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There's More than One Way to Skin a Chimaerin

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Abstract

In two manuscripts published in *Neuron* (Beg et al. and Wegmeyer et al.) and one published in *Cell* (Iwasato et al.), investigators have found that a particular GAP, α -chimaerin, is required in vivo for ephrinB3/EphA4-dependent motor circuit formation.

The diversity of signaling molecules has long fascinated scientists and suggested the possibility that particular proteins might be required for specific cellular events. However, examples of signal transduction proteins acting only during particular cellular events have been few and far between. Now, three papers provide a beautiful in vivo example of a specific function for a particular signaling molecule. Together, the three manuscripts are striking for the extent that they each describe a similar set of results but use independent approaches to do so: one group began by identifying a spontaneous mouse mutant (Iwasato et al., 2007), a second used a yeast two-hybrid screen (Wegmeyer et al., 2007), and a third used a screen for protein-protein interactions (Beg et al., 2007). While each paper offers its own strong evidence for the role of α -chimaerin in mediating ephrinB3/EphA4-dependent motor circuit formation, together these manuscripts provide a compelling and unusually comprehensive picture of an essential role for a particular GAP.

Small GTPases in the Rac, Rho, and Cdc42 families act as molecular switches in signaling pathways, with the GTP-bound “ON” state and GDP-bound “OFF” state. Regulating the cycle between ON and OFF are over 70 small RhoGTPase-activating proteins (GAPs) in the mammalian genome, with 12 known to have specific activity for Rac. Chimaerins are Rho-GAPs with specific activity for Rac that contain a C1 domain that allows them to bind phorbol esters (Yang and Kazanietz, 2007). There are two α -chimaerins and two β -chimaerins made as alternatively spliced products of two genes. α 2- and β 2-chimaerin also contain SH2 domains that enable them to bind phosphorylated tyrosine residues. The α -chimaerin GAPs are expressed in brain and linked to axon guidance (Diaz et al., 2002), semaphorin signaling (Brown et al., 2004), dendritic spine development (Buttery et al., 2006), and the NMDAR (Van de Ven et al., 2005). These properties made chimaerin an interesting candidate for regulating axonal growth in the nervous system.

Work from a number of laboratories has identified a specific role of ephrinB3/EphA4 forward signaling in the formation of the motor circuit between neurons in layer 5 of motor

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cortex and the principle motor neurons in the lumbar spinal cord (Dottori et al., 1998; Coonan et al., 2001; Kullander et al., 2001; Kiehn and Butt, 2003). Mice with mutations to either of these proteins exhibit a hopping rabbit-like gait where both hindlimbs move together—arising from the failure of cortical motor axons to innervate only contralateral motor neurons in the spinal cord and from defects in spinal interneuron axonal projections (Figures 1A and 1C). These defects appear to be due to the failure of these axons to respond to an ephrinB3 repulsive guidance cue along the midline of the spinal cord that allows axons to innervate both left and right motor neurons (Kullander et al., 2001, 2003). Moreover, ephrin-Eph signaling induces growth cone collapse that appears to be mediated via regulation of small GTPase activity (Kullander and Klein, 2002; Cowan et al., 2005; Sahin et al., 2005).

In the manuscript by Iwasato et al. from the RIKEN institute (Iwasato et al., 2007), the authors began by noting a rabbit-like gait in a Chat-Cre line of mice that they generated. Thinking that the insertion of their CRE construct was the cause, they were surprised to learn that the mutation segregated from the CRE, indicating that a spontaneous mutation was causing the phenotype. They determined that the gait defect in the *miffy* mice was likely due to defects in the motor circuit arising from defects in cortical spinal tract projections (CST). This misprojection resulted in an abnormal rhythmic activity of flexors and tensors reminiscent of that seen in mice lacking EphA4. The authors then isolated the *miffy* locus using microsatellites and single-nucleotide polymorphisms (SNPs) to a 3.27 Mb region on chromosome 2. Of the 30 genes in that region, ten were ruled out because mice lacking these genes were known not to have defects in gait. Fortunately for the authors, only one of the 20 remaining genes generated different sized transcript when amplified with RT-PCR— α -chimaerin. The region affected in the *miffy* mutation resulted in a loss of 58 amino acids in both α 1- and α 2-chimaerin and generated proteins that lacked GAP activity in vitro.

