



The Abdus Salam
International Centre
for Theoretical Physics

A GENERAL THEORETICAL FRAMEWORK TO INFER ENDOSOMAL NETWORK DYNAMICS FROM QUANTITATIVE IMAGE ANALYSIS

Authors:

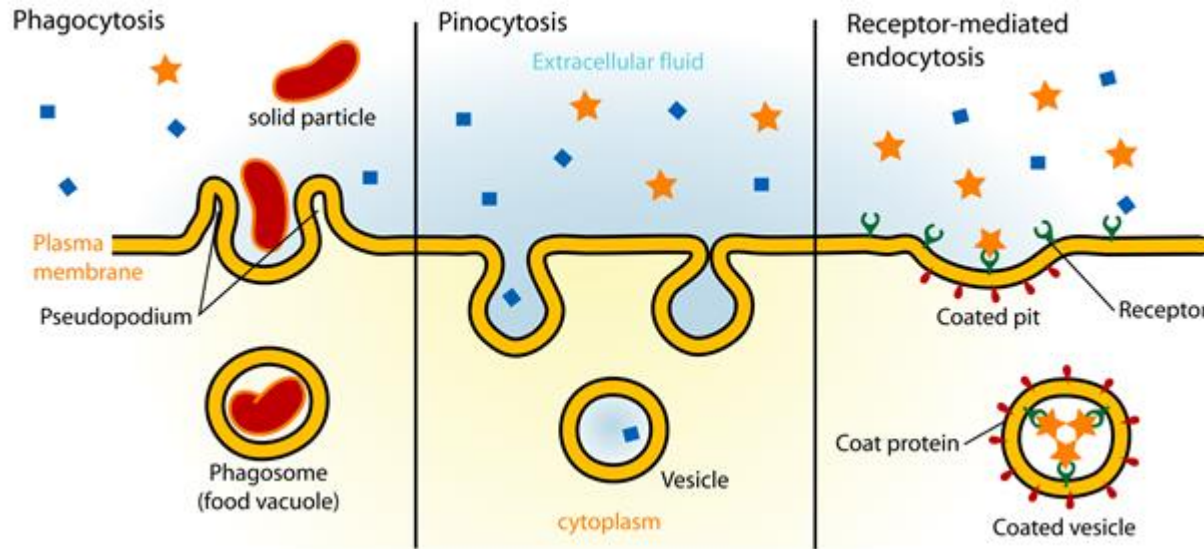
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Paper Review by Ana Milena Forero
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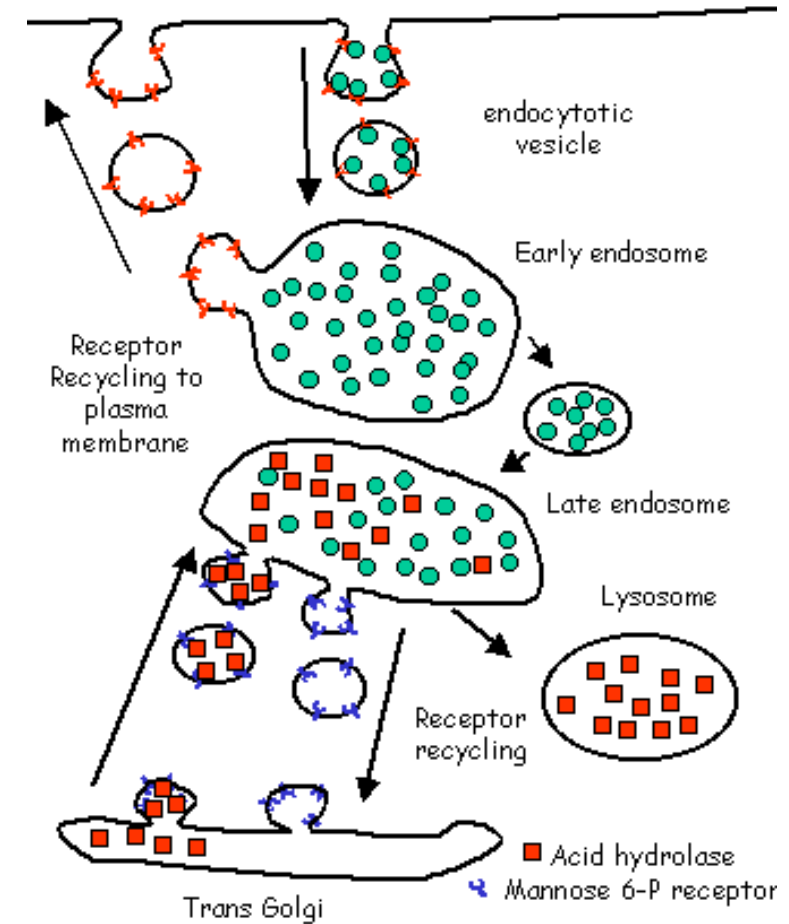
Current Biology 22, 1381-1390, August 7, 2012.

