

日米科学技術協力事業「脳研究」分野
グループ共同研究実施報告書

[研究分野： 2]

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トランスジェニックマウスモデルによるジストニアの病態に関する研究

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5. 研究期間 平成18年 4月 1日～平成21年 3月31日

6. 研究の概要、成果及び意義（1000字）

ジストニアは、不随意かつ持続的な筋肉の収縮と異常姿勢によって特徴づけられる神経疾患で、有病率はパーキンソン病の数分の一と稀ではない。しかし、症状の割には病理学的な変化が殆どなく、何らかの機能的な異常と考えられるなど、謎が多い疾患である。病態生理を調べるためには疾患モデルが必要であるが、なかなか良いジストニアのモデルがなかった。最近、マウントサイナイ医科大学神経学教室の Pullanipally Shashidharan 博士が、ヒト DYT1 ジストニア患者の原因遺伝子である DYT1 を組み込んだ遺伝子改変マウスを開発した。本モデルマウスは臨床症状もあり、良いモデルと考えられるので、Shashidharan 博士と共同研究を行うこととした。

実際は、2006年秋、2007年初夏、2009年冬に、南部 篤と知見聡美博士がマウントサイナイ医科大学神経学教室に滞在し、電気生理学の実験を行った。このマウスは、持続的に回転運動をするなど運動が亢進している。筋電図を記録してみると、主動筋と拮抗筋の同時収縮、持続収縮などジストニアに特徴的な異常な筋活動を示した。覚醒下で大脳基底核から神経活動を記録すると、淡蒼球外節と内節において、バースト発射やポーズ（休止期間）を伴う発射頻度の減少が見られた。大脳皮質運動野を電気刺激すると、淡蒼球外節・内節において、正常例においては観察されない早い興奮とそれに引き続く長い抑制という応答が観察された。また、淡蒼球外節・内節の体部位局在も乱れていた。大脳皮質からの入力によって、淡蒼球内節に生じる長く続く抑制が、視床・大脳皮質を脱抑制することによって、不随意運動が起こっていると考えられた。これらの結果は、Journal of Neuroscience 誌（2008年、28：13967-13977）に発表された。

今後の課題としては、大脳皮質由来の抑制が増強しているメカニズムを明らかにすることが第一である。これが明らかになれば、病理所見が乏しいジストニアの病態の本質に迫れるし、また、何らかの方法で増強された抑制を除いてやれば、ジストニアの治療法に結びつくかもしれない。さらに、全身性ジストニア、とくに DYT1 ジストニアに対して、淡蒼球内節に対する脳深部刺激療法（GPi-DBS）が著効を示し、治療の第一選択となりつつあるが、そのメカニズムは不明である。これら治療メカニズムに関しても、本モデルマウスを使うことにより明らかにしていきたい。

7. その他（実施上の問題点、特記事項等）

多くの実験機材を日本より米国の研究室に持ち込んで実験を行った。このような輸送費も本事業で支出して頂けると有り難い。

米国で動物実験を行う際は、J1 等のビザを必要とされる場合があるようである。今後、注意すべきである。

◎参考資料があれば、添付ください。

Chiken S, Shashidharan P, Nambu A (2008) Cortically evoked long-lasting inhibition of pallidal neurons in a transgenic mouse model of dystonia. J Neurosci 28: 13967-13977

（本研究結果の論文）

モデルマウスの神経活動からジストニアの病態を考える（上記論文の解説）

Cortically Evoked Long-Lasting Inhibition of Pallidal Neurons in a Transgenic Mouse Model of Dystonia

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Dystonia is a neurological disorder characterized by sustained or repetitive involuntary muscle contractions and abnormal postures. To understand the pathophysiology of dystonia, neurophysiological analyses were performed on hyperkinetic transgenic mice generated as a model of DYT1 dystonia. Abnormal muscle activity, such as coactivation of agonist and antagonist muscles and sustained muscle activation, was frequently observed in these mice. Recording of neuronal activity in the awake state revealed reduced spontaneous activity with bursts and pauses in both the external and internal segments of the globus pallidus. Motor cortical stimulation evoked responses composed of excitation and subsequent long-lasting inhibition in both pallidal segments, which were never observed in the normal mice. In addition, the somatotopic arrangements in both pallidal segments were disorganized. Long-lasting inhibition induced by cortical inputs in the internal pallidal segment may disinhibit thalamic and cortical activity, resulting in the motor hyperactivity observed in the transgenic mice.

Key words: dystonia; transgenic mouse model; extracellular recording; globus pallidus; movement disorders; basal ganglia

Introduction

Dystonia is a neurological disorder characterized by sustained or repetitive involuntary muscle contractions and abnormal postures (Fahn, 1988; Fahn et al., 1998; Bressman, 2004). The pathophysiology of dystonia is poorly understood, and no consistent histopathological or biochemical changes associated with it have yet been detected. However, the internal (GPi) and external (GPe) segments of the globus pallidus have been discovered to exhibit decreased and bursting activity during stereotaxic surgery for deep brain stimulation (Lenz et al., 1998; Vitek et al., 1999; Vitek, 2002; Zhuang et al., 2004; Starr et al., 2005; Tang et al., 2007).

Early-onset torsion dystonia, the most common type of primary generalized dystonia, is inherited in an autosomal dominant manner with a penetrance of 30–40% (Kramer et al., 1988; Ozelius et al., 1997a). This dystonia is caused by a 3 bp (GAG) deletion in the DYT1 gene on chromosome 9q34, resulting in loss of a glutamic acid residue (ΔE) in the torsinA protein (Ozelius et al., 1997b). Recently, Shashidharan et al. (2005) generated a transgenic mouse model by overexpression of human ΔE -torsinA using a neuron-specific enolase promoter. These trans-

genic mice developed hyperkinesia and rapid bidirectional circling. They also exhibited abnormal involuntary movements with dystonic-appearing self-clasping of limbs and head-shaking. However, other mouse models have failed to exhibit such severe symptoms (Dang et al., 2005, 2006; Sharma et al., 2005; Grundmann et al., 2007; Zhao et al., 2008).

We therefore further examined the hyperkinesia and dystonic-like abnormal movements in this model by electrophysiological methods in the present study. We first tested whether the transgenic mice share electrophysiological features with human dystonia by recording of the electromyogram (EMG) and neuronal activity. We focused on activity in the GPi and GPe, two important nuclei in the basal ganglia circuitry, because activity change in these nuclei has been reported in human patients. Neuronal activity was recorded in the awake state to exclude effects of general anesthesia on neuronal firing rates and patterns (Bergstrom et al., 1984; Keane and Biziere, 1987; Löscher et al., 1995). We then investigated the responses of GPi and GPe neurons evoked by cortical stimulation to elucidate the neural mechanisms underlying the symptoms exhibited by the transgenic mice. Motor cortical stimulation typically induces triphasic responses composed of early excitation, inhibition, and late excitation in GPi and GPe neurons of normal monkeys and rodents (Nambu et al., 1990, 2000; Yoshida et al., 1993; Chiken and Tokuno, 2003). The origin of each component has been identified, with amplitudes and durations reflecting activity of the corresponding basal ganglia pathways and nuclei. Moreover, in voluntary movements, activity originating in the cortex is transmitted through the basal ganglia circuitry and finally reaches the output station of the basal ganglia (i.e., GPi). Cortical stimulation can mimic information processing through the basal ganglia circuitry (Nambu

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et al., 2002; Tachibana et al., 2008). We then discuss the pathophysiology of dystonia based on our electrophysiological findings.

Materials and Methods

Six transgenic mice (Shashidharan et al., 2005) from 5 to 28 weeks of age were used in the present study. Because ~40% of the transgenic mice displayed dystonic-like behavior, only mice exhibiting such abnormalities on behavioral testing after genotype analyses were used in this study. We also used six age-matched wild-type mice as a control group. The experimental protocols were approved by the Animal Care and Use Committees of the Mount Sinai School of Medicine and the Okazaki Organization of National Institutes, and all experiments were conducted according to the guidelines of the National Institutes of Health *Guide for Care and Use of Laboratory Animals*. Before experiments, the mice were trained daily to acquaint them with handling by humans.

Surgery. The head of the mouse was painlessly fixed in a stereotaxic apparatus while awake based on the method described by Yasoshima et al. (2005). Similar methods are usually used in recording neuronal activity from behaving monkeys. Each mouse was anesthetized with ketamine hydrochloride (100 mg/kg body weight, i.p.) and xylazine hydrochloride (4–5 mg/kg, i.p.) and fixed in a conventional stereotaxic apparatus. The skull was widely exposed, and 10 small stainless-steel screws (0.5 mm in diameter) were attached to the skull as anchors. The exposed skull and screws were completely covered with transparent acrylic resin, and then a small U-frame made of acetal resin for head fixation was mounted and fixed on the head of the mouse.

In three transgenic and three normal mice, bipolar wire electrodes (tip distance, 1–2 mm) made of 125- μ m-diameter Teflon-coated multi-stranded stainless-steel wire (A-M Systems) were implanted in the triceps and biceps brachii muscles to record EMG activity. Wires were passed subcutaneously and connected to connectors attached to the U-frame. Antibiotics and analgesics were injected (intramuscularly) after surgery.

After recovery from the first surgery (2 or 3 d later), the mouse was positioned in a stereotaxic apparatus with its head restrained using the U-frame head holder under light anesthesia with ketamine hydrochloride (30–50 mg/kg, i.p.). A part of the skull in one hemisphere was removed to access the motor cortex, GPi, and GPe (classically termed the entopeduncular nucleus and globus pallidus in rodents, respectively). Large caudal and small rostral forelimb regions of the motor cortex have been reported in rats, corresponding to the primary and secondary motor cortices, respectively (Neafsey and Sievert, 1982; Neafsey et al., 1986; Liang et al., 1993; Rouiller et al., 1993). We inserted two pairs of bipolar stimulating electrodes (tip distance, 300–400 μ m) made of 50- μ m-diameter Teflon-coated tungsten wires into the motor cortex, one into the caudal forelimb region and the other into the orofacial region. These regions were confirmed by observation of body part movements evoked by intracortical microstimulation (ICMS) (train of 10 pulses at 333 Hz, 200 μ s duration, up to 30 μ A). Stimulating electrodes were then fixed therein using acrylic resin.

