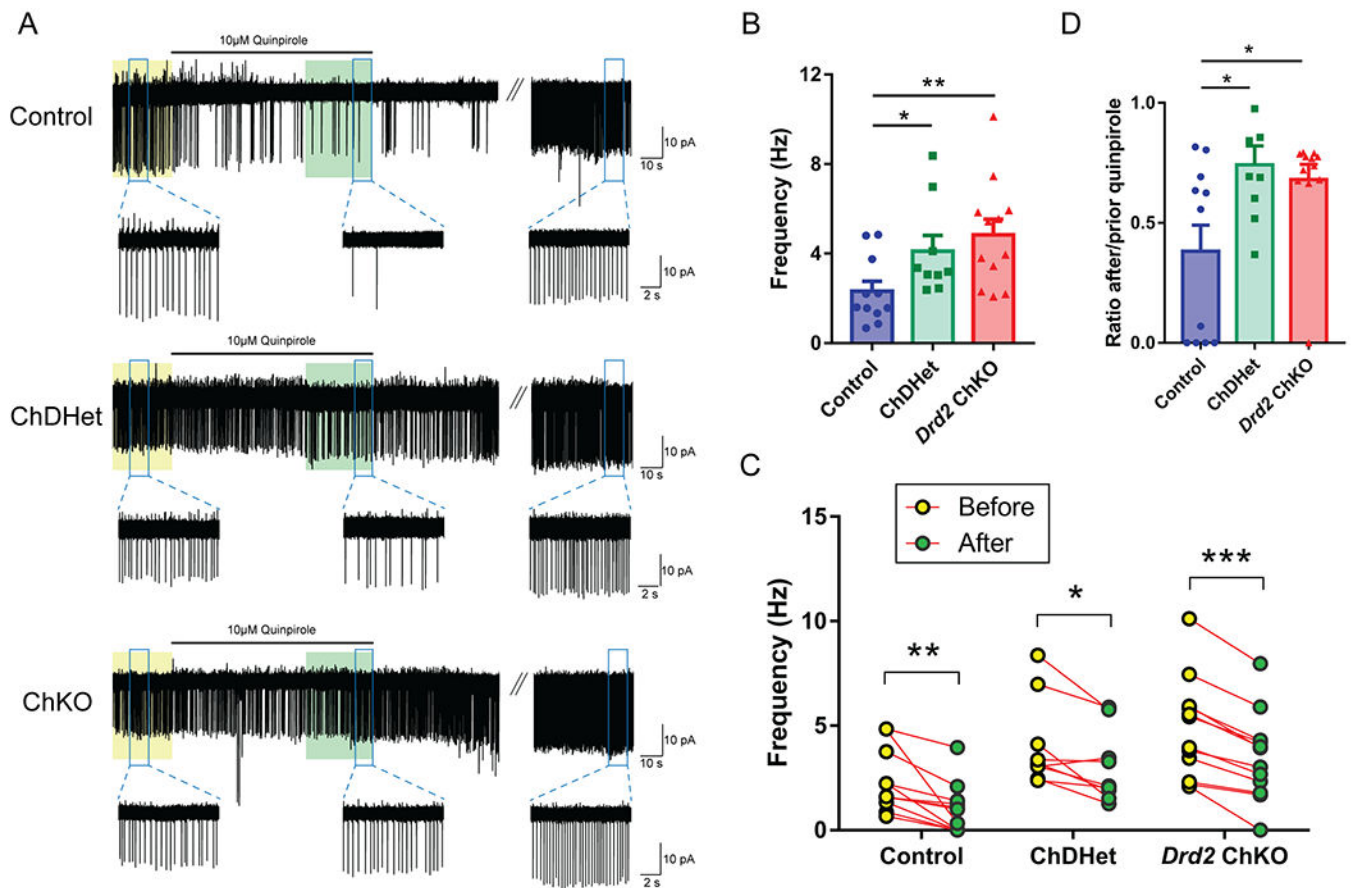


Figure 2. Electrophysiological characterization of D2R function in *Drd2* ChKO mice. (A) Representative recording traces before and after the bath application of the D2R agonist quinpirole (10 μ M, 90 seconds). The yellow area indicates the 30 seconds used to determine spontaneous firing property before drug application. The green area indicates 30 seconds used to determine the frequency after 60 seconds of drug application. (B) Increased spontaneous firing frequency of ChIs in *Drd2* ChKO and ChDHet compared to control mice before quinpirole treatment. (C) All genotypes showed a reduced firing rate after the quinpirole application. (D) *Drd2* ChKO and ChDHet mice showed significantly increased frequency ratios of after/prior quinpirole over the control mice. Vertical bars represent means \pm standard errors. * p 0.05, ** p 0.01, *** p 0.001.

