Single-Molecule Fluorescence and in Vivo Optical Traps: How Multiple Dyneins and Kinesins Interact

Benjamin H. Blehm and Paul R. Selvin*

Physics Department and Center for Physics of the Living Cell, University of Illinois at Urbana-Champaign, 1110 West Green Street, Urbana, Illinois 61802, United States

1. INTRODUCTION

Kinesin and dynein walking on microtubules are the two main drivers of long-distance intracellular transport in a huge variety of systems, from neurons to melanophores. These motors, however, are oppositely directed, with (most) kinesin driving cargos toward the plus ends of microtubules whereas dynein drives cargos toward the minus ends. There are only two types of dynein, cytoplasmic and axonemal, with only cytoplasmic dynein being used for organelle transport. In this review, when we use the term dynein, we are referring to cytoplasmic dynein. Dynein is generally associated with a large multisubunit complex, dynactin, in vivo, which appears to be necessary for many types of transport. Kinesins make up a large family of motors involved in organelle transport, ranging from conventional kinesin (kinesin-1), which is a typical processive, plus-end-directed kinesin, to NCD, a nonprocessive, minus-end-directed kinesin. In addition to dynein and kinesin, there is a third motor, myosin, which walks on actin. Often, myosin is also present on the cargo, and the cargo is made to switch between microtubules and actin; the latter is often for final placement of the cargo.

In this review, we describe experimental systems at multiple levels of complexity, including single-motor-type in vitro assays, multimotor in vitro assays, purified-organelle in vitro assays, and finally in vivo cellular assays (Figure 1). This spread of experiments allows an unprecedented view of the transport complex, as kinesin and dynein can be observed with differing components of the transport complex (i.e., different levels of accessory proteins) and in different environments. Through the

Special Issue: 2014 Single Molecule Mechanisms and Imaging

Received: October 7, 2013
1.1. Kinesin and Dynein Interaction: Tug-of-War versus Coordinated Model

An initial question is why multimotor models are needed. After all, a single motor type is all that is needed for transport in one direction. Most motors appear to be recruited to cargos by specific binding factors, so the cell can control the presence of motors on a specific cargo. However, it is known that, in many systems, both kinesins and dyneins are simultaneously present on cargo. Often, seemingly erratic up-and-back behavior is observed. How multiple motors, and different motor types, interact and are regulated is fundamental to understanding intracellular transport. (For excellent reviews covering intracellular transport, see refs 1, 5, and 9.)

There is currently a wide array of models describing the interaction between kinesin and dynein. In this review, the term interaction means any interplay between kinesin and dynein dynamics, such as through a cargo, not necessarily a direct, physical interaction. These models typically fall into two main categories: coordinated motion, which involves a secondary protein or complex that controls the states of kinesin and dynein, regulating their activity and determining the cargo’s directionality on the microtubule, and tug-of-war motion, which postulates that kinesin and dynein interact directly by force transductions through the cargo that determine directionality (Figure 2). Historically, the definitions of coordinated versus tug-of-war motion have varied somewhat. Today, however, there is general agreement. Coordinated motion typically involves only one particular type of motor being active at any time (kinesin or dynein). Tug-of-war models have several possible states, for example, both motors are pulling and the one that is pulling with more force wins out. Another tug-of-war scenario can have the “losing” motor come off the microtubule or stay bound but walk or diffuse backward. It is

The simplest level of complexity is a single motor with a cargo or label attached and a microtubule track in an in vitro environment. This has been the predominant type of experiment in the study of molecular motors. It has revealed their stepping behavior, stall force, and other characteristics. However, it has little to say about motor–motor interactions. (B) Complexity can be increased by adding extra motors, either multiple kinesins, multiple dyneins, or kinesin and dynein. This is the most basic way to study motor–motor interactions and has been used to study the cooperativity of groups motors. Knowing the absolute number of each motor can be difficult. (C) Adding in accessory proteins and parts of the transport complex, such as dynactin, is the next level of complexity. How accessory proteins and signaling molecules (such as cAMP or a kinase) modulate kinesin–dynein interactions can be studied in this system. (D) The living cell is the most complex system in which to study motor–motor interactions. Cellular gradients, accessory proteins, microtubule-associated proteins (MAPs), organelles, and filament meshes are just a few of the things present that could affect transport. This complexity makes it very difficult to isolate specific causes of transport behavior but also allows for the study of motor–motor interactions in their native settings.

Combination of measurements at all of these levels of complexity, the ability to parse out the function of parts of the transport complex, and reconstitute it in vitro, becomes a real possibility.

In addition, new techniques, from in vivo optical trapping, to high-resolution imaging, are discussed. Such techniques allow for the detailed examination of all of these systems in multiple domains: force, orientation, position, and velocity, among others. They will also allow for the development and testing of theoretical models that describe intracellular transport and multimotor interactions. This review is organized such that, after one has read the first section for an overview, each section can be read more or less independently.
possible that which set of motors “wins” depends on the particular number of the motors pulling and that number might be regulated. In this review, determination of cargo directionality by strain sensitivity is the definition used for tug-of-war motion. A tug of war can lead to stalling (e.g., yeast dynein and mammalian kinesin, as discussed later), inefficient motility, or highly efficient motility (mammalian kinesin and mammalian dynein) depending on motor properties.\textsuperscript{10} Coordinated motion would be any other type of regulatory mechanism of cargo directionality that prevents motors from being simultaneously active (the existence of some external “coordinator” outside the motors and cargo). Higher-order mechanisms could exist that modulate both of these models.

For many years, a coordinated model was popular because a tug-of-war model seemed unable to explain organelle motility. This is because, if both dynein and kinesin were active simultaneously, one would expect the organelle to stall (not move) quite often. Whenever both motors became active, the organelle would stop, and it would only restart when only either kinesin or dynein remained active. Such a scenario assumed that simultaneously active kinesin and dynein would act as anchors against one another. This sort of behavior can be seen in experimental systems with yeast dynein and mammalian kinesin.\textsuperscript{11} However, this particular case is an artificial system: yeast and mammalian motors are not designed to work together.

Recent theoretical arguments\textsuperscript{10b,c} and several experiments\textsuperscript{7d,12} have shown that a tug of war can lead to motility without constant stalling and with efficient directional switching. For example, in 1992, Vale et al. showed that attaching kinesin and dynein to a surface and laying down microtubules on them caused bidirectional gliding of the microtubules with reversals in direction occurring routinely (Figure 3A,B).\textsuperscript{12c} In a somewhat different arrangement, Blehm et al.\textsuperscript{12a} and DeBerg et al.\textsuperscript{13} attached kinesin and dynein to a polystyrene bead and watched it walk on a microtubule. They showed similar reversals and bidirectional motion, with saltations and directional switching (Figure 3C). The fact that directional switching can be seen in both of these systems without any external coordinating complex is strong support for a local tug of war, although it is difficult to say how similar other transport properties are between in vitro and in vivo systems.

These experimental results are supported by several theoretical articles showing that, by varying motor properties, such as stall force, on/off rate, and velocity, among others, different directionalities and types of motility can be engineered.\textsuperscript{5b,c,14} Interestingly, the theoretical results showed that, by having a detachment force (the force needed to pull a motor off the microtubule) that is small compared to the stall force (the force needed to prevent the cargo from proceeding), tug-of-war interactions could occur that result in minimal stalling. Tug-of-war events would happen quickly, with one set of motors quickly detaching while the other took control and transported the cargo. However, a large detachment force relative to the stall force would lead to a situation with both motors attached to the cargo and microtubule and no motility occurring.\textsuperscript{5b}

Bidirectional switching of purified organelles without any cytoplasmic signaling factors (instead of isolated motors, as discussed previously) has also been observed, further adding to the tug-of-war hypothesis.\textsuperscript{7d,15} Organelles were purified from living cells, with a small complement of motors still attached. When placed on microtubules, they exhibited bidirectional, saltatory motion similar to that predicted by tug-of-war models.\textsuperscript{7d,13}

![Figure 3](dx.doi.org/10.1021/cr4005555 / Chem. Rev. XXX, XXX, XXX−−XXX)
and similar to that seen in the cell. This bidirectional motility of cargos in vitro shows that cytoplasmic factors are not necessary for directional switching, although a large array of accessory proteins could still be attached to the transport complex, making a firm conclusion impossible to reach.

