

# Activity and Distribution of Paxillin, Focal Adhesion Kinase, and Cadherin Indicate Cooperative Roles during Zebrafish Morphogenesis

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We investigated the focal adhesion proteins paxillin and Fak, and the cell-cell adhesion protein cadherin in developing zebrafish (*Danio rerio*) embryos. Cadherins are expressed in presomitic mesoderm where they delineate cells. The initiation of somite formation coincides with an increase in the phosphorylation of Fak, and the accumulation of Fak, phosphorylated Fak, paxillin, and fibronectin at nascent somite boundaries. In the notochord, cadherins are expressed on cells during intercalation, and phosphorylated Fak accumulates in circumferential rings where the notochord cells contact laminin in the perichordal sheath. Subsequently, changes in the orientations of collagen fibers in the sheath suggest that Fak-mediated adhesion allows longitudinal expansion of the notochord, but not lateral expansion, resulting in notochord elongation. Novel observations showed that focal adhesion kinase and paxillin concentrate at sites of cell-cell adhesion in the epithelial enveloping layer and may associate with actin cytoskeleton at epithelial junctions containing cadherins. Fak is phosphorylated at these epithelial junctions but is not phosphorylated on Tyr<sup>397</sup>, implicating a noncanonical mechanism of regulation. These data suggest that Fak and paxillin may function in the integration of cadherin-based and integrin-based cell adhesion during the morphogenesis of the early zebrafish embryo.

## INTRODUCTION

The morphogenesis of embryos depends both on interactions between cells and their surrounding extracellular matrix (ECM) via integrin complexes and on direct cell-cell

interactions via cadherins. Although these interactions are perceived as discrete adhesion systems, they may interact with each other during morphogenesis. The integrin family of transmembrane receptors physically connects the actin cytoskeleton of the cells to the ECM at focal adhesions, and thus, mediates migration and adhesion in many cells (reviewed in Turner, 2000a; Schwartz, 2001). The importance of focal adhesions in early development is well illustrated by mouse embryos deficient in fibronectin or focal adhesion components, including focal adhesion kinase (FAK), paxillin, or integrins. These embryos die early in development (days 5–10) with mesodermal defects (Furuta *et al.*, 1995; Georges-Labouesse *et al.*, 1996; Yang *et al.*, 1999; Hagel *et al.*, 2002).

Cadherins are prominent in adherens junctions and desmosomes used in the formation of epithelial tissues. They are transmembrane proteins, which form homophilic interactions between neighboring cells and connect the plasma membrane with the actin cytoskeleton. They are involved in signaling, polarization of epithelia, cell sorting, and probably cell migration (reviewed in Gumbiner, 2000). Cadherins are also thought to regulate many aspects of morphogenesis:

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Abbreviations used: CAK- $\beta$ , cell adhesion kinase- $\beta$ , which is also known as protein tyrosine kinase 2 $\beta$  (PTK2 $\beta$ ), and focal adhesion kinase 2 (FADK 2); ECM, extracellular matrix; EVL, enveloping layer; FAK, mouse, *Xenopus*, or human focal adhesion kinase protein; Fak, zebrafish focal adhesion kinase protein; all numbering of Fak residues is according to the known phosphorylated human sites; *fak*, mRNA or DNA focal adhesions kinase gene; hpf, hours postfertilization; PGMT, postgastrula mesoderm transition; TEM, transmission electron microscopy.

mouse embryos deficient in N-cadherin (Cdh2) develop mesodermal defects such as irregularly shaped somites and an undulating notochord and die at day 10 (Radice *et al.*, 1997). Mosaic expression of a dominant-negative cadherin in *Xenopus* embryos results in failure of the expressing cells to intercalate during notochord morphogenesis (Delarue *et al.*, 1998). In N-cadherin mutants (*parachute*) of zebrafish (*Danio rerio*), convergent cell movements in the neural tube are severely compromised (Lele *et al.*, 2002).

The integrin-mediated and cadherin-mediated adhesion-modulating systems may interact. Somitogenesis in vertebrate embryos involves presumptive somite border cells with cadherin-based adhesion apically (Linask *et al.*, 1998) while forming integrin-based focal adhesions to the ECM basolaterally (Hens and DeSimone, 1995; Henry *et al.*, 2001). In addition, mesoderm migrates as a multilayered coherent mass held together by cadherins in *Xenopus* embryos (Winklbauer *et al.*, 1996). It seems plausible that the mechanism that coordinates these activities may involve proteins that participate in both processes.

Paxillin functions as an adaptor protein coordinating the activities of many focal adhesion proteins (Turner, 2000b). Thus, paxillin is in a position to play a role in the integration and regulation of adhesion and signaling, yet little is known regarding its function during embryogenesis (Turner, 1991; Hagel *et al.*, 2002). FAK (abbreviated "Fak" in zebrafish) is a nonreceptor tyrosine kinase that binds paxillin and is a well-characterized component of focal adhesions. Integrin-mediated FAK-autophosphorylation of Tyr<sup>397</sup> generates a binding site for Src, which in turn phosphorylates a number of other tyrosine residues on FAK, notably Tyr<sup>576/577</sup> in the kinase domain and Tyr<sup>861</sup> in the proline-rich domain (Schlaepfer and Hunter, 1998; Cary *et al.*, 2002). Consequently, the phosphorylation state of Tyr<sup>397</sup> is considered indicative of FAK's activity. FAK has been localized at sites of strong cell-ECM adhesion in embryos. Hens and DeSimone (1995) and Polte *et al.* (1994) showed that in *Xenopus* and mouse embryos FAK is prominent at somite borders and in the brain. We have shown it to be prominent in somite borders and in the notochord cells in zebrafish embryos (Henry *et al.*, 2001).

To elucidate the roles of cell-cell and cell-ECM adhesion in morphogenesis and to characterize these important regulators of adhesion, we have cloned cDNAs representing *paxillin* (this study) and two isoforms of *fak* (Henry *et al.*, 2001; this study). We examined the differential phosphorylation of tyrosine residues in Fak in the context of possible interactions with paxillin and cadherins in the developing fish embryo. We found that Fak is dynamically phosphorylated after gastrulation when the somites form and the notochord extends the body-axis. We found that a novel pattern of Fak phosphorylation occurs in the epithelial enveloping layer (EVL) at epithelial junctions where Fak colocalizes with paxillin and cadherins. Our analysis of the expression and phosphorylation dynamics of these proteins, combined with ultrastructural observations, provides insight into the mechanisms of morphogenesis in the EVL, somites, and notochord.

## MATERIALS AND METHODS

### Cloning of Paxillin cDNA

Polymerase chain reaction (PCR) was used to isolate a probe for zebrafish *paxillin*. The probe was a 603-base pair DNA fragment

amplified from a zebrafish gastrula stage cDNA library (kindly provided by Dr. Randall T. Moon, University of Washington, Seattle, WA) by using 5'-TTC CCT GCT GAA GAA CCT-3' and 5'-CTC GCA GTC GTC ACA GCG-3' as primers. A 2566-base pair cDNA of *paxillin* was isolated from a zebrafish gastrula stage library provided by Drs. Thierry Lepage and David Kimelman (University of Washington, Seattle, WA). The GenBank accession number for the full-length zebrafish *paxillin* sequence is AY144487.

### Cloning of fak1b cDNA

Sequences of *fak1b* were initially identified by a comparison of the PCR products from a zebrafish PAC genomic library generously supplied by Dr. Bruce W. Draper (Frederick Hutchinson Cancer Research Center, Seattle, WA), by using specific and degenerate primers for the amino acid sequences "PPTANLDR" and "EYVP-MVKEVG" in the FAT domain. Degenerate sequences were designed by CODEHOP (Fred Hutch Cancer Research Center and <http://bioinformatics.weizmann.ac.il/blocks/codehop.html>). The *fak1b* PCR product was used to isolate clones of *fak1b* from the Lepage/Kimelman gastrula zebrafish library, which was cloned into a Uni-ZAP XR unidirectional vector (Stratagene, San Diego, CA). To determine the specificity of the FAK C-20 antibody (Santa Cruz Biotechnology, Santa Cruz, CA), which we have previously demonstrated recognizes Fak1a in zebrafish (Henry *et al.*, 2001), we expressed Fak1b and assayed cross-reactivity by Western blot. The *fak1b* clone was subcloned into a pGEX-5X-3 vector (Amersham Biosciences, Piscataway, NJ) and transcribed and translated using PROTEINscript-PRO (Ambion, Austin, TX). The 5' end of *fak1b* was sequenced from a TA-clone derived from a 5' rapid amplification of cDNA ends (RACE) cDNA library generously supplied by Chris J. Thorpe (University of Washington, Seattle, WA), by using SMART RACE (BD Biosciences Clontech, Palo Alto, CA) and 5'-CGTCA-CATTCTCATACACTT-3' as the reverse *fak1b* primer. The GenBank accession number for the full-length zebrafish *fak1b* sequence is AY196213.

### Mapping of paxillin and fak1b

The meiotic mapping of *paxillin* was performed as in Henry *et al.* (2001). Genomic DNAs from the mapping panels were amplified using primers from the 3'-untranslated region of *paxillin* to give a 342-base pair fragment (forward *paxillin* +53 CCTGTGGCCG-TAGTGAGCA, reverse *paxillin* -394 AGACATGATACGGCCGAG-GAAGAA). Genomic DNAs from the mapping panels were similarly amplified using primers from the 3'-untranslated region of *fak1b*.

### Whole-Mount In Situ Hybridization

Single in situ hybridizations were performed as in Henry *et al.* (2001). The antisense in situ probes were synthesized to hybridize with the 3'-untranslated region of the *paxillin*, *fak1a*, and *fak1b* transcripts. Sense controls showed no specific staining (our unpublished data). Images of stained embryos were prepared using Photoshop 7.0 (Adobe Systems, Mountain View, CA).

### Western Blot Analysis

Methods were similar to those in Henry *et al.* (2001). For each stage to be analyzed, embryos were partially de-yolked by poking the yolk ball with sharp forceps and gently squeezing the embryo to force out the majority of yolk. Deflated embryos were suspended in ice-cold embryo medium (Westerfield, 1995) with 10 mM orthovanadate and one tablet of protease inhibitors/5 ml medium (double the recommended dose of Mini Complete tablets; Roche Diagnostics, Indianapolis, IN), settled on ice, and boiled in 2× reducing SDS-PAGE sample buffer at a concentration of 1 embryo/μl. Ten microliters of embryo extracts was run per lane in a 10% gel (Invitrogen, Carlsbad, CA), blotted, and processed for enhanced

chemiluminescence detection as in Henry *et al.* (2001). Primary antibodies were used diluted at 1:1000, except for anti-paxillin, which was used at 1:5000, and secondary antibodies were used at 1:5000. To detect weak signals during epiboly, films were exposed to the blots for 7 to 10 times longer than for postbud stages. Films were scanned using an Umax 6500 scanner and prepared using Photoshop 7.0 (Adobe Systems). For blots that were probed with multiple antibodies, membranes were stripped using Restore Western blot stripping buffer (Pierce Chemical, Rockford, IL) and redetected with secondary antibodies between probes to ensure complete removal of primary antibodies. Antibodies used were polyclonal antibodies to the C-terminal end of FAK (C-20; Santa Cruz Biotechnology) and affinity-purified polyclonal antibodies to the SH2 binding sites pY<sup>397</sup>FAK, pY<sup>576</sup>FAK, pY<sup>577</sup>FAK, pY<sup>861</sup>FAK, and pY<sup>925</sup>FAK (BioSource International, Camarillo, CA), which are conserved in human, mouse, and chick. The antibody against paxillin was a monoclonal to human paxillin (Transduction Laboratories, Lexington, KY).

