



# Understanding Kidney Filtration via Electrokinetic-Hydrodynamic Concepts:

## A Foundry Guided Study

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### Introduction

The human kidney filters blood through nephrons, maintaining waste removal and overall homeostasis. Filtration begins in the glomerulus, where blood pressure drives water and small solutes such as ions, glucose, and urea across a filtration barrier into Bowman's capsule, forming filtrate. This filtrate then moves through the renal tubules, where essential substances are selectively reabsorbed and additional wastes are secreted, ultimately producing urine. To model this process, electrokinetic-hydrodynamics (EKHD) describes fluid flow and electrically driven solute transport across the filtration membrane, and when combined with the Renaissance Foundry Model (RFM), provides a simplified mathematical framework for understanding electroosmotic and electrophoretic transport within the glomerulus.

#### Mathematical Approach

A simplified EKHD-based model represents filtration as the combined effect of pressure-driven flow and electroosmotic transport, with an additional ion concentration term to capture chemical influence. The filtration rate is expressed as a linear combination of these factors, and total filtrate volume is obtained by integrating this rate over time. This approach captures the coupled roles of pressure, electrical potential, and ion exchange in kidney filtration while remaining a first-order approximation suitable for conceptual modeling.

First Set of the P.I.T:  $F(t) = aP(t) + bE(t) + ck[C(t)]$

### Prototype Of Innovative Technology

#### The P.I.T

Ion exchange is incorporated into the model through a time-dependent term,  $I(t)$ , which represents the influence of key physiologically relevant ions such as  $\text{Na}^+$ ,  $\text{K}^+$ , and  $\text{Cl}^-$  on the filtration process. These ions play a critical role in transport across the glomerular filtration barrier due to both their concentration gradients and their interactions with the charged components of the membrane, particularly the glomerular basement membrane (GBM). To capture this effect in a simplified manner, ion exchange is assumed to depend directly on ion concentration, where  $I(t) = k[C(t)]$ , and  $[C(t)]$  represents the overall ion concentration while  $k$  is a proportionality constant. A second constant,  $c$ , is introduced to quantify how strongly ion exchange contributes to the overall filtration rate. Incorporating this into the filtration model yields the expression:

$$F(t) = aP(t) + bE(t) + ck[C(t)]$$

For simplicity and clarity, the constants  $c$  and  $k$  are combined into a single parameter,  $d$ , allowing the model to be rewritten in a more compact and interpretable form. In this formulation, filtration is described as the combined contribution of pressure-driven flow, electroosmotic transport, and ion concentration effects, consistent with electrokinetic-hydrodynamics (EKHD) principles. The rate of filtrate formation is then expressed as:

$$\frac{dV}{dt} = aP(t) + bE(t) + d[C(t)]$$

Here, the pressure term represents hydrostatic forces driving fluid out of the glomerular capillaries, the electroosmotic term captures fluid motion influenced by charge interactions across the filtration membrane, and the concentration-dependent term reflects the overall chemical influence of dissolved ions on transport behavior. The total filtrate volume over time is obtained by integrating the filtration rate, giving:

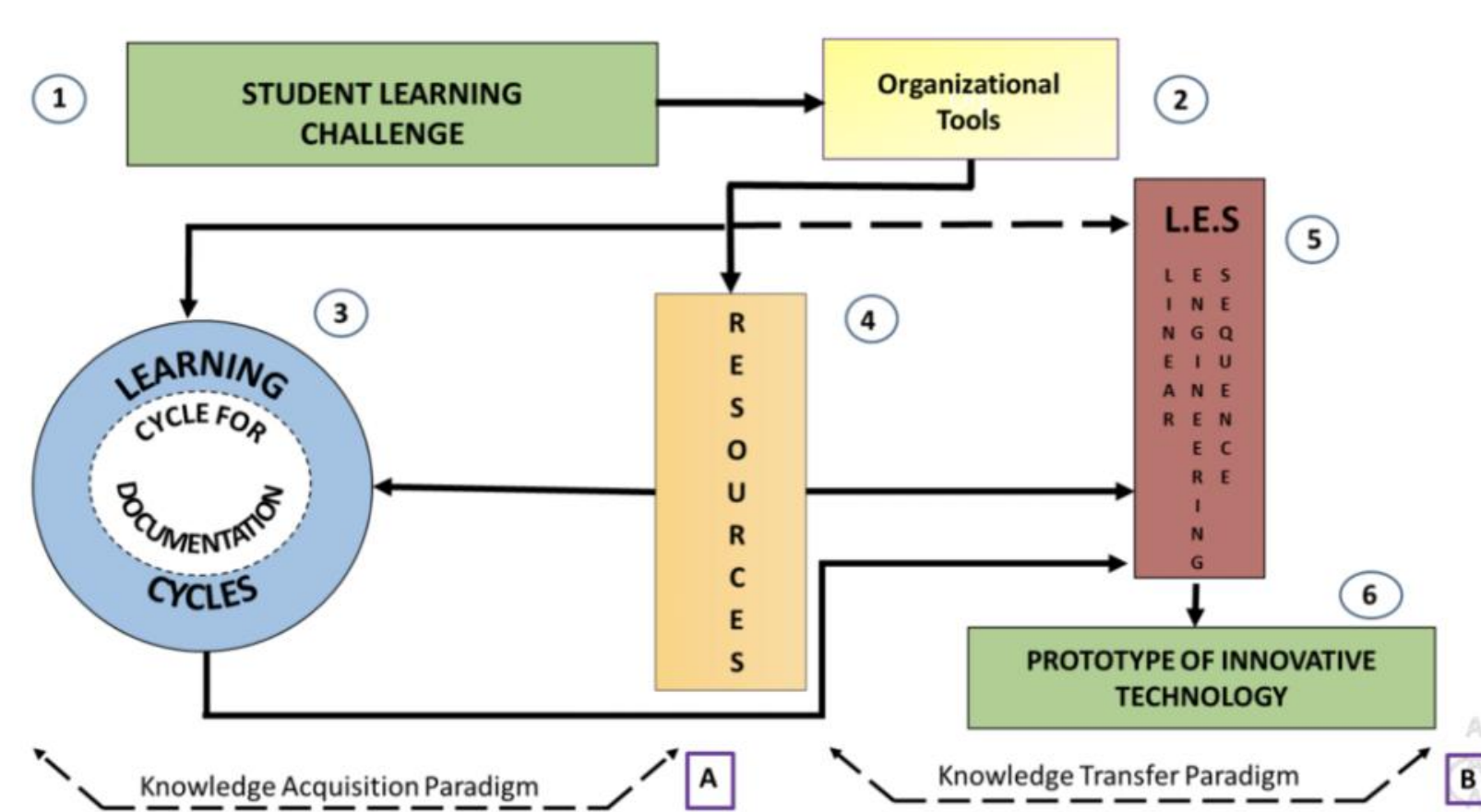
$$V(t) = \int (aP(t) + bE(t) + d[C(t)]) dt$$

This model provides a simplified, first-order representation of kidney filtration that captures the essential coupling between pressure, electrical forces, and chemical composition. While it does not explicitly model individual ion-specific reactions, equilibrium dynamics, spatial variation, or nonlinear transport, it serves as a conceptual EKHD-based framework that highlights the dominant mechanisms governing glomerular filtration while remaining mathematically accessible and suitable for future refinement.

#### Observations & Limitations

The proposed model is a simplified, first-order approximation that assumes linear relationships between filtration rate and contributing variables. There are a variety of factors this model does not account for due to its mathematical limitations, such as spatial variation, nonlinear transport, diffusion, and selective permeability. This is a conceptual framework, derived from EKHD principles, and not a fully predictive physiological mathematical model.

### The RFM



#### RFM Mapping to Kidney Microfiltration

##### Challenge:

Understand glomerular filtration and model electroosmotic effects using EKHD within the RFM framework.