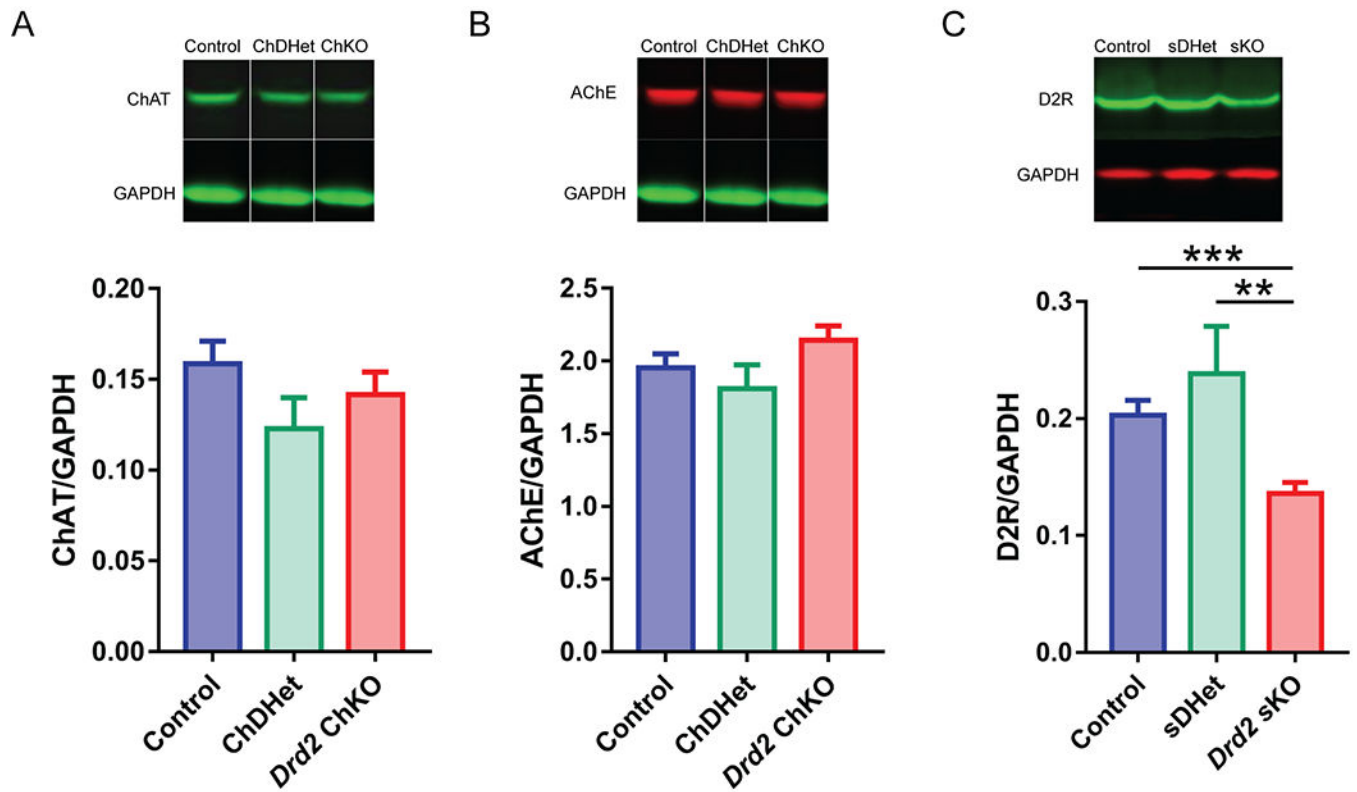