The Tyr<sup>397</sup> antigenic SH2 binding sites of zebrafish Fak1a and Fak1b are identical with that of human FAK (see Figure 2; Dre Fak1a: Henry *et al.*, 2001 and GenBank AAK31154; Dre Fak1b: GenBank AY196213). In comparison, the sequence at this site in human cell adhesion kinase- $\beta$  (CAK- $\beta$ ) is dissimilar at four amino acids and becomes very dissimilar as the length of the peptide is extended. BioSource International shows that their human pY<sup>397</sup>-FAK peptide, but not their human pY<sup>397</sup>-CAK- $\beta$  competes with their pY<sup>397</sup> affinity-purified polyclonal antibody (<http://www.biosource.com/>; catalog no. 44-624). Thus, their antibody to pY<sup>397</sup>-FAK is specific for FAK. The CAK- $\beta$  gene in zebrafish has not yet been identified.

The Tyr<sup>925</sup> SH2 binding site is identical for human FAK and zebrafish Fak1a and Fak1b (see Figure 2). The anti-pY925 antibody, however, is weak and only lightly stains zebrafish proteins on Western blots and does not stain whole mounts. The SH2 binding sites for Tyr<sup>576/577</sup> and Tyr<sup>861</sup> have one or two conservative amino acid differences between humans and zebrafish. In contrast, the site for human CAK- $\beta$  Tyr<sup>576/577</sup> is very different in charge, and CAK- $\beta$  has no Tyr<sup>861</sup> SH2 binding site (see Figure 2). The anti-pY576/577 and anti-pY861 antibodies gave strong signals in both Western blots and whole mounts of fish embryos.

The polyclonal antibody to the C-terminal end human FAK (C-20; Santa Cruz Biotechnology) was shown by us to recognize the expressed *Dre* Fak1a protein (Henry *et al.*, 2001). By Western Blot analysis of the in vitro-expressed proteins, we have shown that the FAK (C-20) antibody does not recognize with *Dre* Fak1b and confirmed that it recognizes *Dre* Fak1a (our unpublished data). The C termini of human CAK- $\beta$  do not significantly match that of human FAK (see Figure 2). Thus, all of our antibodies except that to C-20 FAK recognize both Fak1a and Fak1b and not CAK- $\beta$ . The antibody to C-20 Fak recognizes only Fak1a.

### Whole-Mount Antibody Immunofluorescence

Whole-mount immunostaining was used to determine the subcellular localization of various antigens. In addition to the antisera used in Western blot analysis, we used rabbit anti-laminin and anti-fibronectin (NeoMarkers, Fremont, CA), mouse anti-pTyr (pY99; Santa Cruz Biotechnology), and rabbit anti-PanCadherin (Sigma-Aldrich, St. Louis, MO), which was used previously by Bitzur *et al.* (1994) on zebrafish. With the exception of experiments with the anti-FAK polyclonal antibody, the embryos were dechorionated and fixed with 4% paraformaldehyde in phosphate-buffered saline (PBS) with 10 mM orthovanadate for 24 h at 4°C. All embryos were rinsed with PBDT (1% bovine serum albumin, 1% dimethyl sulfoxide, and 0.1% Triton X-100 in PBS pH 7.3) and then blocked in PBDT with 5% bovine serum albumin at room temperature for 2 h. Embryos were incubated with primary antibodies diluted at 1:200 overnight at 4°C. Embryos were washed three times in PBDT, incubated 2 h with secondary antibodies (goat anti-mouse Alexa-

488, goat anti-rabbit Alexa-568; Molecular Probes) at 1:250 dilution at room temperature, and washed three times in PBDT, mounted, and viewed using a Radiance 2000 confocal microscope (Bio-Rad, Hercules, CA) fitted with 20 $\times$  0.8 numerical aperture dry and 63 $\times$  1.3 numerical aperture water immersion objectives. Microscopic data were processed and analyzed using ImageJ 1.27z (<http://rsb.info.nih.gov/ij/>) and prepared for publication using Photoshop 7.0 (Adobe Systems). For actin-staining experiments, Alexa-568-conjugated phalloidin (Molecular Probes) was added to the secondary incubation at 1:100. For negative control experiments, diluted primary antibody was heat inactivated for 5 min at 70°C before use. No specific staining was observed in any negative control. Embryos stained with anti-FAK were fixed in 2% trichloroacetic acid as in Henry *et al.* (2001).

### Detection of Apoptosis by Terminal Deoxynucleotidyl Transferase dUTP Nick-End Labeling (TUNEL)

Apoptotic cell death in the EVL of normal embryos was detected using the TUNEL technique. Embryos were fixed in 4% paraformaldehyde overnight at 4°C, made permeable by washing 3  $\times$  15 min and 1  $\times$  5 min in PBS + 1% Triton X-100, biotinylated with dUTP according to the Roche protocol, and labeled nuclei detected using streptavidin-Alexa488 (Molecular Probes).

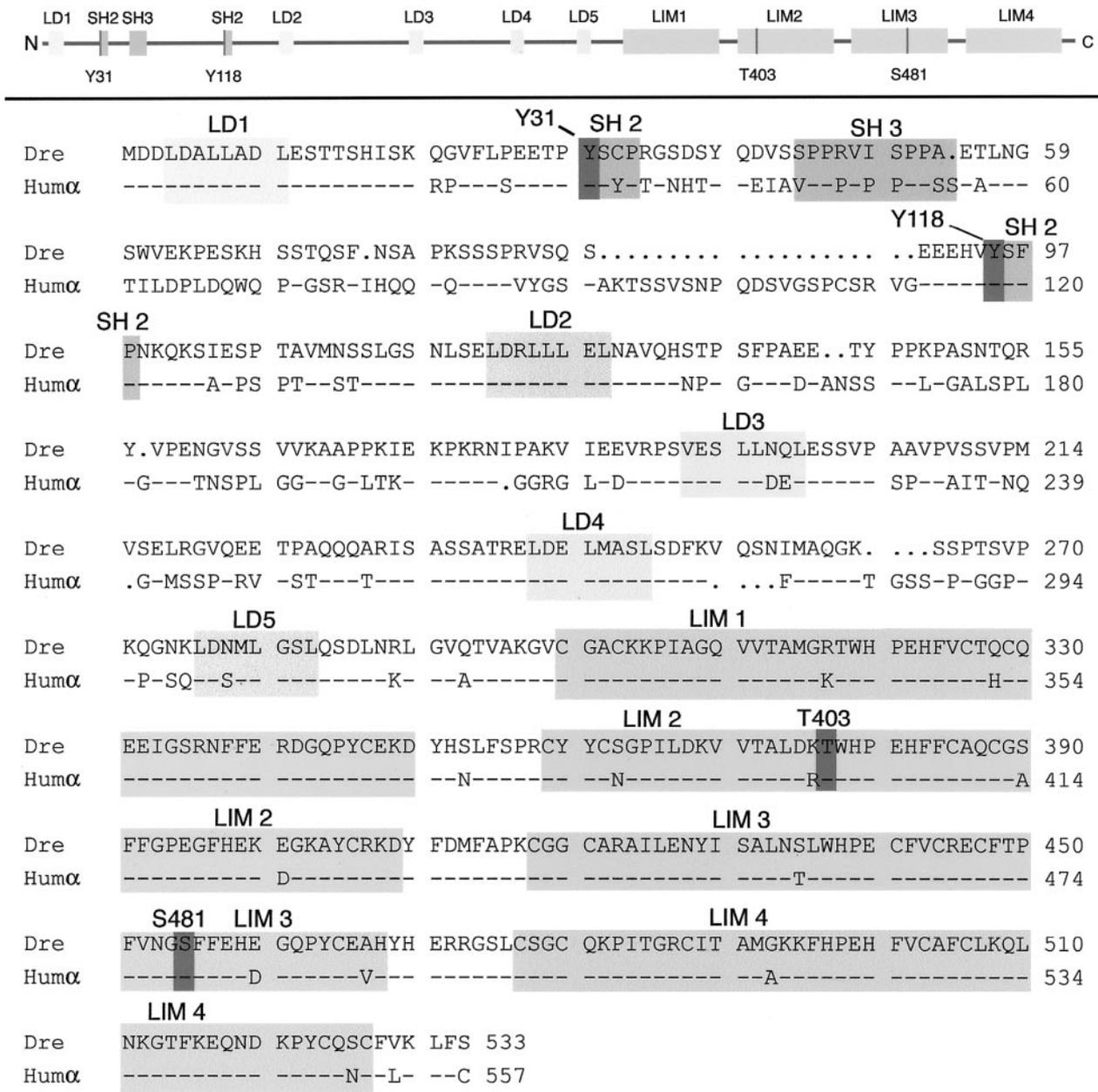
### Transmission Electron Microscopy (TEM)

Embryos were dechorionated and fixed in Karnofsky's fixative (2% paraformaldehyde, 2.5% glutaraldehyde, 5% sucrose, 0.1% CaCl<sub>2</sub>, in 0.2 M cacodylate buffer pH 7.2) overnight at 4°C. For TEM embryos were postfixed in 4% osmium tetroxide (Electron Microscopy Sciences, Fort Washington, PA) in 0.2 M cacodylate buffer for 1 h and then dehydrated through graded ethanols, embedded in LX-112 resin overnight, and polymerized at 70°C. Blocks were sectioned, stained with uranyl acetate and lead citrate, and viewed on a Phillips CM100 transmission electron microscope. Images were prepared using Photoshop 7.0 (Adobe Systems).

