

# Intrinsic programs regulating dendrites and synapses in the upper layer neurons of the cortex

Beatriz Cubelos<sup>1,2</sup> and Marta Nieto<sup>1</sup>

<sup>1</sup>Centro Nacional de Biotecnología; CSIC; Campus de Cantoblanco; <sup>2</sup>Centro de Biología Molecular Severo Ochoa; CSIC-UAM; Madrid, Spain

**Key words:** cerebral cortex, *Cux2*, *Cux1*, *Cut11*, *Cut12*, upper layer, dendrite, spine, synapse, synaptogenesis, *Xlr*, mental retardation

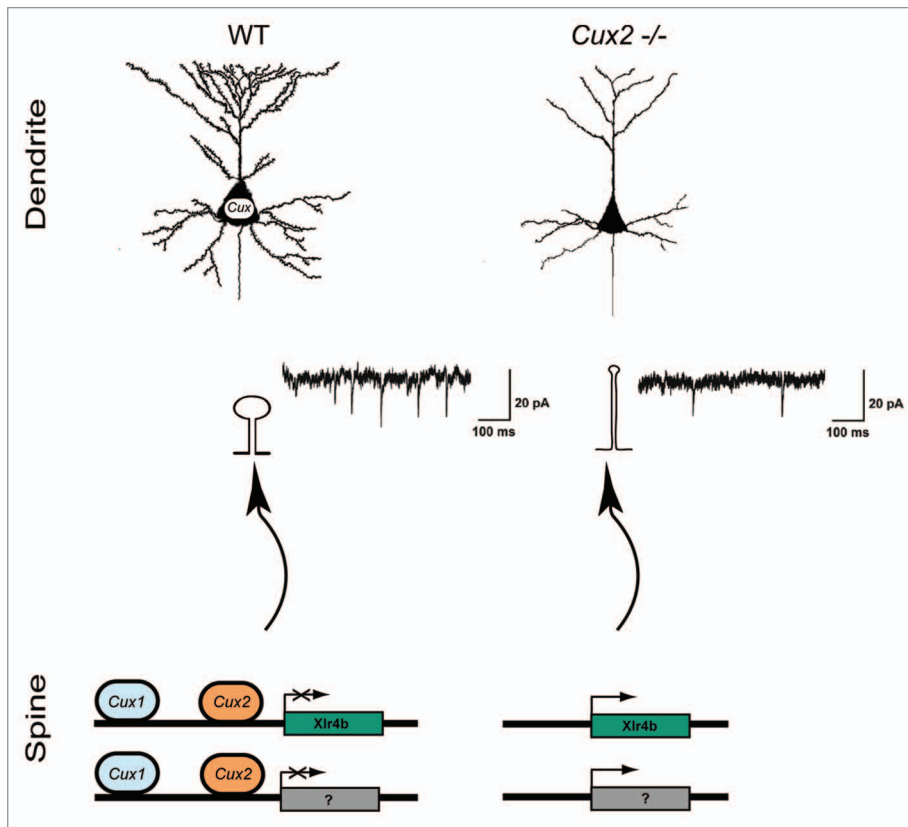
Dendrites and spines are key regulators of neuronal function often affected in cognitive disorders. Neuronal subclasses are characterized by a wide range of dendritic morphologies that aid their specific functions. However, how subclass-specific dendritic trees arise during vertebrate development remains largely unknown. We have recently reported that the restricted expression of *Cux1* and *Cux2* genes in the upper layers of the cerebral cortex determines the specific morphology of dendrites and spines and the function of these neurons. Since *Cux* genes are the vertebrate homologs of *Drosophila Cut*, which specifies the dendritic morphologies of certain sensory neuron populations, our findings suggest that mechanisms of dendrite differentiation are conserved between *Drosophila* and mammals, which had yet to be demonstrated. Importantly, we found that *Cux* genes not only modulate dendritic branching, but also dendritic spine morphogenesis, the functional synapse and cognition. Dendritic spine stabilization was partly mediated by direct repression of genes of the *Xlr* family, previously implicated in cognitive defects in a model of Turner syndrome. Hence, our work indicates that neuronal subclass specific determinants may intrinsically affect synaptic activity beyond expected. The functions of *Cux1* and *Cux2* were additive and complement each other to establish the final pattern of the dendritic tree and the number and strength of the synapses. This work unravels novel mechanisms of dendritogenesis and synaptogenesis and illustrates how regulating dendritic structures contributes to the specialization of upper layer neurons. It will be interesting to dissect how these mechanisms regulate cortical activity, area specialization and cognitive functions.

