The Ras1–Mitogen-Activated Protein Kinase Signal Transduction Pathway Regulates Synaptic Plasticity through Fasciclin II-Mediated Cell Adhesion

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Ras proteins are small GTPases with well known functions in cell proliferation and differentiation. In these processes, they play key roles as molecular switches that can trigger distinct signal transduction pathways, such as the mitogen-activated protein kinase (MAPK) pathway, the phosphoinositide-3 kinase pathway, and the Ral–guanine nucleotide dissociation stimulator pathway. Several studies have implicated Ras proteins in the development and function of synapses, but the molecular mechanisms for this regulation are poorly understood. Here, we demonstrate that the Ras–MAPK pathway is involved in synaptic plasticity at the Drosophila larval neuromuscular junction. Both Ras1 and MAPK are expressed at the neuromuscular junction, and modification of their activity levels results in an altered number of synaptic boutons. Gain- or loss-of-function mutations in Ras1 and MAPK reveal that regulation of synapse structure by this signal transduction pathway is dependent on fasciclin II localization at synaptic boutons. These results provide evidence for a Ras-dependent signaling cascade that regulates fasciclin II-mediated cell adhesion at synaptic terminals during synapse growth.

Key words: mitogen-activated protein kinase; Ras; neuromuscular junction; internalization; cell adhesion; synapse plasticity

Synapse formation and modification are highly complex processes that include the activation of gene expression, cytoskeletal reorganization, and signal transduction activation (Koh et al., 2000; Lee and Sheng, 2000). A pathway involved in these processes is the Ras–mitogen-activated protein kinase (MAPK) signal transduction cascade (Lowy and Willumsen, 1993; Mazzucchelli and Brambilla, 2000). Ras proteins are highly localized in developing and adult brains (Leon et al., 1987), and maintenance of long-term potentiation is critically dependent on MAPK activation (English and Sweatt, 1997). Mutations in genes encoding members of the MAPK pathway, such as MAPK kinase (MEK), Ras–guanine nucleotide-releasing factor, and H-Ras, cause defects in learning and long-term potentiation (Brambilla et al., 1997; Atkins et al., 1998; Manabe et al., 2000).

In Aplysia, ApMAPK, the homolog of P44/42 extracellular signal-regulated kinase (ERK), plays a major role in long-term facilitation (LTF) (Bailey et al., 1997). LTF elicits translocation of activated ApMAPK into the neuronal nucleus and the internalization of ApCAM, a homolog of neuronal cell-adhesion molecule in mice, and fasciclin II (FasII) in flies (Mayford et al., 1992). Mutations in MAPK or MAPK phosphorylation targets in ApCAM block internalization of ApCAM, preventing synaptic growth (Bailey et al., 1997; Martin et al., 1997).

The Drosophila neuromuscular junction (NMJ) is a powerful system to understand the mechanisms underlying synaptic plasticity. Larval NMJs continuously increase in size to compensate for muscle growth during development. This form of plasticity is controlled by electrical activity and FasII-mediated cell adhesion (Budnik et al., 1990; Schuster et al., 1996a,b; Koh et al., 1999). Discs-Large (DLG), a member of the postsynaptic density-95 protein family, associates with the synaptic cytoskeleton, where it clusters FasII (Thomas et al., 1997). With increased electrical activity, Ca2+/calmodulin protein kinase II (CaMKII) phosphorylates DLG, decreasing its FasII-clustering ability and promoting NMJ growth (Koh et al., 1999).

The studies in Aplysia raise the possibility that the MAPK pathway may also regulate the synaptic localization of FasII. Ras1, the Drosophila homolog of N-, K-, and H-Ras, is activated upon GTP binding (Gaul et al., 1992) and activates the phosphoinositide-3 kinase (PI3-K) (p110), Ral–guanine nucleotide dissociation stimulator (GDS), and MAPK pathways (Fig. 1A) (Bergmann et al., 1998). As in mammals, in Drosophila Ras1 activity can be manipulated by mutations in ras1 and by ras1 transgenic variants. Substitution of glycine-12 to valine (Ras1V12) renders a constitutively active form of Ras1, and all three pathways can be activated. Additional mutations in the effector loop of Ras1V12 can result in the activation of a single pathway, because the interaction between Ras1 and the other two effectors is blocked (Fig. 1A) (White et al., 1995; Rodriguez-Viciana et al., 1997; Bergmann et al., 1998; Karim and Rubin, 1998; Halfar et al., 2001).