Background

Endocytosis



The different types of endocytosis.
Source: wikipedia

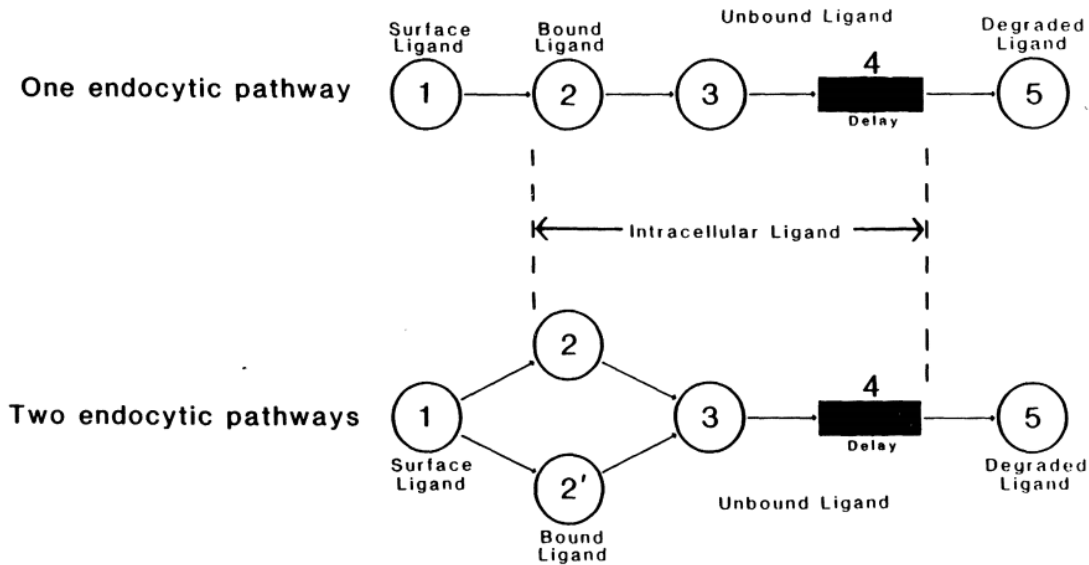


Endocytic pathway

Source: Secretion and Endocytosis (ubc.ca).

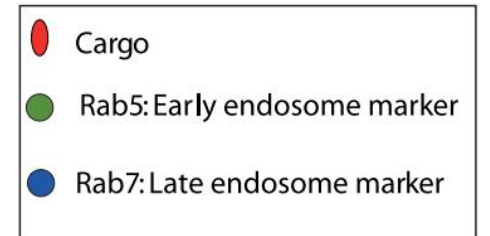
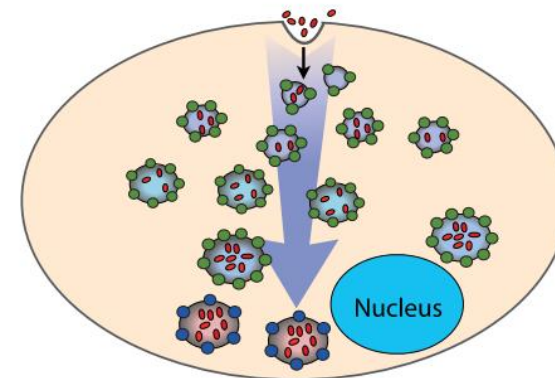
https://www.zoology.ubc.ca/~berger/B200sample/unit_9_secretion/secretion_endocytosis.htm

Problem to address



Compartmental models of asialoglycoprotein endocytosis in rat. [1]

“Endosomes form a dynamic network of membranes undergoing fusion and fission, therefore continuously exchanging and redistributing cargo.”

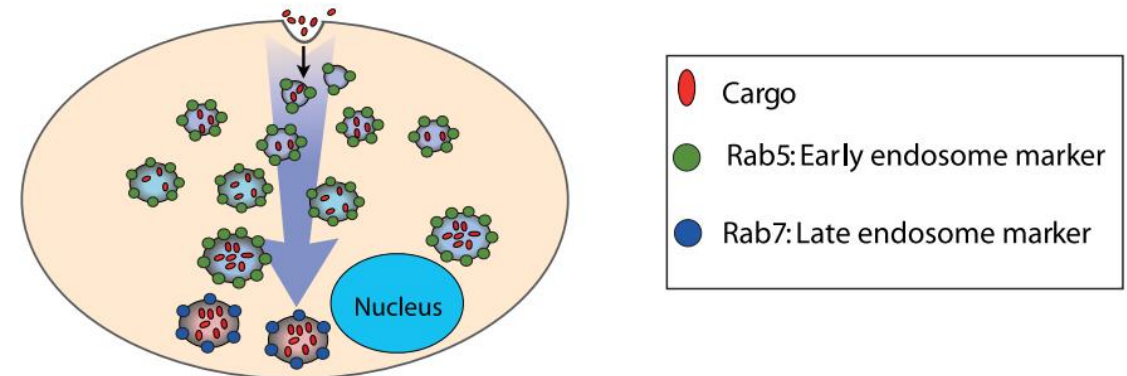


[1] R. Klausner et al., *Mathematical modeling of receptor-mediated endocytosis*. New York (1985), pp. 259-277

Problem to address

- The authors propose to study the macroscopic kinetic properties of the endosomal network from the study of microscopic processes at the level of **individual endosomes**, by means of a combination of theory with quantitative experiments.