Evidence for a tug of war also comes from work on vesicle fission, in Dictyostelium and rat liver cells, both in vitro and in vivo. In the case of fission, it is clear that a tug of war is happening because pulling by both motors is causing the endosome to stretch. In addition, in vivo trapping in both Dictyostelium and human epithelium cells has suggested a model in which dynein remains attached to the microtubule during plus- and minus-end-directed transport whereas kinesin is active during plus-end-directed motion. This “dynein-drag tug-of-war model” (Figure 8B) posits that, during kinesin-driven motion, dynein is dragged backward along the microtubule, effectively reducing kinesin’s stall force. However, during dynein-driven motion, kinesin is detached from the microtubule.

Coordinated motion also has a large amount of support. A huge array of accessory proteins affect intracellular transport, there are known regulatory factors that bias directionality, and there is a lack of competition between opposite-directed motors. The first two methods of coordination, accessory proteins and regulatory factors, do not necessarily exclude a tug of war—they could potentially modify the way the local tug of war works—but a lack of competition between motors seems to directly contradict any tug-of-war model. If the motors were not pulling against one another, how could a tug of war be occurring? For example, in many situations, eliminating a motor reduces motility in all directions, implying that there is a coordination factor that requires both motors to be present to initiate motility. If a tug of war were occurring, eliminating one motor would naively be expected to increase the motility of the opposite motor (eliminating kinesin, for example, would increase dynein-driven motion). When dynein or dynein function was disrupted in Drosophila embryos, plus-end-directed motility was adversely affected: Decreasing dynein-driven motion negatively affected kinesin-driven motion, opposite what would be expected in a tug of war. However, it is possible that impairing a motor in one direction could impair motion in all directions because of the presence of obstacles in vivo.16

Other experiments in Ustilago maydis (yeastlike fungus) and Xenopus melanophores have shown that down-regulating dynein or kinesin-driven motion had no effect on the opposite motor’s motility. This result clearly indicates that the motors are not interfering with one another. In vivo optical trapping also provides support for coordinated-motion models. In Drosophila embryos, organelles that are detached from microtubules using an optical trap tend to move in the same direction as they had been moving when they reactivated. This could indicate that only one set of motors is active at a time during transport, which clashes with the idea of both motors being active simultaneously, that is, a tug of war.

It is also clear that cells must have a way to regulate cargo directionality in the cell and that higher-order mechanisms other than motor-copy number and tug of war between motors might regulate transport. A comparison of the predictions of unregulated tug-of-war models (transport models in which only tug of war regulates directionality) to in vivo transport behavior revealed discrepancies, indicating that additional levels of regulation are required on top of tug of war. For example, changes in motor-copy number in Drosophila embryos had minimal effect on transport behavior, indicating that a mechanism other than tug of war regulates transport. In vivo trapping work with Chlamydomonas also indicated that coordinated motion occurs during intragelellar transport. In this case, large groups of motors of one type appeared to work together—stall forces of 50 pN were generated!—with no tug of war occurring. The generation of these large forces, with rare directional changes and saltations (commonly seen in other systems), suggests coordinated motion, with minimal competition between motor types. In the same study, the knockout of one motor was found to have no effect on transport by the other motor, again opposing tug of war.

Fu and Holzbaur examined another mammalian system, namely, mouse neurons, and found evidence for coordinated motion. They showed that phosphorylation of an adaptor protein, the JNK interacting protein 1 (JIP1), acted as a molecular switch to control the direction of axonal amyloid precursor protein (APP) cargo transport, involved in Alzheimer’s disease. When JIP1 was unphosphorylated, dynein was bound to the microtubule and kinesin was not; after JIP1 phosphorylation, the opposite was true, providing a clear example of coordinated motion, as kinesin and dynein were never simultaneously active.

Characteristics of the tug-of-war and coordinated-motion models are now being merged into more sophisticated models in which transport is regulated at the level of motor properties such as stall force, release force, microtubule binding and unbinding rates, and the relationship between load and motor velocity. These models assume a local tug of war in which motors engage stochastically with the microtubule, with random binding and force events determining directionality but motors rarely engaging in a prolonged tug of war. Instead, their properties are such that, when one set of motors has an upper hand, the other motors stop interfering with transport, either by unbinding or by simply getting pulled along behind the “winners”. These more complex models postulate an interplay between local tug-of-war interactions on the cargos and larger regulatory events, such as changes in motor number, motor properties (through phosphorylation or accessory protein binding), or even microtubule modification.

Conclusion: Evidence for local tugs of war occurring in most transport systems is very strong. Similarly, regulation of motility and directionality at higher levels than a local tug of war has been demonstrated in several systems. Some of the current debate between which model is correct is based on different groups focusing on different behaviors. In vitro systems with just kinesin and dynein clearly display tug-of-war behavior, but this simple behavior cannot explain all transport behavior in vivo. Most systems might have a tug-of-war method for regulating directionality, with a higher level of regulation and control set on top that controls motor number, type of motor, phosphorylation of motor(s), and so on. In addition, the diversity of transport systems also seems to indicate that both coordinated-motion systems and tug-of-war systems exist, with some model systems (Chlamydomonas, Ustilago maydis, APP transport) showing mostly coordination whereas others (A549 cells, Dictyostelium phagosomes, and others) show clear evidence of tug-of-war behavior.
2. CONTROLLING THE TRANSPORT COMPLEX FROM THE BOTTOM UP

This section describes techniques that have been used to study kinesin–dynein interactions in vitro, how these techniques work, and what results have been obtained using them. We have organized the discussion from the simplest, two- or three-component systems, to the most complex systems, consisting of large constructs or entire purified organelles.

Various motors and accessory proteins associated with the transport complex have been purified and are being added piecemeal to in vitro systems to study their effects in isolation. By altering the motor(s), accessory proteins, and cargo types, specific interactions between components of the transport complex can be observed. Tug-of-war interactions of groups of kinesin, dynein, and kinesin and dynein have been observed using this type of assay. These assays have strongly demonstrated that pairs of motors (kinesin and mammalian dynein or kinesin and yeast dynein) undergo a tug of war in vitro without external signals or cofactors. In addition, when cofactors are examined in vitro with a motor (e.g., dynactin–dynein), these cofactors modulate motor properties, which then could influence the tug of war between motors. Finally, different teams of a single motor type (teams of kinesin-only, dynein-only, or NCD-only) have been shown to have different cooperative behaviors.

Kinesin-only teams appear to be poor cooperators, particularly on fixed surfaces (that exist on polystyrene beads, for example). In contrast, dynein and NCD appear to be particularly good at sharing the load equally between motors to generate forces greater than that of a single motor. This impacts dual-motor transport in that teams of dynein would apparently be able to overwhelm kinesin in a tug of war, another method of modulating transport behavior.

2.1. Mixing Multiple Motors in Vitro: Tug-of-War Motion, Cofactors, and Teams

Single-motor behavior for kinesin and dynein has been individually well characterized using various fluorescence and force spectroscopy techniques. Stepping behavior, force–velocity curves, and interactions of various structural elements in the motors have been observed. More remains to be done, but observations of the interplay between multiple motors in simplified in vitro environments have also started to reveal interesting information about the motors.

The most basic dual-motor experiments began with gliding assays, in which kinesin and dynein were attached to a surface and the gliding of microtubules over the surface was observed. In 1992, Vale et al. showed that coupling kinesin and dynein through a microtubule led not to stalled motion but instead to bidirectional motility of the microtubule, with stochastic directional switching (Figure 3A). The motor–motor force interactions through the microtubule affected motor binding and unbinding events, leading to the hypothesis that the motors’ mechanical properties, as opposed to outside activating factors, played a key role in determining directionality. In addition, motor density on the surface tightly regulated the directionality of the microtubules. That is, more kinesin led to more plus-end-directed motion (Figure 3B), and the velocity of microtubule gliding was decreased when opposite-polarity motors were present. This is clear evidence that a tug of war was occurring.

Gliding assays have also shown how two different kinesin motors (OSM-II and kinesin-2 from C. elegens) interact to drive plus-end-directed intraflagellar transport (IFT). These results support previous in vivo data indicating that OSM-II and kinesin-2 are simultaneously active during plus-end-directed IFT. When the ratios of the two motors attached to the surface were varied, the motors were able to continuously vary the velocity of the gliding microtubules between the velocities of the individual motors. This finding strongly indicated that, during IFT, the motors were undergoing a mechanical competition to drive plus-end-directed motion. Therefore, a coordination mechanism beyond mechanical coupling through the cargo is unnecessary for two different kinesin motors to drive plus-end-directed motion, that is, a tug of war can also drive interactions between motors going in the same direction.