A Drosophila MAPK gene, rolled (rl), with homology to P44/44-ERKs has also been identified (Biggs et al., 1994; Oellers and Hafen, 1996). The functions of MAPK in Drosophila have been investigated in eye and wing development (Bergmann et al., 1998;
nuclei persists. NMJs and at the muscle surface is severely reduced, but the signal at the throughout the muscle surface. In the concentrated at synaptic boutons and muscle nuclei as well as at low levels

Ras1, but not in others. Substitution of serine-17 into asparagine (Ras1N17) has been reported in some studies to result in a dominant-negative form of Rubin, 1998). Substitution of glycine (Ras1V12G37), or tyrosine-40 to cysteine (Ras1V12C40) results in the activation of a single pathway (Bergmann et al., 1998; Karim and glycine (Ras1V12G37), or tyrosine-40 to cysteine (Ras1V12C40) results in the activation of a single pathway (Bergmann et al., 1998; Karim and

MAPK signal transduction pathways and Ras1 expression at the Drosophila larval NMJ. We

immunoblotted sequentially with anti-FasII and anti-DLG PDZ (1:2000 dilution). Bands were visualized with peroxidase-conjugated second-
motor neurons and muscles. C, D. Anti-Ras1 immunoreactivity at the larval body wall muscles of wild type (C) and rasP37603 mutant (D). Note that in wild type, immunoreactivity is concentrated at synaptic boutons and muscle nuclei as well as at low levels throughout the muscle surface. In the ras mutant, immunoreactivity at NMJs and at the muscle surface is severely reduced, but the signal at the nuclei persists. E, F. High-magnification view of synaptic boutons double-stained with anti-HRP (red) and anti-Ras (green) in wild type (E) and rasP37603 mutant (F). Scale bar, 20 μm.

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**MATERIALS AND METHODS**

**Flys.** Flies were reared in standard conditions between 22 and 25°C. We used the following strains: a hypomorphic mutation in ras1, ras1P5763 (Schnorr and Berg, 1996), upstream activation sequence (UAS)-RafP19, UAS-RalA124, and UAS-Ras85DV12, obtained from the Bloomington Stock Center (Bloomington, IN); UAS-Ras1WT/CyO, UAS-Ras1585D12S5/CyO. UAS-Ras1585D12G37, UAS-Ras1585D12C40, UAS-Ras1585D117, and hTom/CyO, obtained from Bergmann et al. (1998); and hTom/+ (Oellers and Hafen, 1996), obtained from Dr. L. Zipursky (University of California, Los Angeles, CA). We also used the severe hypomorph fasIIP64, reported to contain ~10% FasII levels (Schuster et al., 1996a), and dglG122, which expresses DLG at very low levels and contains a deletion of the guanylate kinase domain (Woods et al., 1996), and the Calh liver strains C380 and Sca-Gal4 (presynaptic) and BG487 (postsynaptic) (Koh et al., 1999). The wild-type strain Canton-S (CS) was used as a control.

**Immunocytochemistry.** The immunocytochemical procedure for body wall muscle preparations was described by Thomas et al. (1997). The following antibodies were used for this study: rabbit and rat anti-DLG, (1:40,000 and 1:500, respectively) (Thomas et al., 1997), rabbit anti-Ras (1:200; Calbiochem, San Diego, CA), rabbit anti-Drosophila melanogaster (Dm)ERK-A (Biggs and Zipursky, 1992; Gabay et al., 1997b), mouse anti-diphospho-MAPK monoclonal (DPMAPK) (1:20; Sigma, St. Louis, MO) (Gabay et al., 1997a), mouse anti-FasII ID4 monoclonal (1:2 dilution; gift from Dr. C. Goodman, University of California at Berkeley, Berkeley, CA), rabbit anti-FasII (1:3000; Thomas et al., 1997), and anti-HRP-FITC (1:400 dilution; Sigma). As secondary antibodies, we used FITC- or Texas Red-conjugated donkey anti-rabbit IgG, donkey anti-rat IgG, or donkey anti-mouse IgG (1:200 dilution; Jackson ImmunoResearch, West Grove, PA).

The number of type I boutons at muscles 6 and 7 (abdominal segment 2) was counted under an epifluorescence microscope at 63× magnification in preparations stained with FITC- or Texas Red-conjugated anti-HRP to stain the presynaptic terminal. The intensity of FasII and DpMAPK immunoreactivity at synapses of CS, hTom/+ , and hTom/+ was determined as in Thomas et al. (1997) using the NIH Image program (version 1.57). Larvae used for intensity analysis (Table 1) were processed simultaneously for immunocytochemistry, and confocal images were acquired under identical conditions. Briefly, a line beginning from the center of a synaptic bouton and extending through the outer limit occupied by DLG immunoreactivity was traced. Next, the maximum intensity along the line (on a linear relative scale of 0–255) was determined using the plot profile function of NIH Image. Four measurements at right angles were taken for each bouton, averaged, and expressed as a percentage of maximum relative intensity.

**Western blots and immunoprecipitations.** Dissected body wall muscles (without CNS) were homogenized in Drosophila buffer 2% buffer (50 mM Tris-Cl, pH 6.8, 25 mM KCl, 2 mM EDTA, 0.3 μM sucrose, 2% SDS) in a 75°C bath. The homogenate was prepared for SDS-PAGE and separated on an 8% gel. Blots were probed with anti-DmERK-A (1:10,000) or anti-DpMAPK monoclonal (1:500) peroxidase-conjugated secondary antibodies and enhanced with chemiluminescent reagents (Amersham Biosciences, Piscataway, NJ). Coimmunoprecipitations were performed essentially as described by Thomas et al. (1997). Briefly, 10 dissected body wall muscle preparations (consisting of body wall muscles and CNS) were homogenized in 100 μl of radioimmunoprecipitation buffer containing protease inhibitors at 4°C. After centrifugation at 3000 × g for 5 min, the supernatant was precleared with preimmune serum and protein A+G beads for 1 hr. The cleared homogenate was then incubated with rat anti-DLGP512 (5 μl of crude serum) at 4°C for 1 hr. Immunoprecipitates were collected with protein A+G-Sepharose, separated in a 7.5% SDS-PAGE gel, and immunoblotted sequentially with anti-FasII and anti-DLGP512 (1:2000 dilution). Bands were visualized with peroxidase-conjugated secondary antibodies and enhanced with chemiluminescent reagents (Amersham). Quantification of band intensities was performed by scanning the radiographic film on a linear response scanner (UMAX-Powerlook II, UMAX, Dallas, TX). The intensity of the bands was measured by using the NIH Image 1.54 software for densitometric analysis of one-dimensional gels.