## RESULTS

### Zebrafish Paxillin Is Conserved

The functional domains of paxillin are highly conserved. The amino acid sequence of zebrafish paxillin is 75% identical compared with human  $\alpha$ -paxillin (Figure 1). In the amino half of human paxillin, there are five conserved leucine-rich motifs (LDxLLxL), termed LD domains, which bind to a number of other proteins, including FAK, vinculin, and p95PKL, and indirectly to the Rho family of GTPases (Tong *et al.*, 1997; West *et al.* 2001; Brown *et al.*, 2002; Roy *et al.*, 2002). These LD domains are conserved in zebrafish paxillin. Paxillin's carboxyl terminus consists of four tandemly repeated LIM domains essential for localization to focal adhesions, which show 94% identity between the fish and human sequences. Important sites of phosphorylation (Tyr<sup>31/118</sup>, Thr<sup>403</sup>, and Ser<sup>481</sup>) and the surrounding SH2 binding sites are also conserved (Figure 1, dark highlights). Less well conserved is a proline-rich SH3 binding site near the N terminus. The nuclear export signal overlapping the LD2 domain in *Xenopus*, human, and chicken paxillin (Lxx-LxxLxLxL; Ogawa *et al.* 2001) is conserved in the zebrafish sequence as well. However, we do not detect paxillin in the nuclei of cells under our conditions. Notable differences between the zebrafish and human sequences include the 21 amino acid insertion in the human sequence between LD1



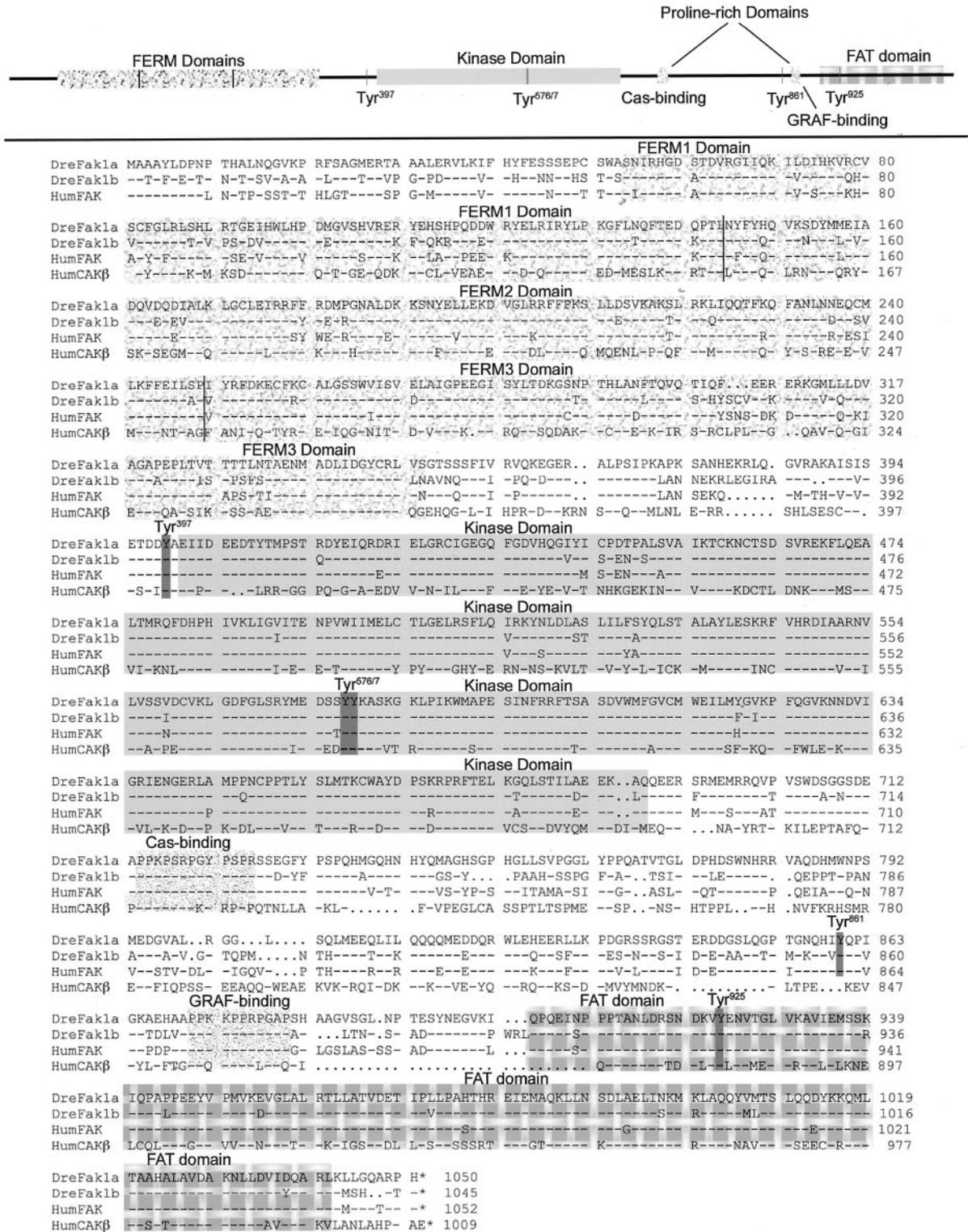
**Figure 1.** Zebrafish paxillin is conserved. The LD and LIM domains, and the SH2 and SH3 binding sites are shaded and labeled. Typically, phosphorylated amino acid residues are shaded and marked as Y31, Y118, T403, and S481. Dre is the zebrafish *D. rerio* sequence and Hum $\alpha$  the  $\alpha$ -paxillin isoform of human. *D. rerio* protein and cDNA sequences are in GenBank AY144487.

and LD2, and considerable sequence variability in the N-terminal region of the protein, which is also less conserved in other vertebrates.

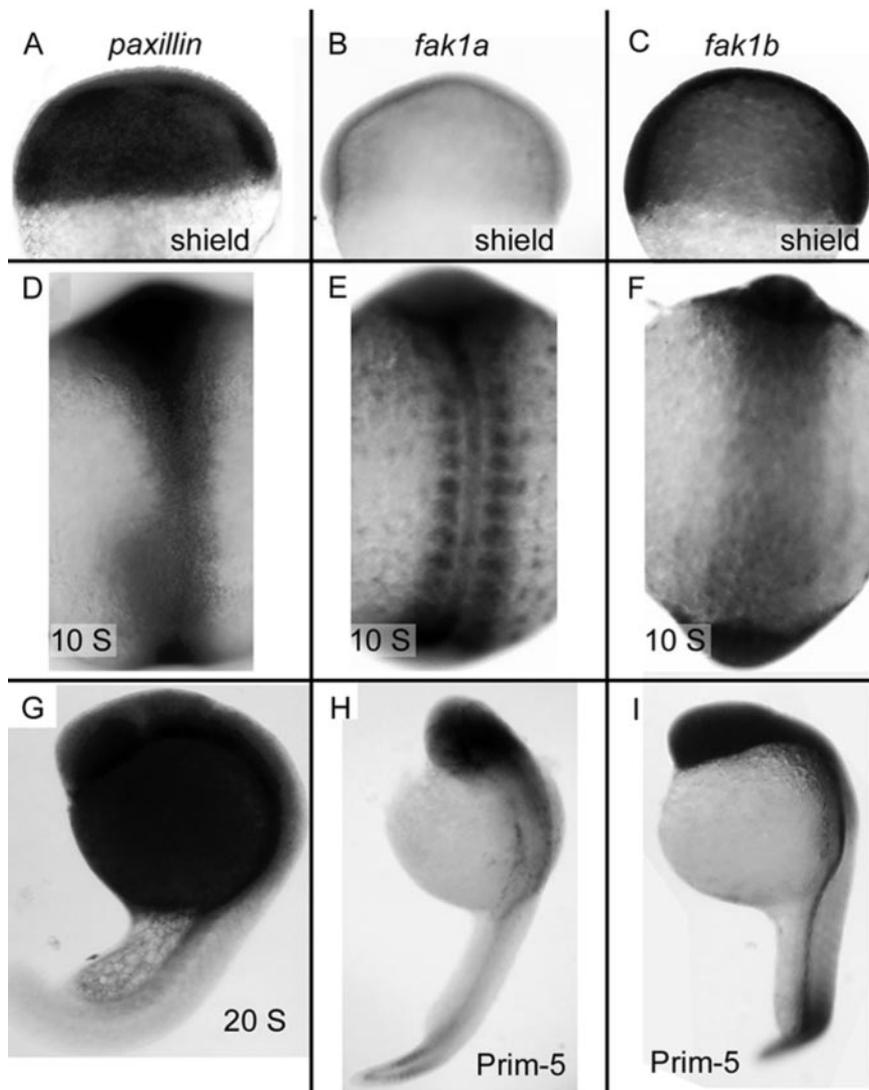
Zebrafish *paxillin* maps to linkage group 5 (LG5) with a high statistical significance (courtesy of Yi-Lin Yan and John Postlethwait, University of Oregon, Eugene, OR). It placed 3.05 cR from Z6323 with a Z6323 LOD of 14.1. Analyses of apparent orthologs show no synteny with human *pavilion*, which maps to human chromosome 14.

**Zebrafish Has Two fak Genes, *fak1a* and *fak1b***

We have cloned the duplicated *Fake* gene, *fak1b*, which is very similar to the previously reported *fak1a* (Henry *et al.*, 2001) and, like *fak1a*, is more similar in derived amino acid sequence to mammalian FAK than to CAK- $\beta$ . The amino acid sequence of zebra fish Fak1b is 69% identical compared with zebra fish Fak1a or with human FAK1 (Figure 2). The kinase domain is very highly conserved being 94% identical to zebra fish Fak1a and 93% identical to human FAK1 (Figure



**Figure 2.** Zebrafish Fak1b is conserved with respect to zebrafish Fak1a and human FAK1 and differs significantly from human CAKβ. The FERM, kinase, and the FAT domains are shaded and labeled. The SH2 binding domains have their tyrosine residues darkening. The proline-rich Cas- and GRAF-binding domains are labeled. Dre designates the zebrafish *D. rerio* sequences and Hum the human sequences. Dre Fak1a is in Henry *et al.* (2001) and GenBank AAK31154; Dre Fak1b in GenBank AY196213; Hum FAK1 from GenBank Q05397; and Hum CAKβ from GenBank Q14289.



**Figure 3.** Expression of *paxillin*, *fak1a*, and *fak1b* mRNAs in the zebrafish embryo. (A–C) Shield stage embryos (6 hpf) hybridized with *paxillin* (A), *fak1a* (B), and *fak1b* (C) probes. Animal pole is up. (D–F) Ten-somite embryos (14 hpf) hybridized with *paxillin* (D), *fak1a* (E), and *fak1b* (F) probes. Presumptive head is up. (G) A 20-somite embryo (19 hpf) hybridized with a *paxillin* probe (G). (H and I) Prim-5 embryos (24 hpf) hybridized with *fak1a* (H) and *fak1b* (I) probes. Head is up and pointing left.

2, shaded area). The focal adhesion-targeting domain (FAT) of zebra fish Fak1b is also highly conserved, 93% with zebra fish Fak1a and 92% with human FAK1 (Figure 2, cross-hatched). The proline-rich Cas binding domain (amino acids 714–724; Figure 2) is identical between all three FAK sequences. The FERM2 phosphotyrosine binding domain, NYFY at amino acids 144–147, is identical for Fak1a and Fak1b and similar to Hum FAK (NFFY) (Garcia-Alvarez *et al.*, 2003). Of importance, for our observations, is that the SH2 binding sites surrounding Tyr<sup>397</sup> and Tyr<sup>925</sup> are identical for zebrafish Fak1a, Fak1b, and human FAK1 and that the SH2 binding sites for Tyr<sup>576/577</sup> and Tyr<sup>861</sup> are identical except for a few conservative changes (Figure 2, highlighted). Although the SH2 binding sites are similar for the zebrafish and human FAK proteins, there are significant differences between them and the SH2 binding sites of human CAK- $\beta$  (Figure 2). Those areas that differ most significantly between zebrafish Fak1a, Fak1b, and human FAK are the N-terminal ends, two areas of the FERM3 domain, and the areas between the proline-rich Cas binding domain and

the FAT domain. These areas are not usually conserved between species.