“Endosomes form a dynamic network of membranes undergoing fusion and fission, therefore continuously exchanging and redistributing cargo.”



[1] R. Klausner et al., *Mathematical modeling of receptor-mediated endocytosis*. New York (1985), pp. 259-277

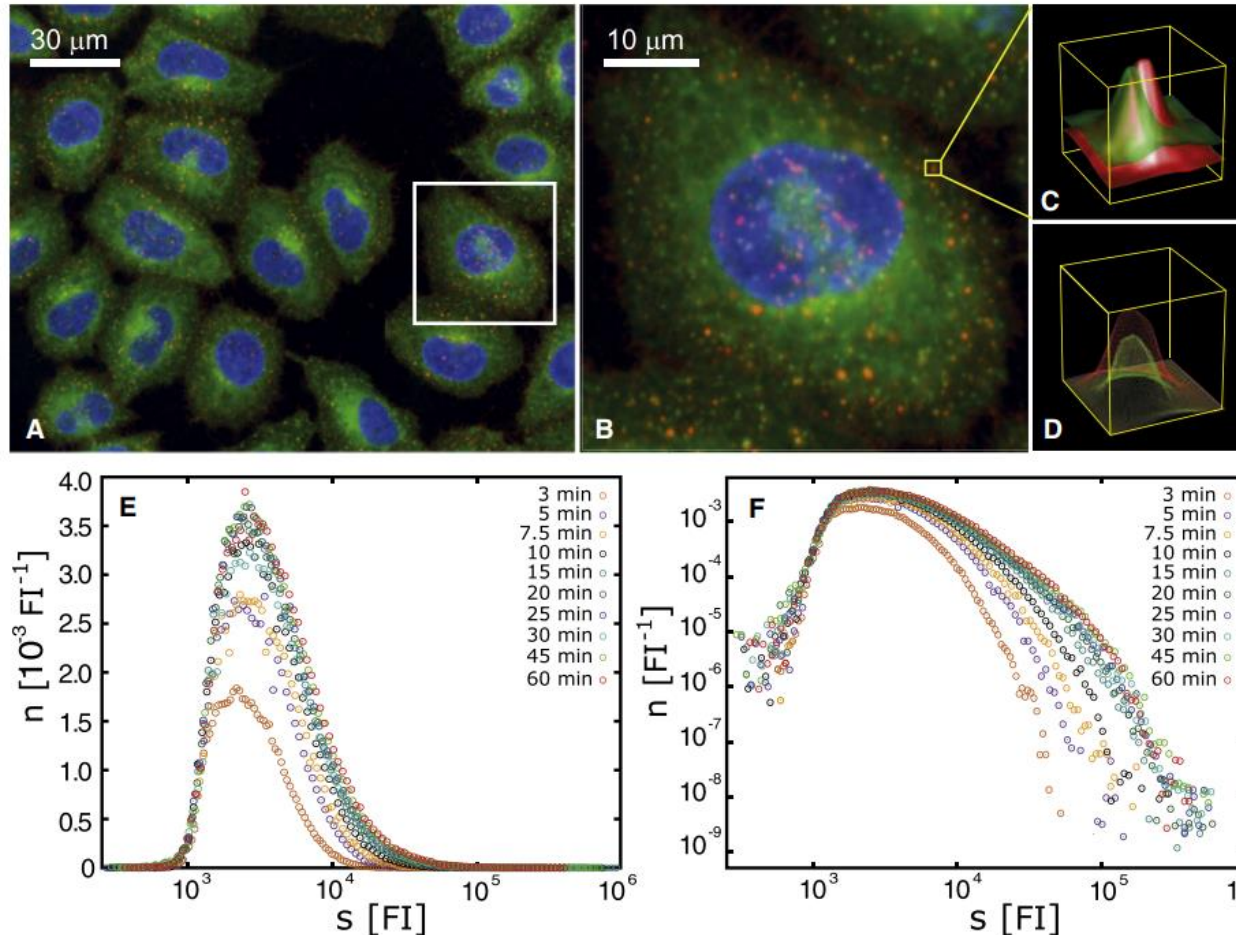
Endosomal Trafficking (Foret et al., 2012)

Results

To study the flux of cargo through the early endosomal network

- Quantitative Image Analysis of the Endosomal Network.
- Theoretical description of Endosomal Dynamics.
- Entry-Fusion-Exit model (EFE).
- Comparison of the Model Predictions with the Experimental Data.