**Statistical analysis.** The Mintab program (Minitab Inc., State College, PA; www.minitab.com) was used for statistical analysis. A two-sample Student’s t test was used to determine differences between samples. Numbers represent mean ± SEM throughout.
RESULTS

Ras1 is expressed at presynaptic terminals and is involved in the regulation of bouton number

To determine the presence of Ras1 at larval NMJs, we used an anti-peptide antibody generated using a conserved Ras1 sequence (Sawada et al., 1989). We found that Ras immunoreactivity was concentrated at type I synaptic boutons and in the nuclei of muscle cells (Fig. 1C,E). Low levels of immunoreactivity were also observed throughout the surface of the muscle membrane. The immunoreactivity at synaptic boutons and at the muscle surface was specific, because in the severe hypomorph rasP5703, Ras immunoreactivity was dramatically reduced (Fig. 1D,F). In contrast, immunoreactivity at the muscle nuclei was similar to wild type, suggesting that the nuclear staining might be cross-reactivity or that the mutant protein is still able to localize at this site.

To investigate the role of Ras1 at the NMJ, we manipulated Ras1 activity at presynaptic and postsynaptic terminals by using the rasP5703 allele and by expressing transgenic wild-type and mutant Ras1 variants using the UAS/Gal4 system (Brand and Perrimon, 1993). For the transgenic experiments, we expressed Ras proteins at both presynaptic and postsynaptic sites by using the Gal4 drivers C380 and Sca-Gal4 (presynaptic) and BG487 (postsynaptic) (Koh et al., 1999). Synaptic morphology was examined by labeling presynaptic arbors using anti-HRP, an insect neuronal marker. We found that decreased levels of Ras1 in the hypomorph rasP5703 mutant resulted in a significant reduction in the number of type I synaptic boutons compared with wild type (Fig. 1B). In contrast, overexpressing transgenic wild-type Ras1 (Ras1WT) at both presynaptic and postsynaptic cells or at presynaptic cells alone caused an increase in the number of boutons (Fig. 1B). An even greater increase was seen by expressing a constitutively active Ras1, Ras1V12 (Fig. 2A). These results suggest a Ras-dependent signal transduction pathway in the regulation of synaptic bouton number. The effect in Ras1WT flies also implies that the endogenous pathway, which presumably is activated by Ras during synapse growth, is overstimulated by increasing levels of wild-type Ras. This hypothesis was confirmed by examining the levels of MAPK activation (see below).

To determine which of the signal transduction pathways known to be activated by Ras1 is involved in the regulation of bouton number, we expressed different constitutively active Ras1V12 variants that selectively activate one of the following pathways: Ral-GDS, PI3-K, or MAPK at the NMJ (Fig. 1A). We found that expression of Ras1V12S35, which activates the MAPK pathway only, induced a striking increase in the number of type I boutons at muscles 6 and 7 compared with controls expressing Ras1WT and compared with wild-type larvae (Fig. 2). This increase was indistinguishable from the increase in bouton number observed in Ras1V12. In contrast, expression of Ras1V12G37 (which activates the Ral pathway), Ras1V12C40 (which activates the PI3-K pathway), and Ras1N17 (which has been demonstrated to block Ras1 activation in certain cases (Feig and Cooper, 1988) but not in others (Malumbres and Pellicer, 1998; Marais et al., 1998)) did not alter NMJ morphology compared with RasWT (Fig. 2A). However, expression of these transgenes did result in a significant increase in bouton number compared with wild-type controls, suggesting that these pathways may also influence NMJ growth or that the activation of the MAPK pathway is not completely eliminated in these Ras variants. Thus, activation of the MAPK pathway appears to be most effective in increasing bouton number, and activating this pathway alone (Ras1V12S35) mimics the effects of constitutive Ras activation (Ras1V12).