Zebrafish *fak1b* maps to linkage group 19 (LG19), which is a duplicate of LG 16 containing *fak1a* (see AAK31154 and AY196213). Both LG19 and LG 16 share synteny with the distal tip of human chromosome 8, the site of the human FAK locus.

#### *paxillin*, *fak1a*, and *fak1b* mRNAs Are Expressed throughout Embryogenesis

We previously found that *fak1a* mRNA is segmentally expressed in the paraxial mesoderm and is regulated at least in part by the Notch signaling pathway (Figure 3E; Henry *et al.*, 2001). We compared the expression of *paxillin* and *fak1b* with that of *fak1a*. In situ hybridization reveals that *paxillin* mRNA is expressed ubiquitously throughout the embryo at the shield stage (Figure 3A), when very little *fak1a* mRNA is detected (Figure 3B; Henry *et al.*, 2001). The expression of *fak1b* is similar to *paxillin* at shield stage (Figure 3C). At the

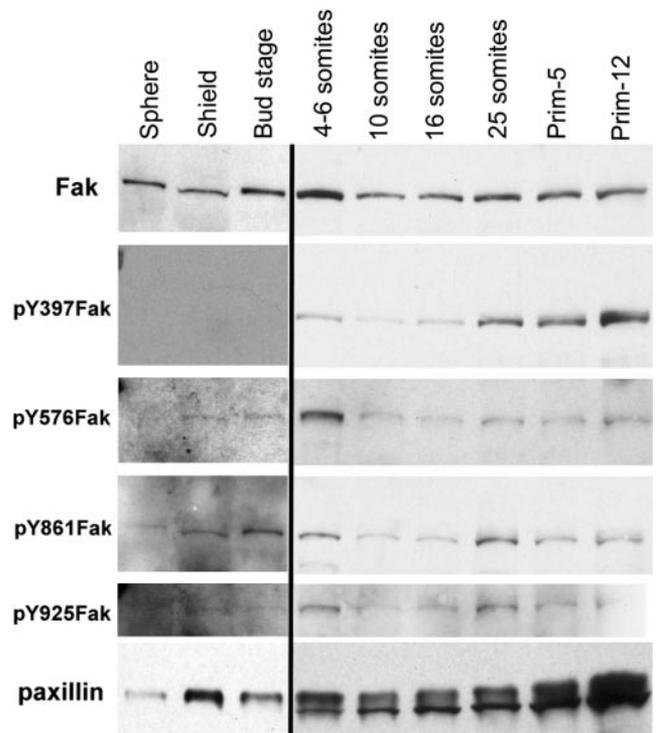
10-somite stages, *paxillin*, *fak1a*, and *fak1b* mRNAs accumulate in the axial region, the head, and the tail. However, in the paraxial region *paxillin* and *fak1b* are expressed homogeneously (Figure 3, D and F, respectively), whereas *fak1a* is expressed segmentally (Figure 3E). At the 20-somite stage, *paxillin* is detected strongly in the head and EVL surrounding the yolk, and less strongly in the tail (Figure 3G). In 20-somite embryos, *fak1a* and *fak1b* are both expressed strongly in the head region and in the dorsal axis, but not strongly in the EVL (Figure 3, H and I). *fak1a* is also segmentally expressed in the newly formed somites in the tail (Figure 3H). The mRNA expression is reflected in the protein expression (see below; Figures 5 and 9).

We also examined the expression of *paxillin* mRNA in the *fused somite* class of mutations, which are believed to disrupt Ephrin and Notch signaling and in which expression patterns of *fak1a* mRNA are disrupted (Henry *et al.* 2001, 2002; Holley *et al.*, 2002). Expression of *paxillin* is ubiquitous throughout the paraxial mesoderm in *fused-somite* (*fss/Tbx24*), *beamter* (*bea*), *deadly seven* (*des/notch 1*), and after-eight (*aei/DeltaD*) embryos and is indistinguishable from wild-type embryos (our unpublished data). Thus, the transcriptional regulation of *paxillin* is markedly different from *fak1a*, in that only *fak1a* is segmentally expressed and regulated by the Notch pathway.

#### Expression and Phosphorylation of Focal Adhesion Proteins Change Dramatically after Gastrulation

FAK is phosphorylated at several SH2 binding sites, both because of autophosphorylation in response to integrin signaling, and because of the activity of other kinases such as Src (Hanks and Polte, 1997). We used several affinity-purified rabbit antibodies that recognize conserved phosphorylated SH2 binding sites from human FAK to analyze the phosphorylation of the zebrafish Fak epitopes in whole-embryo extracts. The SH2 binding sites of both Fak1a and Fak1b are identical for the zebrafish and human sequences flanking Tyr<sup>397</sup>, and those sequences flanking Tyr<sup>925</sup>. The sequences flanking Tyr<sup>576/577</sup> and Tyr<sup>861</sup> have just one or two conserved changes between humans and zebrafish (see MATERIALS AND METHODS). Several lines of evidence indicate that zebrafish CAK- $\beta$  is not recognized by these antibodies. The comparable Tyr<sup>397</sup> and Tyr<sup>576/577</sup> sequences in human CAK- $\beta$  are very dissimilar, the Tyr<sup>861</sup> SH2 binding site does not exist, and binding of the antiserum to the FAK Tyr<sup>397</sup> peptide is not competitively inhibited by CAK- $\beta$  peptides (see MATERIALS AND METHODS).

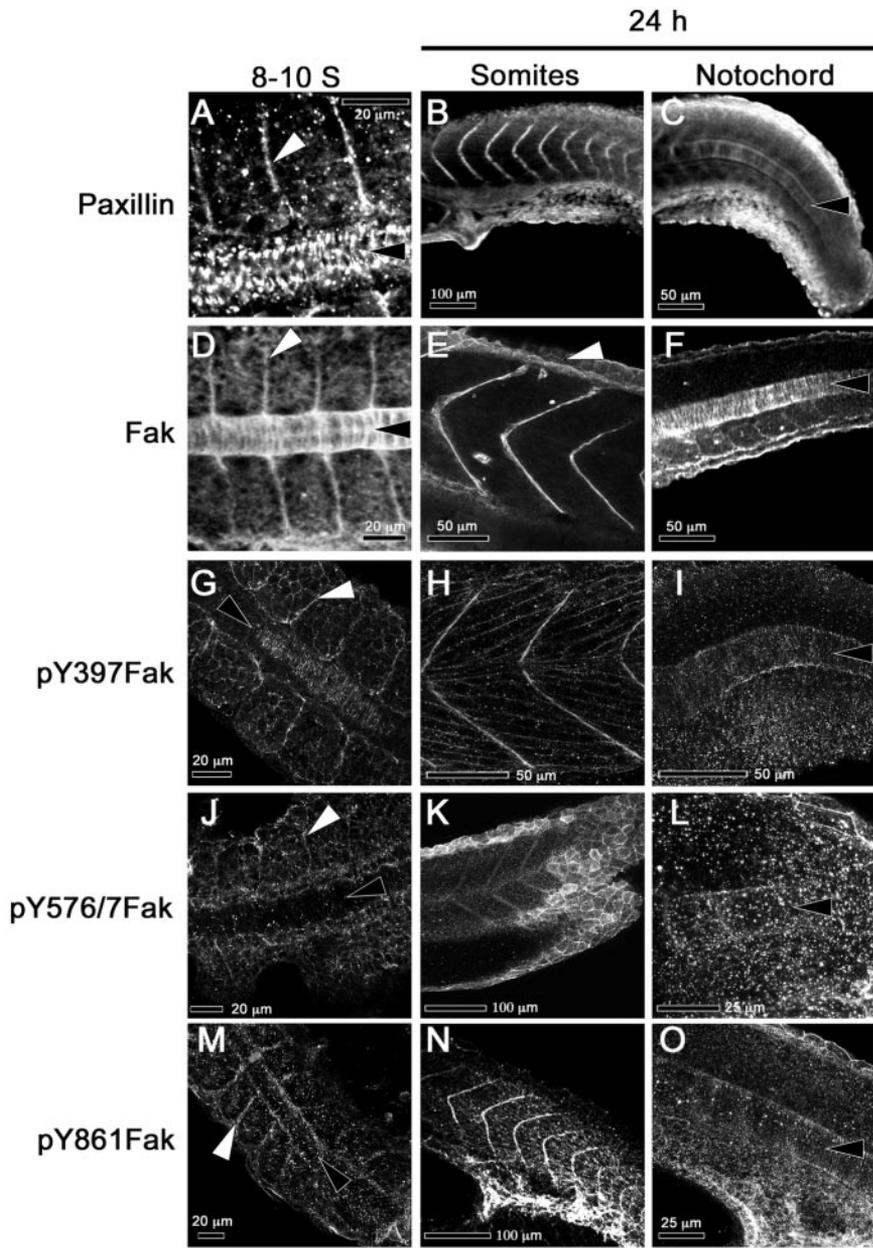
Dramatic increases in the abundance and phosphorylation of Fak and the abundance of paxillin occur during somitogenesis (Figure 4). In Figure 4, this increase is underrepresented because of the long exposures used to detect signals during epiboly and gastrulation (first three lanes). During epiboly and gastrulation, no Fak phosphorylation is detectable on Tyr<sup>397</sup> even after long exposures (Figure 4, lanes 1–3). However, the phosphorylation of other tyrosine residues of Fak, including Tyr<sup>576/577</sup>, Tyr<sup>861</sup>, and Tyr<sup>925</sup>, are just detectable in long exposures (Figure 4, lanes 1–3). By the beginning of somitogenesis, the Tyr<sup>397</sup> residue of Fak is increasingly phosphorylated. The phosphorylation of all four tyrosine residues of Fak increases during somitogenesis. This pattern is consistent with our previous observations (Henry *et al.*, 2001) and will be revisited later in rela-



**Figure 4.** Phosphorylation of Fak and abundance of both paxillin and Fak increase at the end of gastrulation. Each lane represents Western blots from extracts of ten embryos probed with either a polyclonal antibody to the C termini of Fak, or affinity-purified polyclonal antibodies to the phosphorylated SH2 binding sites of Fak, or a monoclonal antibody to paxillin as described in MATERIALS AND METHODS. To maximize detection of antigens before the onset of somitogenesis, the exposures were 7 to 10 times that used for the later stages.

tionship to spatial patterns of Fak phosphorylation in the EVL and at somite boundaries.