Dendritic branches and spines are key regulators of neuronal function. Number, growth and orientation of dendritic branches modify the way information is integrated and select axonal targets.<sup>1,2</sup> Dendritic spines are specialized structures that modulate the activity, strength and stability of the synapse.<sup>3-5</sup> Both structures are often altered in number and morphology in mental disorders.<sup>6-8</sup> From the times of Cajal we learned to identify the myriad of different types of neurons by their stereotyped dendritic morphologies,

and we now begin to understand how these dendritic patterns serve their specialized functions.<sup>1</sup> However, how subclass-specific neuronal features are defined during development is largely unknown. Based in studies on *Drosophila*, dendritic architecture is thought to be instructed by the selective expression of transcription factors (TFs), but few of such TFs have been described in vertebrates.<sup>2,9</sup> Moreover, the possibility that these subclass specific intrinsic factors may affect the formation of dendritic spines and synapses has not been proposed or explored.

*Cux1* and *Cux2* are two transcription factors selectively expressed in the pyramidal neurons of the upper layers (II, III and IV) of the mouse cortex.<sup>10,11</sup> These neurons have elaborated dendritic morphologies and profusion of spines<sup>12</sup> that allow them to integrate intracortical circuits involved in the higher cognitive tasks of the brain. They might be considered as a late evolutive addition, since they appear in mammals and are highly represented in the primate cerebral cortex, especially in humans.<sup>13</sup> We have recently found that *Cux* genes exert an intrinsic control of the dendritic structures of the upper layer neurons of the mouse cerebral cortex (Fig. 1).<sup>15</sup> Our knockout and knockdown studies demonstrate that the homeobox *Cux1* and *Cux2* are early regulators of dendrite branching in a cell autonomous manner. *Cux* genes are the vertebrate homologs of *Drosophila* homeobox *Cut*, which specifies the dendritic morphology of certain sensory neurons.<sup>14</sup> Our findings support the existence of conserved mechanisms of dendritic differentiation between flies and mammals. They may also imply that the activity of *Drosophila Cut* in specifying simpler neuronal types might have been co-opted during cortical evolution to generate the more complex neurons of mammals. But perhaps more unexpected, we found that *Cux1* and *Cux2* instruct also genetic programs that control the number and morphology of the dendritic spines. In the absence of *Cux* genes, the dendritic spines adopt a more immature morphology, with longer necks and smaller heads. Correspondingly, electrophysiological recordings show reduced number and strength of the synapses. A few other TFs, such as MEF2, have been previously implicated in activity dependent spine formation and synaptogenesis,<sup>8,16,17</sup> but these mechanisms apply to most neuronal populations. The implication of *Cux1* and *Cux2* on neuronal plasticity and in normal brain function remains to be understood. Why is it important to selectively restrict or promote synapse formation by intrinsic factors? Do presynaptic axons need this constrains to define their connectivity?

\*Correspondence to: Marta Nieto; Email: mnlopez@cnb.csic.es  
Submitted: 06/18/10; Accepted: 06/18/10  
Previously published online  
[www.landesbioscience.com/journals/cib/article/12755](http://www.landesbioscience.com/journals/cib/article/12755)  
DOI: 10.4161/cib.3.6.12755



**Figure 1.** *Cux* genes control dendrite branching and synaptogenesis. *Cux1* and *Cux2* regulate neuronal differentiation and control intrinsic mechanisms of dendrite development, spine formation and synaptic function in upper layers in the cortex. Upper part: Dendritic patterns in WT and *Cux2*<sup>-/-</sup> pyramidal neurons of the upper layers. Lower parts: Downregulation of *Xlr3b* and *Xlr4b* gene expression by *Cux* proteins contributes to dendritic spine differentiation. *Cux1* and *Cux2* bind and regulate different regions in the *Xlr4b* locus. Miniature excitatory postsynaptic current (mEPSCs) from layer II and III pyramidal cells of *Cux2*<sup>-/-</sup> mice were reduced in amplitude and frequency.

