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Totally asymmetric simple exclusion process simulations of molecular motor transport on random networks with asymmetric exit rates

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Using the totally asymmetric simple-exclusion-process and mean-field transport theory, we investigate the transport in closed random networks with simple crossing topology—two incoming, two outgoing segments, as a model for molecular motor motion along biopolymer networks. Inspired by in vitro observation of molecular motor motion, we model the motor behavior at the intersections by introducing different exit rates for the two outgoing segments. Our simulations of this simple network reveal surprisingly rich behavior of the transport current with respect to the global density and exit rate ratio. For asymmetric exit rates, we find a broad current plateau at intermediate motor densities resulting from the competition of two subnetwork populations. This current plateau leads to stabilization of transport properties within such networks.

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I. INTRODUCTION

Transport of matter and energy is essential for maintaining complex structures. A constant driving can create and maintain the order in man-made structures as well as in living organisms. In fact, efficient transport is crucial for people living in modern societies: food, building components, energy, and people themselves are transported daily between rural and urban areas. Similarly, on the subcellular level, nutrition and molecular building blocks are transported from the periphery of the cell to its core [1–3]. Motor proteins move actively along the filament network, which also acts as cytoskeleton of the cell [4–8]. In both cases, maintaining a robust transport is crucial, in particular with respect to variations in density. In many transport systems, the transport is carried out by unidirectional movement of carriers (e.g., trucks, molecular motors, etc.), along networks consisting of linelike pathway segments and crossings. Such unidirectional particle motion has been modeled by the well-known totally asymmetric simple exclusion process (TASEP) [9–11], which has recently been applied to transport along cytoskeleton assemblies [12,13]. Originally formulated for one-dimensional systems [14], TASEP was extended to networks by defining the dynamics at crossings. These previous works discuss in detail the role of network structure in the modeling of complex traffic problems [12,13].

Yet, recent experimental observation of molecular motor motion highlights the nontrivial dynamics of molecular motors at crossings: specific types of motors were found to switch between segments with specific probabilities [15–19]. This preference at crossings might crucially determine the transport properties of the network. Earlier TASEP simulations of large networks [12,13] used symmetric exit rates, and only recent simulations of a single crossing show the heterogenous motor density that can arise from a preference in outflow directions among the crossing filaments [20,21].

Here, we study the role of asymmetric exit rates on the transport behavior of large networks using the TASEP model. Inspired by in vitro experiments, in which typically arrays of cytoskeleton filaments are created by attachment to a surface, we focus on a network topology of intersecting lines in a two-dimensional plane: in this topology intersections consist of a single crossing of two filaments. We show that interesting heterogeneous network densities and network transport arises from the local asymmetric exit rates. These exit rates, together with the global motor density crucially determine the transport capacity of the network. We identify four regimes of current-density behavior: A low-density regime controlled by the relative exit probabilities and network topology, a jamming regime with heterogeneous density and resulting complex interplay of subnetwork currents, a saturation regime, and a high-density regime with local dynamics and homogenous density. The extent of each regime and the actual value of the current is found to strongly depend on the relative exit rates at crossings.

II. MODEL DESCRIPTION

We model cytoskeleton assemblies with a two-dimensional network of intersecting segments. Inspired by in vitro experiments, where the cytoskeleton is typically grown by sedimenting filaments on a glass plate [Fig. 1(a)], we create a network topology by projecting lines with random positions and directions on a square [Fig. 1(b)]. The part of a line (filament) between two crossings is defined as a segment. Every segment has 1D structure consisting of L lattice sites. Each site can accommodate maximally one motor. The motors can move from site to site along one direction only, which is defined randomly, and once defined, is fixed during the entire simulation run. For simplicity we define a uniform length L for all segments; we found that this produces more robust results while not affecting the general transport behavior through the network [22]. We introduce periodic boundary conditions by connecting the beginning and end of every line at the boundaries of the square, thus creating a closed network. The resulting network is characterized by the number of crossings $N_c$, the number of segments $N_s = 2N_c$, and the total number of sites $M_s = (2L - 1)N_s$. The network is subsequently populated with motors at a density $\rho$, defined as the ratio of the total number of motors and the total number of sites $M_s$.