Recording of activity of pallidal neurons. After full recovery from the second surgery, the mouse was positioned in a stereotaxic apparatus with its head restrained using the U-frame head holder in the awake state. For single-unit recording of GPi and GPe neurons, a glass-coated Elgiloy-alloy microelectrode (0.8–1.5 M Ω at 1 kHz) was inserted perpendicularly into the brain through the dura mater using a hydraulic microdrive (Narishige Scientific Instrument) with local application of lidocaine on the dura mater for anesthesia. The target area was 1.1–1.4 mm posterior and 1.4–2.2 mm lateral to bregma for the GPi, and 0.3–0.8 mm posterior and 1.6–2.4 mm lateral for the GPe (Paxinos and Franklin, 2001). Signals from the electrode were amplified and filtered (0.2–5 kHz). The GPi and GPe were identified by the depth profile of the microelectrode penetrations and their sustained spontaneous firing. Waveforms of the action potentials were continuously monitored with an oscilloscope. Unit activity was isolated and converted to digital pulses using a homemade time-amplitude window discriminator. Then the digital pulses were sampled at 2 kHz using a computer. Spontaneous discharges and responses to the cortical electrical stimulation (200 μ s duration single pulse; 20–50 μ A strength) through the electrode implanted in motor

cortex were recorded. EMG activity was also amplified, filtered (0.25–1.5 kHz), and sampled at 20 kHz using a computer.

Data analysis. Spontaneous discharge rates, interspike intervals (ISIs), and autocorrelograms (bin width of 0.5 ms) of the neurons were calculated from continuous digitized recordings for 30 s. Firing patterns of recorded neurons were classified into three types: regular, bursting, and irregular nonbursting.

Whether a firing pattern was regular was judged by autocorrelograms. The mean and SD of autocorrelation coefficients between 0.9 and 1.0 s (200 bins), which were sufficiently far after time 0, were calculated as control values because the autocorrelogram was flat during this period. Peaks and troughs of autocorrelation were judged significant if the coefficient during at least two consecutive bins (1 ms) exceeded the confidence limits ($p < 0.005$, one-tailed t test) (Abeles, 1982; Karmon and Bergman, 1993; Tachibana et al., 2008). The firing pattern of neurons was judged to be regular if at least two cycles, consisting of a pair of the significant peak and trough, were observed.

Bursts were detected by the “Poisson surprise” method (Legéndy and Salcman, 1985; Aldridge and Gilman, 1991; Wichmann and Soares, 2006). It was assumed that spikes in a certain interval followed a Poisson distribution. This method calculated the probability (P) that a given spike train would be found. First, three spike trains shorter than the mean ISIs in the normal mice (i.e., GPi, 19.8 ms; GPe, 18.3 ms) were detected. The Poisson surprise value, which is defined as $-\log_{10} P$, was maximized by adding additional intervals after the initial interval or deleting first intervals from the initial interval. The spike train was judged to be a burst if it included at least three spikes and if its surprise value was >3.0 . We judged the firing pattern of a neuron to be bursting when at least two bursts were observed in 30 s of data. Distribution of numbers of bursts is shown in supplemental Figure 1, A and C (available at www.jneurosci.org as supplemental material), and the cutoff value of two bursts in 30 s was arbitrarily determined. The firing pattern of the neurons was judged to be irregular nonbursting if firing was neither regular nor bursting.

Pauses of spikes were detected using the method of Elias et al. (2007), whose principles are similar to the algorithm for surprise burst detection described above (Legéndy and Salcman, 1985). First, ISIs >250 ms were detected in the spike train. The Poisson surprise value was maximized by adding additional intervals (less than five) before or after the initial interval. The final interval was considered a pause if its interval was >300 ms and if its surprise value was >3.0 . Finally, two adjacent pauses were merged into one continuous pause if they were separated by one spike. We judged a neuron to be a neuron with pauses if at least one pause was observed in 30 s of the data. Distribution of numbers of pauses is shown in supplemental Figure 1, B and D (available at www.jneurosci.org as supplemental material), and the cutoff value of one pause in 30 s was arbitrarily determined.

Responses to cortical electrical stimulation were examined by constructing peristimulus time histograms (PSTHs) (bin width of 1 ms) for 100 stimulus trials. The mean value and SD of the discharge rate during the 100 ms period preceding onset of stimulation were calculated for each PSTH and considered the baseline discharge rate. Changes in neuronal activity in response to cortical stimulation (i.e., excitation and inhibition) were judged significant if the discharge rate during at least two consecutive bins reached a significance level of $p < 0.05$ (one-tailed t test) (Nambu et al., 2000; Kita et al., 2004, 2005, 2006; Tachibana et al., 2008). The latency of each response was defined as the time at which the discharge rate first exceeded this level.

Histology. In the final experiment, several sites of neuronal recording were marked by passing cathodal DC current (15 μ A for 15 s) through the recording electrodes. The mice were anesthetized deeply with sodium pentobarbital (80 mg/kg, i.p.), and perfused transcardially with 0.1 M phosphate buffer (PB), pH 7.3, followed by 10% formalin in 0.1 M PB, and then 0.1 M PB containing 10% sucrose. The brains were removed immediately and saturated with the same buffer containing 30% sucrose. They were cut into frontal 50- μ m-thick sections on a freezing microtome. The sections were mounted onto gelatin-coated glass slides, stained with 0.7% neutral red, dehydrated, and coverslipped. The sections were observed under a light microscope, and the recording sites

were reconstructed according to the lesions made by current injection and traces of electrode tracks. The sites of stimulation in motor cortex were also examined histologically.

Results

EMG activity

The activities of the triceps and biceps brachii muscles, the extensor and flexor muscles of the forelimb, were simultaneously recorded in three normal and three transgenic mice (Fig. 1). Patterns of EMG activity during voluntary forelimb movements in the transgenic mice did not appear to differ from those in the normal mice during most of the time of recording (Fig. 1*A,B*). In all three transgenic mice, however, the triceps and biceps muscles were sometimes coactivated during forelimb movements (Fig. 1*C*). Sharp EMG activities in the triceps and biceps muscles in these mice were synchronized with each other (Fig. 1*C*); such synchronization was never observed in the normal mice. Moreover, sustained muscle activity that lasted >10 s was frequently observed when the transgenic mice stopped movement (Fig. 1*D*). The sustained muscle activity was sometimes accompanied by coactivation of the triceps and biceps muscles (Fig. 1*D*).

Locations of stimulating electrodes in motor cortex

The caudal forelimb and orofacial regions were successfully identified in all normal and transgenic mice by ICMS. The caudal forelimb region was found 0.3–1.3 mm anterior and 1.3–1.8 mm lateral to bregma in both normal and transgenic mice. Stimulation applied to more medial sites induced movements of the vibrissae. The orofacial region including jaw, lip, and tongue representations was located more rostrally and laterally than the forelimb region, at 1.7–2.6 mm anterior and 1.8–2.5 mm lateral to bregma in both normal and transgenic mice. In some cases, fingers or wrist movements were evoked by stimulation applied to more medial sites, corresponding to the rostral forelimb region. The threshold for evocation of movement by ICMS in the caudal forelimb and orofacial regions of the transgenic mice was 6–30 μ A, and similar to that in the normal mice. These somatotopic arrangements were consistent with those previously reported in rats (Donoghue and Wise, 1982; Neafsey et al., 1986). These observations indicate that the somatotopic arrangement and excitability of motor cortex in the transgenic mice did not differ from those in the normal mice. Movements evoked by single-pulse stimulation (up to 70 μ A) through the implanted stimulating electrodes were restricted to the forelimb (with stimulation in the forelimb region) or face (with stimulation in the orofacial region), suggesting that current spread to neighboring regions was negligible in daily experimental sessions.

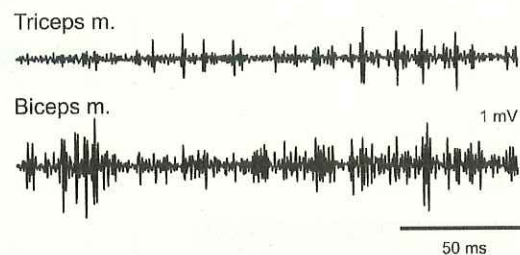
Spontaneous activity of GPi and GPe neurons

The spontaneous activity of 94 GPi and 70 GPe neurons in six normal mice and 90 GPi and 204 GPe neurons in six transgenic mice was recorded. GPi and GPe neurons in the normal mice fired continuously at high discharge rate >50 Hz, as shown in traces of digitized spikes (Fig. 2*A1,B1*). Autocorrelograms indicated that these neurons fired irregularly (Fig. 2*A2,B2*). However, GPi and GPe neurons in the transgenic mice fired at low frequency with bursts and pauses (Fig. 2*A3,B3*). The mean discharge rates of GPi and GPe neurons in the transgenic mice were significantly lower than those in the normal mice (Table 1; Fig. 2*A5,B5*) ($p < 0.001$, t test). Discharge patterns also differed in the transgenic mice. Bursts (indicated by thick black lines in Fig. 2*A3,B3*) and pauses (indicated by thick white lines) were frequently observed in GPi and GPe neurons of the transgenic mice.

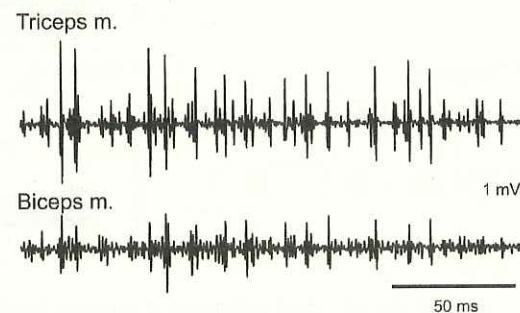
A Normal



B Transgenic



C Transgenic



D Transgenic

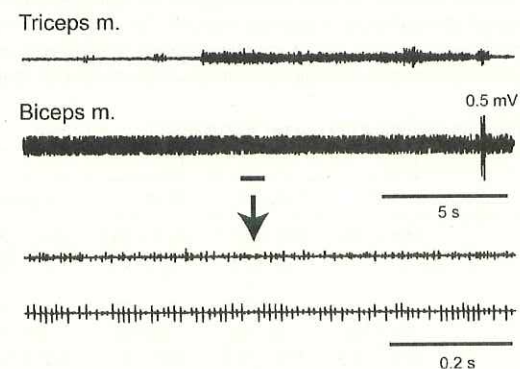


Figure 1. EMG activity of triceps and biceps brachii muscles. *A*, EMG activity in a normal mouse during voluntary forelimb movements. *B–D*, EMG activity in a transgenic mouse. EMG activity during voluntary forelimb movements did not appear to differ from that in the normal mice during most recording time (*B*). However, sharp EMG activities in the triceps and biceps muscles were sometimes observed, synchronized with each other (*C*). Sustained muscle activity that lasted >10 s was frequently observed (*D*). The underlined portions of the top two traces are expanded in the bottom two traces.

