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## Function and regulation of the Arp2/3 complex during cell migration in diverse environments

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### Abstract

As the first *de novo* actin nucleator discovered, the Arp2/3 complex has been a central player in models of protrusive force production via the dynamic actin network. Here, we review recent studies on the functional role of the Arp2/3 complex in the migration of diverse cell types in different migratory environments. These findings have revealed an unexpected level of plasticity, both in how cells rely on the Arp2/3 complex for migration and other physiological functions and in the intricate modulation of the Arp2/3 complex by other actin regulators and upstream signaling cascades.

### Introduction

Actin polymerization drives the morphological changes that allow cells to undergo dynamic processes, such as division, phagocytosis, and migration. The formation of new actin filaments from actin monomers is regulated by three classes of nucleating proteins, including the formins, tandem-monomer-binding nucleators, and the Arp2/3 complex [1–3]. Formins, such as mDia1 and 2, generate unbranched filaments by stabilizing actin trimers and promoting elongation through associations with both the growing barbed end and actin-bound profilin. The tandem-monomer-binding nucleators, including Spire, cordon bleu, and leiomodin, form unbranched filaments by bringing together actin monomers, via a series of monomer-binding WH2 domains, in a configuration that mimics the stable actin trimer. The Arp2/3 complex is unique in that it nucleates branched actin filaments.

The first-identified actin nucleator, the Arp2/3 complex has been studied extensively using *in vitro* reconstitution assays, biochemical and structural analyses, and *in vivo* genetics-based functional experiments [4,5]. The Arp2/3 complex consists of seven subunits, two of which (Arp2 and Arp3) are actin-related proteins that serve as a nucleus for the new actin

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filament. Other Arp2/3 complex subunits bind to existing actin filaments to generate a branch at a  $\sim 78^\circ$  angle from the mother filament [6–8]. Nucleation by Arp2/3 is regulated by many proteins, most prominently the Wiskott-Aldrich syndrome family of nucleation promoting factors (NPFs), such as WASP, N-WASP, Scar/WAVE, and WASH. The WASP family NPFs share a consensus WCA domain in their C-termini that binds monomeric actin through the WH2 domain (W) and the Arp2/3 complex through the cofilin homology region and acidic tail (CA) [9,10]. Thus, in a spatiotemporally-regulated manner, these NPFs facilitate the transition of the Arp2/3 complex from its splayed, inactive conformation to its closed, active conformation and also supply an actin monomer to form the growing barbed end. The WISH/DIP/SPIN90 proteins are NPFs that activate the Arp2/3 complex without binding to F- or G-actin, promoting the formation of unbranched filaments that may serve as the seed for subsequent branch nucleation [11].

Cells must be able to rapidly modify their actin network to adapt to their surroundings. It is well-established that cells can switch between multiple modes of migration depending on their gene expression, the dimensionality of their environment (1D vs 2D vs 3D), and the extracellular matrix (ECM) stiffness and composition [12–24]. This is accomplished in part by complex Arp2/3 regulation combined with the multitude of other mechanisms that promote actin nucleation during cell migration, allowing for tight spatiotemporal control of the formation of different actin architectures. Furthermore, applied mechanical resistance alters the actin network density, geometry, power and stiffness, suggesting that variations in force may influence the organization of actin filaments *in vivo* [25]. Geometric constraints also dictate the structure of the actin network [26–28]. Nonetheless, variability in experimental conditions and the intricacy of the underlying regulatory mechanisms has made it difficult to dissect the role of Arp2/3 in cell migration. This review will focus on recent advances in our understanding of Arp2/3-mediated actin polymerization during cell migration and the mechanisms by which cells fine-tune their actin networks to adapt to internal perturbations or extracellular environments.

### Studies of Arp2/3 Subunit-Disruption in Migration

Because of its central role in the actin polymerization required for cell migration, endocytosis, and other vital processes, functional studies of the Arp2/3 complex were initially hindered by the lethality of Arp2 or Arp3 null cells and animals [29,30]. Early studies with Arp2/3 complex subunit knock-down or NPF inhibition supported the notion that the Arp2/3 complex mediates actin polymerization during lamellipodia formation and migration in cells such as fibroblasts, lymphocytes, mammary carcinoma cells, and the amoeba *Dictyostelium discoideum* [31–42]. Improvements in genetic techniques over the past decade have allowed for more direct tests of Arp2/3 function in migration in a variety of cell types (Table 1).