##### Approach:

**Tools:** Nephron physiology + EKHD concepts (charge, flow, transport)

**Cycle:** Question → Physiology → EKHD concept → Model → Interpret → Refine

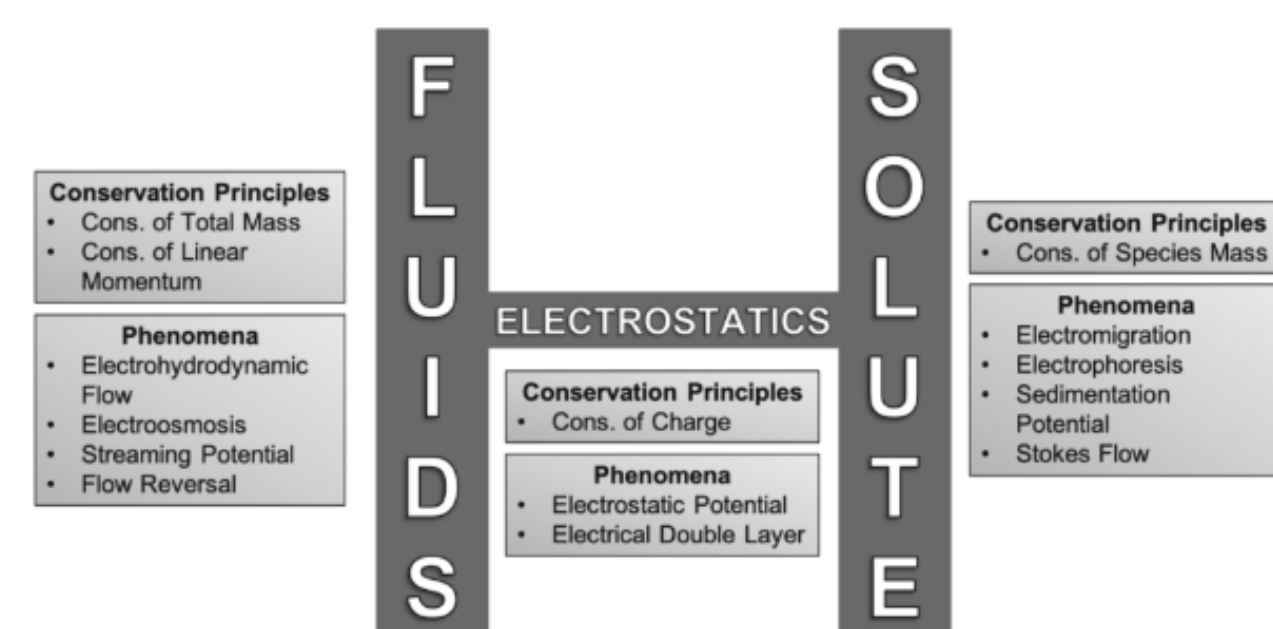
**Resources:** Filtration structure, EKHD framework, Calculus I-II

##### Application:

Link electrostatic potential → fluid flow → solute transport; analyze how parameter changes affect filtration.

##### Output:

Simplified EKHD-based mathematical model + conceptual flowchart.