Results: Quantitative Image Analysis of the Endosomal Network



Quantitative analysis of Confocal Microscopy Images of HeLa Cells expressing GFP-Rab5 and internalizing LDL.

GFP-Rab5c

LDL (Low-density lipoprotein) as endocytic marker. Cargo.

Nuclei

“Using the vesicle recognition algorithm, the software detects the Rab5- and the LDL-positive endosomes and quantifies their characteristics (position, area, elongation, fluorescence intensity, etc.)”.

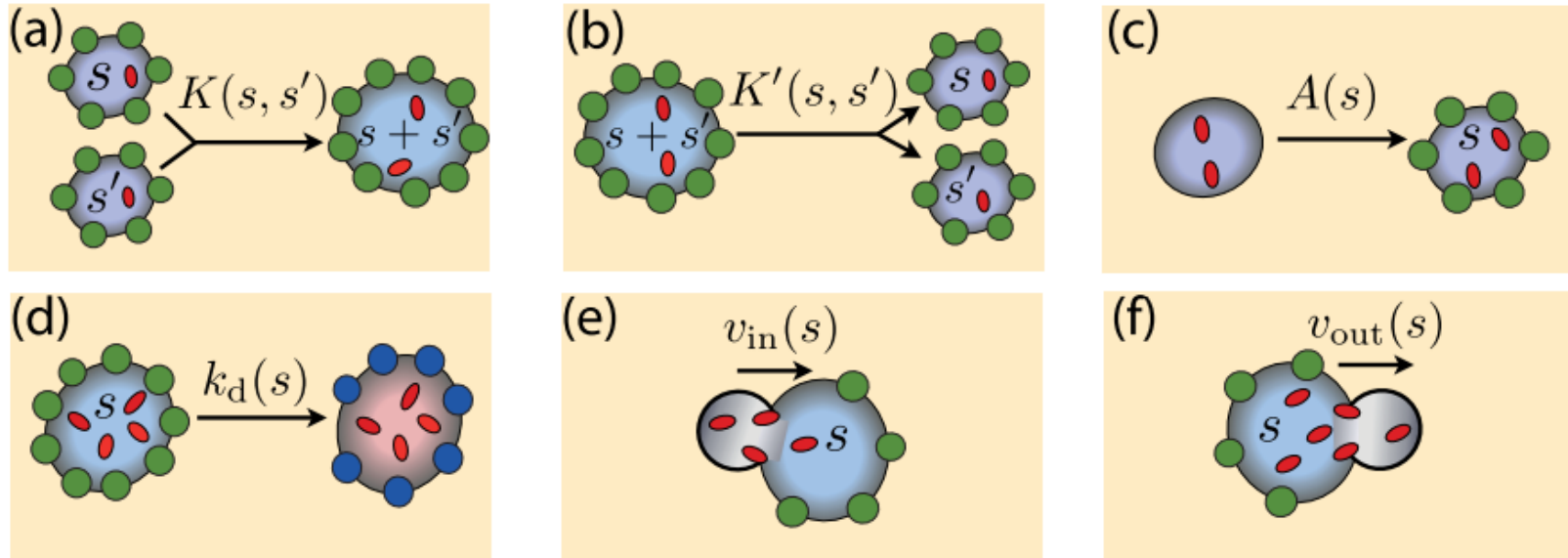
Number of cargo carrying EE per cell:

$$N = \int_0^{\infty} n(s) ds$$

Total LDL fluorescence in the endosomes:

$$\Phi = \int_0^{\infty} s n(s) ds$$

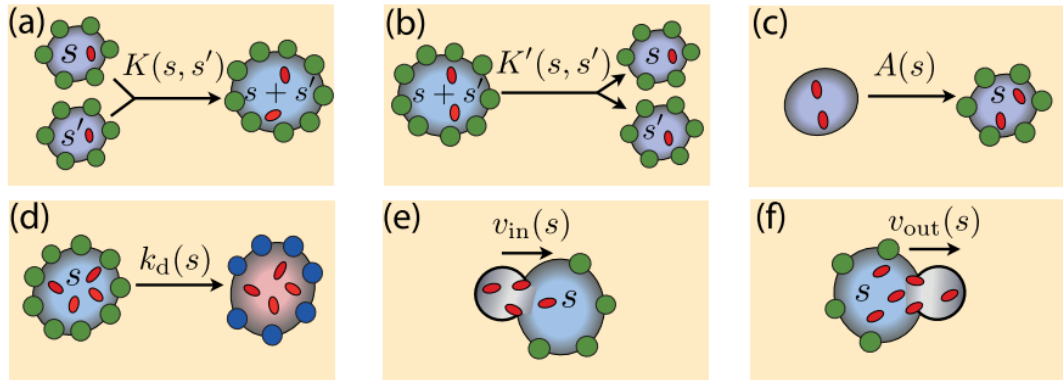
Results: Theoretical description of Endosomal Dynamics