**Fibroblasts**—Our group and others have shown that loss of Arp2/3 activity in mouse embryonic fibroblasts through the use of inhibitors, complex subunit knock-outs, or RNAi results in the loss of lamellipodia [43–47]. Surprisingly, these Arp2/3-deficient cells are still migratory, though with reduced speed, through the generation of formin-containing filopodial structures of bundled actin [45,46]. These actin structures coordinate with myosin

II-mediated contractility in the cortex to drive leading edge extension [44]. Although these studies found similar alterations in F-actin structures at the leading edge, they disagreed on the effect of Arp2/3 complex loss on migration toward chemical cues: our studies showed reduced directional persistence and a cell-autonomous defect in chemotaxis toward growth factors, while Wu *et al* found that Arp2/3 knock-down cells only exhibit a chemotactic deficiency if there is interference by secreted inflammatory cytokines [44,47]. The same group did find that haptotaxis toward surface-bound ECM molecules is impaired in the knock-down cells and later repeated their findings with an Arp2/3 subunit knock-out [43,46]. These discrepancies could be caused by differences in genetic background, the manner by which the cells were produced, the targeted Arp2/3 complex subunit, or experimental conditions.

**Neural and glial progenitors**—Recent studies have examined the role of the Arp2/3 complex in the migration of progenitor cells found in the brain. For example, Arp2/3 is required for the directed migration of neural stem cell-derived oligodendrocyte precursor cells (OPCs) in an electric field, which has implications for neural stem cell transplantation for remyelination [48]. Similar to mouse embryonic fibroblasts, Arpc2-null OPCs were slower, with shorter, filopodial processes than Arpc2-expressing OPCs. We recently found that Arp2/3 is required for neural cell migration *in vivo*: Arpc2-null mice have impaired cortical architecture due to defects in both radial glial cell processes and the migration of neural progenitors along these tracks [49]. *In vitro* experiments showed that the migration defect of neural progenitor cells is particularly pronounced in soft or low-laminin environments that mimic brain tissue.

**Hematopoietic cells**—As opposed to fibroblasts and neural progenitor cells, evidence shows that Arp2/3-mediated actin polymerization inhibits the migration of both dendritic and T cells, which will be discussed below [50,51]. However, a recent study of dendritic cells suggests that successful passage through narrow pores, which is determined by the physical constraints of the nucleus, is associated with an increase in actin polymerization around the nucleus as it reaches the point of constriction [52]. This perinuclear actin, as well as migration under confinement, is reduced by Arp2/3 inhibition or knock-down, but is not required for the confined movement of cells with low lamin levels and softer nuclei, suggesting that actin filaments facilitate migration through narrow spaces by promoting nuclear deformation [52]. In hemocytes, the *Drosophila* macrophage-like immune cells, depletion of Rho1, WASH, or Arp2/3 subunits reduces the formation of cellular protrusions and prevents a subset of migration events during development [53]; similar studies suggest that lamellipodia formation and migration of hemocytes *in vivo* require Scar/WAVE [54,55].

**Epithelial cells**—In MCF10a mammary epithelial cells, inhibition or knock-down of Arp2/3 results in the disruption of lamellipodia and the formation of unstable or bleb-like protrusions, corresponding to a decrease in directional persistence and migratory speed [56]. Furthermore, Arp2/3 inhibition impairs nascent focal adhesion assembly and decreases the coupling between the actin cortex and cell membrane [56].