Molecular motor motion is modeled with the TASEP, in which particles perform unidirectional random sequential hops constrained by the fact that they may not overlap with
each other. During a single update, a motor advances one site forward if the target site is unoccupied; otherwise, no progression occurs. Network sites are updated in random sequential order and one update cycle corresponds to $M_s$ single-site updates.

When a motor arrives at the crossing, it continues with probability $\gamma$ along one outgoing segment and with probability $1-\gamma$ along the other, independently of the incoming segment. As shown in Fig. 1(c), this can account for the local topology of in vitro biopolymer networks, where the preferred filament (overpass filament 1) is lying “on top” of the other filament (underpass filament 2). The probability $\gamma$ thus defined for every crossing is kept constant during the simulation, and is hence a fixed property of the network. Our approach is based on the experimental observation [15] that motors with cargo bead attached are more likely to stay on the overpass filament and change direction when traveling on the underpass filament, making the overpass filament always preferable [23]. For simplicity, we define equal $\gamma$ for all crossings independent of the history of motor motion. For completeness, consequences of history-dependant motor motion are discussed in Appendix C.

Finally, we neglect the possible unbinding of motors, which is observed in Refs. [15,24], and studied in simulations by Neri et al. [13]. Such unbinding weakens the correlations in the traffic flow of the motors. However, it was shown that for motors with attached cargo bead [15] the unbinding probability at the crossings is small or negligible. Neglecting unbinding allows us to focus on the physical effects of jamming and flow [25] and obtain a simple physical picture in terms of $\gamma$. The control parameters of our simulations are thus the exit probability $\gamma$, which due to the symmetry we choose as $0.5 \leq \gamma \leq 1$, and the global motor density $0 < \rho < 1$. We consider relatively large networks with $N_s \sim 500$ and uniform segment length $L = 500$. All data have been averaged over 10,000 cycles in steady state, after discarding the transient (initial $\sim 3000$ cycles).

III. RESULTS

The effect of exit probability $\gamma$ on the network current is shown in Fig. 2, where we plot the total current versus density $\rho$ for various values of $\gamma$ (blue solid curves). For comparison, we also show the mean-field current through a single segment $J_s = \rho (1 - \rho)$ (black dashed line) [26] that exhibits a maximum at $\rho = 0.5$ due to competition of growing density increasing the current and density-induced jamming reducing it. The network current behaves similar to that of a single segment for symmetric exit rates ($\gamma = 0.5$, upper blue solid curve in Fig. 2): it is symmetric around $\rho \sim 0.5$ and deviates from that of the single segment only at densities around 0.5. The situation changes for asymmetric exit probability ($\gamma \neq 0.5$), where the network current becomes asymmetric and progressively reduces and flattens with increasing $\gamma$, until at $\gamma \sim 0.9$ when it becomes largely independent of density. Thus, the asymmetric exit probability reduces the network current and stabilizes it at the same time.

How does this emerging current plateau arise from the interplay of currents in the network? To answer this question