A somewhat more physiological situation, with a bead labeled with kinesin and dynein walking on microtubules attached to a coverslip surface, was also tested. In experiments by Muresan et al. using latex beads coated with kinesin and dynein, the beads always walked in the kinesin direction, that is, bidirectional motility was not observed. However, when kinesin was inhibited (by an antikinesin antibody), dynein would take over, indicating that directionality was regulated by the presence of kinesin. Similarly, DeBerg et al. observed the motility of polystyrene beads coated with kinesin and dynein (without dynactin). They, however, found that the cargos underwent bidirectional, salatory motion (motion with directional reversals and repeated pauses; see Figure 3C). The direction of motion and stall-force behavior could be biased by altering the ratio of dynein and kinesin but only within a fairly tight range. This indicates that one way to regulate directionality in a tug of war is by altering the motor ratio and that the ratio of kinesin to dynein must be tightly controlled if bidirectional motion is to occur.

In vitro stall-force measurements were also made on beads with kinesin and dynein attached by Blehm et al., who used an optical trap to measure stall forces in both the plus and minus directions. A single kinesin and also a single dynein on a bead showed the “normal” stall forces of individual kinesin and dynein (6 and 1 pN, respectively; Figure 8A,C). A bead pulled by kinesin and dynein, however, showed a reduced stall force in the plus direction (that is, with kinesin “winning’) and a stall force equal to or greater than that of a single dynein in the minus, dynein-driven direction (presumably because multiple dyneins were working together; see Figure 8D). These results indicate that, when moving toward the microtubule plus end, kinesin has to drag dynein behind it, with dynein presumably still bound to the microtubule even though dynein is forced to move backwards. When the cargo is moving toward the microtubule minus end, dynein operates freely as a team, presumably with kinesin detached from the microtubule—that is, the kinesin(s) adds no drag. These results provide support for a dynein-drag tug-of-war model (see Figure 8E).

2.1.1. Cofactors. Several elaborations on in vitro motor–cargo systems have also been explored. For example, cofactors, such as the dynactin complex and Lis1, a regulator of dynein, have been added to in vitro systems. These cofactors have been shown to have significant effects on the behavior of dynein alone and have also been suggested to link kinesin and dynein during intracellular transport. Lis1 appears to act as a “clutch”, causing dynein to remain attached to the microtubule for extended periods of time, particularly increasing its binding time (time attached to the microtubule) under load. This behavior would theoretically
help teams of dynein motors apply large forces for the movement of large objects such as nuclei.

Dynactin was shown to increase dynein’s processivity and enhance microtubule binding in vitro, for dynein from yeast and from chick embryo brains. In addition, Ross et al. showed that dynein with dynactin in vitro can undergo bidirectional motion. This surprising result was suggested to be a method of modifying dynein’s properties so as to allow obstacles on microtubules to be bypassed. However, experiments by Ross et al. also showed that groups of dynein–dynactin complexes displayed only unidirectional motion. This finding indicates that multiple dyneins working together (thought to be typical) would not display bidirectional behavior without kinesin present. In addition, it is possible that the plus-end-directed motion of the dynein–dynactin complex is diffusive, although its velocity was ATP-dependent. If the plus-end-directed motion were diffusive, dynactin–dynein complexes could not drive net plus-end-directed motion, indicating that dynactin acts merely as a tether to keep dynein attached to the microtubule. Dynactin’s effects on dynein and bidirectional motility in general are still unclear, although they are definitely significant.

Another cofactor is JIP1, which interferes with kinesin autoinhibition in cultured mouse neurons, thereby activating kinesin when bound to it in vitro. In addition, when p150Glued (a dynactin subunit) binds to JIP1, it counteracts JIP1’s effects on kinesin, inhibiting it again. Thus, it appears that JIP1 and p150Glued can act as a switch to regulate transport. This is a clear case of a cofactor directing coordinated motion. This cofactor can turn kinesin on and off, thereby preventing it from being simultaneously active with dynein.

These experiments show that cofactor modulation of intracellular transport through the alteration of motor properties appears to be a very significant method of transport regulation. It is particularly important for dynein, as there is only one type of cytoplasmic dynein whereas there are many types of kinesins. Different kinesins could therefore be used in different situations in the cell, but a single type of cytoplasmic dynein must play several roles, and cofactors could help modify its behavior. Cofactor modulation is an example of a higher-order method of regulating motion, with a tug of war often occurring between the local motors on the organelle while cofactors modulate the individual motors’ behavior so as to bias the outcome of the tug of war or even, as in the case of JIP1, prevent a tug of war from occurring altogether. 2.1.2. Multiple Motors. The interaction between several motors of the same type is also of interest, as it appears that many cargos carry several kinesins and dyneins. Whether dyneins cooperate with other dyneins, and kinesins with kinesins, will have a large effect on transport. These same-motor interactions could potentially enhance or impair cargo transport by a single motor type. This, in turn, could be an important method of influencing the outcome of a tug of war. For example, it is well-known that increasing the number of dyneins or kinesins can increase processivity under conditions of no load. However, when the motors have to share loads and actively exert force, how well they cooperate is not clear. Several articles have shown that multiple kinesins do not cooperate well whereas multiple dyneins work extremely well as a team. In addition, a minus-end-directed kinesin, NCD, also behaves cooperatively as a group. The cooperativity within groups of a single type of motor now appears to be a major component of the tug-of-war model, with the significant differences between kinesin and mammalian dynein playing a large part in the intracellular tug of war.

Why dynein is more cooperative as a group than kinesin is driven by the fact that a single dynein can function as a gear in response to load, taking smaller steps as the load increases. To show this functionality, Mallik et al. applied force to dynein attached to a bead (using an optical trap in vitro). As the force increased, the step size of dynein decreased (from 32 to 8 nm). This allowed motors with reduced loads to catch up and increase their loads, so as to share the load more equitably. In contrast, kinesins reduce its velocity under load (while its step size remains constant) but not in the same fashion as dynein. Consequently, kinesins do not cooperate well with other kinesins to generate large forces. However, some studies have shown that kinesin can cooperate more effectively at lower velocities, achieved in this case by lowering the ATP concentration. The fact that dynein is more cooperative as a team than kinesin is another high-order method for regulating the tug-of-war interaction between the two motor types. A single kinesin is likely to win out over a few dynein motors, but as the number of dynein motors increases, their ability to cooperate will overwhelm the kinesin, driving motion toward the minus end of microtubules.

Conclusion: Simple in vitro systems consisting of just kinesin, dynein, and cargo, with no external coordinating factors, have shown that bidirectional motility is possible without a coordinating complex and that a tug-of-war mechanism is how motors interact in vitro. Additional studies on groups of either dyneins or kinesins have revealed how they cooperate and the potential impact this will have on a directional tug of war. Kinesin typically wins tugs of war with a dynein, but because dyneins are better at working as a team, enough dynein can overwhelm a kinesin. Finally, the effects of cofactors and how they modulate motor behavior have also been studied, such as the effects of dynactin, Lis1, and JIP1 on dynein, indicating that accessory proteins can significantly modulate motor behavior in vitro. These in vitro experiments have clearly shown that, although a tug of war occurs with just kinesin and dynein present, cofactors and motor-copy number are means of modifying the outcome of a tug of war, sometimes completely eliminating it (JIP1), and are therefore higher-order methods of regulating cargo directionality.

2.2. Beyond Beads: Synthetic Cargos

It is possible that some of the in vitro systems used to study motors create artifacts because of their artificial natures. In particular, beads restrain the bound motors to remain in one spot along the bead surface, whereas on lipid vesicles and presumably organelles in the cell, the free flow of motors allows multiple motors to be recruited to one spot on a vesicle. In addition, the number of motors bound to beads is very difficult to ascertain, as even at a specific concentration of motors, different numbers will bind to different beads and only some of the motors will be bound to microtubules at one time. To overcome these issues, synthetic cargos have been developed to more closely mimic in vivo cargos, and others have been created to control the number and type of motors present.

2.2.1. Motor Diffusion on Cargo Surfaces. The idea that the properties of the cargo influence kinesin cooperativity was demonstrated recently on giant unilamellar vesicles coated with kinesin. Studies in which kinesins were firmly tethered in place (attached to a solid surface, such as a polystyrene bead)
showed poor kinesin-1 cooperativity in vitro. Here again, we mean the ability of motors of a single type to work together (as opposed to kinesin and dynein cooperating with each other). However, on giant unilamellar vesicles, kinesins were recruited to the microtubule-binding site, as they freely diffused around the vesicle until they attached to the microtubule. This increased the number of available motors at the microtubule and enhanced their cooperativity. Other evidence that motor cooperativity can be altered by the cargo’s properties is shown in cooperative pulling of nanotubes, microtubule gliding, and intraflagellar transport. These studies all showed that groups of kinesins would generate forces greater than a single kinesin could against different types of loads (membrane elasticity, magnetic traps, and optical traps, respectively).

