

Actin Localization and Function in Higher Plants

Review Article

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1. Introduction

Eukaryotes from all four kingdoms contain cytoskeletal actin and associated proteins. Functions for actin are well-defined in animals and include cell contraction, intracellular movement, cell morphology determination, cell locomotion, and cytokinesis (ALLEN and ALLEN 1978, SCHLIWA, 1986). In addition, it is now well-established that actin-myosin interaction provides the motive force for cytoplasmic streaming in some lower plants and fungi (for reviews, see ALLEN and ALLEN, 1978, KAMIYA 1981, WILLIAMSON 1980). Evidence for the existence of vascular plant actin and myosin has been accumulating in recent years (reviewed by JACKSON, 1982), although their role in the cell has not been unequivocally defined. This review summarizes current knowledge of actin and associated proteins in plants, considers higher plant actomyosin function, and suggests areas deserving future attention.

2. Lower Plant Actomyosin

Although the remainder of this overview will focus on the occurrence of actin in higher plant cells, a brief look at Characean algae is useful. More is known about actin and its role in cytoplasmic motility in *Characeae* than for any other plants. Light microscopic examination of the giant internode cells of these algae reveals rapid endoplasmic streaming that requires the integrity of cortical filamentous cables (NAGAI and REBHUN, 1966, NAGAI and HAYAMA, 1979 b). Ultrastructurally, these cables consist of 50–100 microfilaments with a

5–7 nm diameter, a double helical structure, and a 36–38 nm repeat (NAGAI and REBHUN, 1966, NAGAI and HAYAMA 1979 a, b, PALEVITZ and HEPLER, 1975). Characean microfilaments reversibly bind mammalian heavy meromyosin (HMM) or S-1 myosin fragments, resulting in arrowhead patterns. All microfilaments in a bundle show uniform polarity, with arrowheads pointing opposite the direction of cytoplasmic flow (KERSEY *et al.* 1976, KERSEY and WESSELS 1976, PALEVITZ and HEPLER 1975, WILLIAMSON 1974). Fluorescence microscopy confirms the presence of actin microfilaments at the interface between stationary ectoplasm and mobile endoplasm which, as suggested by numerous earlier studies (reviewed in KAMIYA 1981), is the site of motive force generation (NOTHNAGEL *et al.* 1981, WILLIAMSON and TOH, 1979, WILLIAMSON *et al.* 1986).

Proteolytic activity in cytoplasmic extracts of plant cells, particularly those possessing large vacuoles, constitutes a problem for biochemical approaches to the study of plant cytoskeletal proteins. It is perhaps for this reason that knowledge about the biochemical properties of plant cytoskeletal proteins is still scarce, at least when compared to animal cells. Cytoplasmic extracts from the Characean alga, *Nitella*, contain an actin-like protein with electrophoretic mobility, solubility properties, and ultrastructure comparable to mammalian actins (ALLEN and ALLEN 1978). A 55.000 MW component copurifies with the 46.000 MW actin from these extracts and may be involved in microfilament bundling (ALLEN and ALLEN 1978, NAGAI and HAYAMA 1979 a, b). Early attempts to identify algal myosin from fractionated extracts elucidated no

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200,000 MW protein in the supernatant, but a Mg^{++} -ATPase in the particulate fraction and a 260,000 MW minor band were noted (ALLEN and ALLEN 1978). Using different extraction procedures, KATO and TONOMURA (1977) described a putative *Nitella* myosin heavy chain of >200 kD MW, which formed bipolar filaments of 0.2 μ m length at low ionic strength: they also observed a Ca^{++} sensitive Mg^{++} -ATPase that was activated by skeletal muscle F-actin. However, these are preliminary and as yet unconfirmed observations, and should be viewed with caution. Endoplasmic organelles frequently show irregularly shaped rod- or horn-like protuberances, and 20–30 nm globular bodies often attached to fine filaments of <4 nm diameter. These structures are believed to be membrane-bound myosin, but direct proof is lacking (NAGAI and HAYAMA 1979 a, b).

The value of Characean algae for the study of actin-dependent processes in plants lies in the fact that reactivatable *in vitro* systems can be obtained with relative ease, and that both they and the intact cells themselves are amenable to a wide variety of experimental manipulations. These include electrical and mechanical stimulation of cells (KAMITSUBO 1980), vacuolar perfusion (ALLEN and ALLEN 1978, WILLIAMSON and TOH 1979, WILLIAMSON 1980), compression, centrifugation, ligation, and mechanical damage (KAMITSUBO 1980, NAGAI and KAMIYA 1977, CHEN and KAMIYA 1975) light microbeaming (KAMITSUBO 1980), and chemical stimulation and inhibition (ALLEN and ALLEN 1978, CHEN and KAMIYA 1975, NAGAI and KAMIYA 1977, NOTHNAGEL *et al.* 1981, WILLIAMSON and TOH 1979). Cell models and *in vitro* cytoplasmic streaming systems were developed using demembrated endoplasmic droplets which continue to show chloroplast rotation and other motility phenomena (HIGASHI-FUJIME 1980). Molecular analysis of *in vitro* motility led to the discovery that the actin cables of mechanically dissected algal cells are capable of supporting the movement of organelles (WILLIAMSON and TOH, 1979, SHIMMEN and TAZAWA 1982), myosin-coated beads (SCHEETZ and SPUDICH 1983, SHIMMEN and YANO 1984), and membranous organelles of *Acanthamoeba* by myosin-I (ADAMS and POLLARD 1986). These studies have provided a current model of Characean cytoplasmic streaming that envisions the motive force for rotational streaming to be generated at the interface between the moving endoplasm and the stationary layer of actin cables attached to cortical chloroplasts. Myosin associated with endoplasmic organelles or soluble myosin is proposed to create a shear force that drives the endoplasm around