A qualitative change in the detection of paxillin also occurs at the beginning of somitogenesis, in the appearance of a doublet separated by an apparent molecular mass of  $\sim 5$  kDa. This doublet does not seem to be due to nonspecific proteolysis, because other proteins detected on the same blot show no sign of degradation. We have not been able to assign the change to a phosphorylation difference: a general phosphatase does not consolidate the bands and both bands are phosphorylated. The faster moving band is probably not the related Hic-5 protein, because Hic-5 is significantly smaller than paxillin (50 compared with 68 kDa), and we have evidence by immunocytochemistry that Hic-5 is not expressed during early somitogenesis, whereas it is during late somitogenesis. In addition, we observed that the slow-migrating band is cytosolic and the faster migrating paxillin band occurs in a low-speed membrane pellet (our unpublished data). Possible explanations for this doublet include expression of a second paxillin gene, alternate splicing, post-translational modifications, and paxillin-specific proteolysis that is not blocked by the protease inhibitors in our extraction buffers. Although we have found no evidence for the existence of a second paxillin gene or splice variants in any



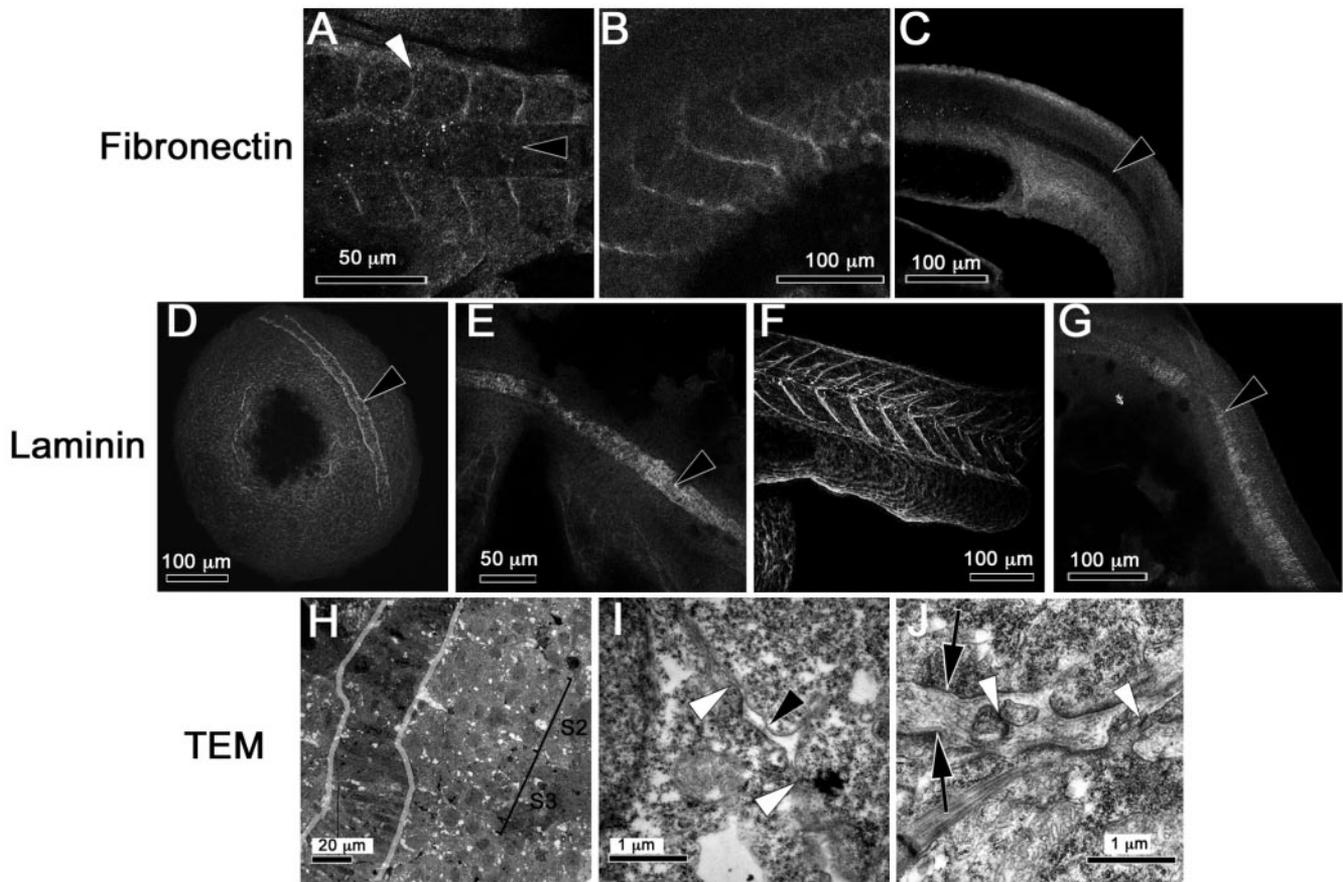
**Figure 5.** Paxillin and Fak are concentrated at somite boundaries and in the notochord, where Fak is phosphorylated at sites of cell-matrix adhesion. The left column of panels shows confocal images of dorsal views of 8- to 10-somite embryos (13.5 hpf, anterior to the left). The middle column shows lateral views of prim-5 embryos (24 hpf, dorsal up) that are focused on the somites. The right column shows lateral views of prim-5 embryos (dorsal up) that are focused on the notochord. The specificities of the antibodies used are indicated at the left. White arrowheads denote somite boundaries and black arrowheads the notochord, except in 5E, where the white arrowhead indicates faint Fak immunoreactivity in the EVL.

of our libraries or in database searches, future completion of the zebrafish genome sequence may resolve this question. Nevertheless, it is clear that a qualitative change is taking place after the onset of somitogenesis. We will designate this period of increased protein synthesis, changes in the patterns of Fak and paxillin synthesis, and increases in the phosphorylation of Fak as the postgastrula mesodermal transition (PGMT).

***Paxillin, Fak, Fibronectin, and Laminin Localize at Boundaries of Somites as They Form***

In zebrafish embryos, somites form by the alignment and polarization of their border cells (Henry *et al.*, 2000). Thus,

we hypothesize that matrix deposition and adhesion at newly formed somite boundaries may be important in the stabilization of somite boundaries. We asked whether paxillin colocalizes with Fak in somites and whether these proteins are found adjacent to the extracellular matrix glycoproteins fibronectin and laminin. Figure 5, A and D, illustrate the abundance of paxillin and Fak in the axial and somitic mesoderm during somitogenesis. Paxillin concentrates at the boundaries of forming somites (Figure 5A, white arrowhead) and in later stages at the boundaries of the chevron-shaped myotomes (Figure 5B). That paxillin protein remains abundant after *paxillin* mRNA has declined suggests that the protein is relatively stable. The pattern of paxillin distribu-



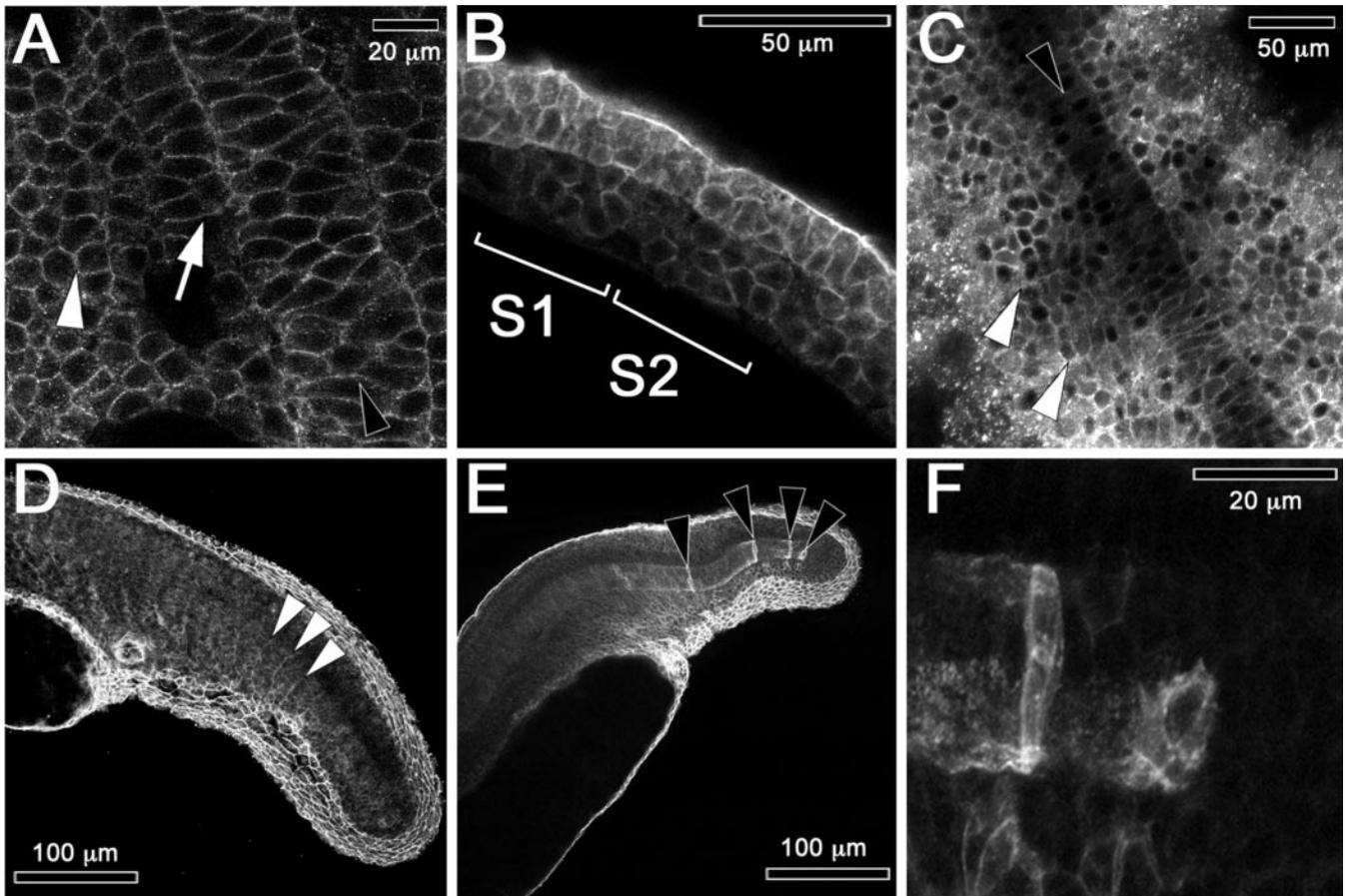
**Figure 6.** Fibronectin is concentrated at somite borders, whereas laminin is concentrated in the forming notochord sheath and at somite borders only after the eight-somite stage. The antigens are given to the left of each row of confocal images. (A) Eight-somite embryo (13 hpf). (B) Prim-5 embryo (24 hpf). (C) Tail of a prim-5 embryo focused on the notochord. (D) Bud-stage embryo (10 hpf), showing staining surrounding the notochord. (E) Eight-somite embryo showing the notochord. (F) Tail of a prim-5 embryo focused on the somites. (G) Prim-5 embryo focused on the notochord. (A, D, and E) Dorsal view, anterior is to the left or left and up. (B, C, F, and G) Lateral view, anterior is to the left. (A–G) Black arrowheads denote the notochord and white arrowheads somite boundaries. (H) Low-magnification TEM of a horizontal section through a three-somite embryo (11 hpf). Two somites are marked (S2 and S3). The notochord is outlined in gray. (I) High-magnification TEM of the section in H focused on a membrane gap with small “island” of matrix forming at the presumptive S2/S3 somite boundary (black arrowhead). Membranes of abutting somite border cells in close contact are marked with white arrowheads. (J) High-magnification TEM of a horizontal section through the somite boundary of a prim-5 embryo (24 hpf) showing rich ECM between somites or black arrows and cellular projections that cross this ECM and form junctions with cells in adjacent somites (white arrowheads).

tion in somites is similar to that described for Fak (Figure 5, D and E; Henry *et al.*, 2001), despite apparent differences in the regulation of these genes at the transcriptional level. All tyrosine-phosphorylated forms of Fak that we are able to detect by immunostaining (pY<sup>397</sup>, pY<sup>576/577</sup>, and pY<sup>861</sup>Fak) concentrate at the somite boundaries during somitogenesis and later stages (Figure 5, G, H, J, K, M, and N). In addition, anti-pY<sup>397</sup>Fak labels the fused myotome cells that extend between somite boundaries (Figure 5H). Fibronectin concentrates at the somite boundaries during somitogenesis (Figure 6A, white arrowhead) and in 24-h embryos (Figure 6B). We did not detect laminin in forming somite boundaries at the eight-somite stage (Figure 6E), but laminin is abundant at the somite boundaries by the 15-somite stage (Parsons *et al.*, 2002) and at 24 hours postfertilization (hpf) (Figure 6F). In TEMs, we saw that by prim-5 (24 hpf) the somites are separated by rich, fibrous ECM (Figure 6J, between the black

arrows), consistent with the presence of both fibronectin and laminin. We also occasionally observe cell processes that cross the somite boundary making adherens and/or gap junctions with the cells of neighboring somites (Figure 6J, white arrowheads). The distributions of fibronectin and laminin are similar to those observed in chick and frog embryos at the somite boundaries (Duband *et al.*, 1987; Krotoski and Bronner-Fraser, 1990) and are consistent with our previous suggestion (Henry *et al.*, 2001) that focal adhesion contacts mediate the stabilization of somite boundaries.