2.2.2. Building a Scaffold to Control Motor Number. A second major concern about bead experiments (and motor experiments in general) is that there is currently no good way to determine the number of motors on the bead or the number of motors that are active. Attempts to determine these numbers have been made by using stall-force assays, with the assumption that stall force increases more or less linearly with motor number (for a single motor type). However, it can be difficult to take stall-force measurements in every instance, as the number of active motors can vary in a single sample with a single concentration of motors, along with a large number of other issues that can occur (see Figure 4A). In addition, some recent studies have shown that single-motor stall forces are not as additive as one might expect, and multimotor assays display a much more complex picture due to interactions between kinesin and dynein affecting the stall force. For example, if a cargo had two kinesins (2 × 7 pN = 14 pN stall force) that were dragging seven dyneins (7 × 1 pN = 7 pN), a net stall force of 7 pN might be measured. This cannot be differentiated from a single kinesin (7 pN) using stall-force measurements. Without knowing the number of active motors present, understanding motor cooperativity and interaction is very difficult.

The first step in elucidating the number of active motors is to control the number of motors present on the cargo. Various scaffolds have been tried to replace beads as artificial cargo (Figure 4B). An early scaffold (Figure 4C), made of protein linkers and “springs”, was built to study the effects of elastic coupling and linker distance on transport by multiple monomeric kinesin motors. In that work, the elastic coupling and linker distance did not play a large role, increasing the velocity and processivity of the kinesin monomers only slightly. Another group used DNA to tether two kinesin monomers together to determine whether the structure of the kinesin outside the motor domain was necessary for processivity. They determined that the neck linker was important (removal abolishes processivity) and that intermotor strain allows processive motion. The native kinesin coiled-coil structure was not necessary for processivity but was the most efficient of all alternatives tested.

2.2.3. Kinesin and NCD on a DNA Scaffold. Whereas previous studies focused on kinesin monomers, a study by Furuta et al. in 2013 focused on dimeric motors, namely, kinesin-1 and NCD, a minus-end-directed kinesin. They used DNA as a scaffold and provided evidence for a “dragging” tug-of-war model, with NCD being dragged rather than dynein. Short pieces of double-stranded DNA with binding-site tags and fluorophores were used to stitch together groups of motors in a line (Figure 5A). Each motor was connected to the ones around it with a short, stiff DNA linker, and the behaviors of the various motor assemblies were observed. The linkers were flexible at the motor attachment site, allowing some freedom of movement for the motors. As the number of motors increased, the processivity of both kinesin and NCD increased, but only NCD’s velocity increased (and then mostly at the transition from one to two motors, with minimal increase thereafter). Force production was similar, as NCD’s stall force was positively correlated with the number of motors whereas kinesin’s showed almost no correlation. Kinesins sometimes cooperated to generate stall forces larger than that of a single kinesin, but they were much less likely to do so than NCD. These experiments indicated that NCD cooperates well in
groups whereas kinesin gains little advantage from increasing motor number.

A final experiment by Furuta et al. tethered kinesin and NCD together with the DNA scaffold previously used to couple together only kinesin or only NCD (Figure 5A). The ratios of the motors (kinesin/NCD) were changed, and the behaviors of coupled plus- and minus-end-directed motors were observed. These experiments showed that the velocity of plus-end-directed motion decreased with increasing numbers of NCD motors, indicating that some sort of tug of war was going on, reducing kinesin’s velocity. It is interesting that NCD, as a weak, cooperative, minus-end-directed motor, behaved so similarly to mammalian dynein when coupled to kinesin. This is evidence that motors with certain properties function together better in tug-of-war situations, as NCD and mammalian dynein behave similarly and both display bidirectional motility when coupled with kinesin.

2.2.4. Kinesin and Yeast Dynein on a DNA Origami Scaffold. A final, more complex, but also more versatile, synthetic cargo also uses DNA, in the form of DNA “origami”. This allowed the construction of more or less arbitrary structures that can be used to determine specific numbers, types, and placements of binding sites on a single cargo, up to 90 unique sites on the DNA construct demonstrated in this article. In this technique, a large, cylindrical scaffold consisting of 12 helical pieces of DNA was the synthetic cargo, and 21-base-pair DNA handles were left at specific sites of the scaffold (Figure 5B). Then, appropriate antihandles could be attached to the motors of interest (in this case, kinesin-1 and yeast dynein). The effects of increasing numbers of kinesin and yeast dynein were observed, and run length increased with motor number, whereas velocity did not change (kinesin) or decreased (yeast dynein) with increasing number. In these experiments, mixed-motor ensembles of kinesin and yeast dynein were routinely immobile; when motile, they moved more slowly than single-motor ensembles, and yeast dynein in general “won” over (mammalian) kinesin in terms of the direction of motility. This is very different from (mammalian) kinesin’s interaction with mammalian dynein.

2.2.5. Issues with DNA Scaffolds and Origami. There is a significant issue with the DNA origami technique, in that, currently, the binding sites available for each motor type are not completely filled. The authors saw about 80% occupancy of each site, leaving uncertainty in the actual number of motors present on the complex. This occupancy issue could be due to the fact the authors attached DNA linkers to the motors (with a SNAP-tag), which were then annealed to the chassis. Therefore the final step was the binding of an oligonucleotide already attached to the motor. When using kinesin and NCD held together by a DNA scaffold, however, Furuta et al. annealed the whole scaffold together and then attached the motors with SNAP-tags, which showed essentially 100% occupancy. Furuta et al. did have difficulty obtaining full occupancy with HaloTags, however. In addition, although the DNA for both of these techniques is commercially available, it still requires design and assembly, which appears to be quite complicated, in particular for the larger origami structures.

2.2.6. Dynein’s Stall Force. Another important note about the experiments by Derr et al. and other experiments is that, in vitro, single yeast dynein and single mammalian dynein have been shown to have very different behaviors. A single yeast dynein has been shown to be a slow, strong (7-pN stall force) motor in vitro, while a single mammalian dynein in vitro has been measured to be weak (1–2 pN) and of a similar speed to mammalian kinesin. Mammalian dynein’s stall force

Figure 5. Synthetic DNA cargos. (A) Schematic representation of DNA–motor construction (not drawn to scale). The typical spacing between motors is 22.7 nm, and the lengths of kinesin, SNAP-tag, and HaloTag are ~17, 4.3, and 4.8 nm, respectively. Note that the DNA was fully ligated and annealed before the motors were attached enzymatically with a HaloTag or SNAP-tag. By altering the number of each tag, the motor ratio could be controlled. Caption and figure adapted with permission from ref 28. Copyright 2006 American Association for the Advancement of Science. (B) Schematic of the 12-helix-bundle chassis structure with 6 inner and 6 outer helices. Each outer helix contains up to 15 optional handles, yielding 90 uniquely addressable sites. Each handle consists of an unpaired 21-bp (~7-nm) oligonucleotide sequence for hybridization to complementary antihandles covalently attached to motors or fluorophores. The inset shows an orthogonal cross section. The chassis is substantially larger and more complex than in the work of Furuta et al., although it is more customizable. (C) Schematic of a chassis labeled with five fluorophores (red) at handle position 14 on each of five outer helices and dynein at handles at positions 1, 5, 9, and 13 on a single outer helix. Oligonucleotide-labeled dynein is also shown. Note that the motors are attached to a piece of single-stranded DNA through a SNAP-tag and then the DNA is attached to the chassis. Figure and caption adapted with permission from ref 11. Copyright 2012 American Association for the Advancement of Science.
and behavior are disputed: Some contend it has a stall force of 5–7 pN.\textsuperscript{39} A substantial comparison of mammalian and yeast dynein will not be undertaken here as it is outside the purview of this review. However, we note that, in the experiments with yeast dynein and mammalian kinesin, it was routine for cargos to be stationary as the motors were engaged in a balanced tug of war, preventing motility.\textsuperscript{15} This might indicate that these motors’ properties are not well balanced for each other. This is in contrast to mammalian kinesin and mammalian dynein, which rarely stall as a result of a tug of war in vitro and appear to have complementary properties: Dynein is dragged by kinesin, but it is weaker and can cooperate in groups as opposed yeast dynein.\textsuperscript{12a,b,26,39}

Conclusion: Synthetic cargos have shown that groups of only kinesin motors can cooperate if they are attached to fluid membranes that allow free motor diffusion.\textsuperscript{41} In addition, synthetic DNA cargos can be constructed that control the number of motors attached (along with allowing detachment with specific signals) and have clearly shown that a tug of war occurs between kinesin and dynein.\textsuperscript{11,28} Interestingly, dynein with different properties (yeast vs mammalian) display extremely different behaviors, indicating that motor properties (stall force, velocity, detachment force) play a significant role in regulating intracellular transport.