![FIG. 2. (Color online) Transport current $J$ dependence on global density $\rho$ for different exit probabilities $\gamma$ obtained from the TASEP simulations. Dashed black curve is shown for comparison with the current through single segment. Data is averaged over 10 000 cycles after waiting for 3000 cycles to reach steady state.](https://example.com/fig2.png)
we investigate the distribution of individual segment densities $\rho_s$. We compare density distributions for symmetric and asymmetric exit probability in Fig. 3. For $\gamma = 0.5$, the density is homogeneous across the entire network: local and global densities largely coincide with only minor variation across segments [Fig. 3(a)]. For $\gamma = 0.8$, in contrast, we find strong inhomogeneity at intermediate global density [Fig. 3(b)]; the asymmetric exit probability leads to redistribution of paths and resulting redistribution of segment densities. Indeed, the asymmetric exit probability redistributes the motors into two subsystems: associated with the vertex exit probabilities, the network can be considered as consisting of two interconnected subnetworks, one consisting of segments with entry probability $\gamma$ and the other consisting of segments with probability $1 - \gamma$. The subnetwork with higher entry probability should populate faster, leading to the heterogeneous distribution of densities shown in Fig. 3(b). This is indeed confirmed in the subnetwork density distributions as shown in Fig. 3(c): at intermediate $\rho$, the density of the high-preferability subnetwork has grown much more, as reflected in the shift of the distribution to higher density. Due to interplay of the individual segments, the density within each subnetwork is heterogeneously distributed.

To link this density evolution to the total current, we plot average densities of the two subnetworks together with the total current in Figs. 4(a) and 4(b). Clearly, the high-preferability subnetwork takes up particles at a faster rate as shown in Fig. 4(b); this is a direct result of the higher entry probability.
The initial slopes of the growing densities in high- and low-preferability subnetworks are $2\gamma$ and $2(1-\gamma)$, respectively, reflecting the two entry probabilities. When roughly half of the network sites are occupied ($\rho \sim 0.5$), high-preferability segments become saturated, and particles are increasingly taken up by the low-preferability segments. This interplay of the two subnetworks leads to stabilization of the total network current: as we will see below, jamming effects in the high-preferability subnetwork lead to an emerging state of the network current: as we will see below, jamming effects in the high-preferability subnetwork lead to an emerging state.
become saturated, and their density does not grow any further [see Figs. 4(b) and 4(c)].

Toward the end of the jamming regime ($\rho = \rho^*_h$) all high-preferability segments and connected vertices are saturated with equal particle density, $\rho^h = \rho_v = 1 - \beta$. At the same time, the low preferability segments are still unjammed. Hence, at $\rho^*_h$, the density difference between the two subnetworks is maximal.

3. Saturation regime

Above $\rho^*_h$, the density in the high-preferability subnetwork no longer increases, and further increase of $\rho$ takes place in the low-preferability subnetwork, where consequently jams start to form. Since high- and low-preferability segments are connected by the same vertices, vertices with saturated uniform density $\rho_v$ control the inflow [$\alpha = \rho_v(1 - \gamma)$] and outflow ($\beta = 1 - \rho_v$) of low-preferability segments. The only regime where the segment density $\rho^i$ can rapidly change while inflow and outflow rates remain constant is during the shock phase [26], when jammed and free flow coexist within one segment and $\alpha = \beta$. Using this equality for the low-preferability segments, we obtain $\rho_v = 1/(2 - \gamma)$. This allows us to calculate the saturating current of high-preferability segments: using $J^h = \beta(1 - \beta)$ with $\beta = 1 - \rho_v$, we obtain $J^h = (1 - \gamma)/(2 - \gamma)^2$. Since in the low-preferability segments, jam and free flow coexist, ($\rho^i$) changes from $\alpha = (1 - \gamma)/(2 - \gamma)$ to $1 - \beta = 1/(2 - \gamma)^2$ with increasing $\rho_v$, corresponding to $1/6$ and $5/6$ for $\gamma = 0.8$, again in agreement with the simulations. During this rapid density change, the low-preferability subnetwork current remains constant $J^l = \beta(1 - \beta)$. The resulting total current is thus $J = 1/2(J^h + J^l) = (1 - \gamma)/(2 - \gamma)^2$ [dashed green line in Fig. 4(a)], independent of $\rho$, manifesting the current plateau.