the cell in a direction determined by actin microfilament orientation. However, the actual distribution of myosin in Characean cells is not known, and the molecular details of this model still need to be elucidated.

3. Higher Plant Actomyosin

3.1. Methods of Higher Plant Actin Identification

Studies on Characean algae set the precedent that allowed researchers to identify actin in vascular plants. The methods used include electron, light, and fluorescence microscopy, inhibitor studies, biochemical characterization, and gene cloning and sequencing.

Early light microscopic studies correlated 0.1–0.2 μ m cytoplasmic fibers with active streaming in the transvacuolar strands of higher plants (KAMIYA 1959, O'BRIAN and THIMANN 1966, PARTHASARATHY and MUHLETHALER 1972). In other cells, cytoplasmic fibrils were frequently found to be oriented parallel to the main axis and to the direction of flow in elongated cells (FRANKE *et al.* 1972, O'BRIAN and McCULLY 1970). Cinematography (MASCARENHAS and LAFOUNTAIN 1972), videotaping (SEAGULL and HEATH 1980 b, PARTHASARATHY 1985), differential interference contrast microscopy (O'BRIAN and McCULLY 1970, PESACRETA and PARTHASARATHY 1984), and computer image enhancement (PARTHASARATHY 1985) have since improved the visualization of fibrils in regions of active streaming and have expanded our knowledge of their disposition. By analogy with the situation in Characean algae (NAGAI and REBHUN 1966), it was suspected that these fibers were composed, at least in part, of actin filaments. However, confirmation of this supposition has met with difficulty.

The earliest evidence for the existence of actin in higher plant cells came from extruded *Amaryllis* cytoplasm which contained 6 nm filaments that specifically bound HMM with a 35 nm periodicity and released it in the presence of ATP (CONDELLIS 1974). Similar results were obtained with glycerinated preparations from *Haemanthus* endosperm cells (FORER and JACKSON 1975, 1976). Fixation and localization of actin microfilaments *in situ* by conventional electron microscopic techniques has proven cumbersome. Osmium tetroxide treatment, routinely used in sample preparation, destroys actin microfilaments under many conditions (MAUPIN-SZAMIER and POLLARD 1978, SMALL 1981, SEAGULL and HEATH 1979). To increase plant microfilament preservation by preferentially stabilizing actin filaments, several approaches have been used, with varying success: HMM binding (CONDELLIS 1974,