2.3. Cellular Organelles in Vitro: Examining Parts of the Transport Complex

Purifying intact organelles with the entire transport complex present and active and examining their behavior is another way to observe transport behavior in a controlled environment. However, it is difficult to ensure that all factors involved in transport are present and functional. Nonetheless, even organelles with part of the transport complex can reveal useful information about intracellular transport. Purified organelles have shown tug-of-war behavior, and the viscoelasticity of the cellular cytoplasm has been found to have minimal effect on organelle transport.\textsuperscript{52,12a,b,50}

The power of examining intact organelles can be seen in that kinesin and cytoplasmic dynein were both discovered as the motors that powered directional transport using this technique.\textsuperscript{51} Organelles have long been purified and studied to see which proteins will copurify with them, in an attempt to determine what proteins are part of the transport complex. One of the first nonmotor parts of the transport complex identified was dynactin, a separate protein from dynein but one that evidently plays an important role in the transport complex. It was examined by in vitro motility assays.\textsuperscript{52} A more recent example is in vitro studies of organelle fission properties. For example, studies on the transition from early- to late-endocytic vesicles have compared the motile properties of the two populations to characterize the difference in proteins present in the transport complex.\textsuperscript{15a} Similarly, Huntingtin protein has been shown to be necessary for dynein-mediated transport in vitro.\textsuperscript{53} Herpes simplex virus transport in vitro was shown to be predominantly mediated by kinesin and associated with the trans-Golgi network marker TGN46.\textsuperscript{54}

Purified organelles can also display motile behavior very similar to that seen in the cell, with bidirectional, saltatory motion.\textsuperscript{76} Some of the first evidence for a tug of war was shown using purified organelles.\textsuperscript{23b} That study showed that the presence of kinesin determined directionality. Only plus-end-directed purified vesicles had kinesin present, whereas both plus- and minus-end-directed vesicles had dynein. Furthermore, when kinesin was inhibited, plus-end-directed vesicles uniformly became minus-end-directed. The study saw no bidirectional movement, but clearly some sort of local tug of war or interaction between kinesin and dynein was occurring, as similar behavior occurred when the motors were bound to beads.\textsuperscript{23b}

Optical trapping of purified organelles has recently been used to measure the behavior of the transport complex without the interference of the highly viscoelastic and complex cellular environment. Organelles purified from a wide array of cells, including Dictyostelium,\textsuperscript{55} A549,\textsuperscript{12a} and mouse macrophages, J774A.1,\textsuperscript{12b} have been trapped. Different methods of purification led to different parts of the motor complex being present; specifically, some of the harsher purification techniques stripped away dynein,\textsuperscript{12a,b} whereas others did not.\textsuperscript{2b} Trapping of these organelles has shown that kinesin’s stall force (with dynein stripped away) is not affected by the remaining transport complex, but kinesin and dynein together on a cargo can interact in surprising ways, leading at times to stretching of the organelle and effectively reducing kinesin’s stall force.\textsuperscript{12a,b} The stretching reveals that kinesin and dynein do engage in a tug of war, as only kinesin and dynein pulling simultaneously would be able to cause the stretching of the organelle.

2.3.1. Effect of Viscoelasticity in a Cell. In addition, in two independent studies, it was found that there is little difference in the forces exerted by purified organelles that display bidirectional motion in vitro and their behavior in vivo.\textsuperscript{12a,b,15b} Given that viscoelastic behavior is extremely different in vitro and in vivo, this is a surprising result. (Viscoelastic moduli of a cell range from $10^2$ to $10^5$ dyn/cm\textsuperscript{2}, leading to viscoelasticities several orders of magnitude higher than that of water, which has a viscosity of 1 cP and essentially no elasticity.\textsuperscript{50–57}) Apparently the high viscoelasticity of the cell has minimal effect on transport.\textsuperscript{12a,15b,50}

The interplay of the components of the transport complex appears to be the major factor determining transport behavior (see also section 3.3). Organelles purified with only one motor behave similarly to single-motor experiments. In contrast, having both motors present on purified cargos can lead to stretching of the cargo and a reduction in kinesin-driven stall forces. Because the simple addition of dynein causes stretching of the cargo and reduces kinesin’s stall force, the motors must be engaging one another in a tug of war. When directional motion occurs with purified organelles, the lack of a difference between their in vivo and in vitro force behaviors also indicates that external factors to regulate motility are not necessary for short-range bidirectional motility. The standardization of purification techniques\textsuperscript{58} and their increasing use is creating a middle ground between in vivo and in vitro studies. Purified organelles analyzed by in vitro techniques might be more suited to exploring the exact components of the transport complex and, thus, lead to clearer conclusions than can be extracted from in vivo studies.

Overall, the preponderance of in vitro data indicates that a tug of war occurs between motors present on in vitro cargos.\textsuperscript{11,12,23b,28} Motors’ properties, particularly those of dynein, are regulated by the presence of regulatory cofactors, and this regulation could potentially influence the outcome of a tug of war.\textsuperscript{52,24a,27b,34,36} In addition, the outcome of the tug of war can be determined by the number of each motor type present and how well these motors can cooperate among

\textsuperscript{dx.doi.org/10.1021/cr4005555 / Chem. Rev. XXXX, XXX, XXX--XXX}
themselves.11−13,23b,26,28 Assays with motor-coated beads,12a,13 DNA scaffolds,11,28 and purified organelles7d,12a,b all support these conclusions.

3. TRANSPORT COMPLEX IN VIVO: EXAMINING ORGANELLES IN LIVING CELLS

Observation of organelle motility in the living cell is a huge area with decades of research behind it. A great variety of techniques exist for observing organelle behavior, from basic light microscopy, phase contrast and differential interference contrast (DIC), to modern fluorescence (total-internal-reflection fluorescence microscopy, TIRFM,59 and fluorescence recovery after photobleaching, FRAP)60 and super-resolution techniques (stimulated emission depletion, STED,61 and structured illumination microscopy, SIM).62 In vivo optical trapping studies have shown that a tug of war is likely occurring7d,12a,b but have also revealed that higher-order mechanisms must exist to regulate and coordinate motor behavior in the cell.7e,20

3.1. Tracking Organelles in Living Cells

Recent discoveries in the area of kinesin−dynein interactions in living cells have revealed a wide variety of behaviors across organelle and cell types. How cells regulate directionality is one of the major questions about bidirectional intracellular transport, and the experiments discussed in this section cover the wide variety of mechanisms observed in the cell: motor concentration, opposite polarity motors, myosin, regulatory factors, and external signaling. Many of these mechanisms might be unique to a specific cellular system, whereas others might be more broadly applied.

3.1.1. Motor Concentration Regulates Tugs of War. During endosome transport in the fungus Ustilago maydis, kinesin-1, kinesin-3, and dynein, have been shown to exhibit a complex interaction that alternates between cooperation and competition.18b Dynein dominates transport near the cell tip, but kinesin-3 takes over the cell tip for long-range transport, necessitating a cooperative “hand-off” between the motors for transport across the cell.18c However, near the plus ends of microtubules, motion driven by kinesin-3 toward the plus end appears to be stopped by a high dynein concentration. The high dynein concentration eventually overwhelms kinesin-3 and prevents it from running endosomes off the plus end. As only minus-end-directed endosomes had dynein present, whereas both plus- and minus-end-directed endosomes had kinesin-3, the presence of dynein apparently regulates directionality in this system.63 This high concentration of dynein is maintained by dynein transport to the plus end of the microtubule by kinesin-1.1,64 This supports a model in which the cell regulates the dynein concentration, thereby regulating overall motion. Locally, at the cargo, the decision of which direction to travel is driven by a tug of war in which net force or motor number is intermittently tested to determine directionality.

3.1.2. Opposite Polarity Motors Are Both Required for in Vivo Transport. In Drosophila S2 cells, it has been shown that opposite-polarity motors must both be present on certain organelles for motility to occur.65 However, it apparently does not matter what the specific motor is.66b During the tracking of fluorescently labeled peroxisomes (GFP was targeted to the organelles), it was observed that knocking out kinesin-1 or cytoplasmic dynein caused all motility to stop. Knocking out kinesin-1 and replacing it with kinesin-3 (specifically unc104 from Caenorhabditis elegans, in this case) restored motility.65b Similarly, replacing dynein with NCD, a minus-end-directed kinesin, would restore motility. (In this case, motility was heavily plus-end-biased, as NCD has different properties than dynein.) If kinesin or dynein was replaced by a nonmotile mutant, motility would not resume. This indicates that active plus- and minus-end-directed motors are simultaneously required during intracellular transport.65b What this means is unclear. It could indicate that there is a coordinating complex that requires the presence of both motors to function.

