

# Towards the Simulation of Bulk Properties via Molecular Dynamics

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# Biomolecular Simulations under Realistic Macroscopic Salt Conditions

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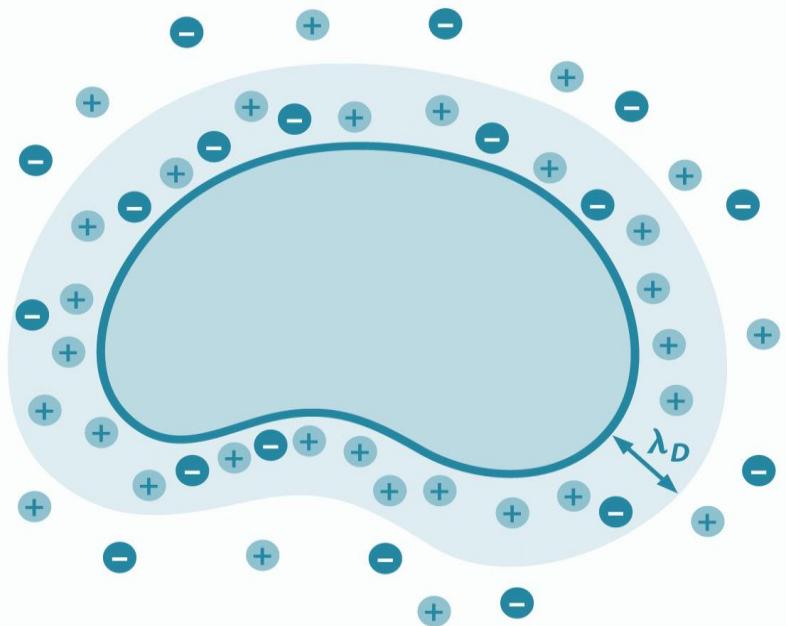
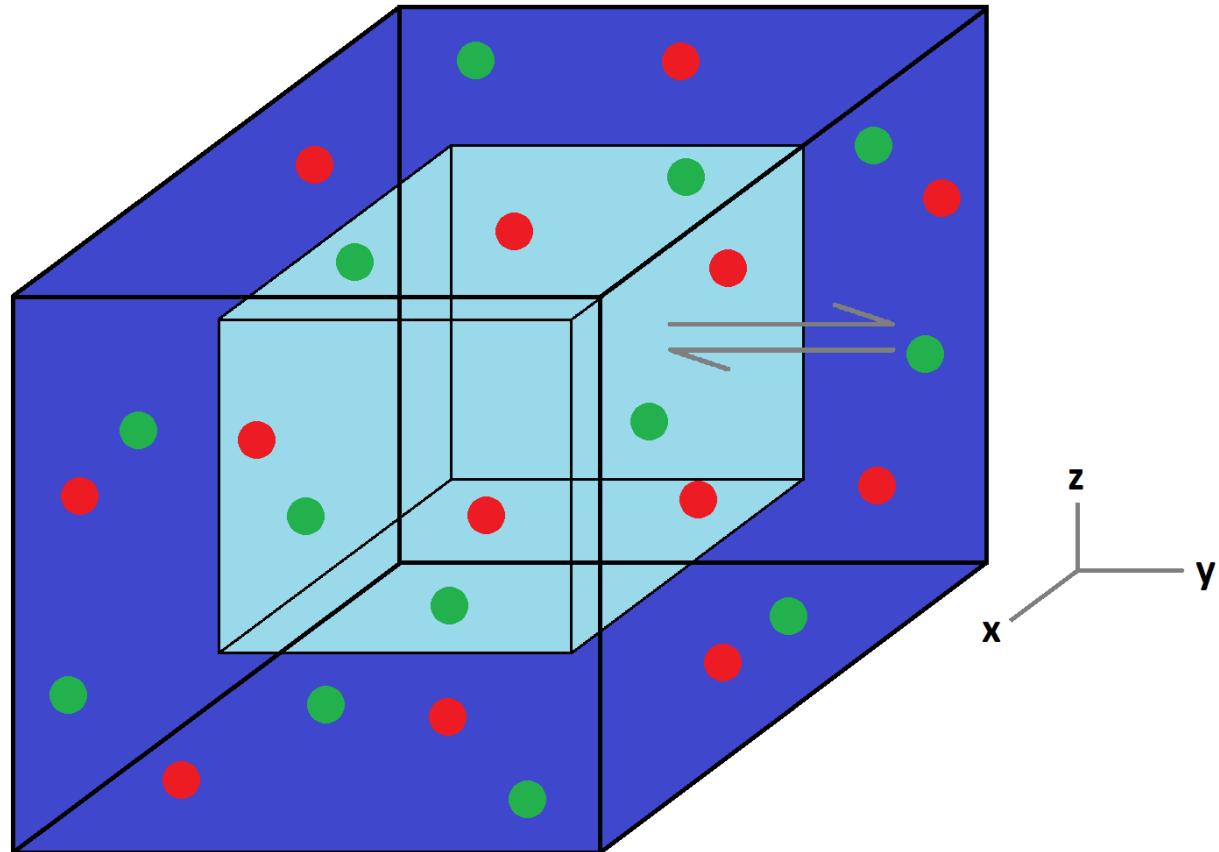


Figure 9.15a Physical Biology of the Cell (© Garland Science 2009)

- Conformations, functions, binding of biomolecules are sensitive to ion content
  - Relevant to pharmacological design
  - Simulations fix salt content and so may not accurately represent actual local environment



■ -- local environment  
(simulation volume)