Another interesting point is the additive and complementary functions of *Cux1* and *Cux2*. *Cux1* and *Cux2* label most neurons of the superficial layers and display overlapping patterns of expression in several areas of the cortex. This indicates that they are likely co-expressed in many upper layer neurons, as we formally corroborated for the neurons of the somatosensory cortex. Initially, this suggested us that *Cux1* and *Cux2* might have redundant functions.<sup>10,18</sup> However, we found that *Cux1* and *Cux2* are complementary but not redundant. Upper layer neurons of the somatosensory cortex of both *Cux1* and *Cux2* single mutants show similar reduction in dendritic complexity and comparable defects in dendritic spine numbers and morphologies. Double loss of *Cux1* and *Cux2* expression induced more dramatic defects in dendrites and spines. Ectopic expression of *Cux1* in cingulate neurons, that normally express *Cux2* but lower levels of *Cux1*, increased branching and reproduced the more complex dendritic morphologies of the somatosensory areas. Therefore, we concluded that the functions of *Cux* genes add to each other to stimulate branching and that it is the combinatorial expression of *Cux1* and *Cux2* that defines the final dendritic pattern.

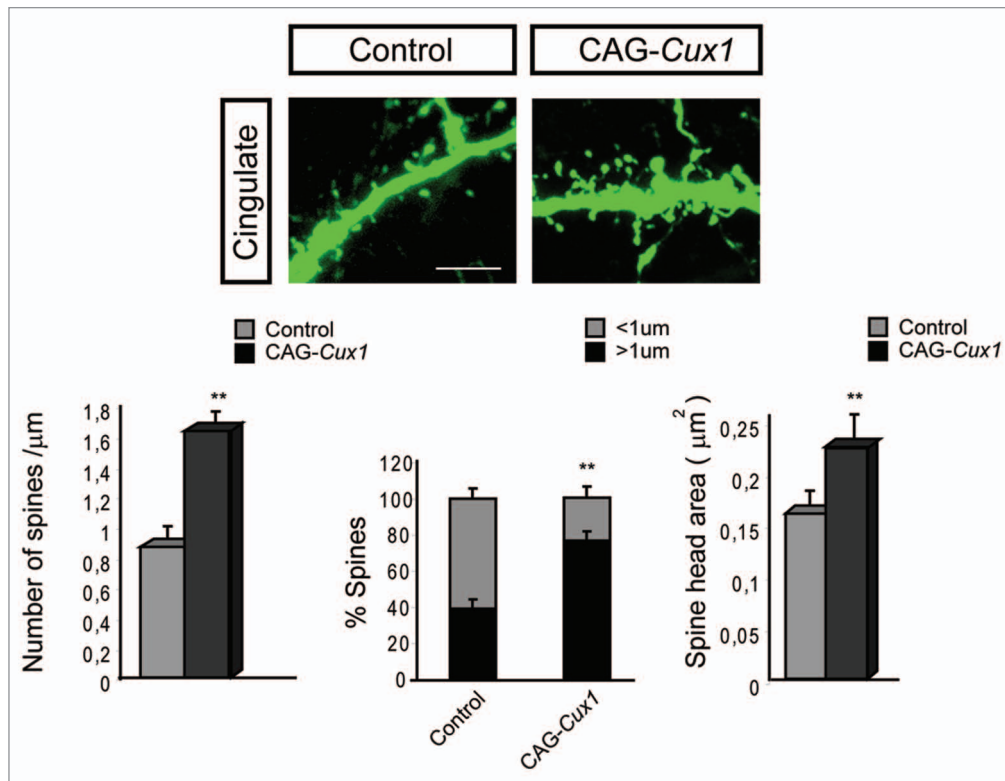
Out of these experiments, it should be highlighted that over-expression of *Cux1* also incremented dendritic spine density and

induced more mature dendritic spine phenotypes (Fig. 2). Thus, the additive functions were revealed also for the intrinsic control of the spine. Altogether, these suggest that discrete differences in the levels of expression of *Cux1* and *Cux2* may modulate dendritic and spine morphogenesis in a dose-dependent manner in subsets of superficial neurons or regionally, in cortical areas. This would refine their functions and establish a fine tuning of their connectivity.