3.1.3. Kinesin and Dynein Interact to Bypass Obstacles. Another potential reason for the simultaneous presence of kinesin and dynein on many cargos is that they might aid in bypassing obstacles on microtubules, such as tau protein patches and microtubule intersections.13,23b,26,28 Assays with motor-coated beads,12a,13 research on nuclear migration in C. elegans has indicated that, even though kinesin is the driver for nuclear movement, the deletion of dynein seriously slows and impairs transport, likely because of the inability to back the nucleus up to bypass obstacles.67 Also, kinesin and dynein are differentially regulated by a microtubule-associated protein, tau. Kinesin generally detaches in the presence of tau-decorated microtubules, whereas dynein reverses at patches of tau. This indicates that dynein and kinesin interact to allow more efficient intracellular transport and could explain why knocking out one motor stops transport in all directions.

3.1.4. Myosin Can Modulate Microtubule-Based Transport. Myosin is often also present on organelles with kinesin and dynein, and its interaction with microtubule-based transport is an open question. Mitochondrial motility on microtubules in Drosophila neurons appears to be inhibited by myosin V in both directions and by myosin VI during retrograde transport, whereas myosin II has no effect on microtubule-based transport. This appears to indicate that myosin inhibits long-range microtubule-based motility and might aid organelle docking and pausing.68 Myosin’s effect on transport is still unclear, but it could have an override effect on kinesin−dynein tugs of war, with myosin perhaps superseding kinesin and dynein to allow switching to actin-based transport.

3.1.5. Regulatory Factors Modulate Transport. A level of complexity typically present only in vivo is external regulatory factors, such as phosphatases and kinases. For example, in the Xenopus laevis melanophore system, melanosomes (pigment granules) can be dispersed or aggregated by external signals.69 Recently, it was shown that this control of directionality is mediated by phosphorylation of the dynein intermediate chain, where phosphorylation of dynein apparently stimulates its activity.70 This type of regulation has also been seen in Huntington’s disease, as one of the downstream effects of pathogenic huntingtin protein is kinesin phosphorylation, which appears to inhibit kinesin.71 In addition, phosphorylation of the scaffolding protein JIP-1 modulates the directionality of amyloid precursor protein (APP) motility in mouse axons. JIP-1 phosphorylation may fully coordinate APP motility, as experiments indicated that kinesin and dynein were not simultaneously active in this system.22b

3.1.6. Controlling in Vivo Motility with External Signals. An exciting new method for studying intracellular transport is the creation of an in vivo transport system that can be induced to move upon addition of an external ligand. This allows much greater control of in vivo transport and control over the motors attached to the tracked organelle. This creates an artificial system where the scientist has control over an
Figure 6. Inducible intracellular motility assays. (A) Assay consists of a fusion construct of PEX, RFP, and FKBP that targets peroxisomes. This PEX−RFP−FKBP construct consists of PEX, a peroxisome membrane targeting signal; RFP, a red-fluorescent protein; and FKBP, a domain that can be cross-linked to an FRB domain using rapamycin analogues. This PEX construct is recruited to peroxisomes, and consequently, (B−D) fusions of FRB with motor constructs (1, 2, 4, 5, 7, and 8) or adaptor protein fragments (3, 6, and 9) are recruited to FKBP and the peroxisomes upon addition of rapalog. This ability to target specific motors to receptors on the peroxisome allows for the control of intracellular motility. Caption and figure adapted with permission from ref 73. Copyright 2010 Elsevier.

external regulatory factor, the motors to be regulated, and the timing of transport. One such system is peroxisomes, which can be targeted by labeling motors with a 93 aa FRB domain (see Figure 6). By addition of an FRB domain to endogenous or exogenous kinesins and dyneins, motility could be stimulated through the addition of rapalog, which cross-links the motor to the peroxisome. As was expected, when FRB-labeled kinesin was present, rapalog led to the rapid in vivo dispersal of peroxisomes, whereas aggregation at the center occurred for FRB-labeled dynein.73 Currently, the effects of rapalog addition are irreversible, and only one binding motif is present, so only one binding event can be controlled (although it was possible to produce bidirectional motility by stimulating kinesin and dynein simultaneously). In addition, the article showed that peroxisomes with only a single motor type attached (kinesin or dynein) underwent motion (the native motors caused only minimal motility). They also showed that kinesin- or dynein-only peroxisomes underwent pauses and changed directions at microtubule intersections, indicating that saltatory motion is not necessarily indicative of a tug of war. The ability to control transport with exo- or endogenous motor proteins is quite promising, as the construction of arbitrary in vivo transport systems could reveal much about intracellular transport.

A similar system was also used to probe transport behavior in dendrites.74 Microtubules are of mixed orientations in dendrites (as compared to axons, where they all run in parallel), and it was unclear how transport to the dendrite was targeted. Recruiting motors to specific cargos showed that dynein preferentially drives cargos to the dendrite but also moves bidirectionally in dendrites, presumably by switching between oppositely oriented microtubules. Because of the small size of the dendrite, this random, bidirectional motion appeared to be capable of maintaining a stable density of cargos in the dendrite. Recruiting kinesin (Kif5 and Kif17, conventional kinesin and another kinesin family member) appeared to increase axonal transport but had minimal effect on dendritic transport.74 In this system, the recruitment of a specific motor type appeared to regulate cargo placement.75 This is a clear system in which both motor types do not appear to be required for transport and could possibly be an example of an exclusionary coordination mechanism. Directionality is controlled by the fact that only one motor type is present at a time.

Conclusion: These in vivo studies have shown how the cell biases motion in one direction or the other. In a couple of cases, the cell appears to completely coordinate motion.22,74 However, in general, the cell appears to modulate a tug of war by modifying the number of motors,63 the properties of the motors (through phosphorylation),71 or even the types of motors (dynein, myosin, or kinesin).68,74 In addition, both kinesin and dynein appear to be necessary for intracellular transport in certain systems,65b although evidence to the contrary has been found in artificial in vivo systems.73 This could be due to the regulation of motor behavior by microtubule-associated proteins such as tau.56 These sorts of mechanisms can determine overall directional bias, whereas a local tug of war between different motor types determines directionality of a specific cargo.

3.2. New Techniques for Tracking Organelles in Living Cells

In addition to standard bright-field and fluorescence techniques, new methods and super-resolution techniques are being applied to the tracking of organelle motility in living cells. These techniques all have their advantages and disadvantages, with many techniques increasing spatial resolution at the expense of temporal resolution or increasing the lifetime or brightness of an imaged molecule at the expense of increasing the difficulty of labeling or interfering with the labeled molecule’s behavior.

The first new technique that is just beginning to be applied to in vivo transport is STED microscopy. This technique uses a focused beam to excite fluorescent molecules and a second donut-shaped beam to stimulate emission around the excited spot at a slightly longer wavelength than normal fluorescence. Ignoring this stimulated emission (cutting out the longer
motor stepping, revealing that motors step similarly in vivo as they do in vitro.\textsuperscript{82} All of these techniques, however, involve serious tradeoffs. Both STED and SIM trade temporal resolution for spatial resolution. STED, being a scanning technique, allows high speed (full frame rate of 28 Hz) scanning of small areas at high spatial resolutions (50 nm), but it is fairly complex to setup and loses time resolution as one scans larger areas, similar to confocal microscopy.\textsuperscript{76,77} SIM, on the other hand, can take large wide-field images at the same rate as smaller image areas, but it is limited to a doubling of resolution and requires significant postprocessing.\textsuperscript{7} New labels can increase photo-stability and brightness but come with their own costs. Most are significantly larger than fluorophores (quantum dots, gold nanoparticles, UCNPs) and present significant difficulties when labeling.\textsuperscript{82} bFIONA is inherently limited to absorbent particles (or organelles).\textsuperscript{174}

3.3. Optical Trapping in Vivo

Although one of the first uses of an optical trap was in a living cell, when Ashkin et al. attempted to measure the forces exerted by mitochondria,\textsuperscript{84} over the next 20 years, trapping was rarely used in the cell. This was because making quantitative force measurements in the cell was extremely difficult because of the complex nature of the cellular environment; the initial article made many assumptions that might or might not have been appropriate when taking in vivo measurements. In the cell, the viscous and elastic properties of the cytoplasm are unknown and can vary both temporally and spatially, making experimental results difficult to interpret. In contrast, in vitro, the spring constant of the trap (the stiffness) can be easily measured, largely because water’s viscous properties are constant and well-known, yielding nanometer and subnanometer results with a millisecond or greater time resolution.\textsuperscript{85}