Table 1. *Actin in higher plants*

Plant	Tissue	Technique	Function	References
Gymnosperms:				
<i>Chamaecyparis</i>	roots	LM/EM/FM	CS	70, 71
<i>Pseudotsuga</i>	roots	LM/FM	CS	70
Angiosperms:				
Monocotyledons:				
<i>Allium</i>	bulb, scale epidermis	EM/FM	CS/SP	67
	cotyledon GMCs	LM/INH	CS/OM/CK	25, 60, 61, 63, 64
	root tips	FM	CS/OM/CK	6, 65
<i>Alstroenemia</i>	pollen tubes	EM/FM	CS/SP	67, 69
<i>Amaryllis</i>	pollen tubes, protoplasts	LM/EM	CS	7
<i>Avena</i>	coleoptile	LM/INH/EM/FM	CS/SP	3, 4, 57, 67, 68, 96, 103
<i>Chamaedorea</i>	stem and root:			
	procambium and vascular tissue	EM	CS	66
<i>Clivia</i>	endosperm	FM	SP/KK/CK	80, 81, 82, 83
	pollen tubes	EM/INH	CS/TG	18
<i>Cocos</i>	stem and root:			
	procambium and vascular tissue	EM	CS	66
<i>Egeria</i>	leaf epidermis	EM/FM	CS (2°)/SP	67
	leaves	LM/FM/INH/EM	CS (2°)/OM	110
<i>Elodea</i>	leaf epidermis	EM/FM	CS (2°)/SP	67
	leaves	EM/FM/LM/INH	CS (2°)/OM	11, 110
<i>Endymion</i>	pollen	FM	CS	28
<i>Haemanthus</i>	endosperm	EM/FM/LM/INH	CS/SP/KK/CK	13, 14, 16, 17, 80, 81, 82, 83
<i>Hordeum</i>	coleoptile epidermis	LM/EM/FM/INH	CS/SP/Gravi	67, 68, 104
	pollen	FM	CS	28
<i>Hydrilla</i>	leaf epidermis	EM/FM	CS (2°)	67
	leaves	EM/FM/LM/INH	CS (2°)/OM	110
<i>Lilium</i>	pollen grains and tubes	LM/EM/FM/INH	CS/SP/TG	18, 27, 67, 69, 74, 76
<i>Limnobiium</i>	root hairs	EM/FM	CS/SP	67
<i>Narcissus</i>	pollen	FM	CS	28
<i>Phleum</i>	root tips	EM	—	98
<i>Prestoea</i>	root: procambium and vasc. tissue	EM	CS	66
<i>Tradescantia</i>	pollen grains and tubes	LM/FM/INH/EM	CS/TG	43, 46, 74
	stamen hairs	LM/EM/FM/INH	CS/SP	10, 43, 67, 98
	male meiocytes	EM	—	26
<i>Triticum</i>	coleoptile epidermis	EM/FM	CS/SP	67
	wheat germ	BC/EM	—	32
	leaf trichomes	EM	—	98
<i>Vallisneria</i>	leaf epidermis	LM/EM/INH	CS (2°)	112
	leaf mesophyll	LM/INH	CS (2°)/OM	33, 95
<i>Zea</i>	genomic library	Mol. Biol.	—	89
	coleoptiles	LM/EM/FM/INH	CS/Gravi	4, 77, 104
Dicotyledons:				
<i>Cucumis</i>	fruit thin-walled parenchyma	EM/FM	SP	67
<i>Cucurbita</i>	fruit thin-walled parenchyma	EM/FM	CS/SP	67
	stem: 1' & 2' vasc. tissue, cambium	EM	CS	66
<i>Fraxinus</i>	stem: active cambium	EM	CS	19
<i>Glycine</i>	seedlings	BC/EM	—	49, 50
	hypocotyls	Mol. Biol.	—	30, 31, 55, 88, 89
<i>Helloborus</i>	pollen	FM	CS	28
<i>Heracleum</i>	petiolar hairs	LM	CS	58
<i>Hibiscus</i>	callus protoplasts	LZR/INH/LM	CS/SP	20
<i>Impatiens</i>	pollen tubes	EM/FM	CS/SP	67, 69
<i>Lepidium</i>	root hairs, root tips	LM/INH/EM/FM	CS/TG/OM/Gravi	22, 23, 27, 102
<i>Lycopersicon</i>	fruit parenchyma and endocarp	BC/EM	CS	99, 100
	stem, petiolar hairs, fruit thin-walled parenchyma	EM/FM	CS/SP	67

Table 1 (continued)

Plant	Tissue	Technique	Function	References
<i>Medicago</i>	suspension culture cells	FM	CW/CS	87
<i>Nicotiana</i>	pith callus	EM/FM	CS/SP	67
	pollen	FM	TG	74
	mesophyll callus protoplasts	LZR/INH/LM	CS/SP	20
	stem: 1' & 2' vasc. tissue, cambium	EM	CS	66
<i>Petunia</i>	pollen grains and tubes	EM/FM	SC/SP/TG	67, 69, 74
<i>Phaseolus</i>	root tips	BC/EM	—	34
<i>Pisum</i>	internode cells	EM/LM/INH	CS	4
<i>Raphanus</i>	root hairs	LM/EM/INH	CS/TG/CW	28, 84, 85, 86
<i>Salix</i>	stem: cambium, 2' vascular tissue	EM	CS	66
<i>Trifolium</i>	seedlings	BC/EM	—	50
KEY:				
Techniques:		Function:		
EM	Electron microscopy	CS	Cytoplasmic streaming	
LM	Light microscopy	SP	Cell shape determination	
FM	Fluorescence microscopy	TG	Tip growth	
INH	Inhibitor studies	OM	Organelle movement	
BC	Biochemistry	KK	Karyokinesis	
Mol. Biol.	Gene cloning and sequencing	CK	Cytokinesis	
LZR	Laser ablation	Gravi	Graviperception	
		CW	Cell wall deposition	

FORER and JACKSON 1976, 1979 a), tannic acid treatment (SEAGULL and HEATH 1979), tissue glycerination (FORER and JACKSON 1979 a), and freeze-substitution techniques (HEPLER and LANCELLE 1986, LANCELLE *et al.* 1986, TIWARI *et al.* 1984). Electron microscopy using these techniques reveals 5–7 nm microfilaments with the morphological characteristics of F-actin (*e.g.* double-helices with a 34–40 nm periodicity) (SEAGULL and HEATH 1979). Mammalian HMM or S-1 fragments will reversibly decorate these microfilaments (SEAGULL and HEATH 1979, YAMAGUCHI and NAGAI 1981).

Recently, fluorescence microscopy, although not having the resolution of electron microscopy, has become increasingly popular for visualizing three-dimensional cytoskeletal arrays in plant cells. Fluorochrome-tagged phallotoxins [fungal metabolites known to bind evolutionarily diverse F-actins (WULF *et al.* 1979, PARTHASARATHY 1985, PESACRETA *et al.* 1982)], and indirect immunofluorescence with antibodies against actin have added to our knowledge of actin arrays in plant tissues (see Table 1). The distribution patterns revealed with these techniques will be discussed in more detail in the next section.