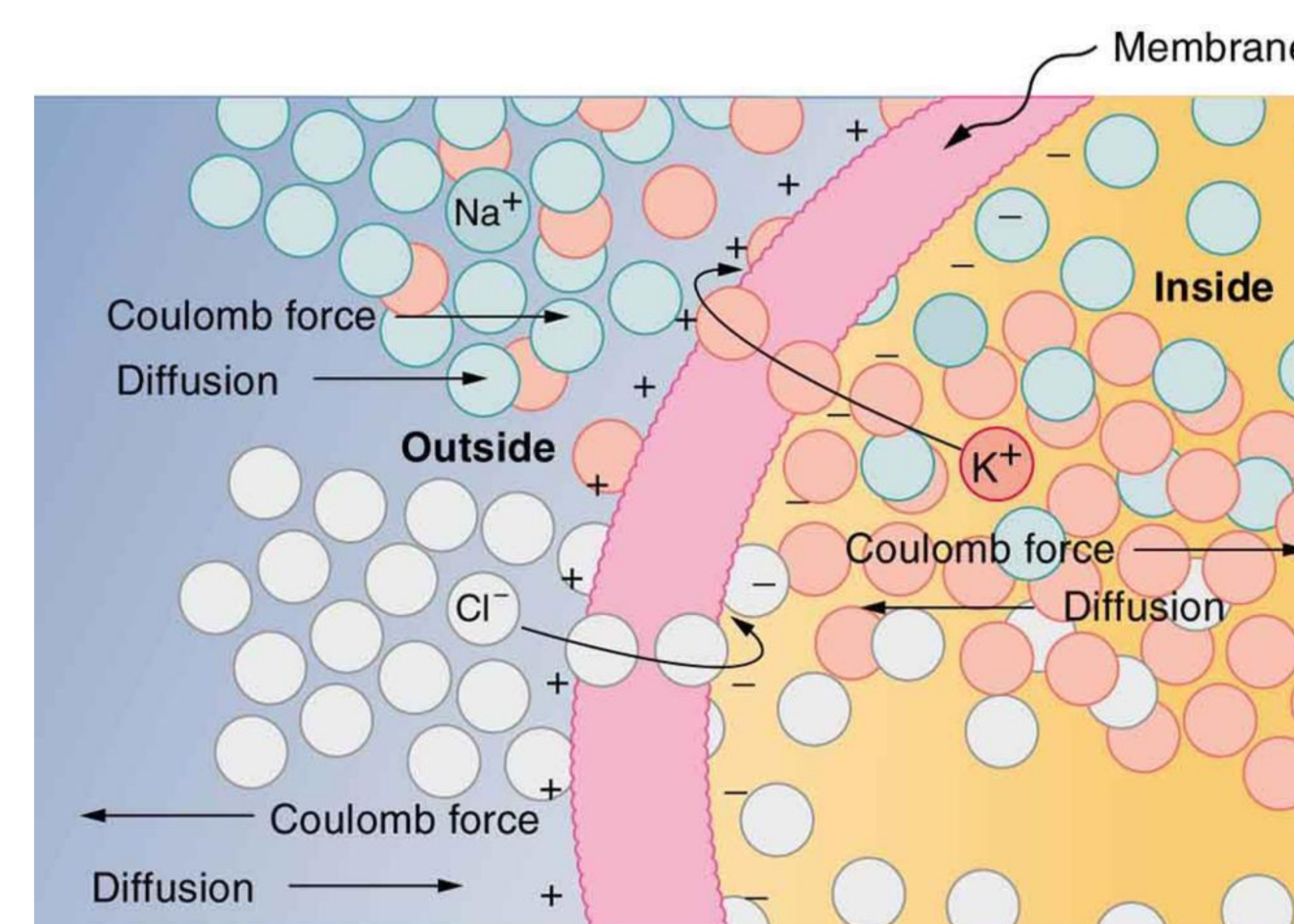
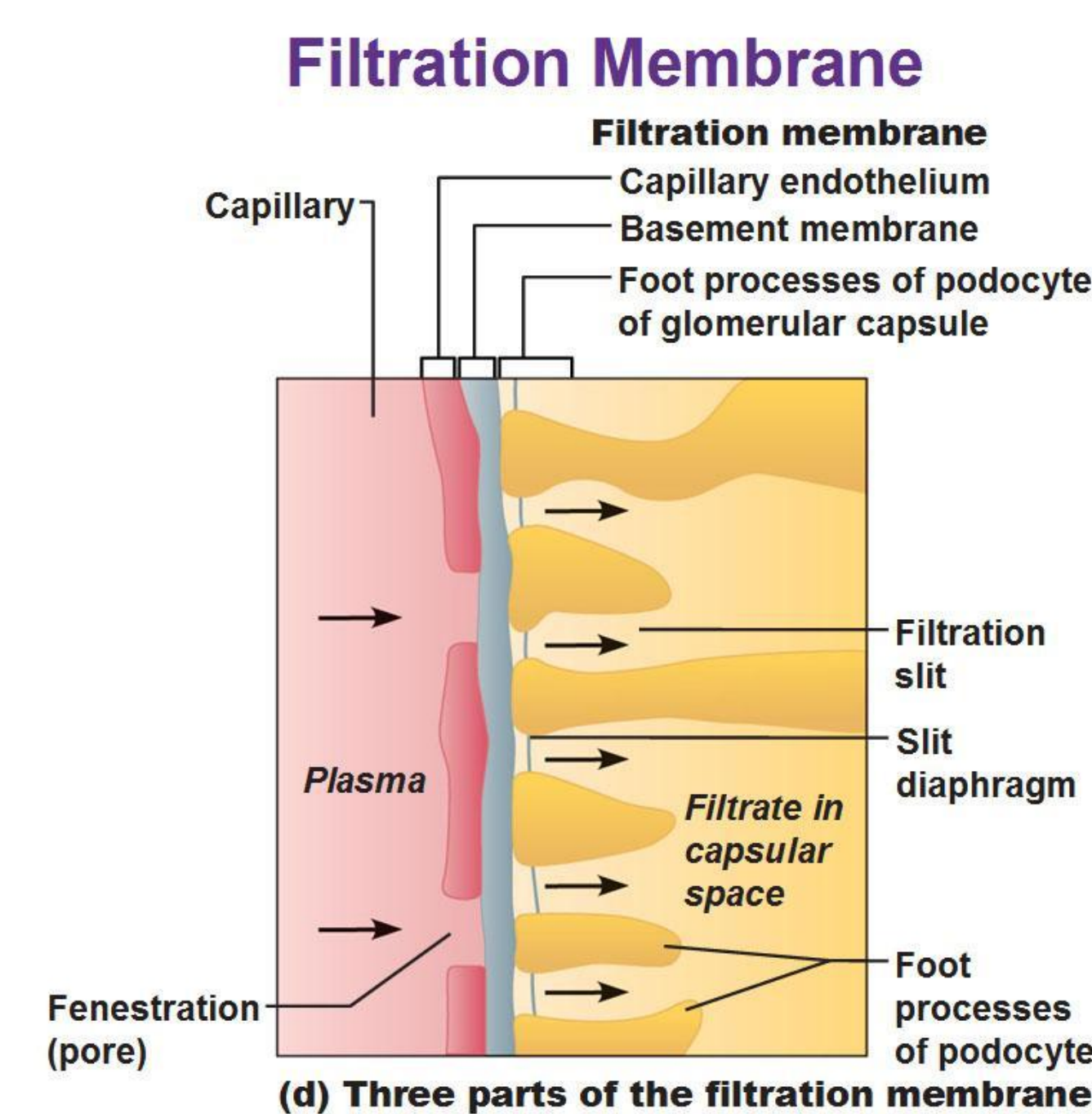