### Ontogeny of Somite Boundaries

Cadherins have been implicated in the morphogenesis of axial mesoderm in mice and frogs (Radice *et al.*, 1997; Delarue *et al.*, 1998) and in the somites of chick embryos (Duband *et al.*, 1987; Linask *et al.*, 1998). In bud-stage fish embryos



**Figure 7.** Cadherin concentrates at plasma membranes in somite and notochord cells. Embryos are labeled with anti-cadherin antibody in these confocal images. (A) Bud-stage embryo (10 hpf, dorsal view, anterior up). Intercalating notochord (black arrowhead), adaxial cells (white arrow), and presomitic mesoderm (white arrowhead). (B) Lateral view of a two-somite embryo (10.6 hpf, anterior to the left, dorsal up). Somites 1 and 2 are labeled as S1 and S2, respectively. (C) Eight-somite embryo (13 hpf, dorsal view, anterior up). White arrowheads denote somite boundaries and a black arrowhead the notochord. (D) Lateral view of the tail of a prim-5 embryo (24 hpf) that is focused on formed somites to the left. White arrowheads mark outlines of recently formed somites. (E) A lateral view of the tail of a prim-5 embryo focused on the notochord and the fin-fold. Black arrowheads mark isolated caudal notochord cells expressing high levels of cadherin. (F) High-magnification view of the embryo in E showing the caudal tip of the notochord.

(Figure 7A) cadherins are concentrated in membranes of the axial mesoderm of the notochord (black arrowhead), and in membranes of the adaxial and paraxial cells of the presomitic mesoderm (white arrow and white arrowhead, respectively). In lateral views of early somitogenesis, we see that cadherins are concentrated in the internal cells of forming somites between the somite boundaries (Figure 7B, see middle of S1 and S2). In dorsal views later in development, cadherins are still present around all cells, but are more concentrated at the somite boundaries (Figure 7, C and D, white arrowheads).

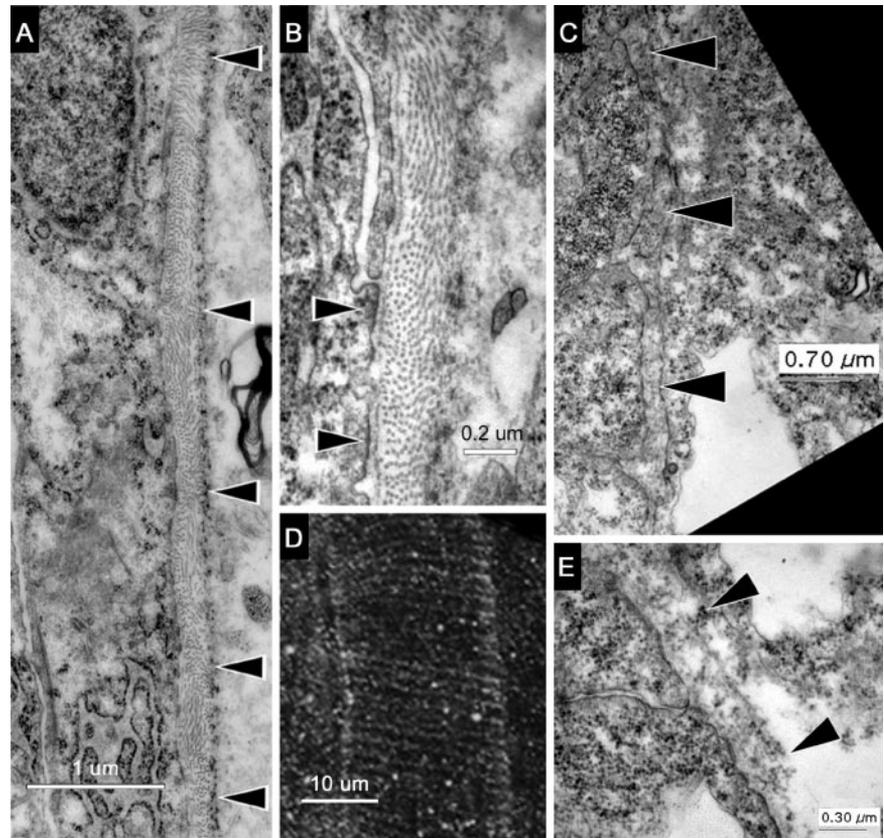
Somite boundaries emerge between mesodermal cells that are in direct cell-cell contact. Wood and Thorogood (1994) suggested the formation of the boundary starts in the lateral somitic mesoderm and then spreads into adaxial somitic mesoderm. In contrast, Henry *et al.* (2000) suggested a random initiation of boundary formation at multiple sites. To test these hypotheses, we examined the ultrastructure of forming somite boundaries of a three-somite embryo where

the location of the forming somite boundaries could be established (Figure 6H, black brackets). At high magnification, we observed, several small “islands” of ECM (Figure 6I, black arrowhead) between stretches of direct cell-cell contact that extend both medially and laterally along nascent somite boundaries (white arrowheads). These small patches of matrix are in stark contrast to a mature somite boundary (Figure 6J), which has rich ECM separating cells of adjacent somites. Our observations suggest that somite boundaries form at multiple sites between adjacent somites.

#### ***Notochord Cells Initially Express Cadherins and Form Circumferential Focal Adhesions after Intercalation***

As previously noted, the earliest adhesion proteins found in the notochord are cadherins that concentrate in the membranes of notochord cells in bud-stage embryos (Figure 7A, black arrowhead). In eight-somite embryos the level of cad-

**Figure 8.** Ultrastructure of the notochord sheath reveals patterns of stress that correlate with rings of phosphorylated Fak. (A) TEM of a horizontal section through the notochord of a prim-5 embryo (24 hpf). Black arrowheads indicate changes in the orientation of fibers in the inner layer. The notochord is to the left of the sheath and somites to the right. (B) High-magnification TEM of this notochord sheath showing sites where the membrane of a notochord cell adheres to the fibrous sheath (black arrowheads). (D) Projection of confocal sections of the notochord of a prim-5 embryo stained with anti-pY<sup>861</sup>Fak shows circumferential striations (anterior up, dorsal to the right). (C) TEM of a horizontal section through the notochord of a three-somite embryo (11 hpf) showing the immature notochord sheath marked with black arrowheads. The notochord cells are to the left of the sheath. (E) High-magnification TEMs of the notochord sheath (black arrowheads) in a three-somite embryo showing only circumferential fiber orientations.



herin detected in the anterior notochord cells (Figure 7C, black arrowhead) becomes reduced compared with posterior regions that are still undergoing intercalation. These observations suggest that cadherins are required primarily during intercalation. By prim-5, high levels of cadherins are detected only in isolated cells in the caudal region of the notochord (Figure 7E, black arrowheads; and F). This residual staining suggests that these cells are still undergoing intercalation.

Paxillin and Fak are expressed abundantly in the cytoplasm of intercalating notochord cells (Figure 5, A, C, D, and F, black arrowheads). Their localization on plasma membranes cannot be verified as distinct from nuclear exclusion of the proteins, which gives the impression of their location in the cortex of the cell. Most obvious, however, is that fibronectin and laminin are absent from the regions where cells are intercalating (Figure 6, A, C, and D, black arrowheads). Instead, laminin is present at the periphery of the notochord (Figure 6, D, E, and G). To assess the interaction of notochord cells with the perichordal sheath, we investigated the pattern of Fak phosphorylation. Figure 5G shows a single confocal section of a 10-somite embryo labeled with an antibody to pY<sup>397</sup>Fak. We can resolve fine circumferential striations where notochord cells are in contact with the perichordal ECM (black arrowhead). Labeling is restricted to the periphery of the notochord. These circumferential striations persist and are shown in a projection of confocal sections comprising the notochord of a prim-5 embryo (24 hpf) stained with anti-pY<sup>397</sup>Fak in Figure 5I (black arrowhead)

and with anti-pY<sup>861</sup>Fak in Figure 5O (black arrowhead). These observations imply a fine structure in the pattern of adhesion to the extracellular matrix that has not been previously described.

To determine what, if any, relationship these circumferential patterns of Fak-immunolabeling have to the ultrastructure of the notochord sheath surrounding the notochord, we used electron microscopy. In the TEMs of prim-5 embryos (Figure 8A), a fibrous sheath denoted by black arrowheads can be seen beside the notochord cells, which are to the left of the sheath. At high magnification, we see what may be focal adhesions where the notochord cell membrane contacts the perichordal sheath (Figure 8B, black arrowheads). The layer of matrix closest to the notochord and these sites of adhesion is laminin-like, based on ultrastructure. The dominant component of the sheath, however, is probably a fibrillar collagen. The orientations of collagen-like fibers occur in alternating groups, which are either perpendicular to the plane of section, and thus, circumferential to the notochord, or else parallel to the plane of section and thus, longitudinal to the notochord. In prim-5 embryos, the distance between groups of circumferential fibers is  $1.56 \pm 0.35 \mu\text{m}$  ( $n = 14$ ) (Figure 8A, black arrowheads), and the distance between circumferential striations of immunostaining for pY<sup>861</sup>Fak is  $1.45 \pm 0.27 \mu\text{m}$  ( $n = 26$ ) (Figure 8D). These are statistically indistinguishable patterns. We hypothesize that the pattern of phosphorylation of Fak represents a spatially periodic pattern of adhesion by notochord cells to the sheath. During notochord elongation, this pattern

of adhesion may generate an uneven mechanical load on the sheath, resulting in the observed alternating pattern of orientation of the collagen-like fibers.