**Cancer cells**—Because of the links to metastatic migration, the role of Arp2/3 has been examined in a number of cancer cell lines. The results are conflicting, which is unsurprising given the tremendous genetic diversity and heterogeneity of cancer cells. Several studies have shown that the Arp2/3 complex or Arp2/3-stimulating factors, such as cortactin, are upregulated in malignant gliomas, and inhibition of Arp2/3 activity reduces lamellipodia formation and invasion [57,58]. Similar trends have been shown for Arp2/3 and its NPFs in models of mammary carcinoma [59–61]. However, other studies of glioma motility concluded that Arp2/3 activity is unnecessary, or even inhibitory, for migration in confined environments [62–65]. The mechanisms of Arp2/3-independent migration are unclear, as studies alternatively conclude that either motility is dependent on formin-mediated actin polymerization or is independent of actin polymerization entirely based on observed migration in the presence of actin depolymerizing agents. Similarly, endocytosis of  $\alpha 5\beta 1$  integrin enhances the invasion of ovarian carcinoma cells into the 3D matrix by promoting the formation of actin spikes via RhoA-mediated formin activity, independent of Arp2/3 [66].

### **Multifaceted Regulation of the Arp2/3 Complex Adapts the Actin Network to Diverse Modes of Migration**

**Proteins that regulate the Arp2/3 complex**—In addition to NPFs and upstream signaling factors, a number of proteins function in the regulation of Arp2/3 activity to control cell migration [10,67–77]. Recent studies have centered on a newly-discovered negative regulator of Arp2/3, Arpin [78]. This protein is recruited to the lamellipod by activated Rac, where it directly inhibits Arp2/3 activity to destabilize protrusions [78,79]. Arpin helps control directional persistence and migration speed by inducing pauses in motility [79]. Subsequent studies showed an inverse correlation between Arpin expression and breast cancer metastasis; low levels of Arpin are also associated with elevated expression of WAVE complex subunits and poor recurrence-free survival [80,81]. Another negative regulator, Gadkin, sequesters the Arp2/3 complex to endosomal vesicles, thereby inhibiting cell spreading and affecting the migration speed of dendritic cells *in vitro* [82,83]. In sum, the numerous potential interactions between Arp2/3 and components of the actin machinery leads to a highly regulated and tunable actin network that can be specifically tailored to a variety of dynamic cellular behaviors.

**Differential activation of NPFs fine-tunes the structure of the actin network**—Evidence suggests that the differential utilization of various Arp2/3-activating proteins, including NPFs, can alter the migratory behavior of cells depending on environmental conditions. For example, recent work has demonstrated that loss of the Scar/WAVE complex in both carcinoma and normal epithelial cells decreases migration in 2D wound healing assays but promotes N-WASP- and Arp2/3-mediated invasion into 3D matrices through the activation of focal adhesion kinase (FAK) at the leading edge [84,85]. Thus, Arp2/3 plays different roles in 2D vs 3D migration depending on whether it is activated by Scar/WAVE or N-WASP. FAK, heavily implicated in cancer cell migration, interacts with Arp2/3, recruiting it to sites of nascent focal adhesions [86, 87]. The interaction between FAK and Arp2/3 also couples cell adhesion to leading edge protrusions and is required for migration of fibroblasts in response to ECM gradients in 2D [87,88]. Similarly, depletion of N-WASP, WAVE1, or

cortactin in fibrosarcoma cells undergoing migration in different environments showed that the regulatory machinery is alternatively activated for 2D vs 3D Arp2/3-based migration, resulting in different morphologies and migratory behaviors [89]. During migration in 3D environments, these cells form thin, dendritic extensions that branch to generate new protrusions that are molecularly distinct from the original, suggesting that the formation of different types of actin structures is coordinated [89].