Despite the progress that has been made in recent years, one needs to ask how reliable are these visualization procedures, and where are their limitations. For example, HMM decoration, although capable of un-

equivocally identifying actin filaments and revealing their subcellular distribution, nevertheless requires a preparatory step, such as glycerination or permeabilization, that may cause rearrangements of actin filaments, and may even induce the polymerization of previously unpolymerized actin. Glycerination also permeabilizes the tonoplast and hence can lead to proteolysis and degradation from the release of vacuolar components. Related problems arise with the use of antibodies or fluorescent phallotoxins for light microscopic visualization of F-actin. Therefore, fixation/extraction protocols required for fluorescent visualization may cause partial destruction, rearrangements, selective extraction, or preferential stabilization of certain cellular components. One has to ask, for example, whether it is only the most stable actin that is visualized, or the actin that is associated with other, more resistant, cellular structures. It is disturbing that, with a few notable exceptions, staining patterns obtained with fluorescent phalloidin are rather diffuse, although one would expect a distinct fibrillar staining based on what is known about phalloidin's interaction with F-actin. In many cases, the techniques used, and the results obtained with fluorescent molecular probes are taken at face-value and are often not evaluated critically enough.

Actin-like proteins have been biochemically identified

and partially characterized in *Phaseolus* root tips (JACKSON and DOYLE 1977), wheat germ (ILKER *et al.* 1979), *Glycine* seedlings (METCALF *et al.* 1980, 1984), and *Lycopersicon* endocarp (VAHEY and SCORDILIS 1980, VAHEY *et al.* 1982). Based on electrophoretic mobility, solubility and immunological and electron microscopic criteria, these proteins show homology to cytoplasmic and muscle actins. All four of the aforementioned plant tissues contain a protein of about 45 kD that comigrates with mammalian actins, while *Phaseolus* (JACKSON and DOYLE 1977) and *Triticum* (ILKER *et al.* 1979) also contain a slower migrating polypeptide. A 58.000 MW protein from *Triticum* copurifies with the 45 kD actin and has properties similar to, and comigrates with, a sea urchin actin binding protein (ILKER *et al.* 1979). Immunologically, these 45 and 58 kD proteins crossreact in Ouchterlony immunodiffusion assays with antibodies against turkey gizzard actin (ILKER *et al.* 1979). Actins from crude cytoplasmic extracts of *Glycine* and *Trifolium*, or from partially purified *Lycopersicon* extract, form 6–7 nm wide filaments at >50 mM ionic strength and reversibly bind HMM or rabbit S-1 fragments in an arrow-head pattern with a periodicity of 28–37 nm (METCALF *et al.* 1984, VAHEY and SCORDILIS 1980, VAHEY *et al.* 1982). *Lycopersicon* actin, which constitutes approximately 6% of extracted proteins, is capable of activating skeletal or tomato myosin Mg^{++} -ATPase (VAHEY *et al.* 1982).

More recently, molecular cloning techniques have aided in the study of the plant cytoskeleton in general, and of plant actin in particular. Actin genes have been cloned from *Glycine max* and *Zea mays* using a *Dicystostelium* actin sequence as a probe (NAGAO *et al.* 1981, SHAH *et al.* 1982, 1983). Soybean contains six or more genes in a multigene family of actin-related sequences (HIGHTOWER and MEAGHER 1985), while maize has at least two actin genes (SHAH *et al.* 1983). Sequences from one maize and three soybean actin genes suggest plant actin genes evolved from a common ancestral gene, with a unique sequence of the N-terminal 9 amino acids and three introns (SHAH *et al.* 1982, 1983, HIGHTOWER and MEAGHER 1986). The deduced amino acid sequence of plant actins shows homology to animal cytoplasmic and muscle actins, yet there is extensive coding sequence heterogeneity among soybean actin genes and between soybean and maize genes (SHAH *et al.* 1983, HIGHTOWER and MEAGHER 1986). The three soybean actin gene sequences differ by 6–9% replacement substitutions (nucleotide changes resulting in amino acid substitutions). Maize and soybean have 8–10% replacement substitutions, while plant actin genes have

diverged 11–15% from their animal counterparts. On the other hand, typical animal-animal replacement substitutions only range from 3–7% (SHAH *et al.* 1983, HIGHTOWER and MEAGHER 1986, VANDEKERCKHOVE and WEBER 1978). The six genes of the soybean actin multigene family fall into three pairs of sequence homology by northern and dot-blot analyses, with one pair highly expressed in seedlings and mature plant organs, constituting >80% of all actin mRNAs (HIGHTOWER and MEAGHER 1985, 1986). It is not known whether actin genes are differentially expressed in plant development or whether different isoforms perform unique cellular functions.