We also studied the effect of constitutively active forms of proteins downstream of Ras: RalA72L, a constitutively active RalA protein (Lee et al., 1996), and RafF179, which activates the MAPK pathway (Brand et al., 1994). We found that in RafF179 there was a dramatic increase in bouton number that was similar to Ras1V12 and Ras1V12S35 (Fig. 2A), consistent with the model that the Ras–MAPK pathway is involved in increasing bouton number. However, constitutively active RalA also enhanced bouton number compared with wild-type controls, although to a lesser extent than Ras1V12, Ras1V12S35, and RafF179, in agree-

Figure 2. Morphology of NMJs in larvae expressing Ras1 variants. A. Number of type I synaptic boutons in wild-type larvae (WT) and in larvae overexpressing wild-type Ras1 (RWT), constitutively active Ras (RasV12; V12), Ras1V12S35 (V12S35), constitutively active Raf (RafF179; Raf), Ras1V12G37 (V12G37), constitutively active Ral (RalA72L; Ral), Ras1V12C40 (V12C40), and Ras1N17 (N17) using the C380 Gal4 driver. B, C, Anti-HRP immunoreactivity in larvae expressing transgenic wild-type Ras1 (RasWT) (B) or the Ras1V12S35 variant (C), which constitutively activates the MAPK pathway. Note that MAPK pathway activation results in a significantly increased number of synaptic boutons. Scale bar, 50 μm.
ment with the idea that activation of other Ras-dependent pathways may also influence synapse growth. Because activation of the Ras–MAPK pathway was the most robust in enhancing bouton number, we centered on the analysis of this pathway.

Similar results were obtained by using either of the presynaptic Gal4 drivers C380 (Fig. 2A) or ScaGal4 alone, or C380 in conjunction with a postsynaptic driver, BG487. In addition, no changes in bouton number were observed when the transgenes were expressed using BG487 alone or in Gal4+/ heterozygotes (data not shown). These results imply that the increase in bouton number is the result of manipulating Ras1 activity at the presynaptic terminals. The increased number of synaptic boutons observed by expressing Ras1V12S35 was not attributable to different levels of expression of the Ras transgenes, as determined by comparing levels of Ras immunoreactivity in the CNS and NMJ in the different Ras transgenic flies (data not shown).

Activated double phosphorylated MAPK is expressed at NMJs, where it regulates the levels of FasII

The above observations suggested that Ras1 is expressed at NMJs, where it can regulate synaptic bouton number through activation of the MAPK pathway. A prediction of this hypothesis is that MAPK also should be expressed at the NMJ. This was tested by using a polyclonal antibody, anti-DmERK-A, against the Drosophila MAPK, Rolled (Biggs and Zipursky, 1992; Gabay et al., 1997b). In addition, we used an anti-MAPK monoclonal antibody, DpMAPK, which recognizes the active, double-phosphorylated form (Gabay et al., 1997a). Western blot analysis of body wall muscle extracts using DmERK-A and DpMAPK antibodies confirmed the presence of a band at ~44 kDa, as expected in Drosophila. The intensity of the DmERK-A band was decreased to 56 ± 13% in the heterozygous hypomorph r10a/+ with regard to wild-type controls, demonstrating the specificity of the DmERK-A antibody (Fig. 3A). Notably, activated MAPK was slightly enhanced in flies overexpressing presynaptic Ras1WT. An additional increase was observed in Ras1V12S35 but not in Ras1V12G37 or Ras1V12C40, corroborating the specificity of RasV12S35 in MAPK pathway activation. Levels of total MAPK protein were similar in all genotypes (Fig. 3B).

Using the DmERK-A antibody, we found that MAPK was expressed at synaptic boutons of the NMJ both diffusely and in immunoreactive hot spots (Fig. 3C). Because the DmERK-A antibody recognizes both inactive and active MAPK forms, we subsequently used the DpMAPK antibody to determine whether it was similarly localized at synaptic boutons. Remarkably, the active, double-phosphorylated MAPK had a more restricted localization than DmERK-A at synaptic boutons, being localized at well defined hot spots (Fig. 3D) that generally colocalized with areas of high DmERK-A immunoreactivity (Fig. 3E). The wider localization of DmERK-A with regard to DpMAPK suggests that active MAPK is selectively recruited to restricted domains at synaptic boutons or that MAPK activation is spatially restricted at the boutons.

In Aplysia, LTF of the gill withdrawal reflex results in an increase in the number of sensory synaptic boutons, an observation that can be replicated in dissociated neuronal cultures by application of serotonin (Mayford et al., 1992). These studies suggest that the decrease of cell adhesion induced by downregulation of ApCAM allows the expansion of sensory processes, resulting in the formation of new synaptic sites. Similarly, at the Drosophila NMJ, manipulations that lead to a decrease in FasII, the homolog of ApCAM, lead to an increase in synaptic bouton number (Budnik et al., 1990; Schuster et al., 1996a; Koh et al., 1999). Notably, in Drosophila, FasII is also required for the maintenance of synaptic boutons. Therefore, increases in bouton number are observed only when FasII is within permissive levels, which are required to maintain the NMJ (Schuster et al., 1996a).

The studies in Aplysia suggest that activation of MAPK during LTF is crucial for the internalization of ApCAM from the surface of sensory neurons and therefore for structural synaptic plasticity during LTF (Martin et al., 1997). In flies, neuronal activity can regulate the localization of FasII by inactivating its clustering at the NMJ by DLG (Koh et al., 1999).