Indeed, the TASEP simulations show plateau-like behavior with current values only slightly higher than the value $(1 - \gamma)/(2 - \gamma)^2$ for $\gamma = 0.8$ in Fig. 4(a) the TASEP simulations plateau value is $\sim 0.141$ while $(1 - \gamma)/(2 - \gamma)^2 = 0.139$. Again, this small discrepancy results from the spread in densities within the subnetworks opposed to the fixed average densities ($\rho^i$) and ($\rho^h$) assumed here.

The transition density $\rho^*_s$ at the start of the saturation regime can be estimated from the requirement that for $\rho \geq \rho^*_s$, all high-preferability segments are jammed; i.e., $\langle \rho^h \rangle = 1 - \beta = \rho_v = 1/(2 - \gamma)$. At the same point all low-preferability segments are still unjammed, $\langle \rho^i \rangle = \alpha = (1 - \gamma)/\rho_v = (1 - \gamma)/(2 - \gamma)$; this leads to the mean density $\rho^*_s = \langle \rho^i \rangle = (\langle \rho^i \rangle + \langle \rho^h \rangle)/2 = 0.5$, which is independent of $\gamma$.

By the end of the saturation regime at $\rho = \rho^*_s$, $\langle \rho^i \rangle$ has increased due to the jamming until $\langle \rho^i \rangle = (\rho^h) = 1/(2 - \gamma)$ and the density is homogenous throughout the whole network. This immediately gives the upper density of the saturation regime: $\rho^*_s = 1/(2 - \gamma)$.

Interestingly, $\langle \rho^i \rangle$ remains much more homogeneous while rapidly increasing than $\langle \rho^h \rangle$ in the jamming regime [see blue and red shades in Fig. 4(b)]. This difference is a direct result of the vertex densities controlling the segment densities [see black curve in Fig. 4(b)]. During the jamming regime the crossing densities $\rho_v$ evolve rapidly, while in saturation regime the crossing densities are saturated and fixed at $\rho_v = 1/(2 - \gamma)$.

\begin{figure}[h]
\centering
\includegraphics[width=\textwidth]{current_plateau.png}
\caption{(Color online) Transport current $J$ dependence on exit probability $\gamma$ for different densities $\rho$ obtained from the TASEP simulations. Data is averaged over 10 000 cycles after waiting for 3000 cycles to reach steady state. Blue (dark gray) curves correspond to $\rho \leq 0.5$, green (light gray) ones to $\rho > 0.5$. The current is then simply $J = \rho(1 - \rho)$, fitting perfectly with both TASEP simulation and iteration method [see dashed green line in Fig. 4(a)].}
\end{figure}

4. High-density regime

For $\rho > \rho^*_h$ the density in both high- and low-preferability segments is given by the outflow rate ($\beta = 1 - \rho_v$) and becomes independent of the exit probability $\gamma$. As a result, all segments behave identically $\rho^h = \rho^i = \rho_v = \rho$, as shown by the perfect overlap and the linear increase with slope 1 of the red (light gray) and blue (dark gray) curves in Fig. 4(b). The current is then simply $J = \rho(1 - \rho)$, fitting perfectly with both TASEP simulation and iteration method [see dashed green line in Fig. 4(a)].

We finally show the explicit dependence of the current on exit probability $\gamma$ in Fig. 5. Interestingly, a similar plateau behavior is observed as for the density dependence, indicating that the current is stabilized also with respect to $\gamma$: green (light gray) curves for $\rho > 0.5$ evidence the emergence of a plateau in $J$ with increasing $\rho$. This plateau corresponds to the high-density regime; hence, its extension can be derived from $\rho^*_s$, and we obtain $1/\rho - 1 \leq \gamma \leq 2 - 1/\rho$. At low densities, on the other hand ($\rho < 0.5$), a somewhat narrower plateau emerges with decreasing $\rho$ and becomes established for $\rho < 0.1$. This rather low density corresponds to the linear regime, i.e., for $\rho < \rho^*_s$. In this case, we can estimate the plateau value from $J = \rho(1 - \rho) - \rho^2(1 - 2\rho)^2$ by neglecting the second term for $\gamma \sim 0.5$. We thus obtain the plateau value $J = \rho(1 - \rho)$, which appears in a narrow range of $\gamma$ close to 0.5; see blue (dark gray) curve for $\rho = 0.1$ in Fig. 5.