3.2. Actin Microfilament Arrays

Plant microfilaments, like microtubules, are organized into distinct arrays that change in distribution during the cell cycle. Several types of interphase microfilament arrays have been reported. Most common are the large cytoplasmic microfilament bundles oriented parallel to the long axis of the cell or the direction of streaming, as in transvacuolar cytoplasmic strands (CLAYTON and LLOYD 1985, PARTHASARATHY and MUHLETALER 1972, PARTHASARATHY *et al.* 1985, PERDUE and PARTHASARATHY 1985, SEAGULL and HEATH 1979, 1980 b, SEAGULL *et al.* 1986, TAKAGI and NAGAI 1983, WHITE and SACK 1986). In addition, other cell types, such as *Haemanthus* and *Clivia* endosperm cells, exhibit a loose cytoplasmic meshwork of microfilaments that intermingles with the interphase microtubule array (SCHMIT and LAMBERT 1985 a, b, 1986 b, SCHMIT *et al.* 1986 a). Recently, a perinuclear microfilament “basket” was described in alfalfa suspension culture cells (SEAGULL *et al.* 1986), *Equisetum* root tip cells (DERKSEN *et al.* 1986), and *Hordeum* and *Zea* coleoptile cells (WHITE and SACK 1986). In these same cell types, as well as in *Raphanus* root hairs and other cells, fine cortical arrays of microfilaments were observed (FRANKE *et al.* 1972, PARTHASARATHY 1985, SEAGULL and HEATH 1979, 1980 b, TAKAGI and NAGAI 1983, WHITE and SACK 1986, YAMAGUCHI and NAGAI 1981, DERKSEN *et al.* 1986). These cortical microfilament arrays are mostly oriented parallel to the cell’s long axis and lie in close apposition to the plasmalemma. Individual microfilaments are sometimes observed in close proximity to, or even parallel to cortical microtubules which are believed to be important in cell wall deposition (SEAGULL and HEATH 1979, 1980 b). Finally, pollen grain of a number of angiosperms contain numerous fusiform, phalloidin-binding bodies ($\sim 2.5 \times 15 \mu\text{m}$) that possibly constitute an actin storage form. These bodies are dispersed into fine fibrils localized near the germination

Table 2. Actin associated proteins in higher plants

Plant	Tissue	Protein	Technique	Function	References
<i>Allium</i>	root tips	myosin	FM/BC	CK	65
		troponin T	FM	—	44
<i>Egeria</i>	leaves	myosin	BC/EM	CS	59
<i>Lycopersicon</i>	endocarp	myosin	BC/EM	CS	99, 100

aperture and in the pollen tube during pollen activation (HESLOP-HARRISON *et al.* 1986).

The distribution of microfilaments during mitosis has long been a controversial topic, and although several microfilament configurations have been described, their functional relevance in mitosis is disputed (for review, see FORER 1978, McINTOSH 1979, PETZELT 1979, PICKETT-HEAPS *et al.* 1982). During prophase in *Haemanthus* and *Clivia* endosperm cells, the cytoplasmic microfilament array gradually redistributes to the cortex (SCHMIT and LAMBERT 1985 a, b, 1986 b, SCHMIT *et al.*, 1986 a). As the mitotic spindle forms in prometaphase and elongates in anaphase, it tends to occupy the cell volume, forcing a rearrangement of the cortical actin network which forms an envelope around the spindle, being stretched until many microfilaments eventually lie parallel to spindle microtubules (SCHMIT and LAMBERT 1985 a, b, 1986 b, SCHMIT *et al.* 1986 a). These findings may help explain early observations on glycerinated *Haemanthus* endosperm mitotic cells which reportedly contain microfilaments between the chromosomes and poles. Most of these microfilaments are oriented parallel to the microtubules and those associated with kinetochore fibers show common polarity (FORER and JACKSON 1975, 1976, 1979 a, b). A localization of microfilaments near spindle fibers has also been observed by fluorescence microscopy in mitotic spindles from alfalfa suspension culture cells (SEAGULL *et al.* 1986), but not in nonglycerinated *Haemanthus* spindles (SCHMIT and LAMBERT 1985 a) or in *Allium* and *Equisetum* spindles (CLAYTON and LLOYD 1985, DERKSEN *et al.*, 1986). Late in karyokinesis increased numbers of microfilaments are observed by fluorescence microscopy in the equatorial division plane. This is the earliest sign of a cytokinetic apparatus (CLAYTON and LLOYD 1985, SCHMIT and LAMBERT 1985 a, b, 1986 b, SCHMIT *et al.* 1986 a).

Cytokinesis in higher plant cells is effected by an organelle known as the phragmoplast which is composed of parallel arrays of microtubules. These presumably guide vesicles to the division plane where they fuse to form a cell plate separating the daughter cells. Fluorescence

microscopy indicates the presence of a diffuse actin network codistributing with phragmoplast microtubules in *Allium* root cells (CLAYTON and LLOYD 1985, PARKE *et al.* 1986), *Equisetum* root tip cells (DERKSEN *et al.* 1986), and alfalfa suspension culture cells (SEAGULL *et al.* 1986). A more distinct microfilament pattern has been observed in phragmoplasts of *Haemanthus* and *Clivia* endosperm cells. In these cells the actin has a definite fibrillar appearance, closely associated with, and parallel to, microtubules (SCHMIT and LAMBERT 1985 a, b, 1986 b, SCHMIT *et al.* 1986 a). When the nuclear envelope reassembles in telophase, the interphase microfilament arrays also reform (SEAGULL *et al.* 1986).