**Balance between Arp2/3 and other actin polymerizing factors**—Studies of cells lacking the Arp2/3 complex have indicated that these cells can still generate protrusive actin networks that sustain migration through other actin nucleators [12,43–48,90]. These studies also showed that different actin polymerizing factors lead to different modes of migration [12,43–48,70,90–92] (Figure 1). A specific example of how the utilization of multiple nucleators contributes to cellular function comes from studies of dendritic cells (DCs) in confining microchannels. DCs have two competing pools of actin: a Cdc42- and Arp2/3-mediated accumulation of F-actin at the cell front that slows motility but is required for micropinocytosis and antigen uptake in immature DCs, and a RhoA- and mDia-mediated F-actin enrichment at the cell rear that is necessary for the fast migration and chemotaxis used by mature DCs to travel to the lymph nodes [51]. Stimulation with LPS, which triggers DC maturation, induces a switch from Arp2/3- to formin-derived actin networks with corresponding changes in migratory behavior [51]. However, previous studies reported that depletion of Arp2/3 activators, including Cdc42 and WASP, impairs migration of DCs to the lymph nodes [93–95]. These contradictions may be due to experimental differences in migratory environments. Similarly to DCs, Arp2/3 activity inhibits T cell migration to promote synapse formation upon high-affinity antigen binding, but is not required for the fast motility of T cells that are unbound by antigen [50]. Homeostasis between Arp2/3- and formin-based polymerization in fission yeast is controlled by competition for actin monomers, ensuring the proper assembly of the contractile ring and endocytic actin patches [96]. These studies and others suggest that the balance between different actin nucleators is critical for specific cell function and can be regulated by extracellular stimuli and intracellular signaling cascades [12,50,51,96].

Although switching between distinct types of actin networks is important for regulating migratory behavior under different environmental conditions, evidence suggests that the interplay between various actin polymerizing factors within the same structure also contributes to the fine-tuning of the actin assembly. For example, formins have been observed in lamellipodia, where they modulate the structure of the dendritic actin network by competing with capping protein for free barbed ends and promoting filament elongation [97–99]. Ultrastructural analysis of cells depleted of the formin mDia1, Arp2/3, or both further suggests that these nucleators are spatially intertwined with distinct consequences on cortical actin structure [90]. Other work has shown that fascin-1-containing bundles of actin may serve as a template for lamellipodia formation [88]. This is supported by the finding, generated by a combination of *in vitro* actin polymerization assays and spatially- and temporally-defined mDia1 inactivation in HeLa cells, that mDia1 stimulates Arp2/3 branching activity by providing mother filaments for nucleation [100]. Furthermore, increases in one nucleating factor can compensate for depletion of the other, supporting the

conclusion that mDia1 and Arp2/3 activities cooperate to influence the impact of the other on actin meshwork structure [100]. Other studies demonstrating that formins can assemble filopodia using dendritic actin generated by the Arp2/3 complex also highlight the cooperative nature of these nucleating factors [101–104].

**Role of profilin as an integrator of multiple actin nucleation factors**—Several recent studies have pointed to a role for the actin monomer-binding protein profilin as a negative modulator of Arp2/3-based actin nucleation. In Arpc2-null mouse fibroblasts, only profilin-bound actin monomers can incorporate into barbed ends, and excess profilin can inhibit Arp2/3-mediated actin polymerization both in wild-type cells and *in vitro* [105]. Profilin is known to enhance formin-mediated barbed end elongation; however, depletion or sequestration of either profilin or Ena/VASP, but not chemical inhibition of formins, which may not be complete, in the Arpc2-null background prevents protrusion formation and spreading [105]. The authors concluded that profilin antagonizes Arp2/3-mediated nucleation and actin branching by facilitating actin polymerization via Ena/VASP, thereby competing away actin monomers from the Arp2/3 complex. In contrast, Ena/VASP has also been proposed to interact with the WAVE regulatory complex to stimulate Arp2/3 activation, enhance migration, and maintain normal lamellipodia formation in *Drosophila* hemocytes and *C. elegans* epidermal cells, suggesting that this family of actin assembly proteins may have alternative functions in diverse cell types or under different conditions [106,107]. Work combining analysis of the contractile ring in *S. pombe* with *in vitro* actin reconstitution assays also shows that profilin inhibits Arp2/3 function, thereby maintaining the balance between formin- and Arp2/3-mediated actin polymerization that is required for cytokinesis [108]. These studies conclude that profilin tunes the actin network through the competition for actin monomers [105,108]; however, another study suggests that profilin competes with formins, VASP, capping protein, and Arp2/3-mediated end branching by binding to the barbed ends of F-actin, thereby regulating the length of actin filaments [109]. Although the mechanisms may be complex, profilin expression is emerging as an important regulator of actin network dynamics during migration by fine-tuning the activities of various actin polymerization proteins.