The major firing pattern of GPi (63%) and GPe (77%) neurons in the normal mice was irregular nonbursting, whereas other firing patterns, such as regular (supplemental Fig. 2, available at www.jneurosci.org as supplemental material) and bursting neurons

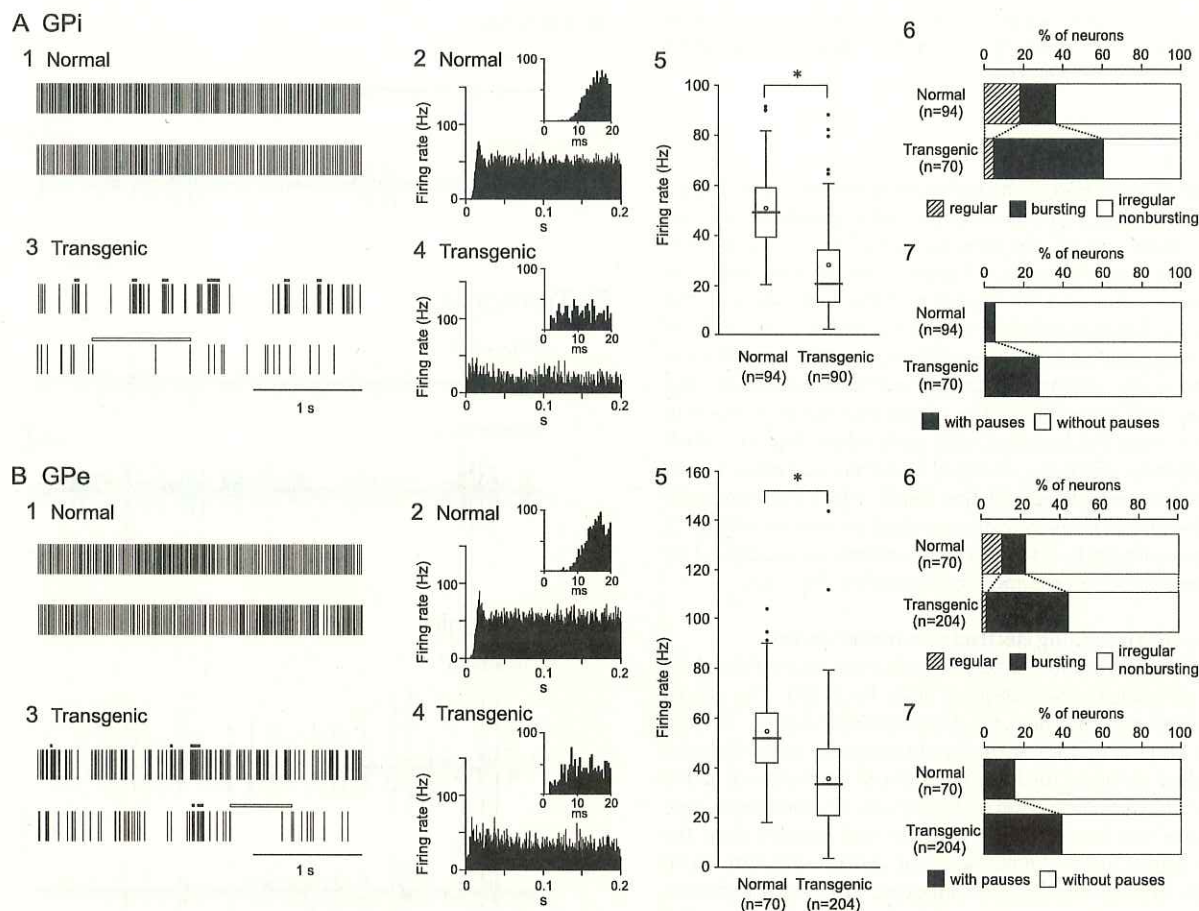


Figure 2. Spontaneous activity of neurons in the internal (GPI) (**A**) and external (GPe) (**B**) segments of the globus pallidus. **1, 3**, Spikes are shown as digital signals in the normal (**1**) and transgenic mice (**3**). Bursts and pauses in the digital signals are indicated by horizontal black thick lines and horizontal white thick lines, respectively. Neither bursts nor pauses are observed in traces of the normal mice exemplified in **A1** and **B1**. **2, 4**, Autocorrelograms with long and short (insets) timescales are shown in the normal (**2**) and transgenic mice (**4**). **5**, Box plots of firing rates in the normal (left) and transgenic mice (right). The boxes are constructed with the top line bounding the first quartile and the bottom line bounding the third quartile. Median and mean values are indicated by a thick horizontal line and an open circle in the box, respectively. The short horizontal lines show the largest and smallest values that are not outliers. Outliers are shown as small closed circles. *Significantly different ($p < 0.001$, t test). **6**, Proportions of neurons classified by firing pattern (i.e., regular, bursting, or irregular nonbursting) in the normal (top) and transgenic (bottom) mice. **7**, Proportions of neurons with or without pauses in the normal (top) and transgenic (bottom) mice.

Table 1. Firing rates of GPI and GPe neurons in normal and transgenic mice

	GPI		GPe	
	Normal	Transgenic	Normal	Transgenic
Total	50.6 ± 15.7 (<i>n</i> = 94)	27.8 ± 19.1* (<i>n</i> = 90)	54.5 ± 16.3 (<i>n</i> = 70)	35.4 ± 19.0* (<i>n</i> = 204)
Regular	60.8 ± 14.8 (<i>n</i> = 16)	70.4 ± 13.8 (<i>n</i> = 4)	61.6 ± 23.8 (<i>n</i> = 7)	62.0 ± 14.9 (<i>n</i> = 3)
Bursting	47.4 ± 16.7 (<i>n</i> = 18)	22.1 ± 12.4 (<i>n</i> = 50)	44.0 ± 8.5 (<i>n</i> = 9)	27.7 ± 14.0 (<i>n</i> = 88)
Irregular nonbursting	48.9 ± 14.9 (<i>n</i> = 60)	30.8 ± 21.7 (<i>n</i> = 36)	55.4 ± 15.7 (<i>n</i> = 54)	40.7 ± 20.1 (<i>n</i> = 113)
With pauses	52.2 ± 11.5 (<i>n</i> = 5)	17.9 ± 6.9 (<i>n</i> = 25)	48.0 ± 15.6 (<i>n</i> = 12)	29.2 ± 16.8 (<i>n</i> = 83)
Without pauses	50.5 ± 16.0 (<i>n</i> = 89)	31.6 ± 21.4 (<i>n</i> = 65)	55.9 ± 16.3 (<i>n</i> = 58)	39.7 ± 19.3 (<i>n</i> = 121)

Values are means ± SD expressed in hertz.

* $p < 0.001$, significantly different from normal (one-tailed t test).

were observed (Fig. 2A6,B6). However, numbers of bursting neurons in the GPI (57%) and GPe (43%) were significantly increased in the transgenic mice ($p < 0.001$, χ^2 test), and numbers of regular and irregular nonbursting neurons decreased. Spike

shapes of bursting and irregular nonbursting neurons in the transgenic mice were compared. Both neurons showed positive-negative deflection with similar peak-to-peak amplitude (mean ± SD in microvolts, average of 10 neurons; bursting GPI, 86 ± 17 ; irregular nonbursting GPI, 79 ± 17 ; bursting GPe, 75 ± 14 ; irregular nonbursting GPe, 72 ± 21) and similar duration (in milliseconds, average of 10 neurons; bursting GPI, 1.6 ± 0.2 ; irregular nonbursting GPI, 1.6 ± 0.2 ; bursting GPe, 1.6 ± 0.2 ; irregular nonbursting GPe, 1.6 ± 0.2), suggesting that both neurons may belong to a single neuronal group. Bursts in the normal and transgenic mice are quantitatively compared in Table 2 and supplemental Figure 1, A and C (available at www.jneurosci.org as supplemental material). Number of bursts in the GPe and total lengths of bursts in the GPI and GPe were increased in the transgenic mice, although the duration of each burst in the GPe was decreased. More GPI (28%) and GPe (41%) neurons exhibited pauses in the transgenic mice than in the normal mice (Fig. 2A7,B7) ($p < 0.001$, χ^2 test). Pauses in the normal and transgenic mice are compared in Table 3 and supplemental Figure 1, B and D (available at www.jneurosci.org as supplemental material). The duration of each pause and total lengths of pauses in the GPI and GPe were increased in the transgenic mice.

Table 2. Burst characteristics in normal and transgenic mice

	GPi		GPe	
	Normal (n = 18)	Transgenic (n = 51)	Normal (n = 9)	Transgenic (n = 88)
No. of bursts during 30 s	6.1 ± 6.2	8.7 ± 8.3	4.1 ± 2.3	16.2 ± 18.5*
Duration of each burst (ms)	33.8 ± 30.8	34.8 ± 27.6	101.8 ± 86.2	42.4 ± 49.1**
No. of spikes during each burst	8.0 ± 5.0	6.3 ± 4.2	15.2 ± 10.4	7.5 ± 4.8**
Intraburst firing rate (Hz)	274.3 ± 80.3	188.7 ± 61.1**	181.6 ± 47.3	200.9 ± 63.5
Total length of bursts during 30 s (% of spikes)	3.2 ± 3.2	8.4 ± 7.3**	5.1 ± 4.7	14.1 ± 16.0*
Total length of bursts during 30 s (% of times)	0.7 ± 0.7	0.9 ± 0.7	1.5 ± 1.5	1.7 ± 1.8
Poisson surprise value for each burst	3.7 ± 0.7	4.0 ± 0.5*	4.4 ± 1.6	4.1 ± 0.8

Values are means ± SD.

* $p < 0.05$ and ** $p < 0.01$, significantly different from normal (one-tailed t test).**Table 3. Pause characteristics in normal and transgenic mice**

	GPi		GPe	
	Normal (n = 5)	Transgenic (n = 25)	Normal (n = 11)	Transgenic (n = 83)
No. of pauses during 30 s	1.2 ± 0.5	2.7 ± 2.0	2.0 ± 1.3	3.5 ± 3.2
Duration of each pause (ms)	410.3 ± 102.0	1074.8 ± 614.8**	418.5 ± 71.4	871.7 ± 689.7*
Total length of pauses during 30 s (% of time)	1.6 ± 0.6	10.0 ± 8.6*	2.8 ± 1.9	10.3 ± 12.7*
Poisson surprise value for each pause	6.2 ± 3.2	4.4 ± 1.9	5.7 ± 2.9	6.1 ± 4.5

Values are means ± SD.