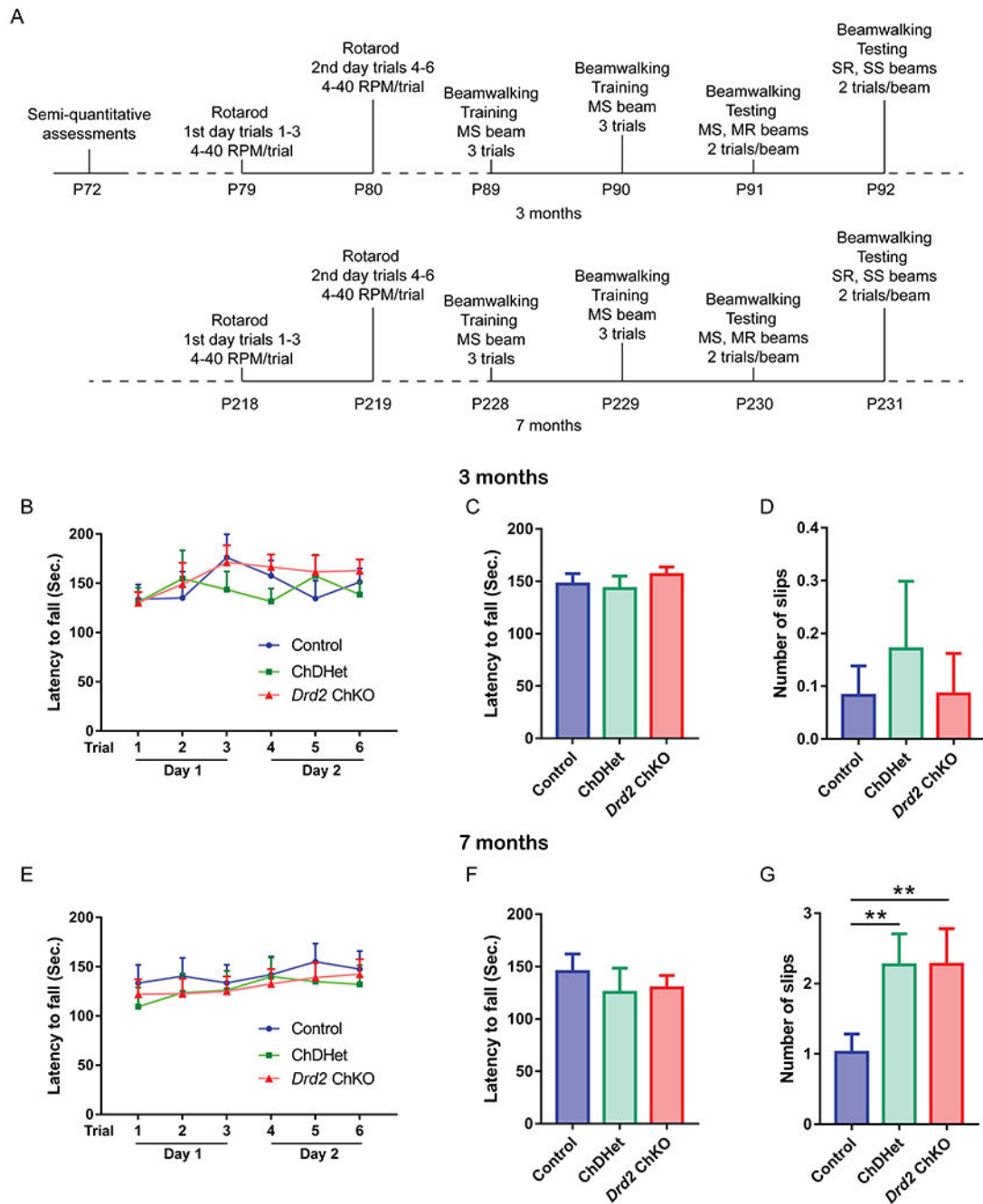