3.3.1. Optical Trap Calibration. In the past few years, in vivo optical trapping has been rejuvenated thanks to new calibration techniques.\textsuperscript{7,12a,b,15b,26,86} As a result, new questions about motor transport in the cell can now be answered.\textsuperscript{87} The spring constant or stiffness ($k$) of an optical trap in a purely viscous medium (i.e., an in vitro optical trap) can be calibrated using a wide variety of methods. All of these methods measure essentially the same thing: the strength of the trap when compared to a known force acting on it. This can be done by measuring the trap stiffness against viscous drag (drag method) or by measuring the trap stiffness against Brownian motion (by either the equipartition method or the power spectrum method).\textsuperscript{88} For a more thorough review of these techniques, see Neuman et al.\textsuperscript{85a}

The most common technique used to calibrate traps in living cells is to estimate the trap stiffness by measuring the index of refraction of the cellular cytoplasm, the index of refraction, and the size of the trapped organelle and then to create a viscous, in vitro environment that replicates these indices of refraction, and measure the trap stiffness in it.\textsuperscript{7,19,26,86} In this index-of-refraction matched environment, standard in vitro calibration techniques are used. Then one creates an organelle size versus stiffness calibration in this in vitro environment. After that, the stiffness of the trap is assumed to be directly related to the size of the organelle being trapped, that is, trap stiffness is linearly related to $r$, the radius of the trapped object (assumed to be spherical). This technique is more straightforward than the actual in vivo calibrations described later, but it suffers from the fact that it relies on many more assumptions. This type of technique and the in vivo calibration technique described below...
appear to measure very similar trap stiffnesses, indicating that this technique might be sufficient for most measurements. Actual results (force measurements) are quite variable between differing calibrations and systems, however, so it is difficult to determine how well various techniques work relative to each other.

For fully in vivo optical trap calibration, the assumptions going into a calibration must be minimized. There are two different in vivo calibration techniques that were developed and tested in vitro and in vivo and are based on the fluctuation dissipation theorem (FDT). Compared to in vitro techniques, they make minimal assumptions about the environment. The minimal assumptions are as follows: The trapped particle’s environment must be locally homogeneous, and the response of the system to a small-applied force must be the same as its response to a similar spontaneous thermal force. These techniques require more complex setups and analysis, but in return, they allow trap calibration in situ, calibrating the trap on each organelle being observed. In addition, these techniques actively measure the local viscoelasticity.

The fundamental issue in vivo is that the trap is in an environment with three variables: the environment’s elasticity, the environment’s viscosity, and the trap’s stiffness (effectively, a second form of elasticity). Both techniques take three measurements, one from passive observation of the trapped object and two from actively applying force to the trapped object: the trapped object’s oscillation in-phase and out-of-phase relative to the trap’s oscillation. The passive measures the combination of the trap’s damping and environment’s damping of the organelle’s thermal vibration. The active measures the trap’s ability to apply force against the local environment. The ratio of the in-phase and out-of-phase measurements corresponds to the relative strengths of the environment’s elasticity and viscosity versus the trap’s stiffness. Because there are three measurements and variables, all three variables can be determined. Therefore, the trap stiffness can be extracted.

The method developed by Hendricks et al. is very similar to the FDT method, with the main differences being in some slightly different mathematical terminology and transformations and the fact that Hendricks et al. performed global fits to the active (or forced) and passive (or thermal) spectra. Fisher et al. applied their equations individually at each frequency. The actual calibration was very similar, with a measurement of the response to an applied force by moving the trapping laser or stage and a measurement of the passive response of the system to thermal fluctuations. These in vivo calibration techniques differ from in vitro methods mainly in adding one extra step for the in vivo case, namely, the active calibration, where the response of the trapped object to the trap’s applied force is tested. This extra measurement allows the measurement of the local viscoelasticity of the trapped object. This then allows the removal of the effect of the local environment on the trap and the calibration of the trap’s stiffness.

3.3.2. In Vivo Optical Trapping Results. The wide varieties of calibration methods and in vivo systems have led to divergent results. This should not be unexpected though, as different types of transport (nuclear, intraflagellar, axonal, etc.) would be expected to have varying properties and purposes.

One of the central questions in the motor field is how many and what types of motors are simultaneously present and active on cargo. Studies on *Drosophila* embryos suggested that there are one or two kinesin motors and one or two dynein motors present and active on most cargos and that the number of motors present on the droplets changes as the embryo develops. Their data also suggested that both motors have a stall force of about 2.6 pN in the cell. They argue for a coordinated-motion model: Motor number had a minimal effect on motility (knocking down kinesin by half), as determined by Western blot and stall-force assays. In addition, droplets going in one direction were more likely to go in that direction again after being pulled off a microtubule by the trap. They inferred from this result that, although motors of both polarities are present on the droplet, only one polarity is active at a time, in opposition to the tug-of-war model, according to which both polarities are simultaneously active. These works all used in vitro calibration techniques dependent on calibrating outside the cell.

Another study on lipid droplets, this time in a human lung cancer cell line, A549, revealed that the stall forces in both directions of transport were similar. This article, by Sims and Xie, also used an in vitro calibration technique. Outward-directed motion, presumably driven by kinesin, had steps around 8 nm in size, whereas inward-directed motion had a variable step size, as has been reported by Mallik et al. for dynein in vitro. In addition, most transport appeared to be driven by one motor, and the fact that the stall forces in both directions were peaked around 7 pN was taken to indicate that dynein and kinesin both have a stall force of around 7 pN. However, Rai and Mallik showed that there were typically 6–10 weak dyneins (1-pN stall force) and 1 kinesin on a typical phagosome in mouse macrophages. This is in agreement with this same group’s previous data on *Dictyostelium*. They also showed that multiple kinesins do not cooperate well during force generation because there were few force events greater than 7 pN in the plus direction in vitro and in vivo. Dynein,
However, appears to be excellent at collectively generating force and appears to form a catch bond (stronger bond as the load increases) to the microtubule under high opposing forces, making it unlikely to detach.\textsuperscript{35} Catch-bond behavior was also seen in \textit{Drosophila} embryos.\textsuperscript{19}

Dynein’s catch-bond behavior was shown by pulling back on groups of dyneins attached to organelles in the cell. The larger the stall force of the cargo (i.e., the more dyneins attached), the longer the dyneins would hold onto the microtubule before release under a superstall force (a force pulling back greater than that required to stall dynein-driven motion). They also observed dynein’s variable step size in the cell and kinesin’s 8-nm step size.\textsuperscript{26} This group calibrated outside the cell. Although catch-bond behavior might look like it would prevent dynein from being dragged backward by kinesin, (dynein-drags model; Figure 8E), it is important to note that the behavior demonstrated here is not what would be expected for an organelle undergoing a tug of war. In the experiments in which catch-bond behavior was observed, a vesicle was moving in the dynein-driven direction when a sudden, large force was applied to it; the vesicle was suddenly jerked backward against the dynein-driven motion. However, in a tug-of-war situation, kinesin would reattach during dynein-driven motion, and the amount of force applied against the dynein would slowly grow as kinesin stopped. This should not initiate catch-bond behavior. Catch-bond behavior would prevent dynein detachment in the case of sudden, large forces and would aid in preventing organelle detachment, but it would not be expected to be significant during typical tug-of-war behavior.

Two groups have also demonstrated optical trap calibration in living cells.\textsuperscript{12a,13b} In these studies, all calibration was done in the cell, accounting for any effects of a viscoelastic environment. Hendricks et al. examined phagocytosed latex beads in a mouse macrophage line, \textit{J774A.1}.\textsuperscript{13b} They discovered that inward- and outward-directed forces were nearly balanced and found that, at high loads (>10 pN), 8-nm stepping occurred, indicating that teams of motors (dyneins, kinesins, or dyneins and kinesins) had correlated stepping (Figure 7).

The other work, by Blehm et al., compared inward and outward forces exerted by lipid droplets in a human lung cancer cell line (A549), phagocytosed polystyrene beads in \textit{Dictyostelium}, and dynein- and kinesin-coated beads in vitro.\textsuperscript{12a} This article indicated that there were typically several dyneins and generally one kinesin on most cargos. This was concluded because the stall force in the minus direction was typically greater than 1 pN: Because a single dynein was shown to have a stall force $\sim$1 pN in vitro, several dyneins were necessary to generate this large force (Figure 8A,B). In the positive direction, the stall force was rarely greater than 6 pN, and hence, only one kinesin was pulling at a time (Figure 8C).