* $p < 0.05$ and ** $p < 0.01$, significantly different from normal (one-tailed t test).

Relationships between firing patterns and firing rates were further analyzed in Table 1. Regular GPi and GPe neurons showed similar firing rates both in the normal and transgenic mice. In contrast, firing rates of bursting and irregular nonbursting neurons decreased in the transgenic mice. These results suggest that reduction of mean firing rates of GPi and GPe neurons in the transgenic mice are caused by both (1) decreased number of regular neurons and (2) reduction of firing rates of bursting and irregular nonbursting neurons. Firing rates were also compared between GPi and GPe neurons with pauses and those without pauses (Table 1). Reduction of firing rates occurred in neurons without pauses as well as in neurons with pauses.

Responses of GPi and GPe neurons to cortical stimulation

Stimulation of the forelimb and/or orofacial region of the motor cortex was applied to 91 GPi and 68 GPe neurons in six normal mice, and induced responses in 59 GPi and 40 GPe neurons (~60%). The typical response pattern of GPi (36%) and GPe (56%) neurons in the normal mice was a triphasic response composed of early excitation, followed by inhibition, and late excitation (Figs. 3A, 4A). Other patterns such as excitation followed by inhibition, inhibition followed by excitation, monophasic or biphasic excitation, and monophasic inhibition were also observed (Figs. 3C, 4C). In most cases, these events ended within 50 ms after cortical stimulation and were not followed by late responses (Figs. 3A, 4A; Table 4).

Cortical stimulation induced responses in similar percentages of GPi and GPe neurons (GPi, 52 of 84; GPe, 111 of 178) in six transgenic mice, although response patterns differed. The most common response pattern of GPi and GPe neurons in the transgenic mice was short-latency monophasic or biphasic excitation followed by long-lasting inhibition (Figs. 3B, left; 4B, left), a pattern never observed in the normal mice. The latency and duration of the early excitation were 4.4 ± 2.6 and 15.3 ± 5.5 ms in the GPi and 4.0 ± 1.6 and 12.4 ± 7.1 ms in the GPe, respectively, whereas the duration of the long-lasting inhibition was 73.7 ± 29.4 ms in the GPi and 66.7 ± 31.3 ms in the GPe. The most common pattern of early response (within 50 ms after cortical stimulation)

was monophasic or biphasic excitation in GPi (62%) and GPe (49%) neurons (Figs. 3C, 4C), and early responses were frequently followed by long-lasting inhibition (GPi, 56%; GPe, 41%) (Figs. 3D, 4D) ($p < 0.001$, χ^2 test). In other GPi (14%) and GPe (30%) neurons in the transgenic mice, cortical stimulation induced a triphasic response similar to that in the normal mice (Figs. 3B, right; 4B, right), although the number of such neurons was decreased (Figs. 3C, 4C) ($p < 0.01$, χ^2 test). The latency and duration of each component were also similar, with the exception of the latency and duration of the late excitation in the GPi and duration of the inhibition in the GPe, which differed significantly (Table 4) (t test).

Relationships between cortically evoked responses and spontaneous firing patterns were further analyzed (supplemental Table 1, available at www.jneurosci.org as supplemental material). The majority of neurons that showed excitation (GPi, 81%; GPe, 70%) to the cortical stimulation in the transgenic mice belonged to bursting neurons, whereas those in the normal mice belonged to irregular nonbursting neurons (GPi, 75%; GPe, 100%). In contrast, neurons with a cortically evoked triphasic response had a tendency to fire in irregular nonbursting manner both in the normal (GPi, 56%; GPe, 92%) and transgenic (GPi, 50%; GPe, 82%) mice. A large number of neurons that showed cortically evoked long-lasting inhibition in the transgenic mice belonged to bursting neurons (GPi, 73%; GPe, 91%) and neurons with pauses (GPi, 42%; GPe, 73%), whereas those neurons were rarely observed in the normal mice.

Somatotopic organization in GPi and GPe

Stimulation of both the forelimb and orofacial regions of the motor cortex was successfully performed for 91 GPi and 68 GPe neurons in six normal mice and 43 GPi and 77 GPe neurons in six transgenic mice. Neurons could be classified into four groups, those with forelimb inputs, those with orofacial inputs, those with convergent inputs from both forelimb and orofacial regions, and those with no responses, based on the responses evoked by cortical stimulation. In the normal mice, small number of GPi (7%) and GPe (10%) neurons responded to stimulation of both forelimb and orofacial regions (Figs. 5A, 6A). In the transgenic mice, the number of neurons with convergent inputs from two regions was significantly increased (GPi, 28%; GPe, 35%; $p < 0.001$, χ^2 test).

The locations of recorded GPi and GPe neurons are plotted using symbols based on cortical inputs in Figures 5B and 6B. In the GPi of the normal mice, the neurons with forelimb inputs were distributed over a wide area of the GPi, although not in the most medial portion of it (Fig. 5B, top). A few neurons with orofacial inputs and with convergent inputs were found in the lateral portion of the GPi. In the transgenic mice, this segregation was not observed (Fig. 5B, bottom). The number of GPi neurons with orofacial inputs and those with convergent inputs increased (Fig. 5A), and they intruded into the central portion of the GPi, although the most medial portion remained unresponsive. Cortical inputs to GPe neurons were mainly examined in the central to lateral portion of the GPe (Fig. 6B). In the normal mice, neu-

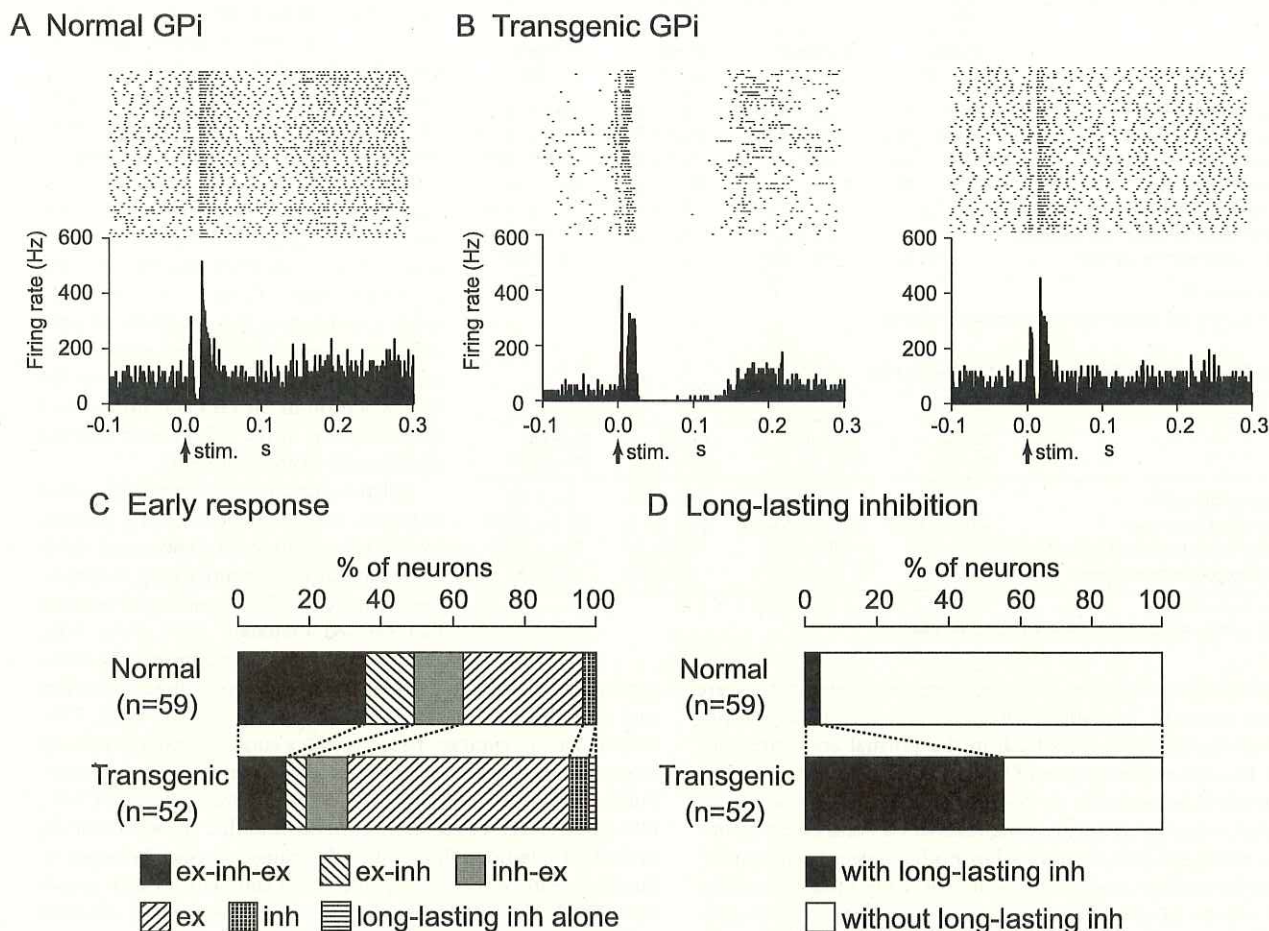


Figure 3. Responses of GPI neurons to cortical stimulation. *A*, Raster and PSTH for the normal mice. Cortical stimuli were delivered at time 0 (arrows). *B*, Raster and PSTH for the transgenic mice. Abnormal responses with long-lasting inhibition (left) and apparently normal triphasic responses (right) were observed. *C*, Proportions of neurons classified based on early response patterns in the normal (top) and transgenic (bottom) mice. ex, Excitation; inh, inhibition. *D*, Proportions of neurons with and without long-lasting inhibition in the normal (top) and transgenic (bottom) mice.

rons with orofacial inputs were located in the ventral portion, whereas those with forelimb inputs were located in the dorsal portion (Fig. 6*B*, top). Neurons with convergent inputs were found in the intermediate portion. In the transgenic mice, this type of segregation was not found. Neurons with orofacial inputs and those with convergent inputs intruded into the dorsal portion, and the three groups of neurons were intermingled (Fig. 6*B*, bottom). Neurons in the most dorsal portion of the GPe were unresponsive in both normal and transgenic mice.