$$\frac{\partial n(s, t)}{\partial t} = \underbrace{\frac{1}{2} \int_0^s K(s', s-s') n(s') n(s-s') ds'}_{(a)} - \underbrace{\int_0^\infty K(s, s') n(s) n(s') ds'}_{(b)} + \underbrace{\int_0^\infty K'(s, s') n(s+s') ds'}_{(c)} - \underbrace{\frac{1}{2} \int_0^s K'(s', s-s') n(s) ds'}_{(d)} + \underbrace{A(s)}_{(e)} - \underbrace{k_d(s) n(s)}_{(f)} - \frac{\partial}{\partial s} (v_{in}(s) n(s)) + \frac{\partial}{\partial s} (v_{out}(s) n(s))$$

Results: Entry-Fusion-Exit model (EFE)

$$J = \int_0^\infty (sA(s) + v_{in}(s)n(s)) ds$$



$$\begin{aligned} \frac{\partial n(s, t)}{\partial t} = & \frac{1}{2} \int_0^s K(s', s-s') n(s') n(s-s') ds' - \int_0^\infty K(s, s') n(s) n(s') ds' \quad (a) \\ & + \int_0^\infty K'(s, s') n(s+s') ds' - \frac{1}{2} \int_0^s K'(s', s-s') n(s) ds' \quad (b) \\ & + A(s) - k_d(s)n(s) - \frac{\partial}{\partial s}(v_{in}(s)n(s)) + \frac{\partial}{\partial s}(v_{out}(s)n(s)) \end{aligned}$$

(c)

(d)

(e)

(f)

$$K'(s, s') = v_{in}(s) = v_{out}(s) = 0$$

$$A(s) = \frac{J}{s_0^2} e^{-s/s_0} \quad k_d(s) = k_d$$

$$K(s, s') = K$$

$$J = \int_0^\infty sA(s) ds$$

$$\begin{aligned} \partial_t n(s, t) = & \frac{K}{2} \int_0^s n(s') n(s-s') ds' - Kn(s) \int_0^\infty n(s') ds' \\ & + \frac{J}{s_0^2} e^{-s/s_0} - k_d n(s) \end{aligned}$$

Results: Entry-Fusion-Exit model (EFE)

$$\partial_t n(s, t) = \frac{K}{2} \int_0^s n(s') n(s - s') ds' - K n(s) \int_0^\infty n(s') ds' + \frac{J}{s_0^2} e^{-s/s_0} - k_d n(s)$$

$$h(x, t) = \int_0^\infty n(s, t) (e^{-xs} - 1) ds \quad \partial_t h = \frac{K}{2} h^2 - \frac{Jx}{1 + xs_0} - k_d h$$

$$h(x, t) = -\frac{\bar{k}(x)}{K} \tanh \left(\frac{t}{2} \bar{k}(x) + \tanh^{-1} \left(\frac{k_d}{\bar{k}(x)} \right) \right) + \frac{k_d}{K} \quad \bar{k}(x) = \left(\frac{2JKx}{1 + s_0x} + k_d^2 \right)^{1/2}$$

Results: Entry-Fusion-Exit model (EFE)

$$h(x, t) = -\frac{\bar{k}(x)}{K} \tanh \left(\frac{t}{2} \bar{k}(x) + \tanh^{-1} \left(\frac{k_d}{\bar{k}(x)} \right) \right) + \frac{k_d}{K}$$