To test this hypothesis, we looked at the sheath of the notochord in three-somite embryos before notochord extension has occurred. The fibers of the matrix surrounding the notochord in the three-somite embryos occur only in circumferential orientation (Figure 8, C and E). We also measured the distance between circumferential striations of pY<sup>397</sup>Fak around the notochord periphery in an embryo that did not yet have a straight body-axis. The distance between striations in the 10-somite embryos in Figure 5G is  $0.63 \pm 0.18 \mu\text{m}$  ( $n = 24$ ), which is significantly smaller than that of the older prim-5 embryo ( $P \ll 0.01$ ) (Figure 8E; see above). Consistent with our hypothesis, the longitudinal collagen fibers indicative of stretching are absent from the notochord sheath in early embryos. Thus, as the notochord elongates, the spacing between striations in Fak phosphorylation expand and the longitudinal collagen fibers occur.

### ***Paxillin, Cadherin, and Noncanonically Phosphorylated Fak Codistribute at the Lateral Membranes of the Enveloping Layer Cells***

The first tissue to develop in the zebrafish embryos is the EVL. The EVL is an epithelial sheet that migrates over the yolk and gastrulating embryo, forming an extraembryonic membrane (Kimmel *et al.*, 1995). The cells of the EVL remain a coherent epithelium that extends and flattens, enveloping the yolk and the rest of the embryo. The mechanism of its migration is not clear (Betchaku and Trinkaus, 1978; Cooper and Kimmel, 1998; Zalik *et al.*, 1999). This epithelium undergoes apoptosis at around prim-15 (30 hpf) shown by TUNEL labeling of apoptotic EVL cell nuclei (Figure 9S).

To better understand the migration of this epithelial EVL, we investigated the distribution and activity of cell adhesion proteins during epiboly of the EVL by using confocal microscopy. Anti-paxillin and anti-cadherin immunolabeling shows concentrations of both proteins at the lateral margins of EVL cells at 50% epiboly (Figure 9, A and D). This codistribution of paxillin and cadherin continues as long as the EVL is present and is shown herein in a six-somite embryo (Figure 9, B and E) and in a prim-5 embryo (Figure 9, C and F). At higher magnification, the paxillin staining (Figure 9C) is often more diffuse than cadherin staining (Figure 9F) and may extend into the cytosol with actin.

The existence of tight and adherens junctions and desmosomes (macula adherens) between cells of the EVL were confirmed by TEM. Tight junctions hold these cells together at their apical-lateral points of contact (Figure 9P, white arrow). Basal to the tight junctions are an adherens junction followed by a row of desmosomes surrounded by dense material that is probably intermediate filaments (Figure 9P, white and black arrowheads). This structure is consistent with tension being applied to these cells as they spread over the embryo.

The surprising observations that paxillin is concentrated at the lateral margins of EVL cells and that some tyrosine residues of Fak, but not Tyr<sup>397</sup>, are phosphorylated during epiboly (Figure 4) led us to investigate the localization of phosphorylated Fak during epiboly and gastrulation. *fak1b*, the dominant isoform of Fak expressed early in develop-

ment (Figure 3, B and C), is not recognized by anti-Fak-C20 and is therefore not shown. However, faint immunoreactivity can be seen in the EVL of prim-5 embryos stained with anti-Fak-C20 (Figure 5E, arrowhead), suggesting that both isoforms of Fak may be expressed in the EVL. The phosphorylated residues of Fak detectable at EVL cell boundaries are pY<sup>576/77</sup>Fak (Figure 9, J–L) and pY<sup>861</sup>Fak (Figure 5, M–O). Phosphorylation on Tyr<sup>397</sup> is not detected at the borders of EVL cells in any stages examined (Figure 9, G–I). Punctate background staining was seen in both cellular and acellular regions of the embryo when using anti-pY<sup>397</sup>Fak antibodies (Figure 9, G–I), but no labeling was ever seen at the lateral membranes of EVL cells. Similar findings are shown in Figure 5 where anti-pY<sup>576</sup>Fak and anti-pY<sup>861</sup>Fak stain the epithelium of EVL cell margins (Figure 5, K, L, and O), but anti-pY<sup>397</sup>Fak does not (Figure 5I). These observations are consistent with the phosphorylation patterns of Fak seen in Western blots before the bud stage (Figure 4). How phosphorylated-Fak functions in the EVL is not yet clear, but it does not seem to depend on stable phosphorylation of Tyr<sup>397</sup>.

Zalik *et al.* (1999) and Betchaku and Trinkaus (1978) have shown that EVL cells contain conspicuous cortical actin belts and that the apical surfaces of the EVL cells are covered with small aggregates of actin coincident with microvilli or ridges. We investigated the relationship between paxillin, cortical belts, and these microvilli. In the EVL at the prim-5 stage (24 hpf), paxillin is abundant (Figure 9R) in areas of actin concentration (Figure 9Q) that is in the cortical belt and microvilli-like extension. One possible interpretation is that paxillin is associated with actin support filaments spreading from adherens or tight junctions.

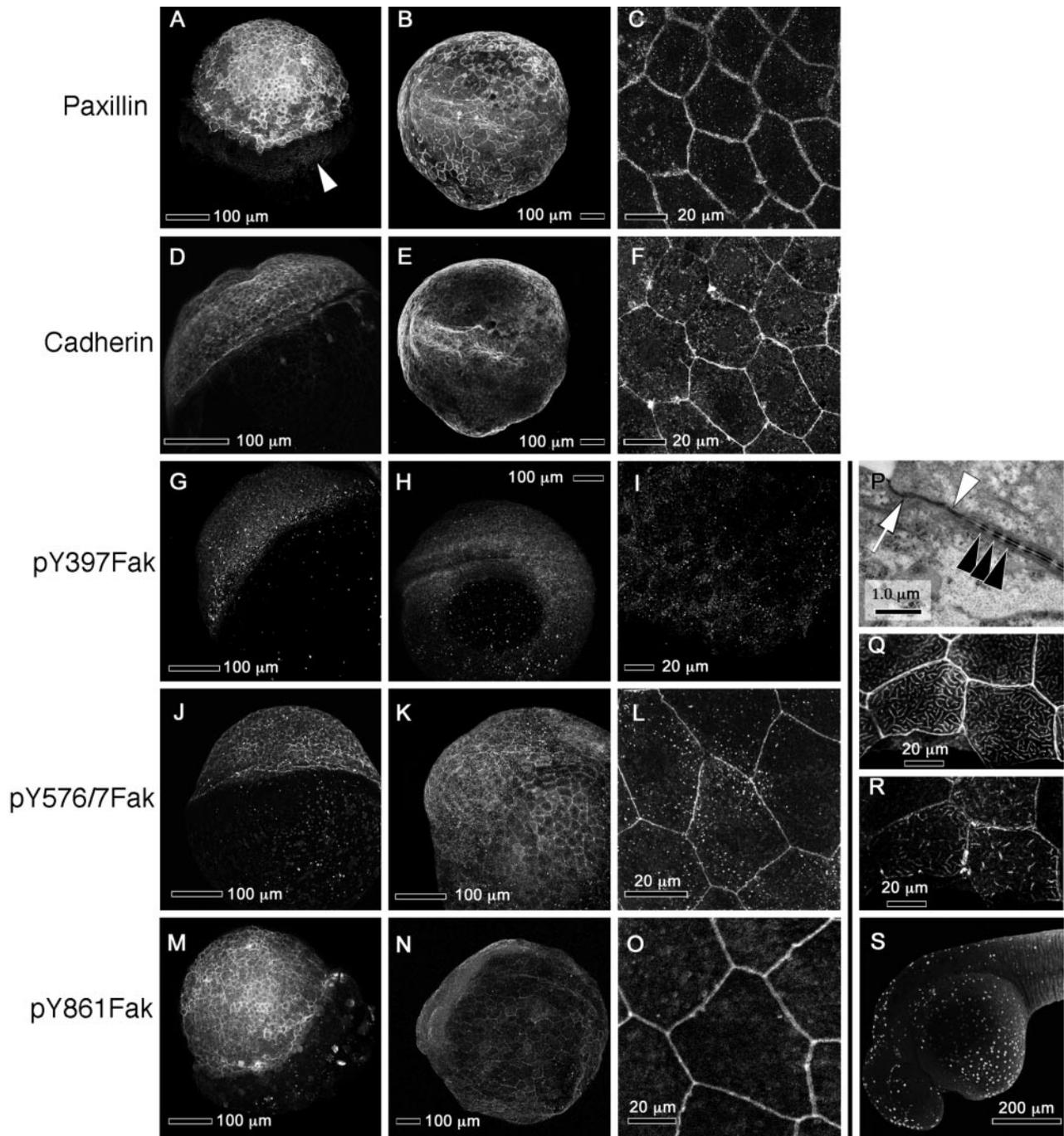
In summary, the EVL migrates as an epithelial layer connected at their lateral margins by tight and cadherin-based adherens junctions. The focal adhesion proteins paxillin and Fak associate near these junctions. The distribution of paxillin, pY<sup>576/577</sup>Fak and pY<sup>861</sup>Fak, cadherin, and actin at the lateral margins of the EVL cells are strikingly similar from 50% epiboly through prim-15, suggesting a cooperation in their functions. The stably phosphorylated tyrosine residues of Fak in the EVL are in the kinase domain, near the Cas-binding site, and in the Map kinase-signaling domain, but not at the autophosphorylation site.

## **DISCUSSION**

Focal adhesion complexes and cadherin-based cell-cell interactions play distinct roles in cell migration and cell adhesion, and therefore in morphogenesis. The regulation of these two modes of cell interaction in embryos is not understood. We have described several tissues where distribution of adherens junctions and/or cadherin proteins overlap with activated focal adhesion proteins and have made some unexpected observations.

### ***Regulation of Focal Adhesion Components***

The transcriptional regulation of the segmentally expressed *fak1a* mRNA was shown to be strongly affected by mutants in Notch signaling (Henry *et al.*, 2001). The expression pattern of the nonsegmentally expressed *paxillin* mRNA is independent of Notch or other signaling that confers segmentation in the paraxial mesoderm (Conlon *et al.*, 1995; Barrantes *et al.*, 1999). Despite these differences in transcrip-



**Figure 9.** Paxillin, cadherin, and noncanonically phosphorylated Fak concentrate at the margins of the EVL cells. (A–O) Each row shows confocal images of immunostaining with an antibody against the indicated target. The first column (A, D, G, J, and M) shows embryos at 50% epiboly (5 hpf). The second column (B, E, H, K, and N) shows embryos at the six-somite stage (13 hpf), and the third column (C, F, I, L, and O) shows high-magnification confocal images of the EVL in prim-5 embryos (24 hpf). (A) White arrowhead shows paxillin on the surface of the equatorial region of the yolk. (B and E) Single embryo double-labeled for paxillin and cadherin. (C and F) High-magnification of a single embryo double-labeled for paxillin and cadherin. (P) TEM of a section through the EVL of a prim-5 embryo showing apical tight junctions (white arrow) and adherens junctions (white arrowhead) and desmosomes (black arrowheads) between lateral membranes of EVL cells. (Q and R) High magnifications of confocal images the EVL of a single prim-5 stage embryo showing phalloidin labeling of actin in Q and paxillin labeling in R. The images are near the apical surface of the cell. (S) TUNEL labeling of EVL nuclei in a prim-15 embryo (30 hpf), showing apoptosis. (A and M) Animal poles are up and tilted toward the viewer. (D, G, and J) Animal poles are toward the upper left and slightly away from the viewer. (B, E, and H) Anterior is to the left and dorsal toward the viewer. (K and N) Dorsal is up. (S) Anterior is to the left and dorsal up. (C, F, I, L, and O) Single confocal sections of the EVL. All other panels are confocal projections of Z-stacks.

tional regulation, paxillin and Fak1a proteins show many similarities in expression pattern at the protein level especially in somitogenesis and notogenesis. Both proteins accumulate at the boundaries of the forming somites where fibronectin is secreted, persist after the formation of the intersomitic borders, and remain through myotome formation. Both proteins are abundant in the notochord and both proteins seem to be participating in the cell-cell adhesion between EVL cells.

