

## Retinoic acid signaling: a new piece in the spoken language puzzle

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# 1 **Retinoic acid signaling: a new piece in the spoken language puzzle**

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12 **Retinoic acid, FoxP2, synaptic plasticity, development, motor skills, striatum, dopamine**  
13 **receptor**

14 **Abstract**

15 Speech requires precise motor control and rapid sequencing of highly complex vocal musculature.  
16 Despite its complexity, most people produce spoken language effortlessly. This is due to activity in  
17 distributed neuronal circuitry including cortico-striato-thalamic loops that control speech-motor  
18 output. Understanding the neuro-genetic mechanisms involved in the correct development and function  
19 of these pathways will shed light on how humans can effortlessly and innately use spoken language  
20 and help to elucidate what goes wrong in speech-language disorders.

21 *FOXP2* was the first single gene identified to cause speech and language disorder. Individuals with  
22 *FOXP2* mutations display a severe speech deficit that includes receptive and expressive language  
23 impairments. The neuro-molecular mechanisms controlled by *FOXP2* will give insight into our  
24 capacity for speech-motor control, but are only beginning to be unraveled. Recently *FOXP2* was found  
25 to regulate genes involved in retinoic acid signaling and to modify the cellular response to retinoic  
26 acid, a key regulator of brain development. Here we explore evidence that *FOXP2* and retinoic acid  
27 function in overlapping pathways. We summate evidence at molecular, cellular and behavioral levels  
28 that suggest an interplay between *FOXP2* and retinoic acid that may be important for fine motor control  
29 and speech-motor output. We propose retinoic acid signaling is an exciting new angle from which to  
30 investigate how neuro-genetic mechanisms can contribute to the (spoken) language ready brain.

31

32 **Main text**

33 **Speech and spoken language**

34 Speech is the primary modality by which humans use language, and human orofacial morphology is  
35 uniquely suited to the production of intricate vocalizations needed for spoken language (Lieberman,  
36 2007). The orofacial musculature is one of the most complex muscle systems in the body and in order  
37 to successfully produce meaningful speech these muscles must be controlled and coordinated in rapid  
38 sequences involving distributed neuronal circuitry. This motor activity is generated in several neural  
39 loops that select appropriate actions and generate the necessary motor patterns. One crucial circuit, the  
40 cortico-basal ganglia loop, sends activity from the motor cortex to the striatum (a component of the  
41 basal ganglia) where activity is integrated. Subsequently, outputs from here modulate activity in  
42 several thalamic nuclei. Activity from the thalamus is then sent back to the motor cortex, where a  
43 specialized population of output neurons organizes the complex thalamocortical inputs (Kravitz and  
44 Kreitzer, 2012; Calabresi et al., 2014). These cortical output neurons send the information, via the  
45 pyramidal tract, to motor neurons directly controlling muscle tissue. These neurons are either located  
46 in the spinal cord (controlling limb and body movements), or in the brainstem's cranial nerve nuclei  
47 (controlling facial and vocal tract movements). An illustration of the cortico-basal ganglia loop (in the  
48 rodent brain) is given in Figure 1A. Proper connectivity within this pathway is necessary to enable the  
49 precise outputs needed for orofacial muscle control.

50 The striatum can be seen as a central hub within the motor pathway, making it one of the most  
51 intriguing regions in which to investigate properties of motor circuitry and orofacial control. Striatal  
52 activity is especially important for fine motor behavior and motor skill learning (Doyon et al., 2003)  
53 and cortical and subcortical circuitry, including the striatum, has been established as highly important  
54 for speech-motor control (Lieberman, 2002). Furthermore, increased activation of the basal ganglia  
55 (which incorporates the striatum) has been shown via functional brain imaging (fMRI) in specific  
56 speech-motor language tasks (Wildgruber et al., 2001; Booth et al., 2007). Lastly, morphological  
57 changes in the striatum have been described in individuals with speech problems such as stuttering  
58 (Craig-McQuaide et al., 2014) and non-fluent aphasia (Ogar et al., 2007).