IV. SUMMARY

Using TASEP simulations and mean-field theory on complex networks, we have studied the transport properties of large networks with asymmetric exit rates. We find that the transport current depends strongly on the relative exit probabilities and the global density that together determine the transport properties and density distribution of the network. Asymmetric...
exit rates lead to strong decrease of the transport current, but at the same time to emergence of a current plateau that stabilizes the current: this plateau makes the current largely independent of variations in particle density. Similar current stabilization is observed with respect to variations in exit probabilities when the density \( \rho \) is rather low or high.

The main mechanism underlying this complex network behavior is the splitting of the network into two subnetworks according to the exit probabilities \( \gamma \) and \( 1 - \gamma \). This splitting into two subnetworks, together with the (random) topology of the network sets the complex transport pathways, which, at intermediate densities, lead to heterogeneous crowding of the network. The resulting four transport regimes—linear, jamming, saturation, and high-density—are characterized by the relative densities of the two subnetworks and the density distribution within each subnetwork. Former is determined by \( \gamma \), latter by the network topology.

The topology and asymmetric exit probabilities used here mimic that of typical in vitro experiments in which molecular motors walking along biopolymer networks exhibit specific preference to proceed along a certain segment at filament crossings. Our work hence provides insight into how local motor dynamics affect the collective transport properties of a biopolymer network. However, important additional effects arise from the unbinding of real motors that is experimentally observed. Such detachment of motors can weaken the correlations and jamming effects that we observe in the simulations.

The work presented here complements simulations of network transport based on random bidirectional \[29\] and guided motion between nodes \[30,31\] that are more suitable for internet traffic and urban transport description. In these studies it was shown that jamming can occur at relatively low densities. In the current work, we have shown that the unidirectional exclusion process together with asymmetric exit rates leads to complex jammed states with heterogeneous steady-state densities and currents that can nevertheless be understood based on the simple interplay of two subnetworks. Due to the generality of the exclusion process in modeling transport and arrest phenomena, our results may provide generic insights into traffic jams and transport capacities of highway networks, biological networks, and other systems with similar unidirectional topology.

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APPENDIX A: ITERATIVE SOLUTION OF MEAN-FIELD MODEL

Here we provide an iterative solution of the mean-field model, which—as we will show below—reproduces the network behavior quantitatively and shows excellent agreement with the TASEP simulations. This iterative method computes steady-state segment currents from the density at the vertices.

For each segment, the inflow rate \( \alpha \) is proportional to the density \( \rho_v \) of the entry crossing and the exit probability, i.e., \( \alpha = \rho_v \gamma \) or \( \alpha = \rho_v (1 - \gamma) \)[26]. On the other hand, the outflow rate \( \beta \) is controlled solely by the availability of an unoccupied exit vertex, so that \( \beta = 1 - \rho_v \), independent of \( \gamma \). The change of density of each vertex is given by particle influx due to the two incoming segments and outflow due to the two outgoing segments, similar to Ref. [12]

\[
\frac{\partial \rho_v}{\partial t} = \sum_{\nu'} J_\nu(\alpha_{\nu'}, \beta_{\nu'}) + \sum_{\nu''} J_\nu(\alpha_{\nu''}, \beta_{\nu''}),
\]

where \( \nu' \) sums over the two incoming segments and \( \nu'' \) over the two outgoing segments. Here, the segment current is \( J_\nu = \alpha(1 - \alpha) \) for \( \alpha < \beta \) and \( J_\nu = \beta(1 - \beta) \) for \( \beta < \alpha \). Since \( \alpha \) and \( \beta \) are themselves related to the vertex density as described above, any change of vertex density will in turn influence the segment currents, which again change the density distribution. We hence use an iterative method to find steady-state currents and vertex densities. By expressing \( \rho_v(t + 1) = \rho_v(t) + \partial \rho_v(t) / \partial t \), we can find steady-state configurations, for which \( \partial \rho_v(t) / \partial t = 0 \) for all vertices (in the case of quantized time steps).