### **Cell Adhesion in Somite Boundary Formation**

Focal adhesions do not act alone in forming somites. Cadherin proteins occur early on the membranes of every cell of the presomitic mesoderm. In early zebrafish embryos, we see a reduction in the abundance of cadherins at the interface with neighboring somite border cells, similar to that in chick embryos where there is strong expression of cadherin proteins in the inner mesenchymal cells of forming chick somites and down-regulation near the somite borders (Duband *et al.*, 1987; Radice *et al.*, 1997). The observations in chicks suggested that the formation of cell-cell adhesions drives the formation of groups of somite cells. In other words, the adhesions of the central mesenchymal cells pull together groups of cells to form discrete somites, segmenting the presomitic mesoderm. We have shown previously that, at least in zebrafish, the central mesenchyme cells are not needed for the formation of somite boundaries (Henry *et al.*, 2000). Herein, we also show with TEM images that a matrix is secreted between somite boundaries in patches as the borders form.

Our observations that phosphorylated Fak and paxillin concentrate at somite boundaries where fibronectin is secreted is consistent with integrin-based adhesion to fibronectin at one edge of these polarized cells (Henry *et al.*, 2001; this study). Laminin is not required because mutations in laminin  $\beta 1$  and  $\gamma 1$  do not disrupt somite boundary formation. Rather, laminin mutations disrupt the formation of the notochord, which is surrounded by laminin, and the differentiation of somites at a stage when we see laminin in somites (*grumpy* and *sleepy*; Stemple *et al.*, 1996; Parsons *et al.*, 2002). We see also in the TEM sections that cell-cell contacts extend across somite boundaries in late-stage embryos, which may contain the cadherin proteins we observe in mature somites and may stabilize these boundaries. Thus, the formation of somites is mediated by both cell-cell (cadherin) and cell-matrix (fibronectin/integrin/Fak) junctions.

### **Cell Adhesion in Notogenesis**

The nature of the cell-cell or cell-matrix interactions involved in the intercalation of notochord cells is obscure (Keller *et al.*, 1989; Winklbauer and Keller, 1996; Henry *et al.*, 2001). We found cadherin proteins and little, if any, matrix material between the intercalating cells of the early notochord. Furthermore, we detected phosphorylated Fak only at the interface between notochord cells and the perichordal sheath, not in the middle of the notochord where the cells are extending processes that intercalate between one another. Thus, integrin-based adhesion to the ECM may not provide traction for the intercalating cells. In zebrafish, the *N-cadherin* (*Cdh2*) gene *parachute* clearly mediates convergence of dorsal cells in the midbrain, a process similar to

notochord cell intercalation (Lele *et al.*, 2002). Zebrafish *E-cadherin* (*Cdh1*) mRNA is present in the midline during early development and could be involved in the ventral notochord cell movements (Babb *et al.*, 2001). We and others (Duband *et al.*, 1987; Krotoski and Bronner-Fraser, 1990; Delarue *et al.*, 1998) have shown that cadherins are present surrounding notochord cells, and we suggest that cadherin-based cell-cell adhesion could provide the major traction force required for notochord intercalation. This idea is consistent with results of mosaic expression of a dominant-negative cadherin construct in *Xenopus* embryos, in which axial mesoderm cells expressing the construct fail to intercalate (Delarue *et al.*, 1998).

Notogenesis involves the deposition of an ECM sheath around the presumptive notochord cells and the intercalation of the notochord cells into a single column of disk-shaped cells that contact the ECM at their peripheries, like a stack of pennies in a tube. The notochord cells then develop large internal vacuoles, the inflation of which results in elongation and stiffening of the notochord. One possible mechanism for this elongation is that the mechanical properties of the sheath and the interactions of the cells with the sheath result in a system that resists lateral expansion, while permitting longitudinal expansion (Adams *et al.*, 1990). Our observations with respect to the ultrastructure of the zebrafish notochord sheath and the pattern of Fak phosphorylation within the notochord are consistent with such a model.

Viewed by TEM, the zebrafish notochord sheath has an extensive fibrillar layer, which seems structurally similar to collagens. Zebrafish notochord cells express type II collagen mRNA (Yan *et al.*, 1995; Lele and Krone, 1997), and types I and II collagen are well-characterized components of the perichordal sheath in other vertebrates (Ghanem, 1996; Zhu *et al.*, 2001). The collagen-like fibrillar matrix is bound on the notochord side by laminin, which is required for notogenesis: mutations in laminin  $\beta 1$  and  $\gamma 1$  block differentiation of the notochord, which is rescued by exogenous sources of laminin (Parsons *et al.*, 2002).

We see a change in the orientation of the collagen-like fibrillar matrix that correlates with notochord elongation and with a change in distance between the circular patterns of Fak phosphorylation on the periphery of the notochord. In the early embryo, the collagen-like fibers within the sheath are consistently circumferential, and circumferential striations in phospho-Fak immunolabeling are close together. Later in development, we see alternating patterns of circumferential and longitudinal collagen-like fibers and the circumferential striations in phospho-Fak immunolabeling are further apart. If the notochord cells were adhering to the fibrillar sheath by periodic junctions such as focal adhesion complexes while the notochord was undergoing expansion by vacuolization, these periodic junctions would move apart and the fibers between the junctions would be stretched longitudinally, like a coil-spring being stretched. We observe this pattern in the perichordal sheath.

### **Focal Adhesion Proteins in the Enveloping Epithelial Layer**

Previous investigations have shown that cadherins are abundant at the lateral margins of EVL cells (Zalik *et al.*, 1999) and are important in maintaining the integrity of the

ectoderm during epiboly in *Xenopus* embryos (Levine *et al.*, 1994, Lee and Gumbiner, 1995). Although Betchaku and Trinkaus (1978) have studied the morphology and ultrastructure of the EVL in the teleost, *Fundulus heteroclitis*, apart from the studies of Zalik *et al.* (1999), little is known about the nature of the EVL in zebrafish.

The vegetal spreading of the EVL is an interesting example of a migrating epithelial sheet. Typically, cells of an epithelium share tight junctions and cadherin-based adherens junctions and desmosomes (reviewed in Jamora and Fuchs, 2002). ECM and integrins are ostensibly absent between neighboring cells in an epithelium. This view of the EVL is supported by the close apposition of the membranes and the absence of detectable ECM in TEMs and high magnification confocal micrographs (Figure 9P). However, we find both paxillin and phosphorylated Fak, hallmarks of integrin-mediated cell-ECM adhesion, concentrated along the lateral membranes of these squamous epithelial cells where cadherin is also present. It is provocative that Fak in the EVL is not detectably phosphorylated on Tyr<sup>397</sup>, but is phosphorylated on Tyr<sup>576/577</sup> and Tyr<sup>861</sup>. McLean *et al.* (2000) showed that v-Src phosphorylates residues in FAK in fibroblasts without phosphorylating Tyr<sup>397</sup> and results in a decrease in the fibronectin-stimulated phosphorylation of FAK. Src phosphorylation of FAK may also drive the deregulation of E-cadherin-associated adhesion in colon cancer cells (Avizienyte *et al.*, 2002). However, the noncanonical pattern of FAK phosphorylation has not been documented in the context of a normal tissue nor its biological significance understood. Tsai and Kinsey (2002) showed that the tyrosine kinase Yes is active in zebrafish embryos during epiboly and thus Yes could potentially contribute to the phosphorylation of Fak1b. Another Src-like kinase, Fyn, is activated by fertilization of zebrafish eggs (Rongish and Kinsey, 2000); however, there is no evidence yet that Fyn is active during epiboly. We propose that, in the epithelium of zebrafish EVL, modulation of the actin cytoskeleton at cadherin-based cell-cell adhesions may involve Tyr<sup>397</sup>-independent phosphorylation of Fak by Src-like kinases such as Yes, although we cannot rule out rapid dephosphorylation of Tyr<sup>397</sup>.

The integration of Fak and paxillin into the protein complex on the cytoplasmic face of EVL cells may be mediated through 1) a direct cadherin-paxillin interaction, because E-cadherin has a potential paxillin binding sequence (Brown *et al.*, 1998), or 2) through an indirect interaction with vinculin, which binds paxillin and thus FAK (Tong *et al.*, 1997). Cell-cell junctions in the corneal epithelium and neuroepithelium also bind paxillin (Sugrue and Zieske, 1997; Chenn *et al.*, 1998).

Several potential roles exist for paxillin and Fak in this migrating epithelial sheet. It is unlikely that Fak in these cells is a signal for cell division because EVL cells divide only once after the 10th cleavage cycle of the embryo (Kane *et al.*, 1992). One possibility is that Fak signaling blocks premature apoptosis (Almeida *et al.*, 2000). Fak also has a FERM domain, which would allow it to interact with molecules such as phosphatidylinositol phosphate kinase type1 $\gamma$  that are involved in membrane recycling (Liddington *et al.*, 2003). Thus, Fak1b may be involved in membrane recycling during the expansion of the EVL epithelial layer to 50% epiboly as it envelops the yolk cell, or in remodeling of

the EVL membrane as it moves over the lower equator of the yolk cells. Paxillin may function to stabilize the microvilli on the apical surface of EVL cells, which in turn might provide storage for membrane and cytoskeletal material needed as the EVL expands over the yolk. Our data support roles for paxillin and Fak in the organization of actin around tight junctions or adherens junctions: functions that would help stabilize or remodel cell-cell borders during the migration of this epithelial sheet.

### The PGMT

The shift in activity patterns of focal adhesion proteins observed during development is of great interest. Dramatic changes in the expression as well as qualitative and quantitative changes in phosphorylation of focal adhesion proteins are indicative of a fundamental change in the adhesive milieu of mesoderm cells at the onset of somitogenesis. Hagel *et al.* (2002) note in the mouse embryo, similar to our observations in the zebrafish embryo, that there is expression of paxillin in extraembryonic membranes early in development, and a dramatic increase in paxillin abundance in the mesoderm after gastrulation. Bitzur *et al.* (1994) also noted in the zebrafish a de novo synthesis of one subtype of cadherin at the end of gastrulation, which persists through somitogenesis. We also observed an increased expression of fak1a mRNA during somitogenesis. The molecular triggers and consequences of this PGMT as well as the interactions between cadherin-based and focal adhesion-based cell adhesion probably play central roles in the morphogenesis of vertebrate embryos. We have only begun to scratch the surface of our understanding of these processes.

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# Activity and Distribution of Paxillin, Focal Adhesion Kinase, and Cadherin Indicate Cooperative Roles during Zebrafish Morphogenesis

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