■ -- nonlocal environment  
(bulk)

● -- positive ion

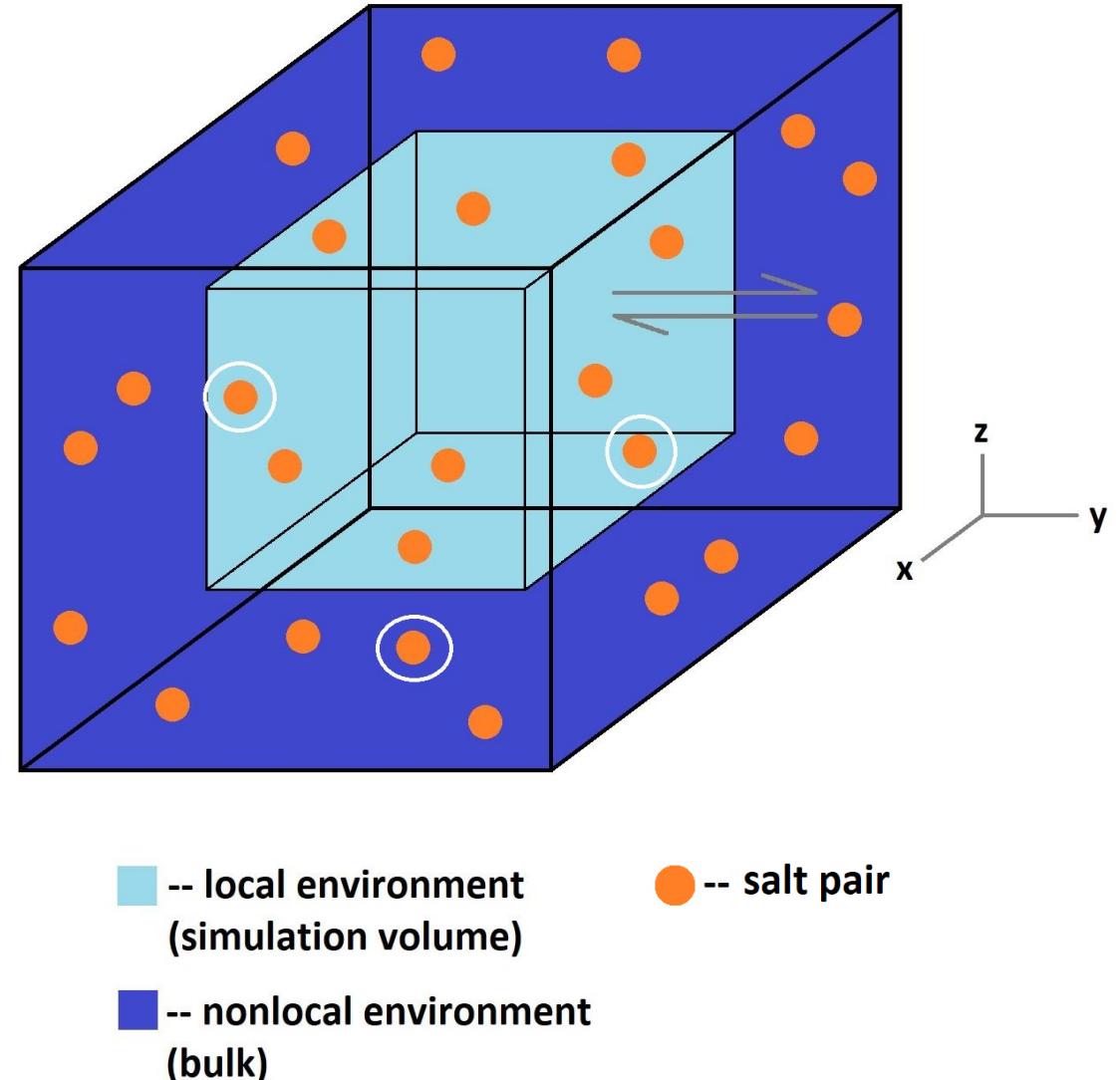
● -- negative ion

# Fluctuations in Local Environment

- How do fluctuations affect processes like folding, screening layer interactions, and binding?
- How do fluctuations affect free energy?