59 The principal cell type in the striatum is the medium spiny neuron (MSN), which makes up  
60 approximately 98% of all striatal cells (Kemp and Powell, 1971; Huang et al., 1992) (for review see  
61 Kreitzer and Malenka, 2008). MSNs can be further divided into two categories of neurons that have

62 different connectivity and opposing functions: dopamine receptor type 1 (D1R) and dopamine receptor  
63 type 2 (D2R) expressing cells (Figure 1A). D1R expressing MSNs connect to thalamic nuclei via the  
64 “direct pathway” which results in excitation of the motor cortex. D2R expressing MSNs form an  
65 “indirect pathway” that connects to the thalamus via multiple subcortical regions leading to inhibition  
66 of the thalamus and thus reduced cortical input (Figure 1A) (Albin et al., 1989; Kravitz and Kreitzer,  
67 2012; Calabresi et al., 2014). This balance between excitation (resulting in more movement) and  
68 inhibition (less movement) is crucial for coordinated motor function (Calabresi et al., 2014) including  
69 fine orofacial motor control.

70 In order to unravel the fundamental components that enable humans to effortlessly use spoken  
71 language, we will need to understand the neuro-genetic mechanisms involved in establishment,  
72 function and maintenance of speech-motor pathways.

73

#### 74 **Spoken language and FoxP2**

75 A breakthrough in speech and language genetics came with the identification of the first gene to cause  
76 a speech/language disorder: *FOXP2* (Lai et al., 2001). Mutations in *FOXP2* were found in a large  
77 pedigree known as the KE family (Hurst et al., 1990; Fisher et al., 1998; Lai et al., 2001). Affected  
78 family members were diagnosed with a severe speech impairment known as developmental verbal  
79 dyspraxia (also known as childhood apraxia of speech; OMIM: 602081) and carried a mutation in one  
80 copy of their *FOXP2* gene. In addition to speech impairments, affected family members demonstrated  
81 receptive and expressive language problems (Watkins et al., 2002a). Although rare, *FOXP2* mutations  
82 have been found in a number of unrelated families and individuals with similar speech/language  
83 phenotypes (MacDermot et al., 2005; Feuk et al., 2006; Shriberg et al., 2006; Lennon et al., 2007; Palka  
84 et al., 2012; Rice et al., 2012; Zilina et al., 2012). For review see Bacon and Rappold, 2012. In depth  
85 investigations of the KE family phenotype indicated a severe impairment in orofacial praxis tasks  
86 (Vargha-Khadem et al., 1995; Lai et al., 2001; Watkins et al., 2002a). In addition, impairments in  
87 language production tasks (e.g. phoneme addition, word repetition) were found between control and  
88 affected individuals (Vargha-Khadem et al., 1995). Different aspects of speech are thus impaired in  
89 KE family members (Watkins et al., 2002a). Orofacial praxis deficits underlie impaired lexicon  
90 building and subvocal (internal) speech representations which can affect irregular verb grammar  
91 (Doyon et al., 2003) and rule based grammar learning (Ullman, 2001). Thus, some of the language  
92 impairments in the KE family could be related to the core speech production deficits observed.

93 FOXP2, and its murine homolog *Foxp2*, are found across many regions of the developing and postnatal  
94 brain (*FoxP2* will be used when referring to both species). Intriguing is the high expression of *FoxP2*  
95 throughout the mouse and human cortico-striato-thalamic motor circuitry (Lai et al., 2003b). During  
96 early development *FoxP2* is broadly expressed in these regions, but in later developmental and  
97 postnatal stages expression becomes more restricted (Figure 1B depicts *Foxp2* expression in the  
98 postnatal mouse brain). In adults, *Foxp2* is limited to deep layer cortical neurons (layer 5 motor cortex  
99 and layer 6 throughout) (Ferland et al., 2003; Morikawa et al., 2009; Hisaoka et al., 2010; Tomassy et  
100 al., 2010; Reimers-Kipping et al., 2011; Tsui et al., 2013). Within the striatum, *Foxp2* is highly  
101 expressed in both types of MSN, though more commonly in D1R MSNs compared to D2R neurons  
102 (Vernes et al., 2011). Commensurate with its expression pattern, imaging studies have shown humans  
103 with *FOXP2* mutations display structural and functional differences in motor areas. Affected members  
104 of the KE family showed structural grey matter volume differences in the motor cortex and striatum  
105 (Watkins et al., 2002b). Furthermore, functional imaging studies showed an underactivation of the  
106 striatum and altered cortical activation (including speech/motor areas such as the left anterior insular  
107 cortex) during word generation and word repetition tasks (Liegeois et al., 2003).