Figure 3. Striatal protein levels in *Drd2* ChKO and *Drd2* sKO mice. Representative western blot images and their quantifications of striatal ChAT (A), AChE (B), and their loading controls (GAPDH) in *Drd2* ChKO mice. Representative western blot images and their quantifications of striatal D2R (C) and their loading controls (GAPDH) in *Drd2* sKO. Vertical bars represent means \pm standard errors.

**Figure 4.**

Impaired motor coordination and balance in *Drd2* ChKO and ChDHet mice. (A) Timeline of the behavior tests. Lack of motor deficits in *Drd2* ChKO and ChDHet mice at 3 months of age on the accelerated rotarod test (B-C) and the beam-walking test (D). Lack of motor deficits in *Drd2* ChKO and ChDHet mice at 7 months of age in the rotarod test. 1-6: trial number. 1-3 trials on the first day and 4-6 trials on the second day. (E-F). *Drd2* ChKO and ChDHet mice displayed excessive slips on the beam-walking test (G). ** $p < 0.01$.

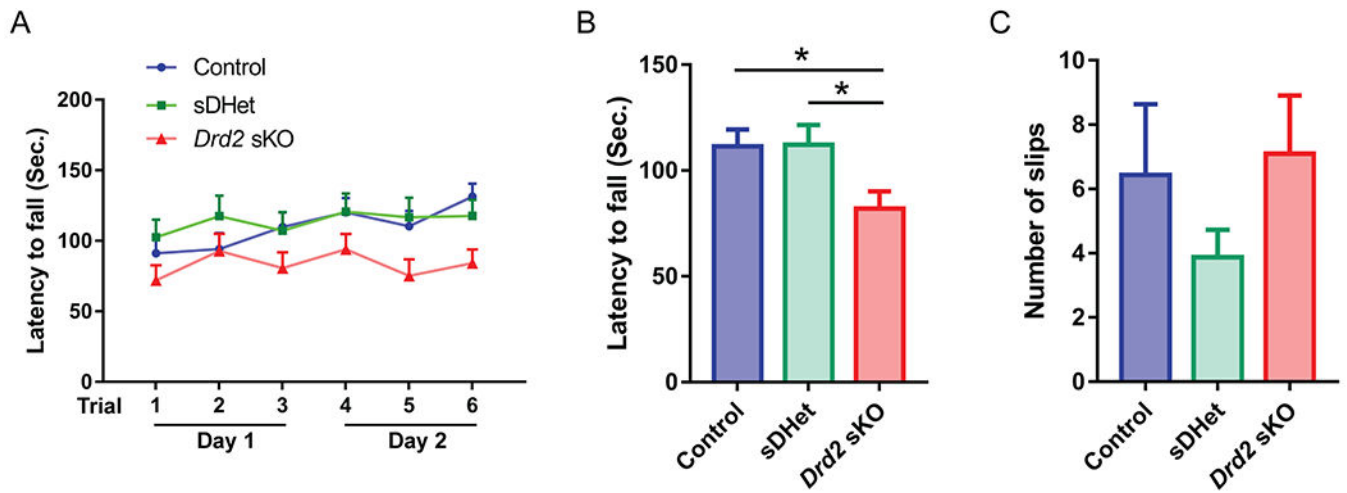


Figure 5.

Impaired motor coordination and balance in *Drd2* sKO mice. (A-B) *Drd2* sKO mice displayed reduced latency to fall in the rotarod test compared with control and sDHet mice. 1-6: trial number. 1-3 trials on the first day and 4-6 trials on the second day. (C) Lack of motor deficits in the beam-walking test in *Drd2* sKO mice. * $p < 0.05$.

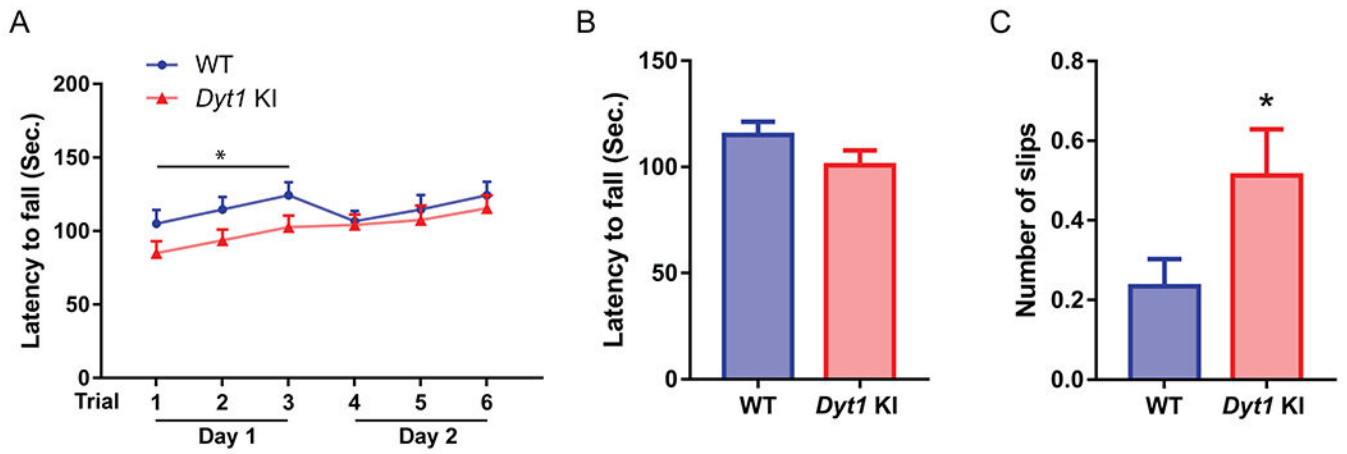


Figure 6.

Impaired motor coordination and balance in *Dyt1* KI mice. (A-B) *Dyt1* KI mice showed motor deficits on the first day but not across all six trials on the accelerated rotarod test. 1-6: trial number. 1-3 trials on the first day and 4-6 trials on the second day. (C) *Dyt1* KI mice displayed excessive slips on the beam-walking test. * $p < 0.05$.