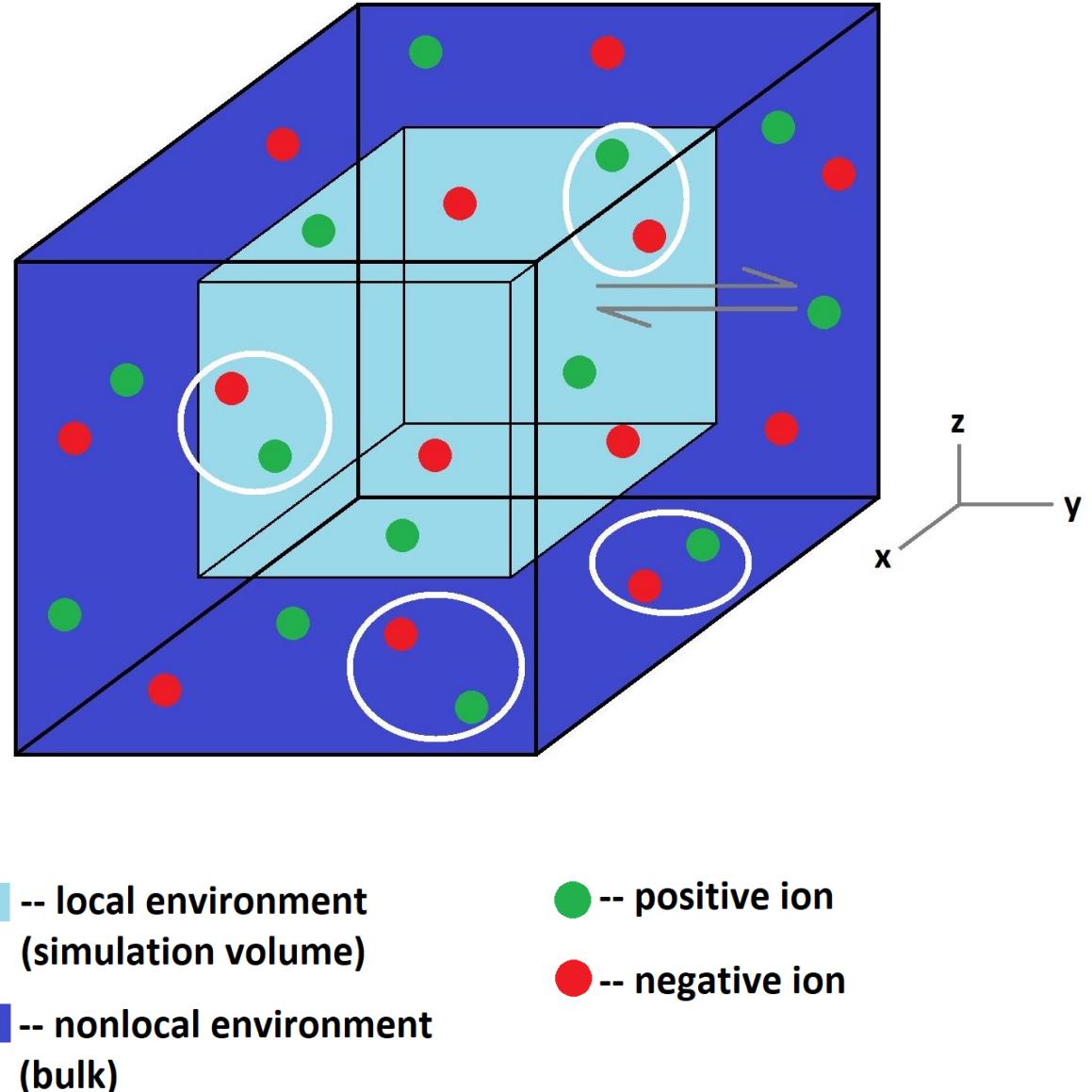
# Single particle model

- One degree of freedom
- Anions and cations fluctuate together
- Simplest but least physical



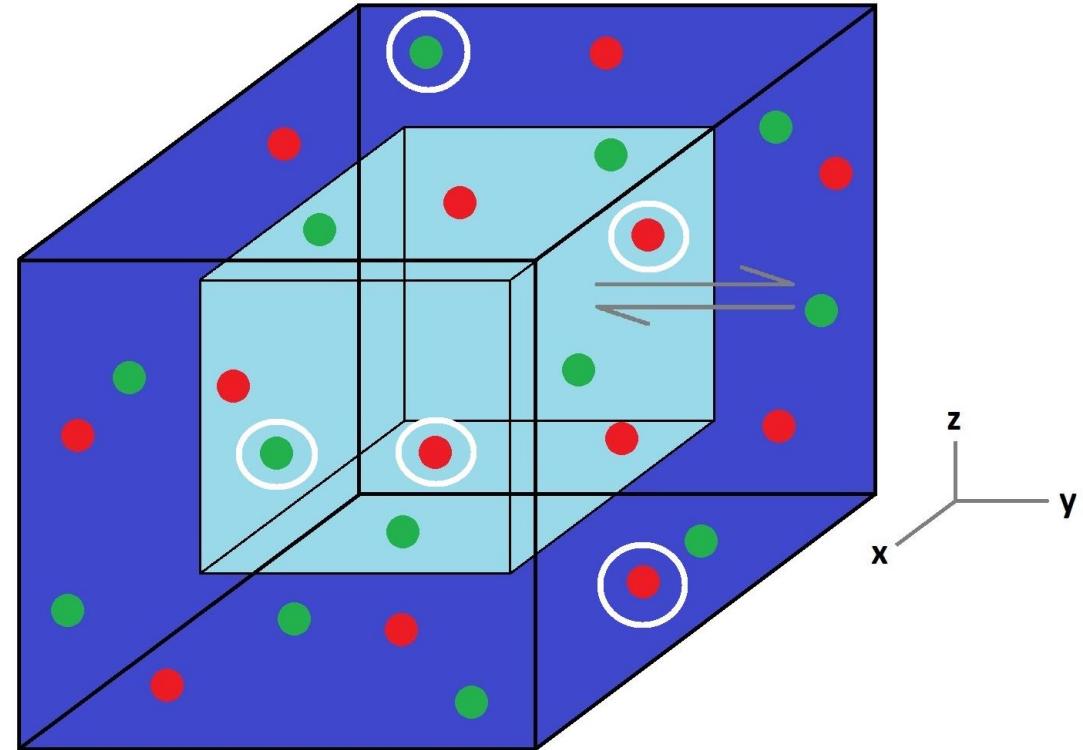
# Salt pair model

- Two degree of freedom
- Anions and cations fluctuate together
- No electrostatic contribution
- Commonly used for computations