The resulting steady-state network current depicted by the brown (light gray) line in Fig. 4(a) reproduces all features of the total current: the initial linear and nonlinear increase, saturation, and the final decrease due to complete jamming of the network. The remaining slight difference between the TASEP simulations and the iterative solution is associated with the approximation \( J_\nu = \rho_v(1 - \rho_v) \), which is strictly true only for infinite segment lengths.

APPENDIX B: LINEAR REGIME LIMIT

In this section we compute the transition density \( \rho = \rho^* \) that bounds the linear regime. The linear regime ends when the density of each subnetwork is no longer linearly proportional to the global density. To pinpoint \( \rho^* \), we determine the transition matrix \( \epsilon \), whose elements \( \epsilon_{ij} \) denote the transition probabilities from vertex \( i \) to \( j \). We define element \( \epsilon_{ij} = 0 \) when vertex \( j \) is not connected directly to vertex \( i \) through one segment, \( \epsilon_{ij} = \gamma \) when \( j \) is connected to \( i \) through a high-preferability segment, and \( \epsilon_{ij} = 1 - \gamma \) when \( j \) is connected to \( i \) through a low-preferability segment. This matrix specifies the transitions between vertices, giving the probability that a particle continues from one vertex to the next; the eigenvector corresponding to eigenvalue 1 of matrix \( \epsilon \) holds the relative distribution of densities \( \rho_v \), among vertices in steady state. Knowing the densities of all vertices we can calculate the densities of every segment \( \rho_v \) through incoming rate \( \alpha = \gamma \rho_v \) or \( \alpha = (1 - \gamma) \rho_v' \) and outgoing rate \( \beta = 1 - \rho_v \). Actually, while using the transition matrix \( \epsilon \) we have assumed that all segments are inflow controlled (\( \alpha < \beta \)) and not jammed, only then the transition rates between vertices will depend on \( \gamma \) only. Indeed at low global density \( \rho \ll \rho^* \), the computed values of \( \rho_v \) are in excellent agreement with the TASEP simulation results, since the transition matrix gives an exact solution of the density distribution among segments as shown in Fig. 6(a). As the first segment becomes jammed, \( \alpha > \beta \), the linear-regime ends and the density distribution is no longer given by the transition matrix; see Figs. 6(b) and 6(c).
The task to find the end of the linear regime is to find the lowest global density $\rho = \rho_s^*$ where $\alpha > \beta$ for one of the segments. For $\gamma = 0.5$ we can find a simple analytical solution, since in this isotropic case all segments and vertices are identical. By solving $\alpha > \beta \Leftrightarrow \gamma \rho_v > 1 - \rho_v$ we get $\rho_v > 2/3$ and find the maximum global density where the linear regime is still valid $\rho_s^* = \gamma \rho_v = 1/3$. For densities $\rho > \rho_s^*$ jamming starts to occur and spread through the network.

For $\gamma \neq 0.5$ the network is anisotropic and the distribution of vertex densities has a finite width, so a simple analytical solution with all $\rho_v$ equal is no longer possible. For every segment within a particular network, however, we can explicitly solve the condition $\alpha > \beta$ using matrix $\epsilon$ and find the global density $\rho_s^*$ where jamming first occurs. Naturally, this transition density depends on the size of the network $N_s$ and the exit probability $\gamma$ for $L \to \infty$ the length of the segment does not play a role. We have averaged the results for $\rho_s^*$ over 200 randomly created networks for every $N_s$ and $\gamma$. We plot the resulting transition density $\rho_s^*$ as a function of $\gamma$ in Fig. 7(a).