We hypothesized that as in Aplysia, FasII levels at the Drosophila NMJ could be additionally regulated by MAPK activity. This was tested by examining the expression of FasII immunoreactivity in the partial-loss-of-function MAPK mutant (r10a/+ ) and in a gain-of-function mutant (d110h/+ ), in which a single amino acid substitution in the kinase domain causes a twofold to threefold increase in levels of MAPK activity (Brunner et al.,

Figure 3. Expression of MAPK at the NMJ. A, Western blot of body wall muscle extracts stained with anti-DmERK-A (left) and reprobed with anti-DpMAPK (middle) after stripping. To show that the blot was completely stripped after staining with DmERK-A, the second lane of the middle panel was incubated with secondary antibody and chemiluminescent reagent before exposure to film. Each lane in the right and middle panels was loaded with an equal amount of protein from body wall muscle extracts (~5 body wall muscle preparations per lane). In the Western blot shown in the left panel, equal amounts of protein (~2.5 body wall muscle preparations per lane) from extracts of wild-type and r10a/+ heterozygotes were loaded, and the membrane was probed with anti-DmERK-A. B, Western blots of body wall muscle extracts from wild type (WT) and from larvae overexpressing Ras1V12S35 (V12S35); wild-type Ras1 (Rwt), Ras1V12G37 (V12G37), and Ras1V12C40 (V12C40) using C380, probed sequentially with anti-DpMAPK and anti-DmERK-A. Equal amounts of protein were loaded in each lane (3.5 body wall muscle preparations per lane). The molecular masses given at the right of the blots are in kilodaltons. C, Anti-DmERK-A immunoreactivity at type I synaptic boutons. D, Antibodies against activated MAPK (DpMAPK) result in highly immunoreactive hot spots at synaptic boutons. E, DpMAPK hot spots colocalize with DmERK-A patches at the boutons, but DpMAPK staining is more restricted than DmERK-A, as shown by merging A and B. Scale bar, 10 μm.
Consistent with our hypothesis, the intensity of FasII immunoreactivity at the NMJ of the MAPK loss-of-function mutant was $135 \pm 3$, an $\sim 30\%$ increase compared with wild-type controls ($106 \pm 3$; Table 1; Fig. 4). In contrast, in the gain-of-function mutant, there was an $\sim 30\%$ reduction in the levels of FasII at the NMJ ($81 \pm 2$; Table 1; Fig. 4). Alteration in MAPK activity levels, however, did not seem to affect the distribution of active MAPK (Fig. 4A–C) or the general morphology of the boutons (Fig. 4G–I), although it did alter bouton number (see below).

Additional evidence that alterations in MAPK activity result in changes in FasII localization at synapses was obtained by performing immunoprecipitation of body wall muscle extracts using anti-DLG antibodies (Fig. 5). FasII is expressed at all neuromuscular junctions, but it interacts with DLG only at type I boutons (Thomas et al., 1997). Therefore, we expected that immunoprecipitation with anti-DLG would result in coinmunoprecipitation of the transmembrane FasII isoform that is expressed at type I boutons. As expected, the amount of FasII at type I synapses was dependent on MAPK activity. Increased MAPK activity in the gain-of-function MAPK mutant resulted in a $27\%$ reduction in the amount of FasII that was coinmunoprecipitated by DLG compared with wild-type controls (Fig. 5). Conversely, reduced MAPK activity in the loss-of-function MAPK allele resulted in a $170\%$ increase in the amount of FasII coinmunoprecipitated by anti-DLG.

We subsequently examined the distribution of active MAPK in relation to FasII to determine whether any change in FasII could be observed at the sites of the synaptic boutons at which active MAPK was concentrated (Fig. 6). As documented by Sone et al. (2000), we found that FasII was not homogeneously distributed at type I synaptic boutons but rather formed an irregular network on their surface, interrupted by nonstaining areas (Fig. 6B). When viewed in rotating Z-series, these nonimmunoreactive areas appeared as little windows through which the opposite side of the bouton was visible. Interestingly, these windows of low FasII immunoreactivity coincided with the highly immunoreactive active MAPK hot spots (Fig. 6B). Thus, the decrease in FasII distribution at synaptic boutons corresponds to sites of active MAPK localization. However, whether there is a causal relationship between active MAPK and the regions of low FasII remains to be established.

![Figure 4](image-url)  
Figure 4. FasII levels at the NMJ are inversely correlated to MAPK activity. A–C, NMJs immunolabeled with anti-DpMAPK antibody in a hypomorphic MAPK mutant ($rl^{10a/+}$) (A), wild type (B), and a gain-of-function MAPK mutant ($rl^{SEM/+}$) (C). D–I, Equivalent views of type I boutons as A–C, but showing anti-FasII labeling (D–F) and anti-HRP labeling (G–I). Note that increased MAPK activity results in decreased levels of FasII and decreased MAPK activity results in increased levels of FasII. Bouton morphology is not affected, as shown by anti-HRP staining. Scale bar, 20 µm.