Of the four major microtubule structures found in higher plant cells (cortical arrays, preprophase band, mitotic spindle, and phragmoplast), only the preprophase band so far has not been found to have associated microfilaments (CLAYTON and LLOYD 1985, PARKE *et al.* 1986).

3.3. Actin-associated Proteins

Whereas the presence of actin in higher plant cells is now well-established, much less is known about the disposition of actin-associated proteins involved in force generation, structural interactions, and actin filament dynamics. Little information is available in part due to the difficulty of applying to plant cells what are routine biochemical approaches for animal cells. Myosin-like proteins have been reported in *Allium* root tips (PARKE *et al.* 1986), *Egeria* leaf extracts (OHSUKA and INOUE 1979), and *Lycopersicon* (tomato) endocarp (VAHEY and SCORDILIS 1980, VAHEY *et al.* 1982) (see Table 2). In some respects these myosins show a similarity to skeletal muscle myosin. High ionic strength extracts of *Egeria* and *Lycopersicon* contain a K^+ -EDTA ATPase activity, which is inhibited by Mg^{++} in tomato (VAHEY and SCORDILIS 1980, VAHEY *et al.* 1982) but not *Egeria* (OHSUKA and INOUE 1979). *Egeria* myosin also shows no p-nitrophenyl phosphatase activity (OHSUKA and INOUE 1979). At low ionic strength

and in the presence of Ca^{++} , tomato myosin contains a Mg^{++} -ATPase that is activated 19-fold by tomato F-actin and 10-fold by rabbit skeletal muscle actin (VAHEY *et al.* 1982). Both of these myosins can be induced to form 0.5–1.0 μm bipolar filaments in a low ionic strength medium (OHSUKA and INOUE 1979, VAHEY and SCORDILIS 1980). Also, *Egeria* myosin can reversibly bind to muscle F-actin with the 40 nm periodicity of projections characteristic of animal myosins (OHSUKA and INOUE 1979). *Allium* root tips contain a protein of 200,000 MW that reacts in Western blots with an antibody against 3 T 3 cell myosin (PARKE *et al.* 1986). This protein comigrates on gels with mammalian myosin heavy chains (PARKE *et al.* 1986) and is similar in size to the ~ 200 kD heavy chain of *Egeria* (OHSUKA and INOUE 1979). Tomato myosin weighs only 100 kD and is believed to be a proteolytic fragment of the actual heavy chain; the authors find no polypeptides >100 kD in polyacrylamide gels (VAHEY *et al.* 1982). Tomato myosin light chains are less well characterized but two, with molecular weights of 16 and 14 kD, have been reported (VAHEY *et al.* 1982). There are no published reports on myosin genes or mRNA's in higher plants. Likewise, ultrastructural studies of myosin are few. Using an antibody against animal myosin, indirect immunofluorescent localization of myosin in *Allium* root tip cells has been achieved (PARKE *et al.* 1986). In interphase, diffuse cytoplasmic staining and aggregates of label were observed, none of which co-localized with actin microfilament arrays. No labeling of preprophase bands or spindles was noted, but a diffuse cytoplasmic staining prevailed in these stages. the cytokinetic phragmoplast is the only structure in which the myosin antibody shows costaining with rhodamine phalloidin-labeled F-actin. Besides myosin, the only higher plant actin-associated protein on which there is some information is a troponin T-like protein, identified in *Allium* root tip cells by immunofluorescent staining with a monoclonal antibody against striated muscle troponin T (LIM *et al.* 1986). Again, the staining pattern does not strictly correspond to actin arrays. This epitope, resembling troponin T, was found associated with interphase cortical microtubules, preprophase band microtubules, mitotic spindle fibers and pole complexes, and with phragmoplast microtubules. Much work is still to be done to identify other actin associated proteins in plants, as there is copious evidence for their occurrence in other kingdoms (for reviews, see KORN 1978, SCHLIWA 1986, STOSSEL *et al.* 1985).

3.4. Actomyosin Function

An involvement of actin in certain plant cellular processes is based on two lines of evidence. The first is localization in cell domains concerned with the execution of motile processes, such as mitotic spindles, phragmoplasts, cell cortex, etc. . . . However, conclusions based solely on this principle of "guilt by association" need to be viewed with caution. The second line of evidence entails the use of pharmacological agents thought to specifically disrupt actin-based processes. Of these inhibitors, the cytochalasins are the most widely used microfilament toxins. These fungal metabolites are thought to disrupt F-actin (MACLEAN-FLETCHER and POLLARD 1980, SCHLIWA 1982), but in a number of systems, especially plants, no such destruction occurs even at concentrations of 1–100 $\mu\text{g}/\text{ml}$ (FORER *et al.* 1972, THOMAS 1978). In fact, cytochalasins are reported to have variable and complex effects on plant cells and it is very difficult to judge whether this is due to differential effects on specific microfilament populations, or nonspecific disruption of other structures and processes (SEAGULL and HEATH 1980 b, WESSELS *et al.* 1971, POPE *et al.* 1979). Phallotoxins are another set of fungal products which reportedly interfere with actin microfilament function by binding to, and stabilizing, F-actin (WIELAND 1977). In many instances the concentrations of these compounds required for disruption of cellular functions are so high that one has to worry about the specificity of the effect observed. On the other hand, it is plausible that the effective intracellular concentrations are much lower since the cell wall may act as a permeability barrier. However, unless these uncertainties are resolved, studies based on inhibitor experiments alone are often difficult to interpret.