For intermediate network sizes in the range $N_s = 3 \times 10^3 - 10^5$, we found that the power-law approximation $\rho_s^* \approx \left(\frac{N_s}{2}\right)^{0.5}$ works relatively well. The prefactor $\rho_s^* = 1/3$ for $\gamma \sim 0.5$ and is changing slightly as $\gamma$ approaches 1 or 0. For $N_s = 500$ and $\gamma = 0.5$ we find $\rho_s^* \approx 0.07$, in very good agreement with Fig. 4(c). In the extreme case $\gamma = 0$ or 1, the transition density becomes simply $\rho_s^* \sim (1/\sqrt{N_v})$.

For large network sizes $N_s \geq 10^3$, $\rho_s^*$ decays slower with respect to a power-law behavior. Additionally for different networks with the same $N_s$ and $\gamma$ the fluctuations of $\rho_s^*$ can be relatively large so any analytical prediction of $\rho_s^*$ will give only its approximate value. If one wants to know the precise value of $\rho_s^*$, one has to explicitly calculate it for every particular network topology using the transition matrix $\epsilon$.

Interestingly, by analyzing the general behavior of $\rho_s^*(N_s, \gamma)$ we conclude that for sufficiently large networks ($N_s \to \infty$) and high anisotropy ($\gamma \neq 0.5$) the transition density $\rho_s^* \to 0$, meaning that local jamming and nonlinear processes start to occur at infinitely small densities.
APPENDIX C: HISTORY-DEPENDENT EXIT PROBABILITIES

In the manuscript, we have concentrated on the case that the exit probabilities at crossings are independent of the history of motor motion; i.e., that they are a fixed property of the network. In this Appendix we show the results for the alternative crossing rule that the choice of the new segment depends on the previous motion. In this case, we introduce the exit probability $\gamma^h$, with which the motor continues along the same line at a crossing; see Fig. 1(b). Consequently, with probability $1 - \gamma^h$, the motor will change direction to the exit segment corresponding to another line, so the choice of the exit segment at the crossing completely depends on where the motor came from.

From the network topological perspective every crossing becomes identical and there is no fixed preferability of one segment over the other. Consequently, for all $\gamma^h$ this situation is very similar to the isotropic case $\gamma = 0.5$ when motors do not have “history” of its past motion. Indeed for all investigated $\gamma^h$, we observe steady-state currents very similar to the previous case $\gamma = 0.5$ as shown in Fig. 8(a). In particular, all current distributions are symmetric with respect to $\rho = 0.5$. In fact, for $\gamma = 0.5$ and $\gamma^h = 0.5$ we get identical transport behavior in the network with current plateau for densities in the range $1/3 < \rho < 2/3$. Increasing $\gamma^h$ above 0.5 makes the equilibration time to reach the steady state longer due to stronger local fluctuations between the segments. We find that these stronger fluctuations also lead to slight decrease of the maximum current; see Fig. 8(b). This current reduction is, however, much weaker (almost negligible, only ~3%) than the one described in the main manuscript for the case of history-independent exit probabilities.

[22] For sufficiently long segments ($L \geq 200$) the density and current of a segment are fully determined by the inflow and outflow at the boundaries and does not depend on $L$.
[23] The choice of the new segment may also depend on the current segment but this is second-order effect comparing to the local topology that we neglect in the current manuscript. Moreover, taking into account switching rules which depends on the incoming segment will make the current and density...
distribution more homogenous but will not add new features; see Appendix C.


[25] To neglect the effect of binding and unbinding one can think not in terms of single motors traveling along the network but rather in terms of the cargos being transported along the microtubules by several motors at once. In this case the chance for the cargo to unbind is relatively small even if one of transporting motor unbinds.