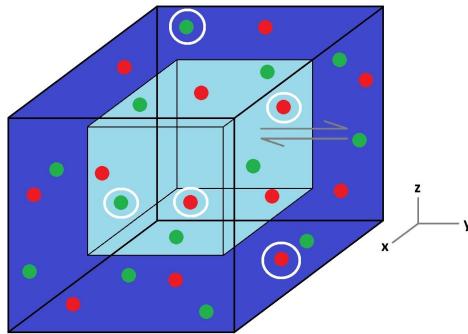
# Independent fluctuations

- Two degree of freedom
- Anions and cations fluctuate independently
- Electrostatic contribution



■ -- local environment  
(simulation volume)  
■ -- nonlocal environment  
(bulk)

● -- positive ion  
● -- negative ion



# Describing independent fluctuations

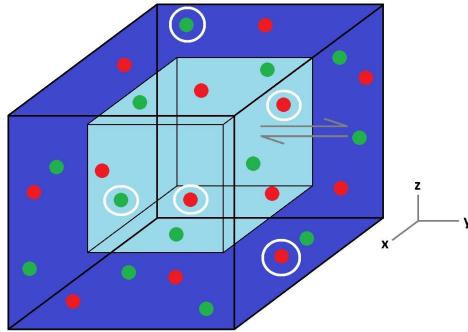
Electrostatic  
component

Two chemical  
components

$$Z = \sum_{N_+, N_-} \frac{V^{N_+}}{N_+!} * \frac{V^{N_-}}{N_-!} * e^{\frac{e^2}{8\pi\epsilon R kT} (Q + N_+ - N_-)^2} * e^{(N_+ + N_-) \frac{\mu}{kT}}$$

Two translational  
components

$N_{\pm}$  = number of  $\pm$  ions  
 $V$  = volume of local environment  
 $Q$  = charge of macromolecule  
 $R$  = radius of local environment  
 $\mu$  = chemical potential  
 $e$  = charge of electron