### Future Work

Future work will focus on giving physical meaning to the model parameters  $a$ ,  $b$ , and  $d$  by connecting them to real chemical and physical properties within the kidney filtration system. In the current model, these constants represent pressure-driven flow, electroosmotic effects, and ion concentration influence, but they are not yet tied to measurable quantities. The next step is to interpret  $a$  in terms of glomerular pressure and membrane permeability,  $b$  in terms of electrostatic potential and fluid mobility, and  $d$  in terms of ion concentration, charge density, and interaction strength with the filtration barrier. This refinement will move the model from a simplified conceptual form toward a more physically grounded framework. The Renaissance Foundry Model will continue to guide this process by structuring how biological understanding, mathematical formulation, and physical chemistry are integrated throughout the project.

At a larger scale, the images below provides a useful visualization of how Coulomb forces and diffusion contribute to ion transport across a charged membrane. The glomerular basement membrane contains fixed negative charges that influence how ions such as  $\text{Na}^+$ ,  $\text{K}^+$ , and  $\text{Cl}^-$  move through the system. Positive ions are attracted toward the membrane, while negative ions may be repelled, creating a selective transport environment. This behavior aligns with the inclusion of the concentration-dependent term in the model, since ion movement is influenced not only by diffusion but also by electrostatic interactions. These interactions can be interpreted macroscopically as a balance between concentration gradients and charge-based forces, both of which contribute to the overall filtration rate.

To further strengthen the model, future work may incorporate ideas from molecular quantum mechanics to better understand ion-membrane interactions at a fundamental level. Charge distribution, potential energy surfaces, and electron density around ions and membrane functional groups could provide insight into why certain ions are preferentially transported or restricted. While the current model treats ion effects in a simplified, averaged way, a quantum-informed perspective could help justify parameter values and improve how electrostatic effects are represented. By linking molecular-scale interactions to macroscopic transport behavior, the model can be refined to better reflect the underlying physics and chemistry governing kidney microfiltration.



### References

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### Conclusion

- The proposed EKHD-based model captures kidney filtration as a coupled process involving pressure-driven flow, electroosmotic effects, and ion concentration.
- Incorporating ion exchange provides a more realistic representation of how charge and chemistry influence transport across the filtration barrier.
- The Renaissance Foundry Model supports a structured approach to connecting physiology, mathematics, and physical chemistry in model development.
- This framework serves as a simplified foundation that can be refined with physically meaningful parameters and extended to better represent real kidney function.

### Acknowledgements

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