![Figure 5](image-url)  
Figure 5. Synaptic FasII levels are altered in a MAPK activity-dependent manner. Body wall muscle extracts from $rl^{10a/+}$, wild-type (CS), $rl^{SEM/+}$, and $dlg^{XI-2}$ were immunoprecipitated with anti-DLG antibody and the Western blots were probed sequentially with anti-FasII and anti-DLG. Note that wild-type, $rl^{SEM/+}$, and $rl^{10a/+}$ have similar DLG levels, but DLG-associated FasII is decreased in $rl^{SEM/+}$ and increased in $rl^{10a/+}$. The molecular masses given at the right of the blots are in kilodaltons.

| Table 1. FasII and DpMAPK immunoreactivity levels at larval NMJs with altered MAPKinase activity |
|----------------------------------|----------------------------------|
| Anti-dpMAPK (n) | Anti-FasII (n) |
| CS | 143 ± 5 (65) | 106 ± 3 (80) |
| $rl^{10a/+}$ | 102 ± 4 (66) | 135 ± 3 (68) |
| $rl^{SEM/+}$ | 194 ± 6 (59) | 80 ± 2 (84) |

Numbers in parentheses represent the number of scored boutons out of $\sim 20$ preparations for each genotype and antibody.
MAPK activity regulates the number of boutons through FasII-mediated adhesion

In Drosophila, as in Aplysia, a decrease in levels of FasII results in an increase in the number of synaptic boutons (Schuster et al., 1996a,b). To demonstrate that activated MAPK can affect FasII levels, leading to an increase in the number of boutons, in this report we used the mutant strains Ras1V12S35 and RafF179, which selectively activate the MAPK pathway (Fig. 1), and quantified the number of boutons in the MAPK mutants. Figure 7 shows the number of synaptic boutons in relation to approximate FasII levels at the NMJ. In a wild-type background (100% FasII levels), enhanced MAPK activity in the gain-of-function allele (rlSEM/+/H11001) resulted in a decrease in FasII levels (Fig. 4; Table 1) that was accompanied by a significant (p < 0.01) increase in the number of boutons (Fig. 7). However, reduction in MAPK activity in the loss-of-function mutant (rl10a/+/H11001) did not significantly affect the bouton number (Fig. 7), although FasII levels were enhanced (Fig. 4; Table 1). Thus, an increase in MAPK activity causes an increase in the number of synaptic boutons, which was accompanied by a decrease in synaptic FasII.

To determine whether the increase in synaptic bouton number observed in rlSEM/+ was likely to be mediated by a decrease in FasII, we examined genetic interactions between fasII and rl by generating fasII rl double mutants. Previous studies have shown that the homophilic cell-adhesion molecule FasII is localized both presynaptically and postsynaptically, where it serves two related roles: it is required for synapse maintenance, and it regulates synaptic growth (Schuster et al., 1996a). The first role was demonstrated by the observation that in fasII-null mutants, synaptogenesis is normal, but motor endings subsequently retract. The second role was established by using fasII mutant alleles with different levels of FasII, in which the motor endings showed an increase in bouton number depending on FasII levels. A decrease in FasII levels to ~50% of normal (e.g., in the heterozygote fasIr70/+ ) results in a striking increase in bouton number, presumably because the strong cell adhesion that stabilizes synaptic endings in the wild type is partially lifted, allowing for sprouting. However, decreases in FasII levels much below 50% (e.g., in fasIr70 homozygotes, which are reported to have on the order of 10% wild-type FasII levels) do not result in an additional increase in bouton number, because FasII-mediated cell adhesion becomes compromised, and this interferes with the maintenance of a large arbor (Schuster et al., 1996a).

To ascertain whether MAPK and FasII function in the same pathway during the regulation of bouton number, we generated the double-mutant combinations shown in Figure 7. The relative positions of the bars in the histograms correspond to the approximate change in FasII levels that MAPK mutants are expected to have (Fig. 4; Table 1) in the different fasII mutant backgrounds. If rl and fasII interact genetically, then the effects that each one has on NMJ growth should be nonadditive. We initially generated the mutant combination fasIIe76/+/H11001; rl10a/+/H11001. We expected that if MAPK and FasII function in the same pathway, then a decrease in MAPK activity should suppress the increased number of boutons resulting from a decreased cell adhesion in the fasIIe76/+ heterozygote. Indeed, we found that the increase in bouton num-

Figure 7. Number of synaptic boutons in mutants with alterations in both FasII and MAPK levels. Background levels of FasII in wild-type (100%), fasIr70 homozygotes (10%), and fasIr70/+ heterozygotes (55%) are indicated on the x-axis. Other bars are positioned on the basis of the influence of the rolled alleles on FasII expression as seen in Figure 4.
ber observed in \( \text{fasII}^{-}\text{79/+} \) heterozygotes was partially suppressed by \( \text{r}^{\text{fl}90/+} \) (Fig. 7). We also generated the combination \( \text{fasII}^{-}\text{79/+}, \text{r}^{\text{fl}2502} \). We expected that if the increase in bouton number observed in \( \text{r}^{\text{fl}2502} \) was attributable to a decrease in FasII at the NMJ, then we should see an enhancement of the increase in bouton number phenotype in \( \text{fasII}^{-}\text{79/+} \), provided that FasII levels do not drop below the levels required to maintain a large arbor. However, we found that in \( \text{fasII}^{-}\text{79/+}, \text{r}^{\text{fl}2502} \) the number of boutons was not significantly different from that in the \( \text{fasII}^{-}\text{79/+} \) heterozygote alone (Fig. 7). The observation that the effects were nonadditive in these double mutants indicates that R1 and FasII interact genetically and act in the same pathway. In addition, we found that the moderate increase in bouton number observed in \( \text{fasII}^{-}\text{79} \) was neither enhanced nor suppressed by changes in MAPK activity. However, the effects of \( \text{rl} \) and \( \text{fasII} \) were nonadditive in every double mutant combination (Fig. 7). The possible interpretation of these results is presented in the Discussion.