# Evaluating Free Energy using SLTCAP

$$F(N_+, \delta N_+, N_-, \delta N_-, Q, V, \mu) = -kT \ln Z$$

$$N_{\pm} = V c_0 * e^{\mp \text{arcsinh}(\frac{Q}{2eVc_0})}$$

$N_{\pm}$  = number of  $\pm$  ions

$\delta N_{\pm}$  = number of fluctuations

$V$  = volume of local environment

$Q$  = charge of macromolecule

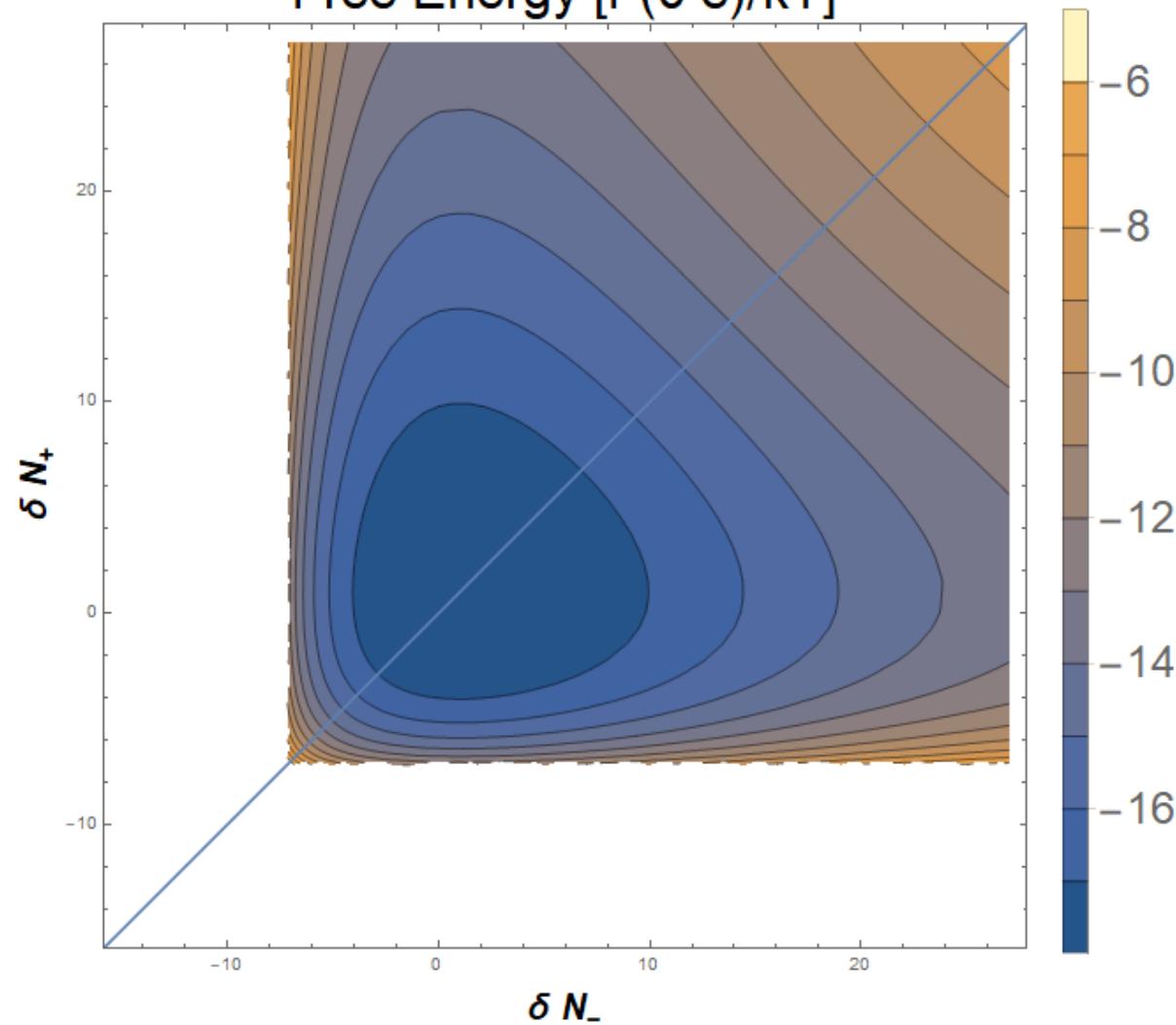
$R$  = radius of local environment

$\mu$  = chemical potential

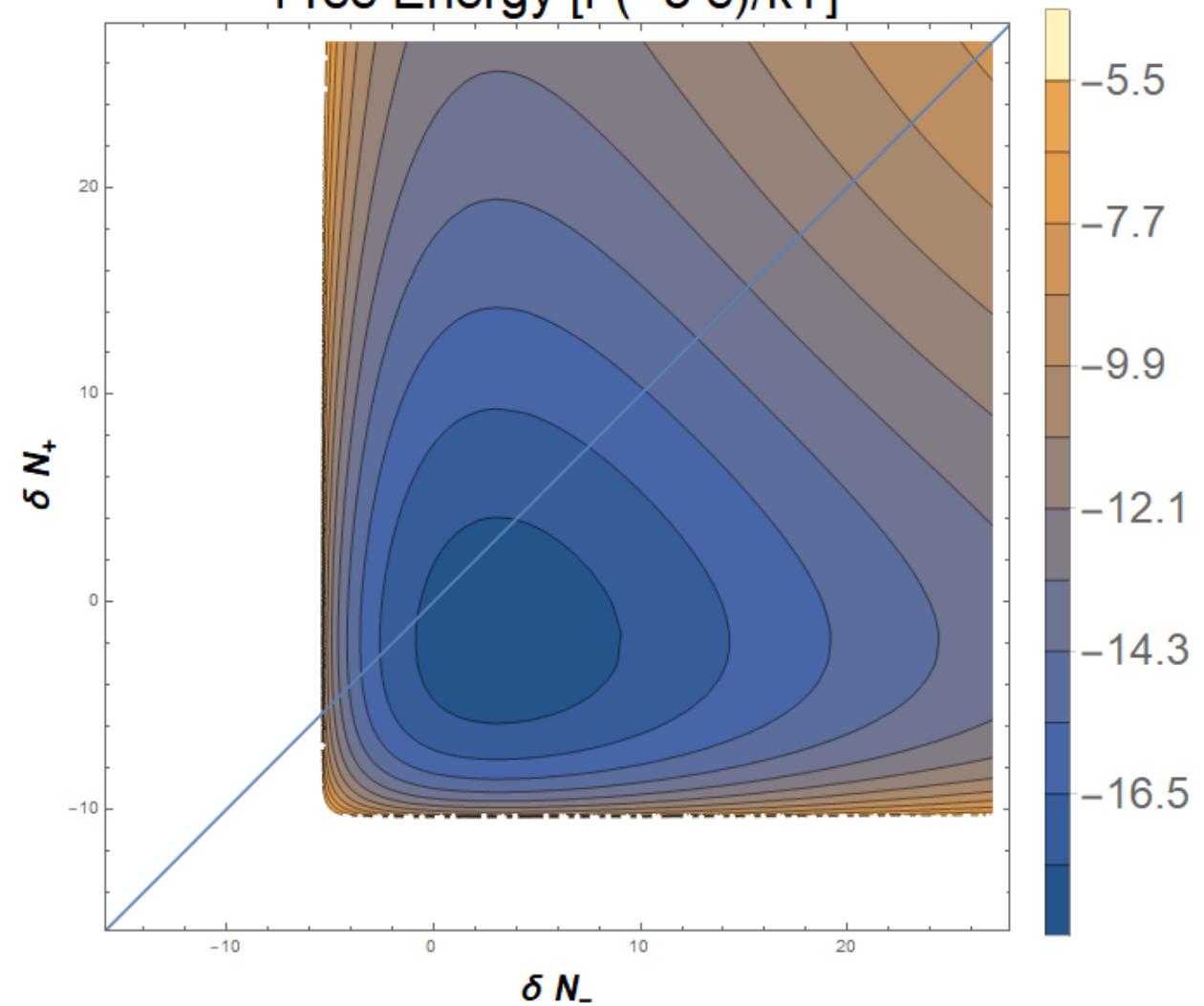
$e$  = charge of electron

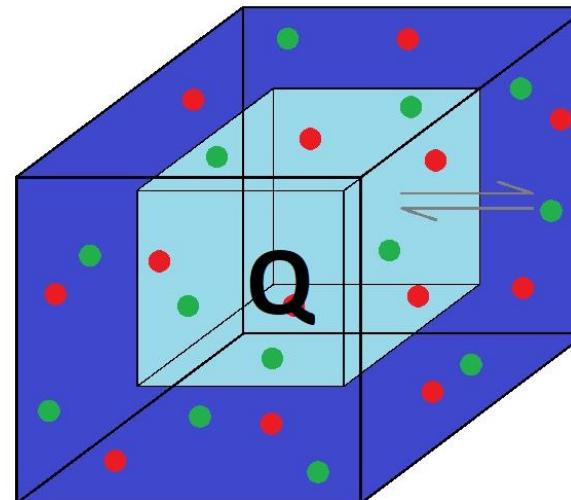
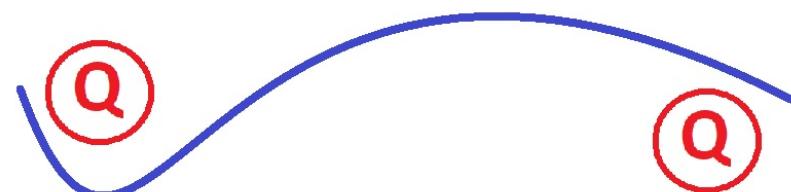
$$F(\delta N_+, \delta N_-, Q, V, c_0)$$