Discussion

The present study characterized the electrophysiological properties of transgenic mice developed to express human ΔE -torsinA. These mice exhibited (1) coactivation of agonist and antagonist muscles and sustained muscle activation, (2) decreased GPI and GPe activity with bursts and pauses, (3) cortically evoked long-lasting inhibition in the GPI and GPe, and (4) somatotopic disorganization in the GPI and GPe. These neuronal abnormalities may be responsible for the behavioral abnormalities exhibited by these mice.

Coactivation of agonist and antagonist muscles and sustained muscle activation in transgenic mice

The triceps and biceps muscles were sometimes coactivated during forelimb movements in the transgenic mice (Fig. 1*C*). Coac-

tivation of agonist and antagonist muscles is common to various types of dystonia (Obeso et al., 1983; Marsden and Rothwell, 1987; Cohen and Hallett, 1988; Berardelli et al., 1998; Farmer et al., 1998; Liu et al., 2004). Sustained muscle activity lasting >10 s was frequently observed in the transgenic mice (Fig. 1*D*). Sustained muscle activity is another important sign of dystonia (Herz, 1944; Yanagisawa and Goto, 1971; Marsden and Rothwell, 1987; Jedynak et al., 1991; Berardelli et al., 1998). It has been suggested that dystonic postures are produced by long periods of continuous EMG activity lasting several seconds (Yanagisawa and Goto, 1971; Berardelli et al., 1998). These observations suggest that the transgenic mice we examined exhibit EMG activity similar to that observed in human patients with dystonia.

Decreased GPI and GPe activity in transgenic mice

Marked reduction of the spontaneous firing rates of GPI and GPe neurons was observed in the transgenic mice (Table 1, Fig. 2). Alteration of firing patterns was also observed in them, including bursting discharges and pauses. Decreased discharge rates and irregularly grouped discharges with intermittent pauses in GPI and GPe neurons have also been observed in patients with generalized dystonia (Vitek et al., 1999; Zhuang et al., 2004; Starr et al., 2005). These findings suggest that the transgenic mice we examined may share neurological abnormalities with dystonia.

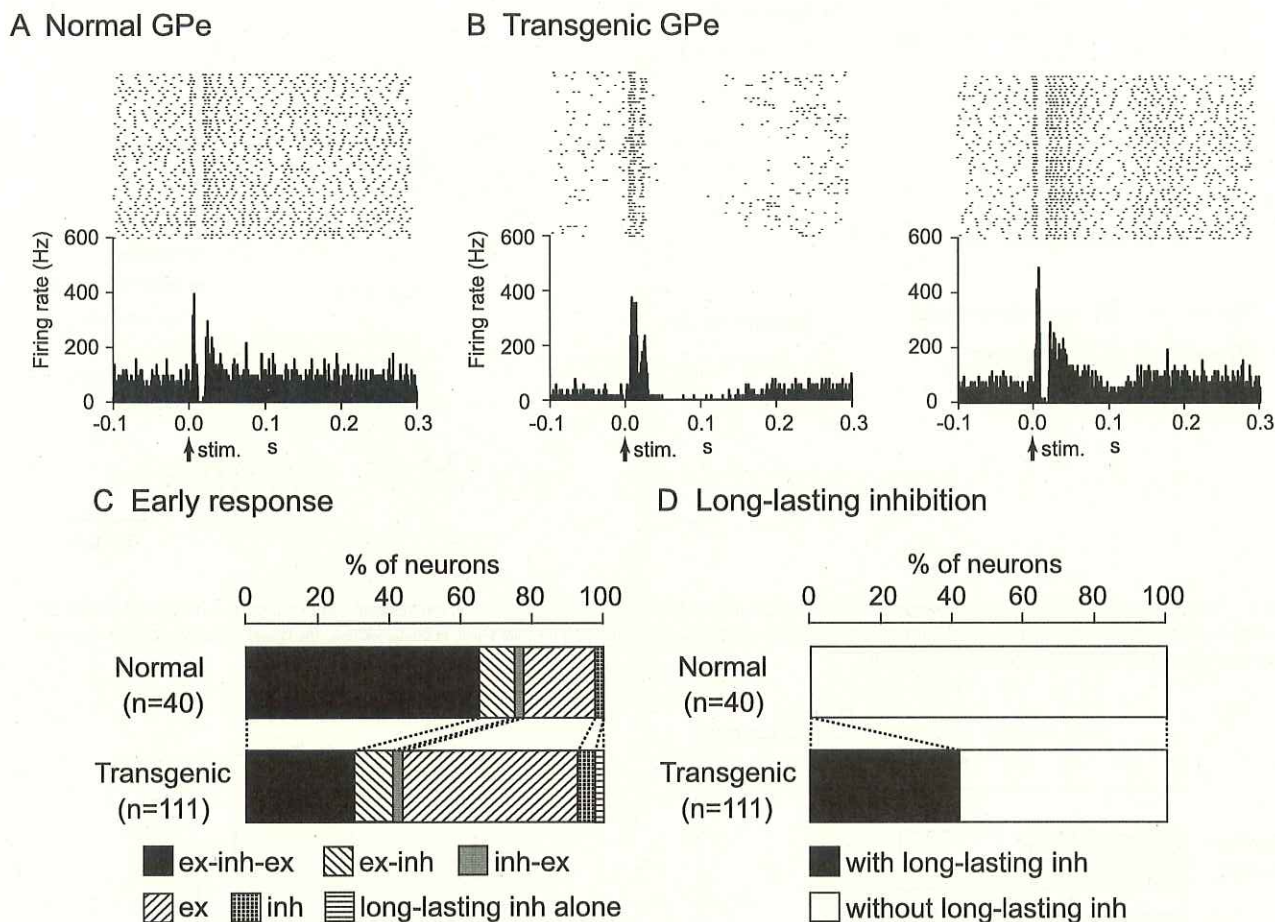


Figure 4. Responses of GPe neurons to cortical stimulation. *A*, Raster and PSTH for the normal mice. *B*, Raster and PSTH for the transgenic mice. *C*, Proportions of neurons classified based on early response patterns in the normal (top) and transgenic (bottom) mice. *D*, Proportions of neurons with and without long-lasting inhibition in the normal (top) and transgenic (bottom) mice.

Table 4. Latency and duration of triphasic responses evoked by cortical stimulation

		GPi		GPe	
		Normal (<i>n</i> = 21)	Transgenic (<i>n</i> = 7)	Normal (<i>n</i> = 26)	Transgenic (<i>n</i> = 33)
Latency (ms)	Early excitation	3.3 ± 0.9	3.5 ± 0.5	4.3 ± 0.8	4.4 ± 1.1
	Inhibition	11.6 ± 2.3	11.1 ± 1.7	11.1 ± 1.5	12.1 ± 2.5
	Late excitation	20.9 ± 2.1	18.6 ± 1.8**	21.1 ± 2.2	21.2 ± 5.4
Duration (ms)	Early excitation	5.2 ± 2.9	5.7 ± 1.3	4.9 ± 1.8	5.3 ± 1.7
	Inhibition	6.1 ± 2.9	6.1 ± 2.0	8.7 ± 3.0	6.8 ± 3.2*
	Late excitation	9.9 ± 5.0	34.6 ± 45.0**	16.8 ± 18.7	26.3 ± 28.5

Values are means ± SD expressed in milliseconds.

p* < 0.05 and *p* < 0.01, significantly different from normal (one-tailed *t* test).

patients. Dystonic hamsters with paroxysmal generalized dystonia also exhibited reduced and bursting GPi activity (Gernert et al., 2002). Firing rates of both bursting and irregular nonbursting neurons decreased, suggesting that reduction of firing rate universally occurred in GPi and GPe neurons. The mechanisms responsible for decreased firing rates may include (1) alteration of membrane properties of GPi and GPe neurons; (2) increased inhibitory inputs to the GPi and GPe, such as GABAergic inputs from the striatum; and/or (3) decreased excitatory inputs to the GPi and GPe, such as glutamatergic inputs from the subthalamic nucleus (STN). Inhibitory inputs from the striatum to the GPi and GPe increase in the transgenic mice as discussed in the next section.

Cortically evoked long-lasting inhibition in GPi and GPe neurons of transgenic mice

To investigate the mechanism of abnormal firing of GPi and GPe neurons in the transgenic mice further, responses evoked by cortical stimulation were observed. In the normal mice, cortical stimulation typically induced triphasic responses composed of early excitation, inhibition, and late excitation in GPi and GPe neurons (Figs. 3*A*, 4*A*). Similar triphasic responses were also observed in the GPi, GPe, and substantia nigra pars reticulata (SNr), another output station of the basal ganglia, in rats and monkeys. The origin of each component has been intensively studied (Ryan and Sanders, 1994; Maurice et al., 1998, 1999; Nambu et al., 2000; Kita et al., 2004; Tachibana et al., 2008). Early excitation is mediated by the cortico-STN-GPe/GPi/SNr pathway, whereas inhibition and late excitation are mediated by the cortico-striato-GPe/GPi/SNr and cortico-striato-GPe-STN-GPe/GPi/SNr pathways, respectively. However, in the transgenic mice we examined, cortical stimulation induced early excitation followed by late long-lasting inhibition in GPi and GPe neurons (Figs. 3*B*, 4*B*). Because the excitability of motor cortex was unchanged in the transgenic mice, these abnormal patterns