Table 1 lists reports of actin occurrence in higher vascular plant tissues, the techniques used for identification, and some proposed microfilament functions. Cytoplasmic streaming occurs in all of these tissues with only a few exceptions (see PARTHASARATHY *et al.* 1985). By extrapolation from the well-documented involvement of actomyosin in Characean algal motility and other lines of indirect evidence, *e.g.*, presence of myosin, orientation of microfilaments parallel to the direction of streaming and its reversible disruption by cytochalasins and phalloidin (BRADLEY 1973, CANDE *et al.* 1973, FORDE and STEER 1976, FRANKE *et al.* 1972, HAHNE and HOFFMAN 1984, HERTH *et al.* 1972, ISHIGAMI and NAGAI 1980, MASCARENHAS and LAFOUNTAIN 1972, PALEVITZ 1980, PARTHASARATHY *et al.* 1985,

PARTHASARATHY 1985, PERDUE and PARTHASARATHY 1985, SEAGULL and HEATH 1980 b, WESSELS *et al.* 1971, WITZMUN and PARTHASARATHY 1985), it seems plausible to suggest a role for actin in cytoplasmic streaming of higher plant cells as well (see JACKSON 1982 for a review).

Another process in which microfilaments are thought to play a role is tip growth of pollen tubes (FRANKE *et al.* 1972, HERTH *et al.* 1972, MASCARENHAS and LAFOUNTAIN 1972, PIERSON *et al.* 1986) and root hairs (HERTH *et al.* 1972). Both processes are disrupted by cytochalasins. This inhibitory action is likely due to a disturbance of the vectorial control of vesicular wall material transport which may be mediated by axial microfilaments.

Organelle movement is a third cytoplasmic event believed to require the presence of microfilaments. Light-induced chloroplast migration (2° streaming) in *Valisneria*, *Egeria*, *Elodea*, and *Hydrilla* has been demonstrated to be dependent on a peripheral microfilament network by cytochalasin-induced disruption of movement, structural evidence, and physiological and mechanical manipulations (ISHIGAMI and NAGAI 1980, PARTHASARATHY *et al.* 1985, TAKAGI and NAGAI 1983, WITZMUN and PARTHASARATHY 1985, YAMAGUCHI and NAGAI 1981). *Allium* guard mother cells (GMC) require microfilaments for nuclear migration following division, as shown by specific disruption with cytochalasin and phalloidin (PALEVITZ 1980).

Many authors posit a role for cytoplasmic microfilaments in cell shape determination (O'BRIEN and THIMANN 1966, PARTHASARATHY *et al.* 1985, PARTHASARATHY 1985, SCHMIT and LAMBERT 1985 a, b, SEAGULL and HEATH 1980 a, b). In plant cells, the cell wall, a semirigid cellulosic matrix, defines the boundaries to be filled by turgid protoplasts. Whereas cortical microtubules are generally believed to play an important role in cell shape determination by controlling the orientation of cellulose microfibril deposition, the involvement of actin is less well understood (HEPLER and PALEVITZ 1974, PALEVITZ and HEPLER 1975, HEATH and SEAGULL 1982). A few reports suggest that cell wall deposition might be mediated by both microtubules and microfilaments (reviewed by HEATH and SEAGULL 1982). Cytochalasin B treatment of *Raphanus* root hairs did not result in a disruption of cellulose microfibril deposition or cortical microtubule arrays but rather led to an overall increase in microtubule lengths and in the number of microtubule-microtubule associations (SEAGULL and HEATH 1980 a). *Allium* guard mother cells treated with cytochalasin B or phalloidin also do not

show specific disruption of microtubule arrays, atypical wall thickenings, or cellulose microfibril orientation (PALEVITZ and HEPLER 1976, PALEVITZ 1980). The observation that normal wall morphogenesis can occur despite the inhibition of microfilament function seemingly refutes the claims of microfilament-microtubule interactions in cell wall formation and cell shape determination. In a unique study using *Hibiscus* and *Nicotiana* protoplasts, either laser microsurgery or cytochalasin treatment led to the reversible disruption of large cytoplasmic strands, a cessation of streaming, and the formation of isodiametric protoplasts (HAHNE and HOFFMAN 1984). These observations are suggestive of a role for cytoplasmic microfilaments in protoplast shape determination, but whether processes of protoplast shape determination influence intact plant cell morphogenesis is poorly understood.