Free Energy [ $F(0 \text{ e})/kT$ ]

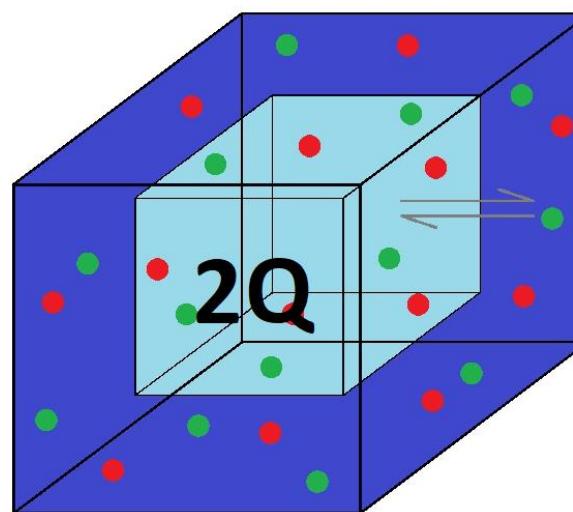
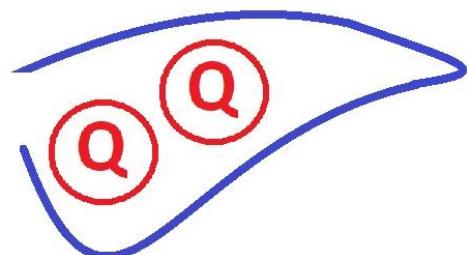
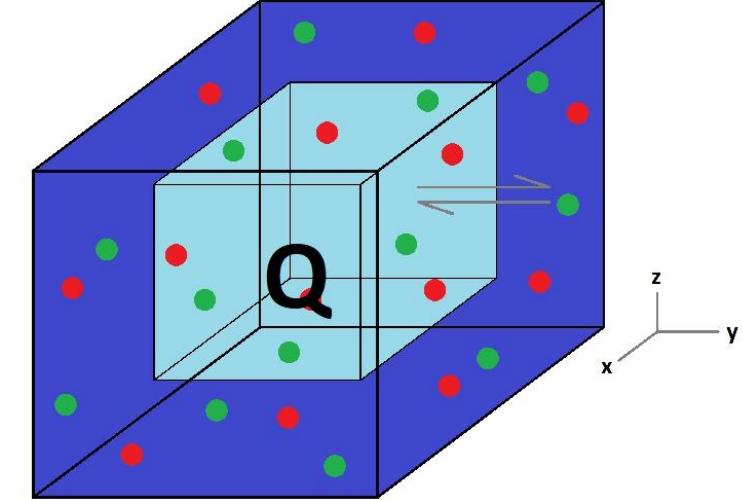


Free Energy [ $F(-5 \text{ e})/kT$ ]