To demonstrate that this mutation was the cause of the defects in motor circuit formation, Iwasato et al. then undertook to rescue the expression of α -chimaerin using a bac-transgenic rescue in the *miffy*^{-/-} mice and attempted to phenocopy their spontaneous mutation by making a targeted deletion of α -chimaerin. Likely due to its low sub-wild-type level of expression, the bac-transgenic gave a partial rescue, but knockout of the locus phenocopied the spontaneous mutation, indicating that *miffy* is likely due to a mutation in α -chimaerin.

Iwasato et al. then began to examine whether α -chimaerin and EphA4 might interact. Immunocytochemistry revealed that α -chimaerin and EphA4 proteins colocalize in developing CST axons in vitro. Moreover, Iwasato et al. were able to demonstrate that α 2-chimaerin and EphA4 coimmunoprecipitate when coexpressed in HEK293T cells, in neurons, and from brain lysates of mice. By making mutations to α 2-chimaerin and EphA4, they found in HEK293T cells that the interaction appears to be EphA4 kinase independent and relies on the C-terminal domain of α 2-chimaerin. The interaction between EphA4 and α 2-chimaerin also appears to be functional: α 2-chimaerin can inactivate Rac downstream of kinase-active EphA4 when these proteins are coexpressed in COS cells and then stimulated with a soluble form of ephrinB3 (ephrinB3-Fc). Finally, the authors use an RNAi knockdown approach to demonstrate that motor cortex axons expressing less α 2-chimaerin are less sensitive to ephrinB3-dependent growth cone collapse.

The two papers published in *Neuron* begin with a more traditional approach, with both groups first demonstrating that $\alpha 2$ -chimaerin or domains of this protein interact with Ephs. Using a yeast two-hybrid screen with $\alpha 2$ -chimaerin SH2, Wegmeyer et al. identified EphA4 and EphB1 as $\alpha 2$ -chimaerin-interacting proteins (Wegmeyer et al., 2007). This group then generated two mice with mutations to the α -chimaerin locus: a conventional knockout and a knockin with a point mutation in the C1 domain. The knockin mutation resulted in a loss of $\alpha 1$ -chimaerin expression and reduction of $\alpha 2$ -chimaerin transcript. Similar to what Iwasato et al. reported, heterozygous animals had no phenotype, and homozygous mice of both genotypes displayed defects in gait. Examination of the CST in both lines of mutant mice revealed increased axonal recrossing and a decrease in the size of the dorsal funiculus. Recording motor output from lumbar dorsal roots revealed a loss of motor coordination that was similar to that seen in the mice generated by the Iwasato et al. group. Neurons from mice lacking α -chimaerin had impaired growth cone collapse following ephrinB3 or ephrinA1 stimulation. Moreover, consistent with the data from all three papers, while loss of α -chimaerins resulted in defects in CST axonal projections, axons that were dependent on EphA4 reverse signaling were unaffected.

Having demonstrated that mice lacking α -chimaerin have defects in the CST projection and neuronal growth cones from these animals fail to respond robustly to ephrin treatment, Wegmeyer et al. next examined how EphA4 might interact with and regulate α -chimaerins. Unlike Iwasato et al., Wegmeyer et al. were not able to coimmunoprecipitate α -chimaerin with EphA4; therefore, they turned to a GST-pulldown assay. Using this assay, Wegmeyer et al. determined that both $\alpha 1$ - and $\alpha 2$ -chimaerin can bind to EphA4 with a complex series of experimental results. $\alpha 1$ -Chimaerin appears to bind EphA4 strongly, in the kinase domain, but does not require the kinase to be active. $\alpha 2$ -Chimaerin, on the other hand, appears to bind to EphA4 both through its SH2 domain and via a second kinase-independent interaction. Because they interact, the authors then asked whether EphA4 might phosphorylate α -chimaerin. Both isoforms of α -chimaerin are phosphorylated in 293FT cells when coexpressed with EphA4, and a mutation of three different tyrosine residues (y202, y303, y333) on $\alpha 2$ -chimaerin reduces its phosphorylation, suggesting that EphA4 phosphorylates these sites. Consistent with a role of EphA-dependent regulation of $\alpha 2$ -chimaerin, EGFP-tagged $\alpha 2$ -chimaerin overexpressed in neurons becomes phosphorylated following soluble ephrinA1 activation of EphA receptors.