108 Converging evidence from *FoxP2* expression pattern studies and phenotypic characterization of human  
109 mutations suggests that *FOXP2* may play an important role in the development of the speech-motor  
110 pathway. The high expression of *Foxp2* in a specific subset of neurons (D1R MSNs) in the striatum  
111 indicates a functional specificity related to motor tasks requiring the striato-thalamic connections of  
112 the direct pathway. Malfunctions within this pathway could ultimately affect aspects of the motor  
113 circuitry related to fine motor control and contribute to the observed speech-motor deficit in humans.

114

### 115 **FOXP2 as a molecular entry point into speech-motor pathways**

116 *FOXP2* is a transcription factor; its molecular function is to regulate the expression of other genes,  
117 switching them on or off in a temporally and spatially controlled manner. *FoxP2* has been shown to  
118 regulate hundreds of different genes involved in processes crucial to brain development and function,  
119 ranging from neurogenesis and migration to neurite outgrowth and synaptic activity (Spiteri et al.,  
120 2007; Vernes et al., 2007; Konopka et al., 2009; Vernes et al., 2011; Devanna et al., 2014). Recently,  
121 evidence has suggested that *FOXP2* regulates a number of genes involved in the retinoic acid (RA)  
122 signaling pathway (Devanna et al., 2014). Retinoic acid is a vitamin-A derivative essential to  
123 mammalian development and disruption of the retinoic acid signaling pathway (caused by genetic

124 disruptions or dietary deficiencies) can have severe consequences during development and adulthood  
125 (Holson et al., 1997; Krezel et al., 1998b)

126 RA induces genetic and morphological changes in cells. When neuronal precursors (cells that generate  
127 neurons during development) differentiate into neurons they switch on genes normally found in mature  
128 neurons, stop dividing and grow long processes known as neurites (Siegenthaler et al., 2009; Korecka  
129 et al., 2013). We previously compared how neuron-like cells with or without FOXP2 responded to RA  
130 and found that cells showed stronger genetic and morphological changes in response to RA if FOXP2  
131 was present (Devanna et al., 2014). In addition we discovered that FOXP2 changed the expression of  
132 retinoic acid receptors - proteins that directly control the cellular response to RA (Devanna et al., 2014).  
133 Of particular interest, FOXP2 upregulated RAR $\beta$  and a number of other genes involved in transport or  
134 modification of retinoic acid were also transcriptionally regulated (e.g. ROR $\beta$ , CRABP II and ASCL1).  
135 These experiments suggest an intriguing link between FOXP2 and the RA pathway, in which FOXP2  
136 seems to contribute to or modify the cellular response to RA.

137 Given the importance of the RA pathway for development, this raises new questions about how FOXP2  
138 might mediate its effects on brain and neural circuit development. Could the relationship between  
139 FOXP2 and the RA pathway be relevant for (1) normal motor circuitry development and function,  
140 and/or (2) effects of FOXP2 dysfunction in patients? To address these questions, we need to understand  
141 how FoxP2 and the RA pathway might interact and in what way FoxP2 mutations might affect the RA  
142 pathway on a cellular, functional and behavioral level.

143

#### 144 **RA, FoxP2 and motor behavior**

145 Retinoic acid is a key compound during embryogenesis, affecting a multitude of critical developmental  
146 pathways. Precise control of RA levels is essential for normal brain development as either an excess  
147 or a deficiency of RA results in widespread adverse effects on the brain.

148 Gestational treatment of rats with excess RA results in behavioral deficits in learning, memory and  
149 motor function (Holson et al., 1997). Rats treated with excess RA displayed poor generalized motor  
150 control including impairments in the 'righting reflex' (the ability to return to upright position), and the  
151 ability to sit only on the back paws. In addition, gestationally treated adult rats showed problems with  
152 learning and memory, such as decreased learning rates in a water filled T maze (Butcher et al., 1972;  
153 Holson et al., 1997). Rats lacking dietary vitamin A (of which retinoic acid is a metabolite) also perform  
154 poorly on motor learning and motor performance tasks (Carta et al., 2006). Furthermore, mice

155 engineered to lack a key facilitator of retinoic acid signaling (the retinoic acid receptor  $RAR\beta$ ) develop  
156 severe locomotion deficits and are highly impaired on motor learning tasks (Krezel et al., 1998a).