## Conclusions and outlook

Recent findings on the function of the Arp2/3 complex, an actin nucleator long thought to play a key role in actin-based protrusive force generation, during the migration of diverse cell types in various experimental settings have been both fascinating and confusing. Underneath the conflicting functional consequences of Arp2/3 inhibition, it is becoming apparent that there is an unexpected level of complexity and plasticity in actin network formation, regulation, and force production to drive cell motility in response to specific geometric or mechanical properties of the environment. Particularly important is the intricate interplay, both competitive and collaborative, between different actin nucleators. Through gene expression and distinct subcellular localizations or activating signaling molecules, maintaining a balance between the Arp2/3 complex, its regulators, and other nucleating proteins is crucial for modulating the dynamic actin structures found during cell migration. In cancer, the mechanistic adaptability of the actin network is influenced not only by environments but also by the genetic heterogeneity of tumor cells, highlighting the

considerable challenge in understanding and preventing cancer cell migration during dissemination and metastasis. Because of the extraordinary plasticity of cell motility machineries, inconsistencies in experimental observations associated with different cell lines or environmental conditions could shed light on the adaptive principles and regulatory complexity of the actin network in cell migration.

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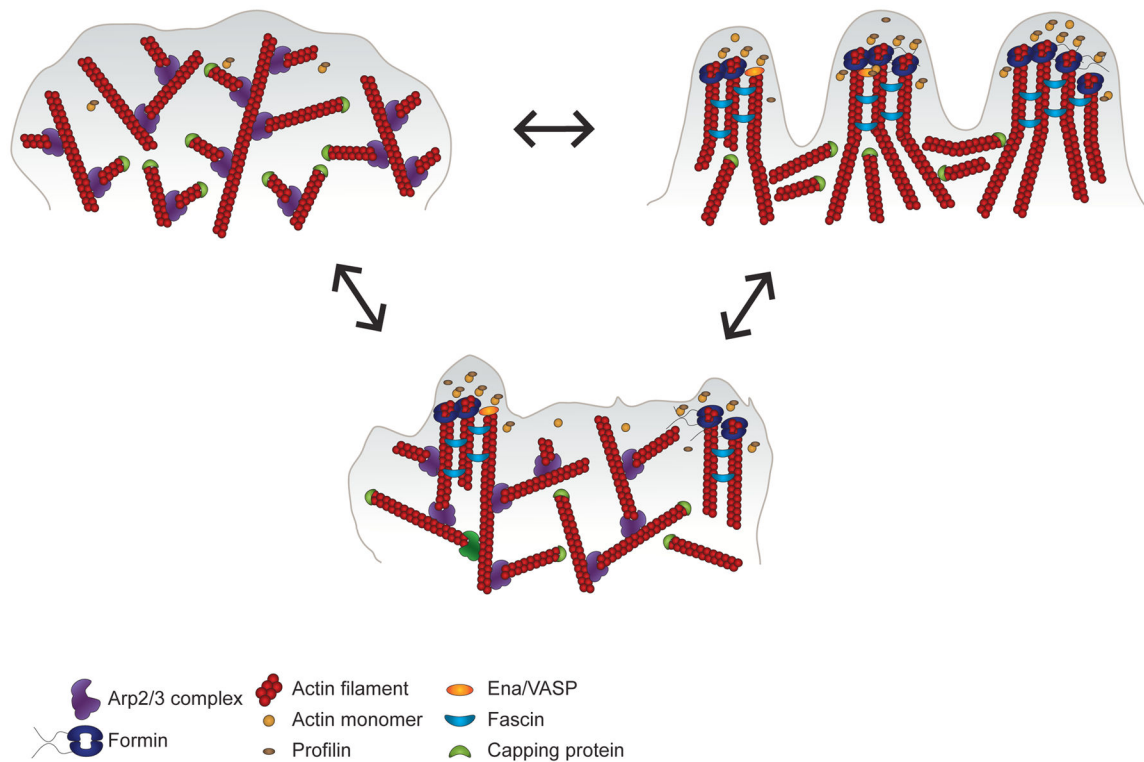
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### Figure 1. Cells use different actin networks for migration