In search for the downstream elements by which *Cux* genes exert their functions we found mechanisms of synaptogenesis key to cognition, including the regulation of NMDAR2B and PSD95.<sup>8,19-21</sup> Directly downstream of *Cux*, we found the chromatin remodeling genes of the *Xlr* family (Fig. 2). These genes were initially identified as upregulated in the *Cux2*<sup>-/-</sup> cortex in a screen of genes differentially expressed. Previous report showed that increased level of *Xlr3b* and *Xlr4b* expression correlated with more acute behavioral inflexibility in a mouse model of Turner syndrome.<sup>22,23</sup> Nothing was known about the functions of *Xlr* genes, but dendrite and spine defects associate to mental retardation and therefore, it seemed possible that these genes were involved in the changes in dendritic structures in the absence of *Cux*. Further research identified *Cux* binding sites in the *Xlr3b* and *Xlr4b* locus and proved that *Cux* proteins bind to these sites *in vivo* and repress *Xlr* expression. The functional demonstration of the direct implication of *Xlr* genes in the control of the synapse was provided by experiments in which RNAs of interference targeting *Xlr* rescued normal spine density and reduced the proportion of long spines upon *Cux* loss of function. Dendritic tree was not affected, proving to be independently regulated. Interestingly, *Cux1* and *Cux2* proteins selectively bound and repressed distinct regulatory regions on *Xlr3b* and *Xlr4b* loci, illustrating the mechanisms conveying the additive functions (Fig. 1, lower parts).

The expression of *Cux2* selectively defines the upper layer of the human cortex.<sup>24</sup> We identified *FAM9A*, *B* and *C*<sup>25</sup> as the closest orthologs of *Xlr* genes in human and found sequences containing *Cux* binding sites in *FAM9A*, *B* and *C* loci that are conserved between primates and humans. *In vitro* ChIP experiments in human cell lines demonstrated binding of *Cux1* and *Cux2* proteins to these regions, indicating that it is possible that similar *Cux* mediated synaptic mechanisms act in humans.

The functions of *Cux* in spine morphogenesis highlight the existence of neuronal subclass specific mechanisms of synaptogenesis that contribute to the establishment of cognitive circuits. Accordingly, we found defects in working memory in *Cux2*<sup>-/-</sup>. Work lies ahead to further investigate these intrinsic mechanisms



**Figure 2.** Dendritic spine formation in neurons of the cingulate cortex is stimulated upon *Cux1* overexpression. Upper parts show representative confocal image of GFP positive spines in the cingulate cortex. These neurons had been electroporated with control or *CAG-Cux1* plasmid. Scale bar represents 5  $\mu\text{m}$ . Lower parts show quantification of dendritic spine number, spine morphology and spine head area. Data in bar graphs depict mean  $\pm$  SD. \* $p < 0.005$ , \*\* $p < 0.001$ , compared with control. This figure is a modification of Cubelos et al.<sup>15</sup>

of synapse regulation. It will likely reveal genes and proteins affected in cognitive disorders and neurodegeneration. In addition, it will be a challenge to investigate how the selective control of dendritic structures and spines by *Cux1* and *Cux2* contribute to laminar, columnar and area connectivity, and ultimately to the establishment of the intellectual capabilities that rest in the cortex.