157 The displayed motor deficits are similar to phenotypes observed in mouse models of *Foxp2*  
158 dysfunction. Mouse models of two well characterized patient mutations of *FOXP2* have been created  
159 that have comparable phenotypes. One mouse model reflects the R553H missense mutation found in  
160 the KE family (Lai et al., 2001). The second mouse model mirrors an early stop codon in exon 7  
161 introduced by a nonsense mutation that leads to a loss of FoxP2 protein in an independent family with  
162 speech/language disorder (MacDermot et al., 2005; Groszer et al., 2008). Mice that have a homozygous  
163 *Foxp2* mutation show severe general motor impairments, reminiscent of animals treated with excess  
164 RA. However these *Foxp2* homozygous mutants do not survive beyond 3-4 weeks after birth, possibly  
165 due to a requirement for *Foxp2* in other organs such as the lungs or heart (Groszer et al., 2008). In mice  
166 where a single copy of *Foxp2* is affected (as per the heterozygous state of the mutations observed in  
167 patients) general motor control is normal but motor learning is impaired (Groszer et al., 2008; French  
168 et al., 2012). This more subtle phenotype closely resembles the motor learning phenotype observed in  
169 RA deprived rats (Carta et al., 2006). For an overview of the different phenotypes exhibited by *Foxp2*  
170 mutation, *RAR* mutation and RA treatment, see Table 1.

171

## 172 ***Foxp2* and RA signaling affect neuronal function**

173 In addition to the behavioral deficits, vitamin A deprivation/supplementation adversely affects striatal  
174 development and function. Cells in the developing lateral ganglionic eminence (the precursor region  
175 of the striatum) do not differentiate into the appropriate neuronal subtypes when RA signaling is  
176 blocked (Toresson et al., 1999; Chatzi et al., 2011). However restoring RA levels rescued this  
177 phenotype and resulted in normal differentiation into appropriate neuronal cell types (Chatzi et al.,  
178 2011). Separately, mice engineered to knockout the retinoic acid receptor  $\beta$  (*RAR\beta*) gene display gross  
179 morphological striatal defects including impaired neurogenesis and deficits in acquiring proper  
180 neuronal identities (Liao et al., 2008). Lastly, chronic postnatal vitamin A supplementation has been  
181 linked to oxidative cell toxicity in the striatum (de Oliveira et al., 2007).

182 *Foxp2* also contributes to striatal cell morphology and function. *Foxp2* mutant neurons exhibit reduced  
183 neurite growth and branching in primary striatal cultures (Vernes et al., 2011) and in the *in vivo* striatum  
184 displays aberrant neuronal activity. Mice with a heterozygous *Foxp2* mutation showed unusually high  
185 activity in the dorsomedial striatum during active motor behavior (French et al., 2012). This suggests

186 striatal cells can no longer properly modulate their activity following input from motor areas when  
187 lacking *Foxp2*. Moreover, the increased striatal activity normally seen when animals perform motor  
188 learning tasks was absent in mutant mice. Instead, a decrease in firing rate was seen, again suggesting  
189 aberrant modulation of responses to cortical and/or thalamic input (French et al., 2012). Additionally,  
190 extracellular measurements on striatal brain slices from heterozygous *Foxp2* mutant animals show  
191 these cells fail to respond to induction of long term depression (Groszer et al., 2008). An inability to  
192 induce long term plasticity (either long term depression (LTD) or potentiation (LTP)) has debilitating  
193 consequences as scaled activity (plasticity) is necessary for circuits to properly regulate their input and  
194 output. Synaptic long term plasticity changes underlie information storage and are necessary for  
195 learning and memory (Novkovic et al., 2015; Zhu et al., 2015). Interestingly, in the striatum, synaptic  
196 plasticity has been strongly linked to motor learning (Dang et al., 2006; Kreitzer and Malenka, 2007).  
197 Defects specifically related to striatal LTD and LTP are known to affect procedural motor learning and  
198 the acquisition of new motor paradigms (Gubellini et al., 2004).

199 Aberrant induction of synaptic scaling has also been found in mice following acute retinoic acid  
200 depletion, which results in a complete lack of hippocampal LTP or LTD (Misner et al., 2001). This  
201 phenotype was specific to RA depletion and was reversible, as vitamin A supplementation rapidly  
202 restored normal synaptic plasticity (Misner et al., 2001). At a molecular level, RA signaling is mediated  
203 by the action of retinoic acid receptors (RARs;  $RAR\alpha$ ,  $RAR\beta$  and  $RAR\gamma$ ) and similar plasticity defects  
204 have been shown for mice lacking  $RAR\alpha$  (Sarti et al., 2012) or  $RAR\beta$  (Chiang et al., 1998).  
205 Hippocampal cells from these mice fail to establish LTD when subjected to low frequency stimulation  
206 – the paradigm necessary to induce LTD in the hippocampus. By contrast, excess RA induced the  
207 reverse effect in cultured hippocampal slices, where increased excitatory activity was observed (Aoto  
208 et al., 2008). It is not yet known if RA signaling affects synaptic plasticity in the striatum. However,  
209 the similarity in synaptic activity phenotypes between *Foxp2*-,  $RAR\alpha$ - and  $RAR\beta$ -deficient animals  
210 (albeit focusing on different brain regions) does indicate these transcription factors may play a role in  
211 similar intracellular pathways regulating neuronal activity and synaptic plasticity.