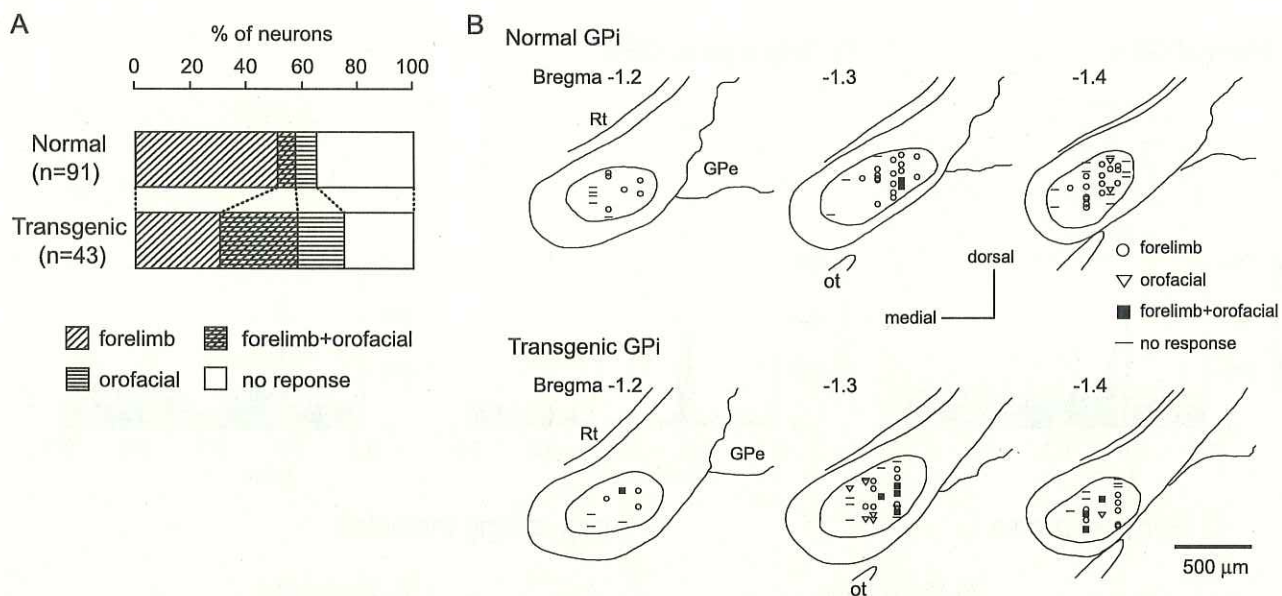


Figure 5. Somatotopic organization in the GPi. **A**, Proportions of neurons classified based on cortical inputs in the normal (top) and transgenic (bottom) mice. **B**, Distribution of recorded GPi neurons indicated by symbols based on cortical inputs. Data from two normal (top) and two transgenic (bottom) mice are shown in frontal sections. The figures in the left top corner represent distance from bregma. Rt, Reticular thalamic nucleus; Ot, optic tract.

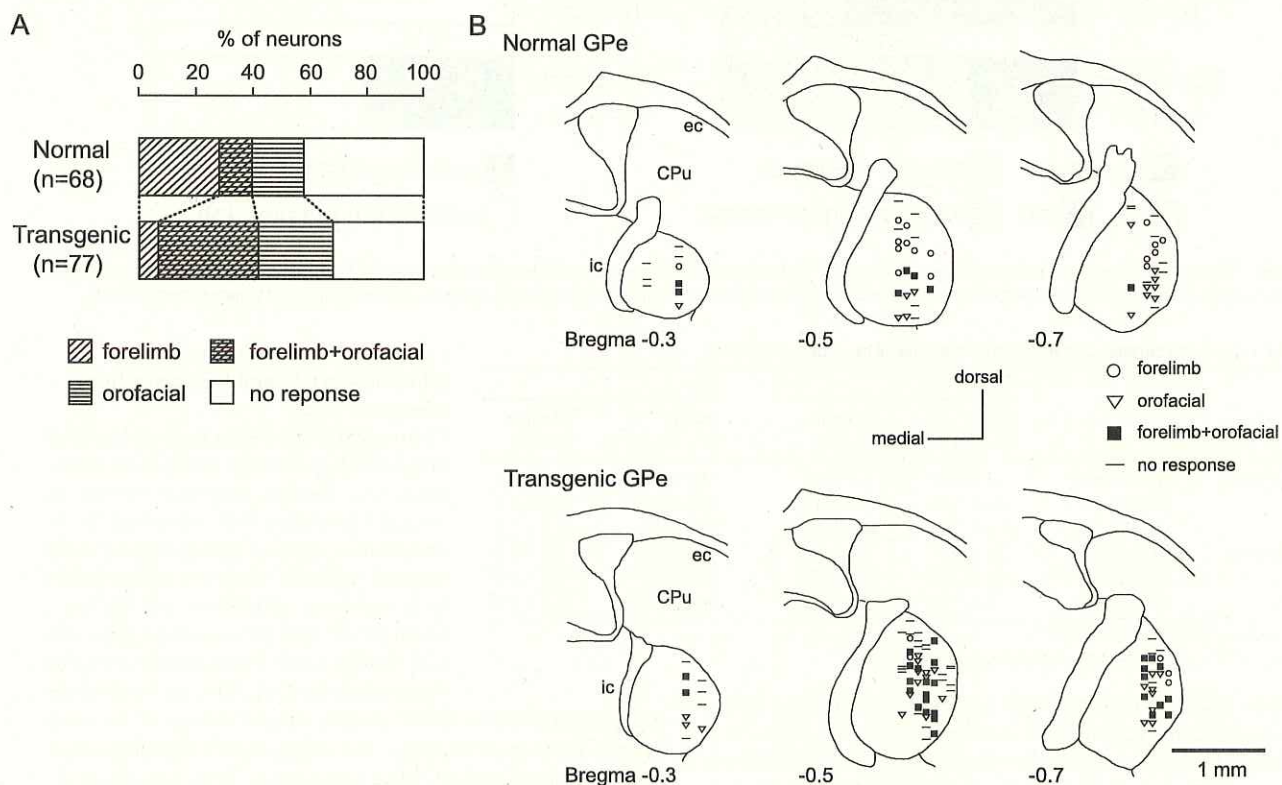


Figure 6. Somatotopic organization in the GPe. **A**, Proportions of neurons classified based on cortical inputs in the normal (top) and transgenic (bottom) mice. **B**, Distribution of recorded GPe neurons. Data from two normal (top) and two transgenic (bottom) mice are shown in frontal sections. CPu, Caudate-putamen; ec, external capsule; ic, internal capsule.

of response may be generated through the cortico-basal ganglia pathways. The early excitation may, at least its early phase, be mediated by the cortico-STN-GPe/GPi pathway, as in the normal mice, because the latency of the early excitation in the transgenic mice was short and similar to that in the normal mice. These

observations also suggest that the activity of STN neurons is unchanged in the transgenic mice. The origin of the late long-lasting inhibition may be (1) increased inhibitory input via the striato-GPe/GPi pathway or (2) decreased excitatory input via the STN-GPe/GPi pathway. The latter explanation seems less likely to be

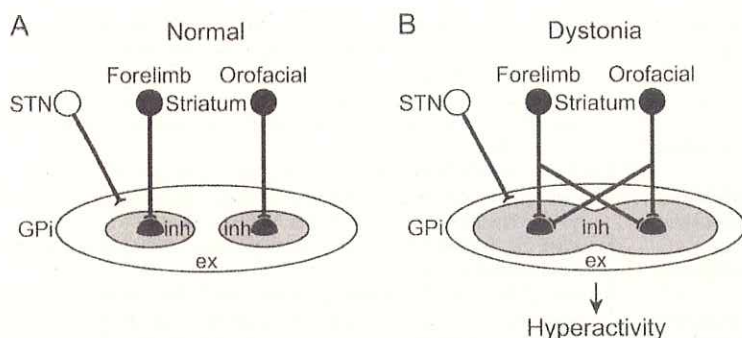


Figure 7. Schematic diagrams showing information processing through the basal ganglia in normal condition (A) and dystonia (B). In dystonia, cortical activation induces strong inhibition over wide areas of the GPI, as well as strong excitation in the thalamus and cortex as a result of disinhibition, resulting in the motor hyperactivity and involuntary muscle contractions.

correct, because activity along the cortico-STN-GPi/GPe pathway appeared to be unchanged, as discussed above. The early excitation sometimes included two excitatory peaks (biphasic excitation) (Figs. 3*B*, 4*B*). Strong inhibition in the GPe evoked by the cortico-striato-GPe pathway might disinhibit STN neurons and produce excitation in the GPi and GPe, corresponding to the second excitation of the early response in the transgenic mice. The above observations, together with positive relationships between cortically evoked responses and spontaneous firing patterns, also suggest that spontaneous excitation in the cortex is transmitted to the GPi and GPe through the cortico-basal ganglia pathways, and induces short-latency excitation and long-lasting inhibition, which might be the origins of bursts and pauses, respectively.

Somatotopic disorganization of GPi and GPe in transgenic mice

Motor territories in the GPI and GPe of monkeys are somatotopically organized, as indicated by their somatosensory inputs, activity during voluntary movements, and cortically evoked responses (DeLong et al., 1985; Yoshida et al., 1993). The present study confirmed the somatotopic organization in the normal mice. In the transgenic mice, however, somatotopic disorganization was observed, and many GPI and GPe neurons received convergent inputs from both forelimb and orofacial regions (Figs. 5, 6). Widened somatosensory receptive fields in pallidal neurons have been reported in patients with generalized (Vitek et al., 1999) and focal (Lenz et al., 1998; Sanger et al., 2001) dystonia. Because the somatotopic arrangement and excitability of the motor cortex in the transgenic mice appeared normal in the present study, information-crossing may have occurred through the cortico-basal ganglia pathways. One explanation for this is that single GPI or GPe neurons receive inputs from more striatal neurons in the transgenic mice than in the normal mice (Fig. 7). This explanation agrees well with the assumption of increased inhibitory input via the striato-GPe/GPI pathway noted in the previous section.

Neural mechanisms of motor hyperactivity in transgenic mice

The GPI, an output nucleus of the basal ganglia, is composed of GABAergic inhibitory neurons and fires at high frequency in normal states. Its target structures, such as the thalamus and frontal cortex, are thus continuously inhibited. Striatal inputs reduce GPI activity in temporal manner, excite thalamic and cortical neurons via disinhibition, and finally release appropriate movements with appropriate timing (Fig. 7A) (Nambu et al., 2000,

2002). However, in the transgenic mice, cortical excitation induced long-lasting inhibition in the GPi, suggesting that even tiny amounts of neuronal activity originating in the cortex are transmitted through the cortico-basal ganglia pathways and finally induce strong and long-lasting inhibition in the GPi (Fig. 7*B*). Moreover, somatotopic disorganization was noted in the GPi and cortical activation induced inhibition over a wide area of this region. Wide areas of the thalamus and cortex are thus activated in uncontrollable manner, resulting in the motor hyperactivity and involuntary muscle contractions observed in the transgenic mice. A similar mechanism may underlie the symptoms of hu-

Relationships with other transgenic models

The transgenic mice investigated in the present study exhibited hyperkinesia and dystonic-like movements. Recently, other mouse models have been developed via overexpression of ΔE -torsinA (Sharma et al., 2005; Grundmann et al., 2007; Zhao et al., 2008) or GAG deletion in the DYT1 gene (Dang et al., 2005, 2006). However, because these models have failed to replicate such severe symptoms, the phenotype of the present transgenic mice may not be attributable entirely to overexpression of ΔE -torsinA. Electrophysiological analyses of other models of transgenic mice may improve our understanding of the pathophysiology of dystonia and enable the development of more effective treatments of it.