$$\bar{k}(x) = \left(\frac{2JKx}{1 + s_0x} + k_d^2 \right)^{1/2}$$

Analytic solution

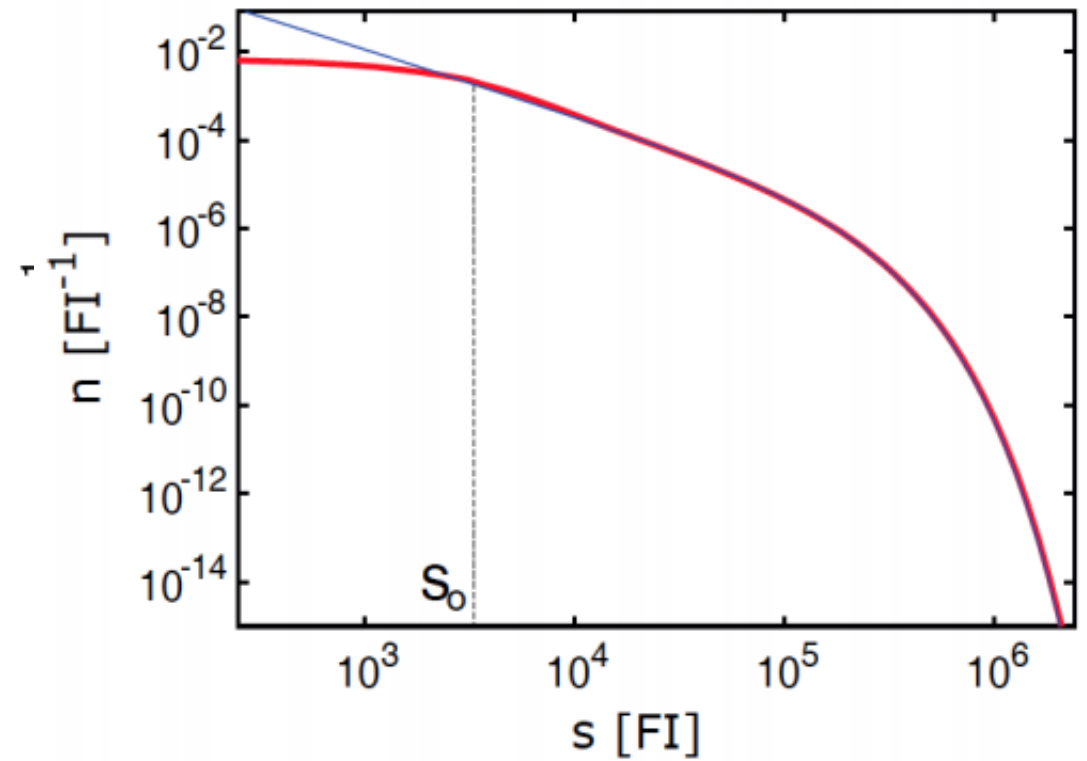
Steady state distribution $n(s)$:

$$h(x, t) \cong -\frac{\bar{k}(x)}{K} + \frac{k_d}{K}$$

$$s \gg s_0 \rightarrow xs_0 \ll 1: \quad \bar{k}(x) = (2JKx + k_d^2)^{1/2}$$

Taking the Inverse Laplace transform:

$$n(s) \cong \left(\frac{J}{2\pi} \right)^{1/2} \frac{e^{-s/s_\infty^*}}{s^{3/2}} \quad s_\infty^* = 2JK/k_d^2$$



Numerical and analytical solutions of the kinetic equation at steady state regime.

Results: Comparison of the Model Predictions with the Experimental Data

$$h(x, t) = -\frac{\bar{k}(x)}{K} \tanh\left(\frac{t}{2}\bar{k}(x) + \tanh^{-1}\left(\frac{k_d}{\bar{k}(x)}\right)\right) + \frac{k_d}{K}$$

$$\bar{k}(x) = \left(\frac{2JKx}{1+s_0x} + k_d^2\right)^{1/2}$$

Time dependent properties of $n(s, t)$ for large s :

$$t \ll \frac{1}{k_d}, \text{ i.e., } k_d \approx 0$$

$$xs_0 \ll 1$$

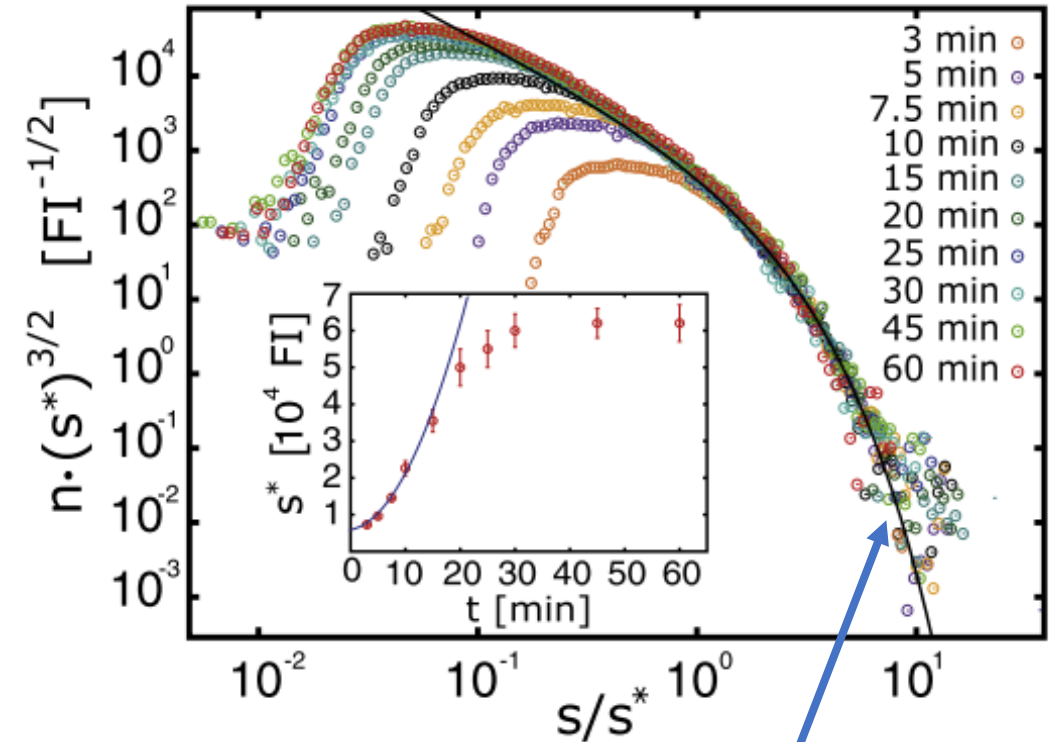
$$h(x, t) \cong -\left(\frac{2Jx}{K}\right)^{\frac{1}{2}} \tanh\left(\sqrt{3s^*(t)}x\right)$$

$$s^*(t) = \frac{JKt^2}{6}$$

$$xs^* \ll 1 \quad h(x, t) \cong -\left(\frac{6Js^*(t)}{K}\right)^{\frac{1}{2}} \frac{x}{1+xs^*(t)}$$