**DISCUSSION**

Previous work in *Drosophila* demonstrates that synapse stability and synapse expansion during muscle growth are regulated by changes in FasII expression at presynaptic and postsynaptic membranes and that FasII expression is in part controlled by electrical activity (Schuster et al., 1996a). One mechanism through which electrical activity alters FasII levels is by regulating its synaptic clustering via CaMKII-dependent phosphorylation of DLG (Thomas et al., 1997; Koh et al., 1999). In this report, we provide an additional mechanism by which the levels of FasII at the presynaptic terminal are modified: the activation of the Ras–MAPK pathway. This redundant mechanism may serve the differential regulation of FasII localization at the presynaptic and postsynaptic site or may represent FasII regulation in response to different signals. Whereas activation of CaMKII is elicited by an increase in electrical activity, activation of the MAPK pathway may be triggered by activity or by an as yet unknown but different signaling mechanism.

Studies in *Aplysia* (Bailey et al., 1992, 1997; Martin et al., 1997) indicate that activity-dependent endocytosis of ApCAM results in an increase in the number of synaptic contacts during long-term facilitation (Bailey et al., 1992). Studies by Martin et al. (1997) suggest that ApMAPK is likely to induce ApCAM internalization in a process that depends on ApMAPK activity in dissociated neurons. However, its involvement in the intact organism has not been tested.

In this study, we used *Drosophila* larval neuromuscular synapses to determine the involvement of the Ras–MAPK pathway in the regulation of synaptic FasII levels and in morphological synaptic plasticity. We demonstrate that both Ras and MAPK are expressed at the NMJ, where they regulate presynaptic expansion. We also showed that this regulation is accomplished by altering FasII levels at synaptic boutons.

Ras proteins are conserved from yeast to humans (Cohen, 1997). In *Drosophila*, Ras1 has been involved in several processes, including the mechanisms of cell proliferation, eye development, and apoptosis (Grether et al., 1995; Bergmann et al., 1998; Kurada and White, 1998; Malumbres and Pellicer, 1998; Prober and Edgar, 2000). In this study, we used a ras hypomorphic mutant and anti-Ras antibodies to determine that Ras1 is specifically expressed at the larval NMJ. Although Ras1 immunoreactivity at synapses and muscles was severely reduced in ras1 hypomorphic mutants, nuclear staining persisted.

*Drosophila* MAPK has also been studied intensively for its role in cell proliferation and apoptosis. In particular, antibodies directed against the activated form of MAPK (DpMAPK) have shown it to be present in proliferating tissues, such as embryos and imaginal disks (Gabay et al., 1997b). In this study, we used two antibodies to demonstrate the synaptic localization of MAPK at the NMJ, an antibody that recognizes all forms of the MAPK Rolled (DmERK-A) and an antibody that exclusively labels active, double-phosphorylated MAPK (DpMAPK). Interestingly, although both antibodies labeled synaptic boutons, their distribution was not identical. In particular, the antibody against active MAPK-labeled hot spots was more restricted in its localization than general MAPK staining. This suggests that active MAPK is recruited to specific domains within the synaptic bouton or that MAPK activation occurs at discrete regions within the boutons. Interestingly, the same domain that is occupied by active MAPK has lower levels of FasII, consistent with the idea that MAPK activation might be involved in the downregulation of FasII. A recent report (Sone et al., 2000) suggested that the regions of low FasII concentration correspond to the active zone, suggesting that active MAPK is localized to the active zone. The localization pattern of Ras1 and MAPK at synapses is also consistent with the localization protein 14-3-3, another protein that has been involved in the Ras1–*Drosophila* Raf–MAPK signal transduction pathway (Chang and Rubin, 1997; Zou and Cline, 1999).

**Ras1–MAPK signal transduction pathway regulates the number of synaptic boutons**

In this study, we found that expression of constitutively active Ras (Ras1V12) drastically increased the number of synaptic boutons. This change was indistinguishable from the increase in boutons observed in the Ras1V12S35 variant and the constitutively activated Raf\(^{179}\), suggesting that these changes were induced by activation of the MAPK pathway. Consistent with these results was the observation that a hypomorphic mutation in ras1, ras1\(^{7703}\), had the opposite phenotype, a decrease in bouton number, and that a gain-of-function mutation in \( \text{rl} \) led to an increase in bouton number. Our finding that Ras1V12 and Ras1V12S35 elicited identical phenotypes at the NMJ is consistent with findings in other tissues, such as in the retina, in which the epidermal growth factor receptor–Ras1 pathway is involved in photoreceptor survival (Bergmann et al., 1998), or in the wing disks, where the Ras pathway is involved in hyperplastic growth (Karim and Rubin, 1998).