The Wegmeyer et al. study implicates the adaptor proteins Nck1 and Nck2 as potentially important in the ephrin/EphA4/ $\alpha 2$ -chimaerin pathway by showing that these proteins can interact with full-length $\alpha 2$ -chimaerin, but not with $\alpha 1$ -chimaerin or mutant forms of $\alpha 2$ -chimaerin that lack an SH2 domain. The presence of Nck2 negatively regulates the RacGAP activity of only $\alpha 2$ -chimaerin in the presence of EphA4. While interpretation of these results was limited because the authors could not control the expression of each protein, the results demonstrate that EphA4, $\alpha 2$ -chimaerin, and Nck2 can act in a concerted fashion on Rac activity.

The third paper, by Beg et al. from Columbia University (Beg et al., 2007), uses affinity chromatography with solubilized neuronal membrane proteins on an immobilized recombinant $\alpha 2$ -chimaerin SH2 domain and GST-pulldown assays in HEK293T cell lysates

to identify a number of phospho-tyrosine proteins as potential $\alpha 2$ -chimaerin interactors. One of these interacting molecules was EphA4. The SH2 domain of $\alpha 2$ -chimaerin fused to GST was able to pull down EphA4 and EphB1, but not TrkA. The SH2 domain interaction required EphA4 kinase activity and likely occurred via interaction with the juxtamembrane tyrosines. While the authors could not demonstrate that EphA4 and $\alpha 2$ -chimaerin coimmunoprecipitated, they did note that coexpression of EphA4 and $\alpha 2$ -chimaerin led to $\alpha 2$ -chimaerin phosphorylation. In contrast to the experiments from Wegmeyer et al., Beg et al. found that the $\alpha 2$ - but not the $\alpha 1$ -chimaerin interaction with EphA4 required EphA4 kinase activity.

To examine the in vivo significance of $\alpha 2$ -chimaerin, Beg et al. used a gene-trap approach that enabled them to substantially reduce $\alpha 2$ -chimaerin expression while leaving $\alpha 1$ -chimaerin expression at nearly wild-type levels. The authors first asked whether axons from these mutant animals would still respond to ephrinA5-Fc treatment normally and display growth cone collapse. Consistent with a role of $\alpha 2$ -chimaerin alone in EphA-mediated growth cone collapse and reports from Iwasato et al. using siRNA, axons from motor cortex explants of mice lacking $\alpha 2$ -chimaerin were significantly less responsive to ephrinA5 than controls.

Beg et al. then demonstrated that the loss of $\alpha 2$ -chimaerin generated a phenotype strikingly similar to that seen in animals lacking EphA4 and to the reports by the other two groups examining the $\alpha 1$ - and $\alpha 2$ -chimaerin mutant animals. Gene-trap $\alpha 2$ -chimaerin mice showed defective gait, increased CST axon crossing, and increased crossing of spinal interneurons. Then, in an elegant series of experiments, Beg et al. demonstrated a genetic interaction between EphA4 and $\alpha 2$ -chimaerin by showing that double-heterozygous mice ($\alpha 2$ -chimaerin^{+/-}, EphA4^{+/-}) also have enhanced spinal interneuron axon crossing compared to the single-heterozygous mutants and similar to that seen in the $\alpha 2$ -chimaerin mutant. These experiments provide strong genetic evidence for EphA4 and $\alpha 2$ -chimaerin interaction.

Each group conducted a similar series of experiments to demonstrate that the loss of α -chimaerin generates a phenotype similar to that seen with loss or disruption of EphA4. In the end, these three papers paint a compelling picture of a molecule that is essential for a particular aspect of EphA4 receptor signaling (Figure 1). Overall, the data from these three papers fit together nicely, though there are a few places, largely in the biochemical analysis, where there is some disagreement. In most cases, these differences are most likely due to differences in methodology. In any case, the major findings are in agreement. Whether other GAPs and GEFs act so specifically remains to be determined; meanwhile, these papers provide a comprehensive and exciting picture of how a widely expressed signaling molecule can be essential for particular events. There are a number of new questions that are raised by this work. Is axon guidance generated by a balance between GAP and GEF activity? How often do specific GAPs act in particular cells? Do they act sequentially within a signaling cascade? If high levels of specificity do exist, rather than expecting that loss of signaling proteins will generate dramatic phenotypes, we will have to re-examine previously generated models for more specific and subtle phenotypes that point to the essential function of other widely expressed signaling proteins.

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