Taking the Inverse Laplace transform:

$$n(s, t) \cong \left(\frac{6J}{Ks^*(t)^3}\right)^{1/2} e^{-s/s^*(t)}$$



$$n_0 \left(\frac{s}{s^*}\right)^{-3/2} e^{-s/s^*}$$

Results: Comparison of the Model Predictions with the Experimental Data

$$h(x, t) = -\frac{\bar{k}(x)}{K} \tanh\left(\frac{t}{2} \bar{k}(x) + \tanh^{-1}\left(\frac{k_d}{\bar{k}(x)}\right)\right) + \frac{k_d}{K}$$

$$\bar{k}(x) = \left(\frac{2JKx}{1+s_0x} + k_d^2\right)^{1/2}$$

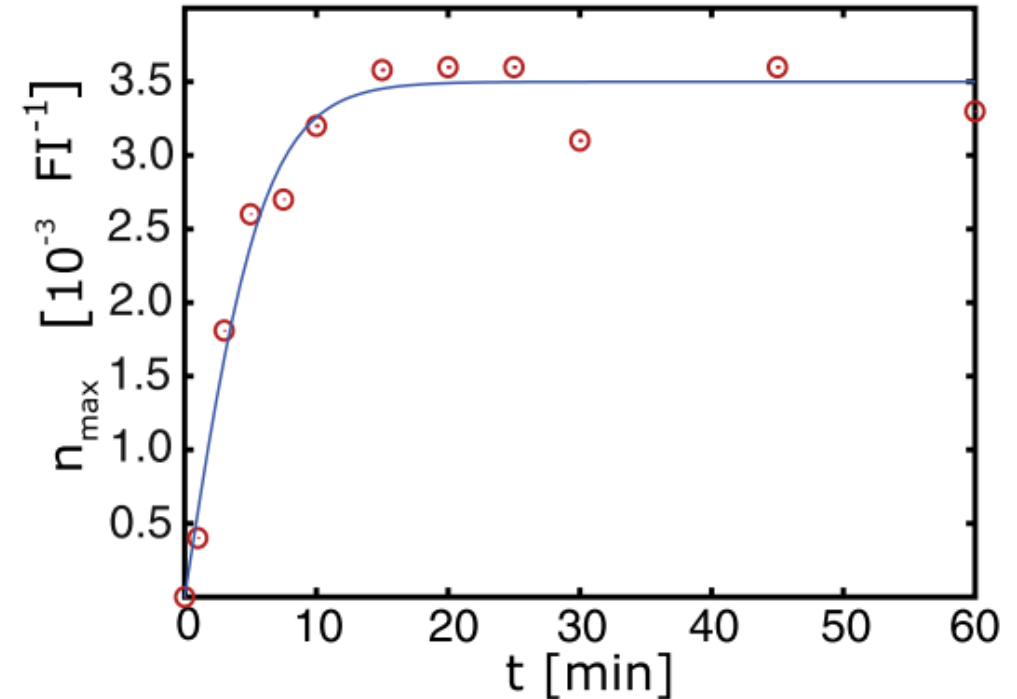
Time evolution of $n(s, t)$ for s small:

$$t \ll \frac{1}{k_d}, \text{ i.e., } k_d \approx 0 \quad s \ll s_0$$

$$h(x, t) \cong -\left(\frac{2J}{K}\right)^{\frac{1}{2}} \left(\frac{x}{1+s_0x}\right)^{\frac{1}{2}} \tanh\left(\left(\frac{3s^*(t)x}{1+s_0x}\right)^{\frac{1}{2}}\right)$$