The plasticity underlying the regulation of the actin machinery allows cells to adapt to diverse migratory environments. The migration of fibroblasts and other cell types on 2D surfaces is characterized by the formation of broad lamellipodia at the leading edge (upper left). These structures are composed of Arp2/3-mediated branched actin networks. In confined environments or in the absence of the Arp2/3 complex, many cell types generate filopodial structures of bundled linear actin filaments generated by formins (upper right). Properties of the extracellular matrix and inputs from upstream signaling pathways can mediate a switch between these different actin machineries. However, it is likely that cells migrating in diverse 3D environments *in vivo* use a combination of actin nucleators in order to fine-tune their cytoskeletal networks (lower center); these nucleators have been shown to both cooperate with and antagonize each other, depending on environmental context. Competition for free actin monomers, influenced by profilin, can modulate the balance between Arp2/3- and formin-mediated actin polymerization. Likewise, competition between formins, Ena/VASP, capping protein, profilin, and even NPFs for free barbed ends can influence the length and structure of the actin filaments. Thus, the migration machinery can be thought of as a spectrum, and cells can shift along this continuum from one mode to the next depending on regulatory protein concentrations, genetic factors or environmental cues. It should be noted that this Figure does not include contractility- or pressure-based modes of migration, such as blebbing or lobopodia.

Table 1

Functional consequences of Arp2/3 depletion on migration in various cell types and environments

Cell Type	Method of Arp2/3 Depletion	Migratory Environment Tested	Arp2/3 Function or Phenotype	Reference
Fibroblasts	ES cells from Arpc3-null mice were differentiated into fibroblasts	2D closed microfluidic chambers	Null cells lost lamellipodia but migrated by combining formin-based filopodia formation with myosin II-based contractility to advance the leading edge; null cells were unable to migrate directionally during wound healing or chemotaxis toward EGF	Suraneni <i>et al.</i> , 2012; Suraneni <i>et al.</i> , 2015
	Mouse embryonic fibroblasts with combined knock-down of Arpc2 and Arp2 subunits in the <i>Ink4a/Arf</i> <sup>-/-</sup> genetic background; inhibition with CK666	2D continual-flow microfluidic chambers	Knock-down cells lost lamellipodia, showed focal adhesion and spreading defects, and migrated with reduced speed via filopodial structures; knock-down cells were deficient in haptotaxis but showed normal chemotaxis if media was replaced; knock-down cells, via NF- $\kappa$ B activation, secreted growth factors that disrupted chemotaxis in a closed system	Wu <i>et al.</i> , 2012; Wu <i>et al.</i> , 2013
	Mouse fibroblasts with conditional Arpc2 knock-out in the <i>Ink4a/Arf</i> <sup>-/-</sup> genetic background	2D continual-flow microfluidic chambers	Null cells were able to chemotax toward PDGF	Asokan <i>et al.</i> , 2014
	Null cells lost lamellipodia, showed an increase in filopodial protrusions, and migrated with reduced speed		Rotty <i>et al.</i> , 2015	
Neurons and glia	Neural stem cells from Arpc2 knock-out mice were differentiated into oligodendrocyte precursor cells (OPCs)	Oligospheres plated on glass chambers with applied electric fields	Null cells had shorter, filopodial extensions and moved with reduced speed compared to wild-type OPCs; Electric fields enhanced the migration and biased the directionality of wild-type OPCs but did not elicit these responses in null cells	Li <i>et al.</i> , 2015
	Radial glial cells from Arpc2 knock-out mice	Cortical slices were embedded in matrigel for <i>ex vivo</i> imaging	Null cells had disorganized processes that lacked dynamic ruffles and were shorter than control cell protrusions due to frequent retraction; nulls showed altered cell fates due to premature differentiation, increased apoptosis, and decreased proliferation; these defects contributed to abnormal cortical architecture	Wang <i>et al.</i> , 2015
	Neuronal precursor cells from Arpc2 knock-out mice	Wild-type or null neurospheres cultured <i>ex vivo</i> with wild-type brain slices; glass or elastic laminin-coated surfaces <i>in vitro</i>	Null cells could not migrate in the <i>ex vivo</i> system or on soft or low-laminin surfaces <i>in vitro</i> ; null cells lacked lamellipodia but could migrate, with reduced speed, on stiff or high-laminin surfaces	Wang <i>et al.</i> , 2015
Hematopoietic cells	Drosophila pupal macrophages with Arp2 or WAVE RNAi knock-down	2D glass surfaces and <i>in vivo</i> (WAVE knock-down only)	Knock-down cells lost lamellipodia and generate filopodial protrusions; WAVE knock-down cells extend processes in the direction of a chemotactic gradient but fail to migrate	Sander <i>et al.</i> , 2013
	Drosophila embryonic macrophages with	<i>In vivo</i>	Partial failure of knock-down cells to migrate anteriorly during development due to reduced	Verboon <i>et al.</i> , 2015