## References

- van Elburg RA, van Ooyen A. Impact of dendritic size and dendritic topology on burst firing in pyramidal cells. *PLoS Comp Biol* 2010; 6:1000781.
- Parrish JZ, Emoto K, Kim MD, Jan YN. Mechanisms that regulate establishment, maintenance and remodeling of dendritic fields. *Ann Rev Neurosci* 2007; 30:399-423.
- Majewska A, Brown E, Ross J, Yuste R. Mechanisms of calcium decay kinetics in hippocampal spines: role of spine calcium pumps and calcium diffusion through the spine neck in biochemical compartmentalization. *J Neurosci* 2000; 20:1722-34.
- Noguchi J, Matsuzaki M, Ellis-Davies GC, Kasai H. Spine-neck geometry determines NMDA receptor-dependent  $\text{Ca}^{2+}$  signaling in dendrites. *Neuron* 2005; 46:609-22.
- Yuste R, Bonhoeffer T. Genesis of dendritic spines: insights from ultrastructural and imaging studies. *Nat Rev Neurosci* 2004; 5:24-34.
- Dierssen M, Ramakers GJ. Dendritic pathology in mental retardation: from molecular genetics to neurobiology. *Genes, Brain, Behav* 2006; 2:48-60.
- Boda B, Dubos A, Muller D. Signaling mechanisms regulating synapse formation and function in mental retardation. *Curr Opin Neurobiol* 2010; 20:519-27.
- Tada T, Sheng M. Molecular mechanisms of dendritic spine morphogenesis. *Curr Opin Neurobiol* 2006; 16:95-101.
- Chen JG, Rasin MR, Kwan KY, Sestan N. Zfp312 is required for subcortical axonal projections and dendritic morphology of deep-layer pyramidal neurons of the cerebral cortex. *Proc Natl Acad Sci USA* 2005; 102:17792-7.
- Nieto M, Monuki ES, Tang H, Imitola J, Haubst N, Khoury SJ, et al. Expression of *Cux-1* and *Cux-2* in the subventricular zone and upper layers II-IV of the cerebral cortex. *J Comp Neurol* 2004; 479:168-80.
- Zimmer C, Tiveron MC, Bodmer R, Cremer H. Dynamics of *Cux2* expression suggests that an early pool of SVZ precursors is fated to become upper cortical layer neurons. *Cereb Cortex* 2004; 14:1408-20.
- DeFelipe J, Jones EG. *Cajal on the Cerebral Cortex*. New York: Oxford University Press, 1988.
- Hill RS, Walsh CA. Molecular insights into human brain evolution. *Nature* 2005; 437:64-7.
- Gruber WB, Jan LY, Jan YN. Different levels of the homeodomain protein cut regulate distinct dendrite branching patterns of *Drosophila* multidendritic neurons. *Cell* 2003; 112:805-18.
- Cubelos B, Sebastian-Serrano A, Beccari L, Calcagnotto ME, Cisneros E, Kim S, et al. *Cux1* and *Cux2* regulate dendritic branching, spine morphology and synapses of the upper layer neurons of the cortex. *Neuron* 2010; 66:523-35.
- Flavell SW, Cowan CW, Kim TK, Greer PL, Lin Y, Paradis S, et al. Activity-dependent regulation of MEF2 transcription factors suppresses excitatory synapse number. *Science* 2006; 311:1008-12.
- Shalizi A, Gaudilliere B, Yuan Z, Stegmuller J, Shirogane T, Ge Q, et al. A calcium-regulated MEF2 sumoylation switch controls postsynaptic differentiation. *Science* 2006; 311:1012-7.
- Cubelos B, Sebastian-Serrano A, Kim S, Moreno-Ortiz C, Redondo JM, Walsh CA, et al. *Cux-2* controls the proliferation of neuronal intermediate precursors of the cortical subventricular zone. *Cereb Cortex* 2008; 18:1758-70.
- El-Husseini AE, Schnell E, Chetkovich DM, Nicoll RA, Brecht DS. PSD-95 involvement in maturation of excitatory synapses. *Science* 2000; 290:1364-8.
- Ultanir SK, Kim JE, Hall BJ, Deerinc T, Ellisman M, Ghosh A. Regulation of spine morphology and spine density by NMDA receptor signaling in vivo. *Proc Natl Acad Sci USA* 2007; 104:19553-8.

## Acknowledgements

This work was supported by the MICINN grants SAF2008-00211; PIE-200820I166, and a grant from the Spanish Comunidad de Madrid CCG08-CSIC/SAL-3464. B. Cubelos holds a fellowship from the CSIC (JAEDoc2008-020).

21. Edbauer D, Neilson JR, Foster KA, Wang CF, Seeburg DP, Bartterton MN, et al. Regulation of synaptic structure and function by FMRP-associated microRNAs miR-125b and miR-132. *Neuron* 2010; 65:373-84.
22. Raefski AS, O'Neill MJ. Identification of a cluster of X-linked imprinted genes in mice. *Nat Genet* 2005; 37:620-4.
23. Davies W, Isles A, Smith R, Karunadasa D, Burrmann D, Humby T, et al. Xlr3b is a new imprinted candidate for X-linked parent-of-origin effects on cognitive function in mice. *Nat Genet* 2005; 37:625-9.
24. Arion D, Unger T, Lewis DA, Mirnics K. Molecular markers distinguishing supragranular and infragranular layers in the human prefrontal cortex. *Eur J Neurosci* 2007; 25:1843-54.
25. Martinez-Garay I, Jablonka S, Sutajova M, Steuernagel P, Gal A, Kutsche K. A new gene family (FAM9) of low-copy repeats in Xp22.3 expressed exclusively in testis: implications for recombinations in this region. *Genomics* 2002; 80:259-67.