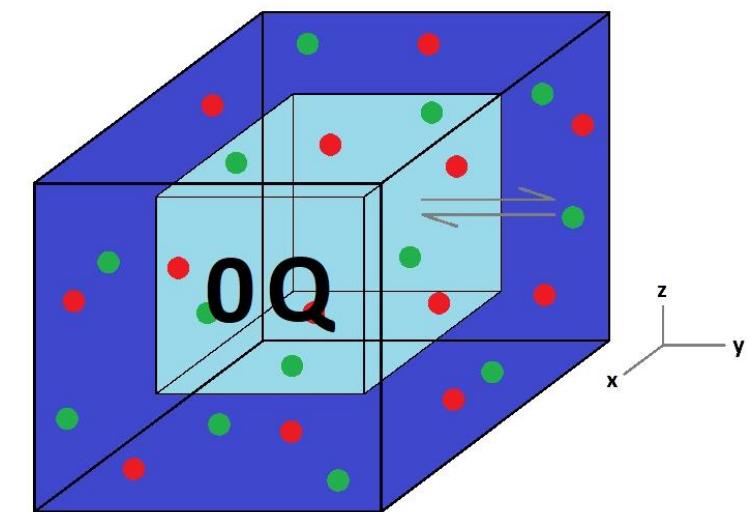




+



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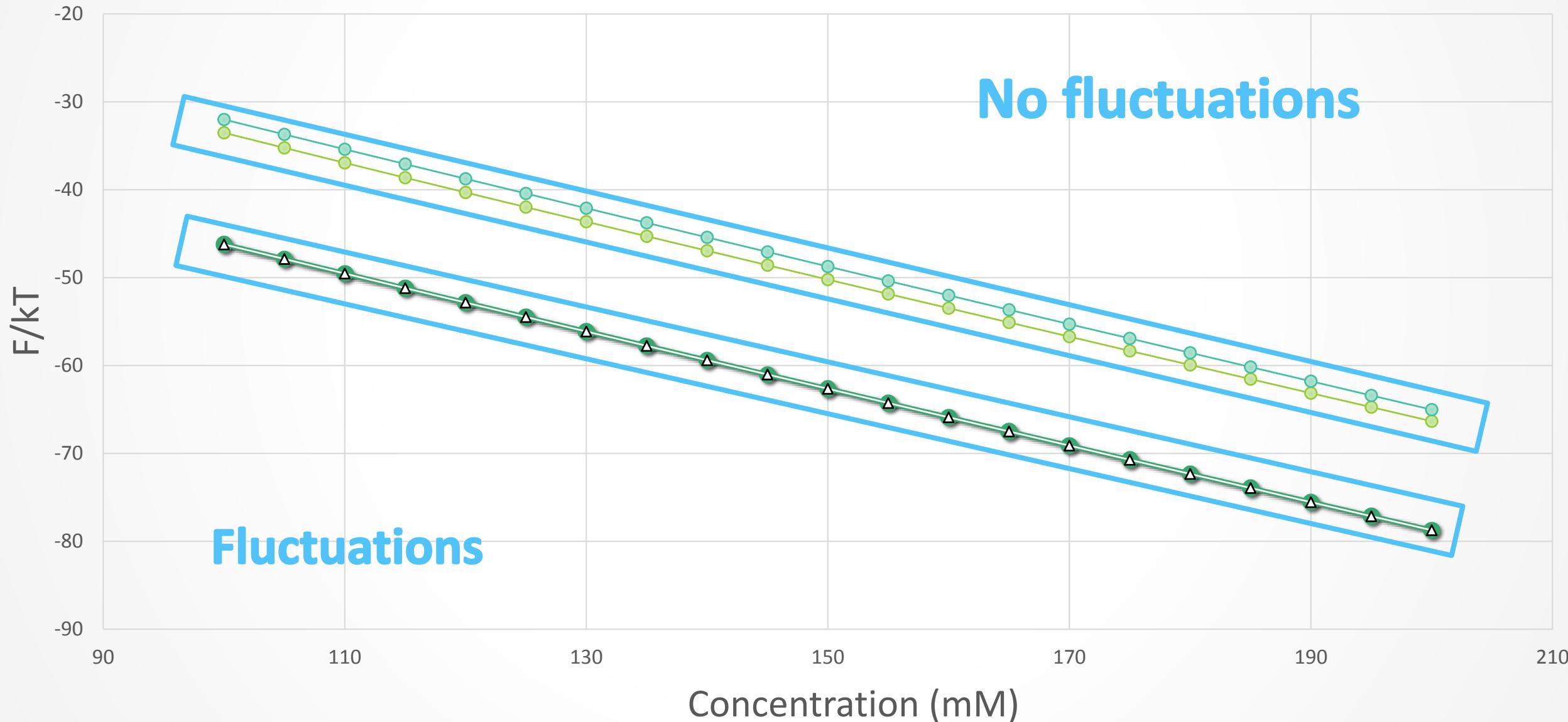
-- local environment  
(simulation volume)

-- nonlocal environment  
(bulk)