Cell Type	Method of Arp2/3 Depletion	Migratory Environment Tested	Arp2/3 Function or Phenotype	Reference
	Arp3 or WASH RNAi knock-down		protrusion size; knock-down cells could migrate posteriorly, but were disorganized	
	OT-I CD8+ T cells with chemical inhibition of Arp2/3 via CK666	Microchannels coated with pMHC antigenic complexes	Inhibition of Arp2/3 did not affect fast T cell migration (unbound by antigen) or exploratory migration during kinapse formation (bound by low-affinity antigen), but impaired T cell deceleration for synapse formation (bound by high-affinity antigen)	Moreau <i>et al</i> , 2015
	Dendritic cells (immature and mature, via LPS induction) with conditional Arpc2 knock-out, Arpc4 siRNA knock-down, or chemical inhibition via CK666	1D confined microchannels; under agarose; collagen gels; <i>ex vivo</i> in ear epidermal sheets	Arp2/3 depletion enhanced migration speed in immature cells and reduced F-actin at the cell front, leading to defective antigen macropinocytosis; Arp2/3 was not required for F-actin enrichment at the cell rear or for chemotaxis in collagen gels or to lymphatic vessels <i>ex vivo</i>	Vargas <i>et al</i> , 2016
	Immature dendritic cells with Arpc4 siRNA knock-down or chemical inhibition with CK666	Microchannels with constrictions 1.5 $\mu$ m	Arp2/3 depletion did not affect normal migration but prevented cells from passing through narrow confinements (3 $\mu$ m or less); Arp2/3 depletion prevented perinuclear actin accumulation and nuclear deformation required for passage through narrow pores	Thiam <i>et al</i> , 2016
Epithelial cells	MCF10A mammary epithelial cells with siRNA knock-down of Arp3 or chemical inhibition via CK869 and CK666	2D coverslips	Arp2/3 depletion disrupted lamellipodia formation, reduced migration speed and directional persistence, and impaired focal adhesion assembly and attachment to the ECM; Arp2/3 depletion also weakened the coupling between the cytoskeletal cortex and plasma membrane	Beckham <i>et al</i> , 2014
Cancer cell lines	Human glioma cell lines U251, LN229 and SNB19 with chemical inhibition via CK666	2D wound healing; transwell invasion through Matrigel-coated Boyden chambers	Arp2/3 inhibition reduced lamellipodia size, disrupted polarity, and reduced migration and invasion in all cell lines; Arp2/3 expression is correlated with malignancy of patient tumors	Liu <i>et al</i> , 2014
	Rat C6 glioma cells or human patient-derived glioma propagating cells with chemical inhibition via CK666	2D surfaces or thin (3 – 7 $\mu$ m), linear laminin tracks	Arp2/3 inhibition blocked 2D migration but enhanced linear/saltatory migration by altering cell morphology, increasing cell velocity, and improving cell alignment in antiparallel arrays	Monzo <i>et al</i> , 2016
	A2780 human ovarian carcinomas with siRNA knock-down of Arpc2 or Arpc3 or chemical inhibition via CK666	2D surfaces or elastic 3D cell-derived matrix	Arp2/3 depletion did not affect $\alpha$ 5 $\beta$ 1 integrin recycling-mediated protrusion morphology, migration, or invasion	Paul <i>et al</i> , 2015

\*For simplicity, this table does not include the many studies involving depletion of Arp2/3 regulators, including NPFs