Notably, expression of Ras variants that activate the PI3-K and Ral signal transduction pathways and a constitutively active Raf may also induce an increase in bouton number that was similar in extent to RasWT and considerably lower than Ras1V12. These results raise the possibility that Ras1V12G37 and Ras1V12C40 may still retain some degree of affinity for Raf or, alternatively, that other Ras-mediated pathways might also influence NMJ development. All known ras genes encode a protein region, the effector loop, that is highly conserved in all species. Mutations in this loop interfere with the ability of Ras to bind to specific effectors without altering its catalytic activity. A series of mutations in the effector loop that allow almost exclusive activation of a single effector have been isolated in mammals. The specificity of these mutants has been tested by *in vitro* binding assays as well as by genetic and biochemical approaches in cell culture (White et al., 1995; Khosravi-Far et al., 1996; Rodriguez-Viciana et al., 1997). In *Drosophila*, a genetic approach has been used to demonstrate specificity. These studies suggest that Ras1V12 and
RasV12S35 phenotypes are emulated by a hyperactivated form of Raf and suppressed by Raf, MEK, and MAPK mutants (Karim et al., 1996; Karim and Rubin, 1998; Halar et al., 2001).

Studies in vertebrate cells and in Drosophila suggest that MAPK regulates levels of synaptic FasII, a cell-adhesion molecule that plays a key role in the maintenance and expansion of NMJs in Drosophila (Schuster et al., 1996a,b). This model was supported by experiments in which only type I synaptic FasII was immunoprecipitated. This was accomplished by using anti-DLG antibodies, because DLG binds directly to FasII at type I boutons but not at other bouton types (Thomas et al., 1997). The immunoprecipitation experiments demonstrated that enhancing the levels of MAPK activity at synaptic terminals resulted in a reduction of type I synaptic FasII. Conversely, decreasing levels of MAPK activity resulted in an increase in type I synaptic FasII levels. These results are in agreement with the studies in Aplysia dissociated neurons, which show that ApMAPK is involved in the internalization of ApCAM (Bailey et al., 1997; Martin et al., 1997).

Additional support for the idea that the changes in bouton number elicited by alterations in RasI and MAPK activity are mediated by alterations in FasII levels was demonstrated by examining the overall expression of FasII in MAPK gain- or loss-of-function alleles, examining the distribution of FasII within single synaptic boutons in relation to active MAPK, and using hypomorphic fasII mutants. The studies with nI mutants demonstrated that there was an inverse relationship between levels of synaptic FasII and MAPK activity. Furthermore, active MAPK localization coincided with regions of the bouton that have no or low FasII levels.

The studies by Schuster et al. (1996a,b) demonstrate two main functions of FasII in the regulation of synapse number. First, FasII is critically required for synapse maintenance: below threshold FasII levels, synaptic boutons are not maintained. Second, FasII operates by constraining synaptic growth, similar to the Aplysia system (Abel et al., 1998). Therefore, a decrease in FasII to a level still sufficient for maintenance results in an increase in synaptic arbor size (Schuster et al., 1996a). On the basis of this model, we propose the following interpretation of our results. The dramatic decrease in FasII levels in the homozygous fasII mutant did not allow any influence of MAPK activity changes on NMJ structure. Similarly, when FasII levels were decreased to approximately one-half the wild-type levels (fasII<sup>79/+</sup>), an increase in MAPK activity did not induce an additional increase in bouton number, probably because an additional decrease in FasII compromises synaptic maintenance, thus preventing NMJ growth. However, the increase in FasII levels induced by a reduction of MAPK activity (nI<sup>+</sup>/+ in a fasII<sup>79/+</sup> background) suppressed the increase in boutons observed in fasII<sup>79/79</sup> alone. This result suggested that MAPK regulates FasII levels and exists upstream of FasII at signal transduction pathways that regulate the number of type I synaptic boutons.

Notably, the hypomorph nI<sup>+/+</sup> had no significant decrease in bouton number, although these mutants had a striking increase in FasII levels compared with wild-type controls. An explanation for this result is that FasII is a homophilic cell-adhesion molecule that is required both in the presynaptic and in the postsynaptic cell for function (Schuster et al., 1996a; Thomas et al., 1997). If the Ras–MAPK pathway functions to regulate FasII at the presynaptic cell, as suggested by our studies with cell-specific Gal4 drivers, then an asymmetric increase in FasII levels in the presynaptic cell alone may not have much of an effect. Previous studies also show that although the NMJ is very sensitive to a decrease in FasII levels, an increase in FasII over wild-type levels does not have much of an effect (Schuster et al., 1996a).

Although our results are consistent with a regulation of FasII-mediated synapse growth by the Ras–MAPK pathway, it is important to note that several other molecules in addition to FasII are involved in the regulation of synapse growth (Torroja et al., 1999; Sone et al., 2000; Parnas et al., 2001). Moreover, several studies suggest that many changes at the fly NMJ are compensated by yet unknown homeostatic mechanisms (Davis and Goodman, 1998). Therefore, further understanding of these regulatory and compensatory signals will be necessary to fully explain our observations.

In conclusion, we have identified a signaling pathway intimately involved in the regulation of synaptic growth at the NMJ. Identification of the mechanisms involved in the activation of this pathway may provide valuable clues toward understanding the plasticity of this synapse.

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