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モデルマウスの神経活動からジストニアの病態を考える

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要旨

ジストニアの病態生理を調べるため、ヒト DYT1 ジストニア患者の原因遺伝子である DYT1 を組み込んだ遺伝子改変マウスの神経活動を調べた。このマウスは、持続的に回転運動をするなど運動が亢進している。筋電図を記録してみると、主動筋と拮抗筋の同時収縮、持続収縮などジストニアに特徴的な異常な筋活動を示した。覚醒下で大脳基底核から神経活動を記録すると、淡蒼球外節と内節において、バースト発射やポーズ（休止期間）を伴う発射頻度の減少が見られた。大脳皮質運動野を電気刺激すると、淡蒼球外節・内節において、正常例においては観察されない早い興奮とそれに引き続く長い抑制という応答が観察された。また、淡蒼球外節・内節の体部位局在も乱れていた。大脳皮質からの入力によって、淡蒼球内節に生じる長く続く抑制が、視床・大脳皮質を脱抑制することによって、不随意運動が起こっていると考えられた。

1、はじめに

ジストニアは、不随意かつ持続的な筋肉の収縮と異常姿勢によって特徴づけられる神経疾患で、有病率はパーキンソン病の数分の一と稀ではない。しかし、症状の割には病理学的な変化が殆どなく、何らかの機能的な異常と考えられるなど、謎が多い疾患である。大脳基底核の神経生理に携わるものとして、ジストニアの病態に関する研究を行いたいと思っていたが、良い疾患モデルがないなど、きっかけがつかめないでいた。

2004年秋にスコットランドで開催された国際大脳基底核学会（IBAGS）に参加したところ、私がニューヨーク大学に留学していた時、ポスドクの同僚であった Margaret Rice 博士が、ジストニアモデルマウスの線条体におけるドーパミン伝達に関する発表を行っていた。彼女によれば、そのモデルマウスは最近、マウントサイナイ医科大学神経学教室の Pullanipally Shashidharan 博士が、ヒト DYT1 ジストニア患者の DYT1 遺伝子を組み込んで開発したもので、

他の遺伝子改変モデルと異なり、臨床症状も出ているとのことであった。

また、私たちの研究チームはサルの実験が中心であったが、2005年より知見聡美博士が加わり、げっ歯類を使った実験も新たに始まることとなった。そこで、Shashidharan 博士とは面識が無かったが、メールを送ったところ、共同研究をしようということになった。とりあえず私たちが米国を訪問するということで、日米脳（日米科学技術協力事業「脳研究」分野）のグループ共同研究に申し込んだ。

実際は、2006年秋、2007年初夏、2009年冬に、私と知見博士がマウントサイナイ医科大学神経学教室に滞在し、知見博士が電気生理学の記録を主に行い、私が手伝うこととした（図1）。Shashidharan 博士の研究室は完全な分子生物学の研究室なので、増幅器、電気刺激装置、マニピュレータ、オシロスコープ、コンピュータなど必要な電気生理の器具を全て、日本より持参した。実験結果の詳細は原著論文（Chiken et al., 2008）を見て頂くとして、本稿ではジストニアの病態を中心に考察したい。ところで、不勉強で行くまで知らなかったのだが、マウントサイナイ医科大学神経学教室はボスが C. Warren Olanow 教授、先代の教授はパーキンソン病の Hoehn-Yahr の重症度分類に名を残す Melvin D. Yahr と、超有名教室であった。

2、実験方法

遺伝子改変技術によりヒト DYT1 患者の DYT1 遺伝子を組み込み、異常 torsinA を過剰発現させたマウス（Shashidharan et al., 2005）から記録を行った。

大脳基底核疾患の際、大脳基底核ニューロンの発射頻度や発射パターンが変化し、これによって病態を説明しようと、議論されてきた。一方、動物実験において、全身麻酔をするとニューロンの発射頻度が減少したり、発射パターンが変わることが知られている。そのため、神経活動を記録し病態を調べるためには、麻酔の影響をできるだけ排除する必要がある。これまでサルを用いて行動時の神経活動を記録してきたので、無麻酔記録については、それなりに熟練していたが、げっ歯類に関しては不案内であった。そこで、げっ歯類から無麻酔記録を既に行っている福島県立医大の小林和人教授、小山純正准教授（現福島大学教授）八十島安伸博士（現大阪大学准教授）に方法を教えて頂き、行うこととした。

図2に示すように、頭部固定と記録用のチェンバーを頭部に固定する手術を、

麻酔下であらかじめ行っておく。また、大脳皮質一次運動野 (Motor cortex) の上肢領域と口腔顔面領域をマッピングしておき、それぞれに刺激電極を埋めておく。このような手術により、動物に痛みを与えることなく、頭部を固定することができ、十分、馴らしておけば、実験台の上で2-3時間は大人しくしてくれる (図2A)。そして、金属電極 (ガラス被覆エルジロイ電極) を、チェンバーを通して、大脳基底核内に刺入し、単一ニューロン活動を記録した (図2B)。記録した部位は、淡蒼球外節 (external segment of the globus pallidus, GPe) と淡蒼球内節 (internal segment of the globus pallidus, GPi) (古典的には、げっ歯類では、それぞれ淡蒼球、脚内核というが、最近では霊長類との相同性を強調して、このように呼ぶことも多い) であるが、両者とも良く似た変化を示したので、以下の実験結果は主に淡蒼球内節について述べる。また、細いワイヤー電極を上腕二頭筋と三頭筋に埋め込み、筋電図の記録も行った。

3、筋電図活動

本モデルマウスは運動が亢進しており、しばしばケージ内を回転運動している (Shashidharan et al., 2005 の Supplementary Material にある動画 <http://hmg.oxfordjournals.org/cgi/content/full/ddi012/DC1> を参照)。回転運動はジストニアらしくないが、時として頸部にジストニア様症状も現れる。

まず、本モデルマウスがジストニアに特徴的な筋電図を示すのか調べた (図3)。正常マウスでは、主動筋と拮抗筋である上腕二頭筋と三頭筋が同時に収縮することはない (図3A)。一方、本モデルマウスでも多くの場合、正常マウスと同じような活動パターンを示した (図3B)。しかし、時として主動筋と拮抗筋が時間的に同期して収縮したり (図3C)、10秒以上にわたるような持続収縮 (図3D) を示した。このような主動筋と拮抗筋の同時収縮や持続収縮は、ジストニアに特徴的な筋電図所見である。

4、大脳基底核ニューロンの自発発射活動

次に電極を脳内に刺入し、大脳基底核のうち淡蒼球外節・内節の活動を調べた (図4)。正常マウスの淡蒼球内節ニューロンは50Hz前後の高頻度で休止期間なく発射をしている (図4A, C)。本モデルマウスでは約28Hzと発射頻度が減少していた (図4B, C)。また、短い時間に複数の発射が見られるバースト発射が増えたり、発射が暫くないポーズ (休止期間) が見られるなど、発射

パターンも変わっていた（図 4 A, B）。このような自発発射活動の変化は、淡蒼球外節でも観察された。

ヒトジストニア患者の淡蒼球外節・内節からニューロン活動を記録したこれまでの報告によれば、発射頻度減少、発射パターンの変化などが観察されている。しかし、多くの場合プロボフォールなどの全身麻酔下で手術を行うので、このような自発発射活動の変化が、麻酔の影響ではないかとの疑問があった。今回、本モデルマウスでもヒト患者と同等なニューロン活動の変化が観察されたことにより、このような異常活動は麻酔のせいではなく、ジストニアの病態と関わっていることが示唆される。

5、大脳皮質刺激に対する応答と体部位局在

大脳皮質を電気刺激（持続時間 0.2ms、強度 20-50 μ A の単発刺激）すると、淡蒼球外節・内節のニューロンは、正常例では、早い興奮、抑制、遅い興奮の 3 相性のパターンで応答する（図 5 A）。これまでの霊長類を用いた実験などから、淡蒼球内節の早い興奮は大脳皮質—視床下核—淡蒼球内節路（ハイパー直接路）、抑制は大脳皮質—線条体—淡蒼球内節路（直接路）、遅い興奮は大脳皮質—線条体—淡蒼球外節—視床下核—淡蒼球内節路（間接路）を介した反応であることがわかっている。

本モデルマウスでは、一部、正常マウスと同じような反応も観察されたが、多くの淡蒼球外節・内節ニューロンにおいて、（2 相性の）早い興奮とそれに引き続く長い抑制が観察された（図 5 B）。このような反応は、正常マウスでは観察されないものである。早い興奮と遅い興奮の由来は、現在のところ不明であるが、早い興奮の少なくとも第一成分は、正常の早い興奮の潜時と同じことから、ハイパー直接路を介していると考えられる。抑制が増強した成因としては、

（1）直接路の興奮性増強、（2）視床下核から淡蒼球外節・内節への興奮性入力への減弱、（3）淡蒼球外節・内節ニューロンの興奮性の低下（例えばイオンチャンネルの変化など）が考えられる。早い興奮がハイパー直接路を介しているとするれば、視床下核から淡蒼球外節・内節への投射も正常と考えられ、（2）の可能性は低い。また、線条体を刺激した予備的な実験でも、通常は観察されない長い抑制が淡蒼球外節・内節に観察されることから、（1）と（3）の可能性が高い。

大脳皮質運動野の上肢領域と口腔顔面領域を別々に刺激し、淡蒼球外節・内

節ニューロンの反応を記録することにより、どの大脳皮質から入力を受けているのか同定することができる（図6）。正常マウスの淡蒼球内節では、多くのニューロンが上肢領域の刺激に反応し、口腔顔面領域の刺激に反応したものの数は少数であった（図6 A, Normal）。また、その両者から入力を受けるニューロンも少数であった。淡蒼球内節での分布を見てみると、上肢領域の刺激に応じるニューロンは淡蒼球内節の中央部に、口腔顔面領域の刺激に応じるものは、わずかに最外側部に観察された（図6 B, Normal GPi）。また、最内側部は、何れの刺激にも反応しなかった。これらから、淡蒼球内節は小さな核であるが、その中に体部位局在があることを示している。