212 The aforementioned plasticity (LTD and/or LTP) deficits in *Foxp2*,  $RAR\alpha$  and  $RAR\beta$  mutant animals  
213 suggests an improper reaction of neuronal circuits to changes in external input. Induction of LTD or  
214 LTP leads to a decrease or an increase, respectively, in the amount of glutamate receptors (of the  
215 AMPA-receptor class) at the synaptic membrane (Seidenman et al., 2003; Briand et al., 2014) (for  
216 review see Luscher and Huber, 2010). This change in AMPA receptor abundance modifies the response  
217 strength of a cell when it is excited. The change in stimulus-response strength is transient, and in time



218 the normal AMPA receptor distribution will be restored, returning synaptic responses to normal levels.  
219 RA treatment of hippocampal cultures has shown an increase of AMPA receptors on the cell surface  
220 (Aoto et al., 2008), but no data on the striatum is currently present. The shared synaptic plasticity defect  
221 following disruption of RA signaling pathways or Foxp2 mutation does suggest that they both may  
222 influence receptor abundance or localization at the synapse in the striatum, an intriguing area for further  
223 study.

224 A thorough investigation of the mechanisms leading to LTD and LTP deficits resulting from RA/RAR  
225 and Foxp2 malfunction will be necessary to understand if they function in the same pathways.  
226 Understanding the molecular mechanisms underlying striatal function, especially related to complex  
227 motor circuitry function, will lead to a better understanding of striatal speech-motor control.

228

### 229 **Molecular links between RARs and FOXP2**

230 RARs canonically function as transcription factors, regulating genes responsible for directing normal  
231 embryogenesis and brain development. Interestingly, FoxP2 and RARs share some of the same target  
232 genes (Balmer and Blomhoff, 2002; Delacroix et al., 2010; Devanna et al., 2014). RARs are highly  
233 expressed in the brain (Krezel et al., 1999) and are present throughout embryonal development  
234 (Mollard et al., 2000), postnatal development (Wei et al., 2011), and in adults (Krezel et al., 1999;  
235 Zetterstrom et al., 1999). Notably high expression of RARs can be found throughout the motor  
236 circuitry, including cortical, striatal and multiple thalamic regions (Krezel et al., 1999) (Figure 1B).  
237 We focus on two key receptors found in the motor circuitry: RAR $\alpha$  and RAR $\beta$ . RAR $\alpha$  is found in layer  
238 5 of the cortex and in the thalamus - both regions that overlap with murine Foxp2 expression (Krezel  
239 et al., 1999; Zetterstrom et al., 1999; Ferland et al., 2003; Lai et al., 2003a; Hisaoka et al., 2010).  
240 Interestingly, Foxp2 only overlaps with RAR $\alpha$  in the motor cortex layer 5, because Foxp2 expression  
241 is largely restricted to layer 6 of other mature cortical areas. RAR $\beta$  is strongly expressed only in the  
242 striatum, another site where Foxp2 expression is highest (Figure 1B). Notably, FOXP2 has been shown  
243 to directly drive RAR $\beta$  expression in cell models (Vernes et al., 2007; Devanna et al., 2014), although  
244 this is yet to be shown in the striatum. This high level of overlap, combined with shared target genes  
245 and molecular interactions, strongly supports interplay between FoxP2 and RARs in motor pathways.