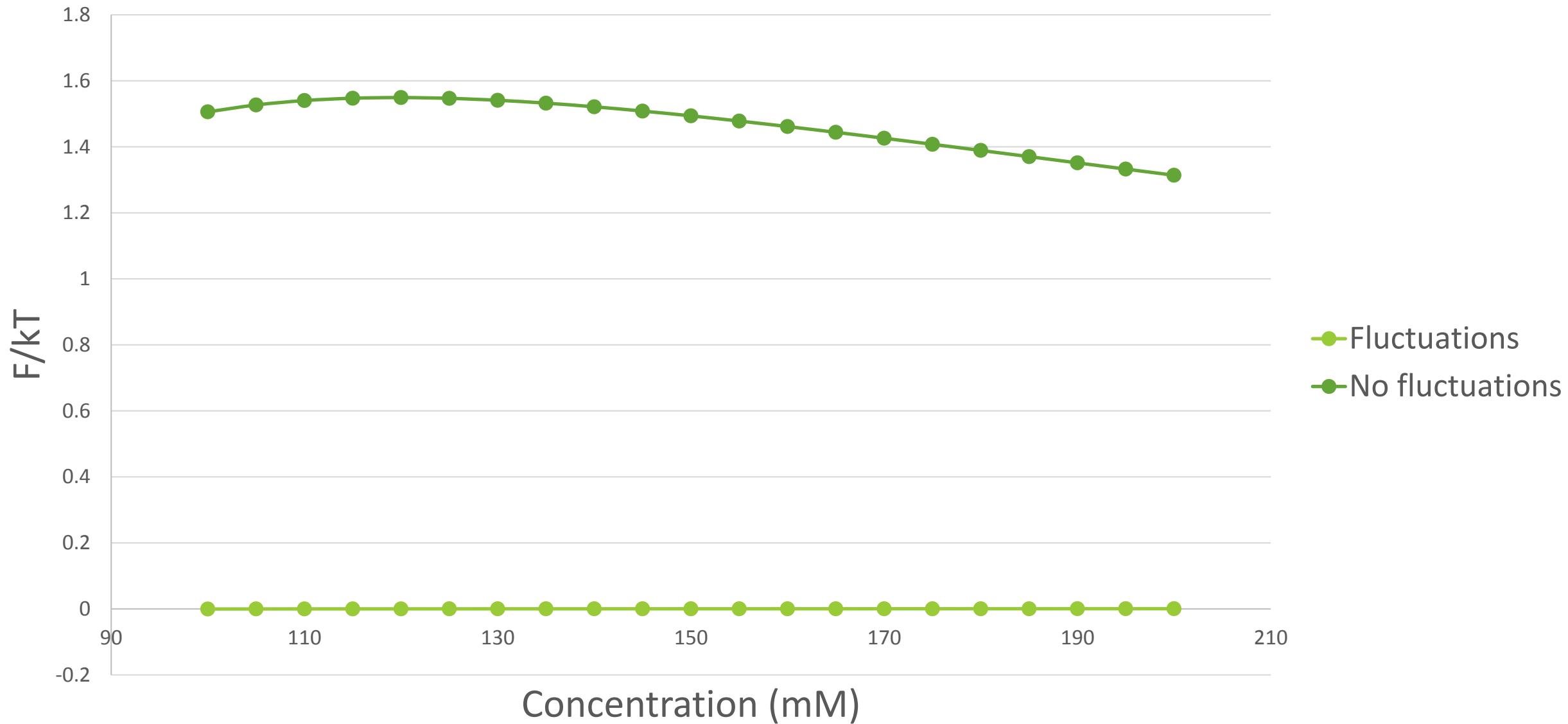
-- positive ion  
 -- negative ion

# Free Energy

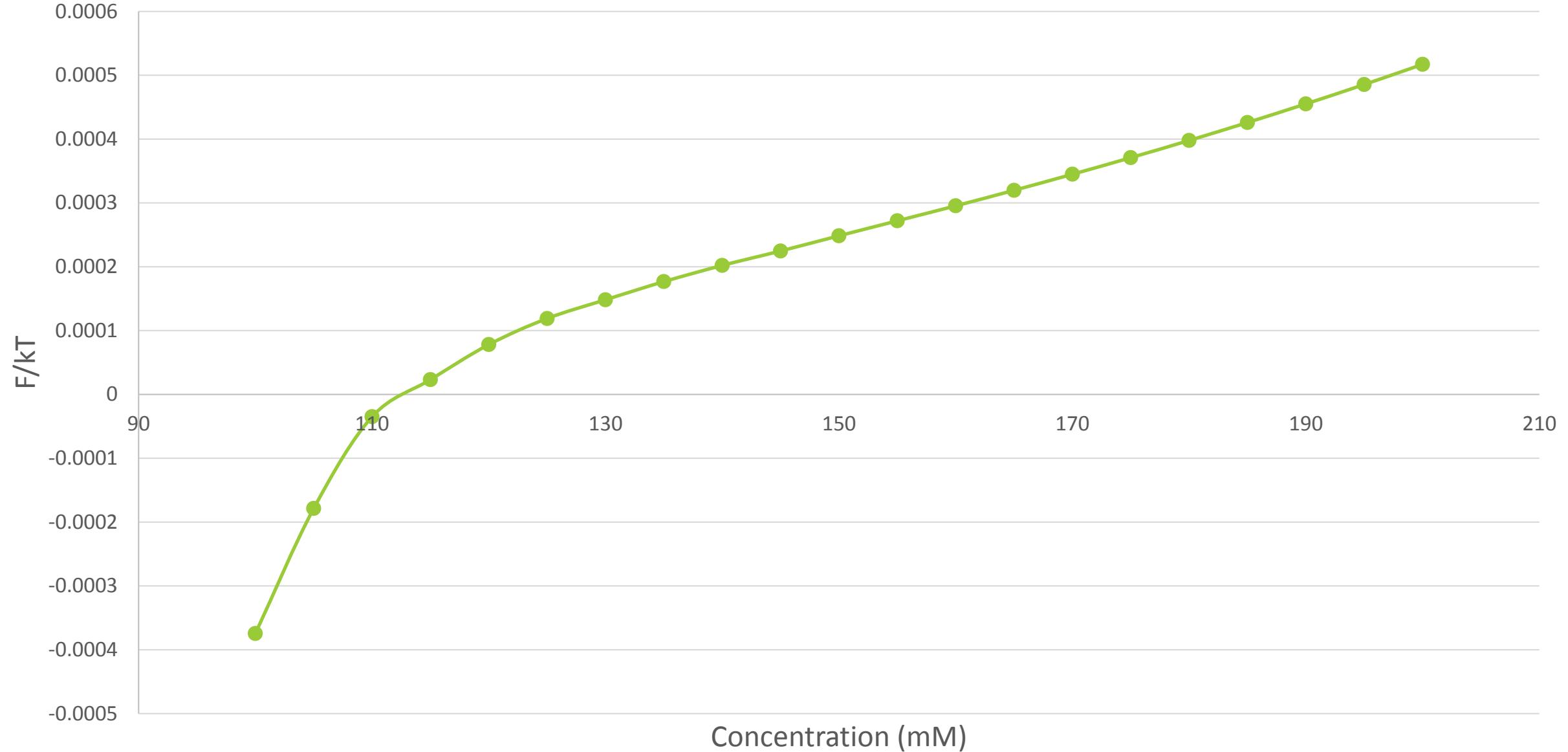
●  $2^*F(-10 \text{ e}): \text{fluc}$  ○  $2^*F(-10 \text{ e}): \text{no fluc}$  △  $F(0 \text{ e}): \text{fluc} + F(-20 \text{ e}): \text{fluc}$  ●  $F(0 \text{ e}): \text{no fluc} + F(-20 \text{ e}): \text{no fluc}$



$$\Delta F [ 2F(-10 \text{ e}) - (F(0 \text{ e}) + F(-20 \text{ e})) ]$$



# $\Delta F_{\text{fluctuations}}$



# Further Research

Explain why  $\Delta F_{fluctuations} \ll \Delta F_{no fluctuations}$

Collaborate to implement model in simulations



# Acknowledgements

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