一方、本モデルマウスを調べてみると、上肢領域の刺激に応答するニューロン以外に、口腔顔面領域の刺激に応じるニューロンも多く観察され、とくに両者から入力を受けるニューロンの数が有意に増えていた（図6 A, Transgenic）。淡蒼球内節での分布を見てみると、上肢領域の刺激に応じるニューロンは依然として核の中央部に存在するが、口腔顔面領域や、両者から入力を受けているニューロンが外側部だけではなく、中央部にも侵入していることがわかる（図6 B, Transgenic GPi）。一方、最内側部は応答を示さない領域が残されていた。このような体部位局在の乱れ、重なりは、本モデルマウスの淡蒼球内節ばかりでなく、淡蒼球外節でも観察された。

6、ジストニアの病態

以上、本モデルマウスの淡蒼球外節・内節において、

- 1) 発射頻度減少、バースト発射や長いポーズなどの発射パターンの変化
- 2) 大脳皮質刺激に対する興奮+長い抑制の反応
- 3) 体部位局在の乱れ

が観察された。

まず、最初に、このうちどれが第一義的な変化か考察してみたい。電気刺激と同様に、大脳皮質の自発神経活動が淡蒼球外節・内節に、早い興奮と長い抑制をもたらしているとすれば、それぞれバースト発射とポーズに対応すると考えられる。また、大脳皮質由来の抑制が増強すれば、淡蒼球外節・内節の発射頻度が減少する。さらに前節において、直接路の興奮性増強によって長い抑制が起こる可能性を考えたが、そのメカニズムのひとつとして、体部位局在が乱れていることから、ひとつの淡蒼球内節ニューロンが、多数の線条体ニューロ

ンから多重支配を受けている可能性がある。したがって、3) → 2) → 1) という因果関係が考えられる。

つぎに、ジストニアの病態について考察してみたい。大脳基底核の機能として、直接路が必要な運動を引き起こし、間接路、ハイパー直接路が不必要な運動を抑制していると考えられている(図7A)。本モデルマウスでは、直接路を介した抑制が増強し、その結果、大脳皮質の小さな興奮が淡蒼球内節を広く、あるいは時間的に長く抑制することになる(図7B)。淡蒼球内節の過剰な抑制は、視床を脱抑制し、大脳皮質を過剰に興奮させ、その結果、不必要な運動が抑制されず、意図しない時間に引き起こされる。このようにしてジストニアでは不随意運動が起こっていると考えられる。また、淡蒼球内節において体部位局在が乱れていることから、例えば上肢の運動を行う際、淡蒼球内節において本来は抑制されない口腔顔面領域も抑制されることになる(図7B)。これによって、意図した体部位以外の運動が起こり、ジストニア患者で観察される筋電図のオーバーフロー現象に相当するのではと考えられる。

今後の課題としては、大脳皮質由来の抑制が増強しているメカニズムを明らかにすることが第一である。これが明らかになれば、病理所見が乏しいジストニアの病態の本質に迫れるし、また、何らかの方法で増強された抑制を除いてやれば、ジストニアの治療法に結びつくかもしれない。さらに、全身性ジストニア、とくに DYT1 ジストニアに対して、淡蒼球内節に対する脳深部刺激療法(GPi-DBS)が著効を示し、治療の第一選択となりつつあるが、そのメカニズムは不明である。これら治療メカニズムに関しても、本モデルマウスを使うことにより明らかにしていきたい。

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図の説明

図 1 マウントサイナイ医科大学の実験室にて。左 2 人目より、筆者、Shashidharan 博士、Rice 博士、一人おいて知見博士。

図 2 実験のセットアップ。

A, 動物の固定方法 ; B, 刺激部位と記録部位を示す模式図。GPe, 淡蒼球外節 ; GPi, 淡蒼球内節 ; Motor cortex, 大脳皮質運動野 ; SNr, 黒質網様部 ; Striatum, 線条体

図 3 上腕三頭筋 (Triceps m.) と上腕二頭筋 (Biceps m.) の筋電図。

A, 正常マウス ; B-D, ジストニアモデルマウス

図 4 淡蒼球内節 (GPi) ニューロンの自発発射活動。

正常マウス (A) とジストニアモデルマウス (B) における自発発射パターンを、パルス列で示す。C, 平均発射頻度の比較。

図 5 淡蒼球内節ニューロンの大脳皮質運動野刺激に対する応答。

正常マウス (A) とジストニアモデルマウス (B) における大脳皮質刺激に対する応答を、刺激前後時間ヒストグラム (peri-stimulus time histogram, PSTH; bin 幅, 1ms; 100 回加算) で示す。

図 6 淡蒼球内節の体部位局在。

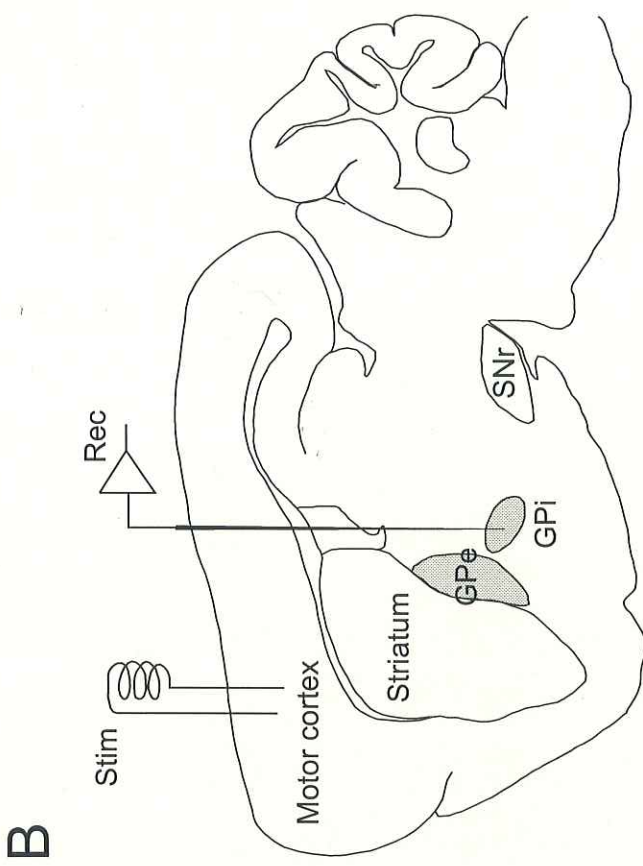
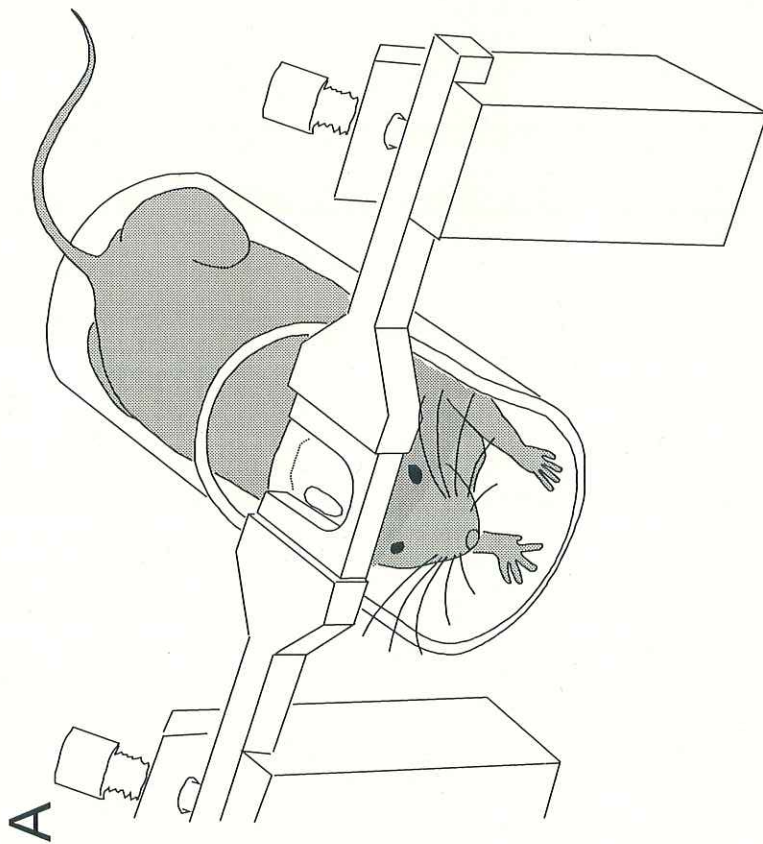
A, 大脳皮質運動野の上肢領域 (forelimb) と口腔顔面領域 (orofacial) から入力を受ける淡蒼球内節ニューロンの割合を正常マウス (Normal) とジストニアモデルマウス (Transgenic) で示す。

B, 大脳皮質運動野の上肢領域と口腔顔面領域から入力を受けるニューロンの淡蒼球内節における分布を正常マウスとジストニアモデルマウスで示す。

図 7 正常 (A) とジストニア (B) の大脳基底核の活動を示す模式図。

STN, 視床下核





A Normal

Triceps m.



Biceps m.



1 mV

50 ms

B Transgenic

Triceps m.



Biceps m.



1 mV

50 ms

C Transgenic

Triceps m.



Biceps m.



1 mV

50 ms

D Transgenic

Triceps m.

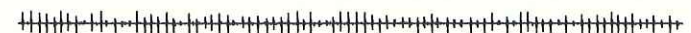


Biceps m.



0.5 mV

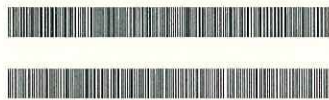
5 s



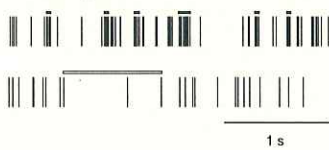
0.2 s

GPI

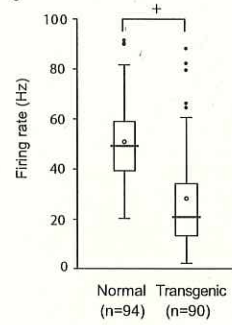
A Normal



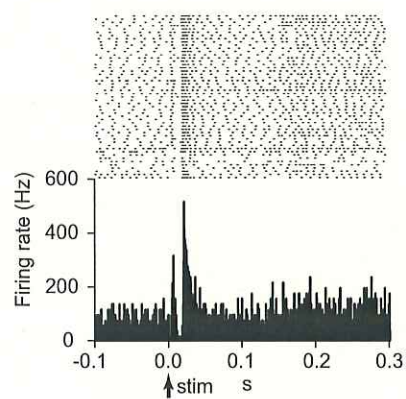
B Transgenic



C



A Normal GPi



B Transgenic GPi