Blehm et al. also found that the plus-end-directed stalls in the cell (i.e., outward or presumably kinesin-driven) looked very similar to the in vitro plus-end-directed stalls for kinesin- and dynein-coated beads (Figure 8D). In particular, a range of stall forces from 1 to 7 pN was found, equal to or more often less than the 7-pN in vitro stall force of a single kinesin alone (Figure 8C). This finding is important, as it indicates that kinesin’s stall force was reduced by the presence of dynein, not by some other.

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**Figure 8.** Dynein drags. (A) In vitro (red) beads with a single dynein have a lower stall force than minus-end-directed lipid droplets in vivo (blue). (B) Beads coated with multiple dynein in vitro have stall forces similar to those of minus-end-directed lipid droplets in vivo. (C) A single kinesin on a bead in vitro has a narrower and higher stall-force distribution than plus-end-directed lipid droplets in vivo. (D) In vitro, beads coated with kinesin and dynein stall at similar forces while walking toward the microtubule plus end as lipid droplets and phagosomes in vivo (less than the stall force of a single kinesin).\textsuperscript{12a} (E) Dynein being dragged by kinesin during plus-end-directed motion appears to be the simplest explanation for why the beads with kinesin and dynein in vitro and organelles in vivo have reduced plus-end-directed stall forces. Adapted with permission from ref 12a. Copyright 2013 National Academy of Sciences.
effect of the cell. This suggests that dynein reduces kinesin’s stall force, perhaps by remaining attached to the microtubule during plus-end-directed motion and acting as a drag (Figure 8E). This behavior could potentially help bypass obstacles, allowing for quicker directional switching or stronger attachment to the microtubule if kinesin detaches. It could also simply be the most efficient way for these two motors to interact to implement a tug of war. Data from both the Blehm et al. and Hendricks et al. articles support a weak 1-pN dynein and a strong 7-pN kinesin in vivo.\textsuperscript{12a,15b} They also strongly support a tug-of-war model, with cooperative dynein working in a group against a single kinesin (or perhaps two kinesins) during intracellular transport.

In contrast, Laib et al. measured extremely high in vivo stall forces in an intraflagellar transport (IFT) system, indicating large, coordinated groups of kinesins or dyneins. They bound a microsphere to the outside of \textit{Chlamydomonas}’ flagella by attaching the sphere to a transmembrane protein undergoing IFT (Figure 9).\textsuperscript{21a} They could then measure the force the molecular motors inside the cell were exerting, even though the bead was outside the cell. This allowed typical in vitro stiffness calibration techniques to work.\textsuperscript{21a} This system appears to be quite different than most others studied, as the forces measured were extremely high, around 60 pN in both directions. This indicates upward of 10 motors cooperating during transport assuming a stall force similar to kinesin’s, 5–7 pN. Using the same technique and system, Shih et al. also saw similar behavior, with a somewhat lower peak stall force of 20–30 pN.\textsuperscript{21b} The reason for the difference in peak forces between these articles was unclear but could potentially be due to the microsphere being nonspecifically bound in the Laib et al. work, whereas Shih et al. specifically bound the bead with an antibody. In addition, temperature-sensitive kinesin-2 and dynein-1b were used to measure the effects of each motor on transport. When kinesin-2 was knocked out, dynein-driven motion continued unaffected, with the typical pauses that accompanied two-motor transport still occurring.\textsuperscript{21c} Similar behavior was observed with dynein-1b.\textsuperscript{21b} This was taken as evidence for reciprocal coordination as opposed to a tug-of-war model; in a tug-of-war model, knocking out kinesin should have improved dynein-driven transport and removed most saltations, and vice versa.\textsuperscript{21a}

The Shih et al.\textsuperscript{21b} and the Laib et al.\textsuperscript{21a} articles seem to show that there are perhaps many ways in which kinesin and dynein can interact. That kinesin (and dynein) could yield up to 60 pN of force indicates excellent cooperativity in their systems, in addition to coordination between the two motors to prevent them from interfering with each other. Other systems seem to indicate poor kinesin cooperativity and rarely if ever saw motor cooperation on this scale.

Conclusion: From the wide variety of in vivo optical trapping results, the different kinesin–dynein transport systems present quite a bit of complexity. All of the different techniques used to observe and measure force in these in vivo systems might or might not be entirely compatible. In addition to differing calibration techniques, some groups looked at maximum force,\textsuperscript{15b,86} some looked at stalls,\textsuperscript{7f,12a,39} and some looked at escape forces (the minimum force required to escape the trap).\textsuperscript{12b} Each of these methods has its merits, as does each cellular system, but it will be difficult to compare and contrast these systems until more data are collected, so it can be understood how and why these systems act so differently. It might be expected that transport systems with different functions and purposes, for example, intraflagellar transport\textsuperscript{21a} as opposed to neuronal transport, would have different characteristics, which could explain much of the variability encountered in different systems.

4. CONCLUSIONS

Recent interest in how kinesin and dynein interact during bidirectional motion has led to many new systems and techniques that can be used to explore this interaction. New data and theoretical works have also led to much more nuanced models of the kinesin–dynein interaction, with evidence for local directional regulation (individual organelles) being mainly through tug of war between motors, whereas intermediate- and large-scale regulation (regional and whole cell) are dominated by other factors (motor number, phosphorylation state, microtubule modifications, etc.).

Motors also seem to be somewhat specialized in working with each other, in that mammalian kinesin and dynein appear to have characteristics allowing them to work together when simultaneously present on a cargo, or at least not interfere with each other to the point of preventing transport.\textsuperscript{7f,12a,15b} Dynein apparently is dragged behind kinesin during some plus-end-

Figure 9. Schematic of the assay developed by Laib et al.\textsuperscript{21a} for the study of the transport of FMG-1-bound beads on the surface of a flagellum. The flagellum of a single \textit{Chlamydomonas} cell is immobilized on a coverslip, and a polystyrene bead held in an optical trap is used to determine the force produced by the motors. Caption and figure reprinted with permission from ref 23a. Copyright 2009 Elsevier.
directed transport, whereas kinesin routinely releases to allow unhindered minus-end-directed transport.\textsuperscript{2,4,12a} This behavior has also been shown with mammalian NCD and kinesin.\textsuperscript{11} This is in contrast to yeast dynein and human kinesin,\textsuperscript{11} which seem to actively work against each other, leading to stalled cargos. Compatible motors seem to be able to coordinate with each other naturally through a local tug of war,\textsuperscript{7d} with weak, cooperative dyneins acting as a foil to strong, weakly cooperative kinesin. Noncompatible motors (yeast dynein and mammalian kinesin) act similarly to early tug-of-war predictions in that both motors appear to be active and neither can win out, leading to stalling and failed transport.\textsuperscript{11}

Whereas evidence in a majority of systems investigated so far seems to point to a local tug of war to determine directionality and motility behavior for individual cargos\textsuperscript{7d,10a,c,12} and general transport bias being regulated by some higher signaling mechanism,\textsuperscript{7d,122,18a,23a,37a} only a few systems have been examined. In particular, there are systems that seem to show strong coordination (no tug of war occurring at all). Intraflagellar transport\textsuperscript{22} and amyloid precursor protein transport\textsuperscript{22} both displayed heavily coordinated motion: kinesin or dynein was active, but rarely, if ever, were they both active simultaneously.

More complex systems, such as systems with more motor types present,\textsuperscript{24} or other specialized types of transport promise to add more complex regulatory mechanisms. Intraflagellar transport systems have already shown this to be the case, displaying very different behavior than that seen so far in other in vivo systems.\textsuperscript{18b,63,64}

New techniques are being developed to count motor number in vitro and observe organelle behavior in the living cell and in various purified in vitro systems. As these techniques improve and more systems are explored, a more complete understanding of how the interplay between motor–motor interactions (the local tug of war) and cellular regulation (motor number, phosphorylation, signaling) controls bidirectional transport will be uncovered.

AUTHOR INFORMATION

Corresponding Author

E-mail: selvin@illinois.edu.

Notes

The authors declare no competing financial interest.

Biographies

Benjamin H. Blehm is currently a postdoctoral fellow at the National Institutes of Health and previously received his Ph.D. at the University of Illinois, Urbana-Champaign. Current research interests include the role of force and tension in mediating cell–microenvironment interactions and developing new optical trapping and imaging techniques.

Paul R. Selvin is a Professor of Physics, Biophysics and Cell and Developmental Biology at the University of Illinois, Urbana-Champaign. He has developed many single-molecule fluorescence techniques, as well as optical trap techniques, with the goal of understanding molecular motors and, more recently, neuronal synapses.

ACKNOWLEDGMENTS

Funding for this work was provided by Grants NIH GM068625 and NSF 0822613

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