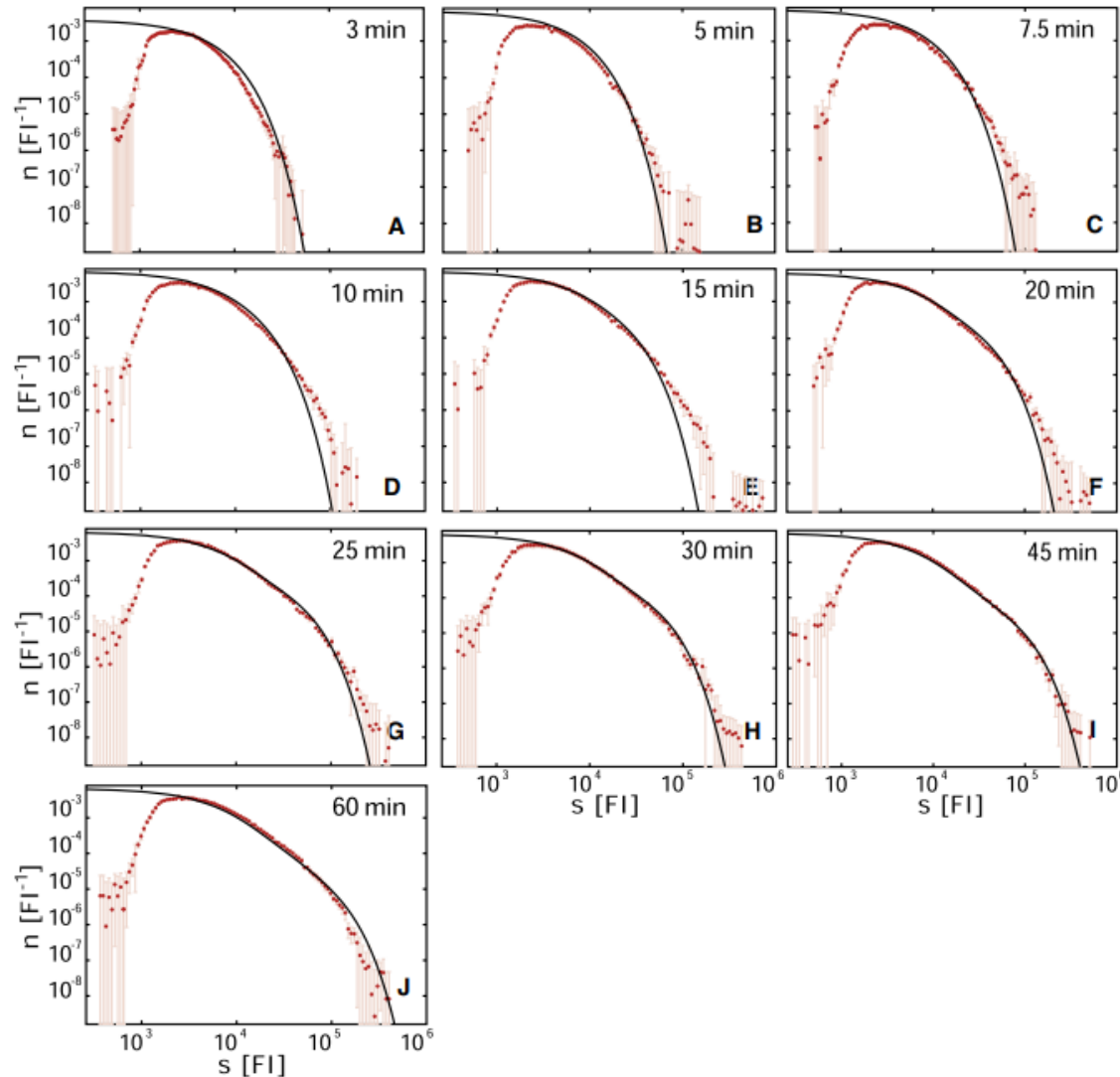
$s \ll s_0 \ll s^*(t)$ and taking the Inverse Laplace transform:

$$n(s, t) \cong \left(\frac{J}{2Ks_0}\right)^{1/2} \tanh(t/\tau) e^{-s/s_0} \quad \tau = \left(\frac{JK}{2s_0}\right)^{-1/2}$$



$$n_{max}(s, t) \propto \tanh(t/\tau)$$

Results: Comparison of the Model Predictions with the Experimental Data



Estimate of the Kinetic Rates for LDL concentrarion of $2.5 \mu\text{g/ml}$

Parameters of
EFE model

Parameter values

K (constant)	$K = 1.6 \cdot 10^{-4} \text{ s}^{-1}$
$(J/s_0^2)e^{-s/s_0}$	$J = 546 \text{ FI} \cdot \text{s}^{-1}$, $s_0 = 3600 \text{ FI}$
k_d (constant)	$k_d = 1.5 \cdot 10^{-3} \text{ s}^{-1}$

They also tested the influence of constant fission rate K' and exit average cargo amount per unit time v_{out} by fitting the data.

$$\frac{f_{fis}}{f_{fus}} = 0.26 \quad \frac{f_{v_{out}}}{f_{k_d}} < 5\%$$

Conclusions



A general theoretical framework to infer endosomal network dynamics from quantitative image analysis.

Three main processes of LDL trafficking and dynamics through the endosomal network: A source of cargo-carrying EE, homotypic EE fusion and endosome conversion.

They suggest that Rab conversion instead of vesicle budding is the principal mode of transport of LDL from early to late endosomes.

Their approach allows to measure kinetic transport rates from a series of still images.