246

### 247 **Concluding remarks**

248 In addition to its canonical role during embryogenesis, studies described here suggest RA signaling  
249 plays a specific role in the development and function of striatal motor circuitry and may link to FoxP2  
250 function. Disruption of the RA pathway results in strikingly similar phenotypes to FoxP2 mutation on  
251 multiple levels, which suggests a potential mechanistic interaction. FoxP2 and RARs can regulate some  
252 common target genes, affect similar cellular phenotypes and show highly overlapping expression  
253 patterns in the cortico-striato-thalamic motor circuitry. In the striatum, aberrant function of Foxp2 and  
254 RA signaling contributes to altered development and, in the case of mutations of mouse Foxp2, altered  
255 synaptic plasticity similar to that seen in the hippocampus of RAR $\alpha$  mutant animals. Given that RAR $\beta$   
256 is predominantly expressed in the postnatal striatum, it seems likely that its disruption will also affect  
257 striatal plasticity, however this is yet to be experimentally determined. Lastly, animals with mutated  
258 Foxp2 or RA signaling defects show comparable motor control/learning impairments. Thus at multiple  
259 levels (molecular, cellular, circuit, and behavioral) there is evidence that interplay between FoxP2 and  
260 RA signaling may facilitate proper development and function of motor circuitry. This evidence from  
261 mice is strengthened by findings in songbirds which show both FoxP2 and RA influence song learning  
262 by acting in circuits that have parallels with human vocal-motor pathways (Haesler et al., 2007; Wood  
263 et al., 2008). In the future it will be of great value to understand if these signaling cascades interact to  
264 influence neuronal mechanisms related to song learning or speech-motor control, and if RA signaling  
265 deficits are involved in aberrant speech-motor development in humans. The capacity for human speech  
266 and spoken language is dependent on multiple molecular and neural building blocks. With the link  
267 between FoxP2 and RA signaling, a new block has been suggested, giving us new opportunities to  
268 investigate the evolution and development of the (spoken) language ready brain.

269

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### 274 **Conflict of interest**

275 The authors report that they have no involvements that might raise the question of bias in the work  
276 reported or in the conclusions, implications, or opinions stated.

277

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487 **FIGURE LEGENDS**

488 **Figure 1. Foxp2 and retinoic acid receptors show overlapping expression patterns in motor**  
489 **associated circuitry.** (A) An overview of the direct and indirect pathways represented in the sagittal  
490 overview showing connectivity between different regions. Dopamine receptor type 1 (D1R) and  
491 Dopamine receptor type 2 (D2R) expressing cells in the striatum are separated to highlight direct and  
492 indirect pathways. (B) Sagittal Schematic of the mouse brain showing that Foxp2, RAR $\alpha$  and RAR $\beta$   
493 are all expressed in motor associated circuitry. RAR $\alpha$  and RAR $\beta$  are expressed in distinct regions, but  
494 each receptor partially overlaps with Foxp2. RAR $\alpha$  and Foxp2 can be found in deep layers of the cortex,  
495 thalamus, subthalamic nucleus (STN), the internal (GPi) and external (GPe) globus pallidus,  
496 cerebellum and olfactory bulbs (OB). Foxp2 and RAR $\beta$  overlap in the striatum. RAR $\alpha$  shows non-  
497 overlapping expression in the hippocampus (hi.), RAR $\beta$  in the hypothalamus (hy) and Foxp2 in the  
498 substantia nigra (SN). Connectivity between regions involved in motor processing (including outputs  
499 to brain stem nuclei and spinal cord) is shown by solid lines. The direct (excitatory) and indirect  
500 (inhibitory) pathways, which are the two outputs from the striatum, are shown by dashed lines.

501 **Table 1:** Overview of phenotypes described in Foxp2 mutation, RAR mutation and RA  
502 excess/depletion treatments. ‘-’ = no effect, ‘+’ = mild effect, ‘++’ = strong effect, ‘N/A’ = not  
503 applicable, ‘NT’ = not tested

504

Deficit		Foxp2 mutation		RA receptor mutation	RA excess/depletion
		homozygous	heterozygous		
Development	Embryogenesis defects	-	-	-	+
	Lethality	++	-	+	++

	Aberrant basal ganglia development	++	+	++	++
<b>Cellular</b>	basal ganglia cell identity defects	<b>NT</b>	-	+	++
	Decreased neurite growth and branching	++	<b>NT</b>	<b>NT</b>	<b>NT</b>
	Aberrant neuronal activity in striatum	++	++	-	+
	Unable to induce LTD	<b>NT</b>	++	++	<b>NT</b>
	Unable to induce LTP	<b>NT</b>	<b>NT</b>	++	<b>NT</b>
	General motor control deficits	++	-	++	++
<b>Behavior</b>	Motor learning deficits	<b>N/A</b> (postnatal lethality)	+	++	++ (postnatal treatment)
	Spatial learning deficits	<b>N/A</b> (postnatal lethality)	<b>NT</b>	+	+

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Figure 1.JPEG