Karyokinesis has also been suggested to be a process requiring microfilament-microtubule interactions. Kinetochore fiber-associated microfilaments could, theoretically, drive chromosome-to-pole movement (FORER 1978, MCINTOSH 1979). This is controversial since cytochalasin treatment or antimyosin antibody injections, in general, do not disrupt chromosome movement (MCINTOSH 1979). Cytochalasin B does not inhibit chromosome movement in *Haemanthus* spindles (SCHMIT and LAMBERT 1985 b). The spindle-associated cortical microfilaments reported in *Haemanthus* endosperm might instead control the lateral movement of interzone microtubules in anaphase observed in these cells; alternatively, they might be involved in vesicle transport on the spindle surface (SCHMIT and LAMBERT 1985 a, SCHMIT *et al.* 1986 a). This cortical framework might also maintain spindle integrity and cell shape during mitosis (SCHMIT and LAMBERT 1985 b, 1986 b, SCHMIT *et al.* 1986 a). Treatment of *Allium* guard mother cells with cytochalasins or phalloidin does not disrupt spindle formation or spindle structure, nor does it interfere with chromosome movement (PALEVITZ 1980), which therefore suggests that microtubule-microfilament interactions are not important in these processes. In contrast to these observations, spindle reorientation in anaphase and telophase of *Allium* guard mother cells is sensitive to microfilament, microtubule, and metabolic inhibitors (HEPLER 1981, PALEVITZ and HEPLER 1974 b, 1976, PALEVITZ 1980). Numerous reports of actin and myosin localization led to speculation on their role in phragmoplast and cell plate formation and function (CLAYTON and LLOYD 1985, PARKE *et al.* 1986, SCHMIT and LAMBERT 1985 a, b, 1986 b, SCHMIT *et al.* 1986 a). Cell plate orientation

in *Allium* guard mother cells has been shown to be cytochalasin sensitive, but attempts to visualize significant numbers of microfilaments by electron microscopy so far have been unsuccessful (HEPLER 1981, PALEVITZ and HEPLER 1974 b, 1976, PALEVITZ 1980). In contrast to the disruption of cell plate orientation, cytochalasin or phalloidin treatment does not disrupt phragmoplast microtubule organization, or the transport of vesicles to the division plane for cell plate formation (PALEVITZ and HEPLER 1974 b, PALEVITZ 1980). These results do not rule out the possibility that phragmoplast actin plays a role in the centrifugal expansion of this organelle or in vesicle fusion at the cell plate. Finally, a cytoskeletal role in graviperception has been suggested based on an apparent structural interaction between amyloplasts and microfilaments in the statocytes of *Zea* and *Hordeum* coleoptiles (WHITE and SACK 1986). Light microscopic observation of these cells suggests that cytoplasmic streaming may contribute to the dynamics of amyloplast sedimentation (SACK and LEOPOLD 1985). Other reports suggest that actin microfilaments are important in generating a polar organization of cellular organelles, specifically the rough endoplasmic reticulum, in presumptive statocytes of cress (*Lepidium*) roots (HENSEL 1985, WENDT and SIEVERS 1986). This polar organization is thought to be crucial to the perception of gravity by these cells, and can be disrupted by cytochalasin treatment (WENDT and SIEVERS 1986, HENSEL 1985). Localization of microfilaments with rhodamine-phalloidin did not reveal any specific interaction with the rough endoplasmic reticulum, but instead displays weak or diffuse cortical staining and perpendicular cytoplasmic bundles (HENSEL 1986).

4. Outlook

This account of higher plant actin and associated proteins makes it apparent that much work remains to be done to understand the role these cytoskeletal proteins play in plant cellular processes. To date, most researchers have relied upon "function by correlation", *i.e.*, actin is localized in motile structures therefore it plays a role in driving the motility. This is a necessary, yet insufficient, step in understanding actin's involvement in cellular motile processes. It is important to complement these observations with physiological and mechanical manipulations, as was done in Characean algae. Similarly, inhibitor studies with, for example, cytochalasins, can provide valuable first insights but are of limited value because of their reported nonspecific effects (WESSELS *et al.* 1971, POPE *et al.* 1979). Phal-

lotoxins, which apparently also disrupt actin-mediated events even when applied to intact cells, should be used to complement experiments employing cytochalasins. The ability to microinject plant cells has been demonstrated (CROSSWAY *et al.* 1986), and injection of antibodies against actin, myosin, and other cytoskeletal proteins will provide inhibitory molecules of higher specificity than pharmacological agents. More effort should be directed towards the development of *in vitro* systems since they bridge traditional biochemistry and *in vivo* observations. Reactivable lysed cells have proven invaluable tools in the study of animal cell motility, and there is reason to believe that this approach will be similarly informative for plant cell motility, once the horrendous problems posed by the presence of vacuoles, and the design of cell milieu-compatible buffers, can be overcome. Other approaches that should prove fruitful in defining the role of actomyosin in the cell cycle are analyses of mutants and genomic manipulation by molecular genetic techniques. Finally, a search for actin-associated proteins in general, and proteins that mediate microtubule-microfilament interactions in particular seems desirable, as these interactions will prove crucial for understanding higher plant cytoskeletal function. The study of the cytoskeleton of higher plant cells is just beginning. However, one can be satisfied with the progress that has been made to date, and can look forward to significant contributions in the near future.

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Note Added in Proof:

Rhodamine-phalloidine localization of F-actin in the preprophase band of *Allium* root tip cells (PALEVITZ, BA (1987) *J Cell Biol.* 104: 1515-1519) and *Daucus* suspension culture cells (TRAAS JA *et al.* (1987) *J Cell Biol* 105: 387-395) has recently been reported in the literature.

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