

# *Differential geometry, topological invariant and machine learning approaches to virus dynamics*

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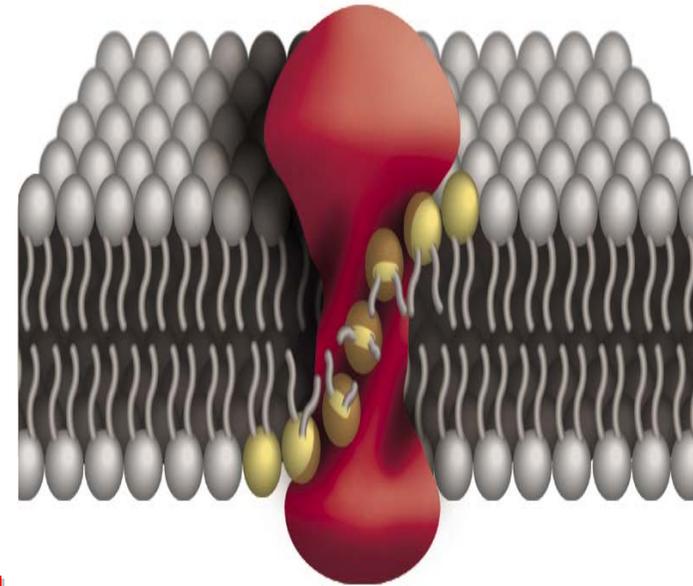
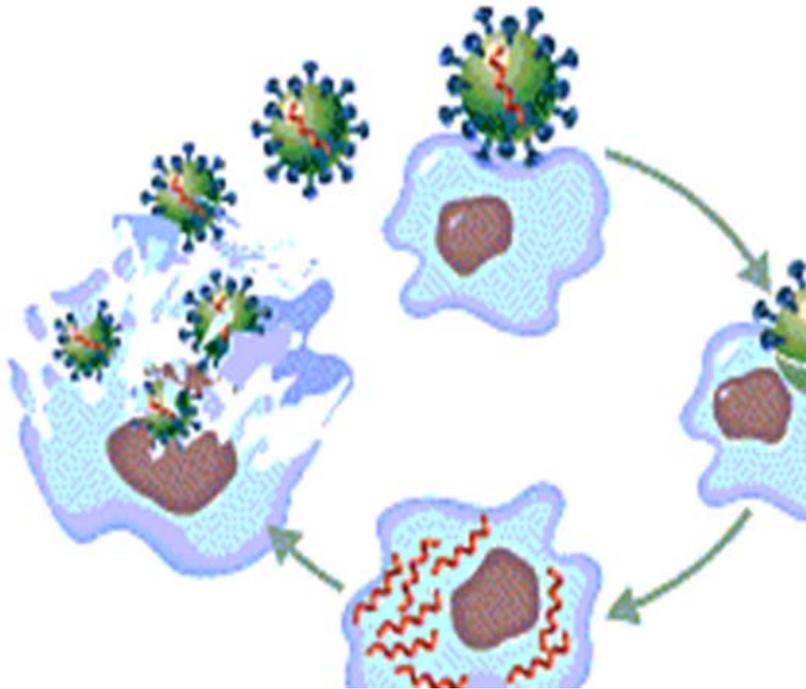
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# Differential geometry, topological invariant and machine learning approaches to

- Understand molecular mechanism of virus life circles
- Develop visual-analytic methods for virus infection prevention
- Extract biological functions and properties from dynamic data



## Challenges

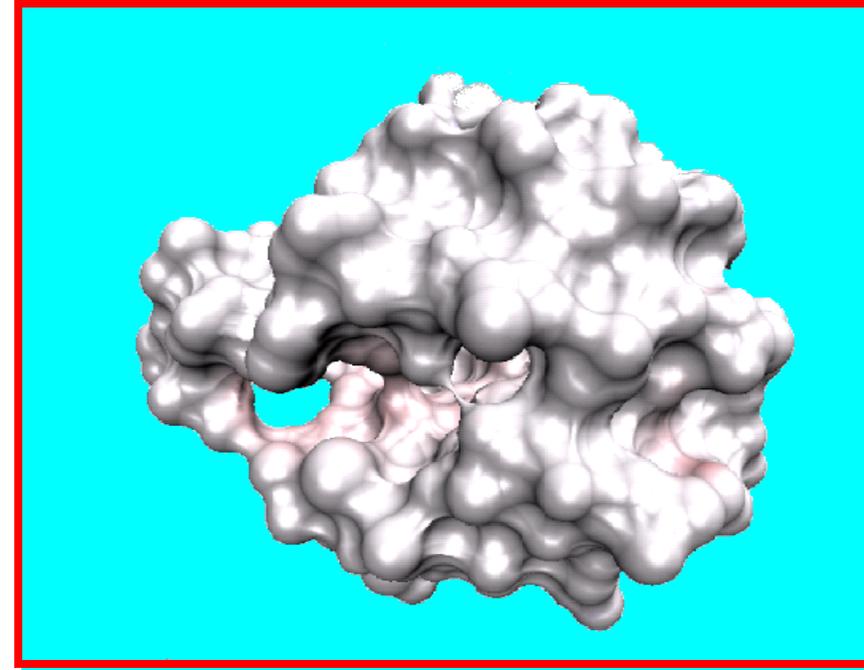
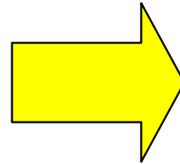
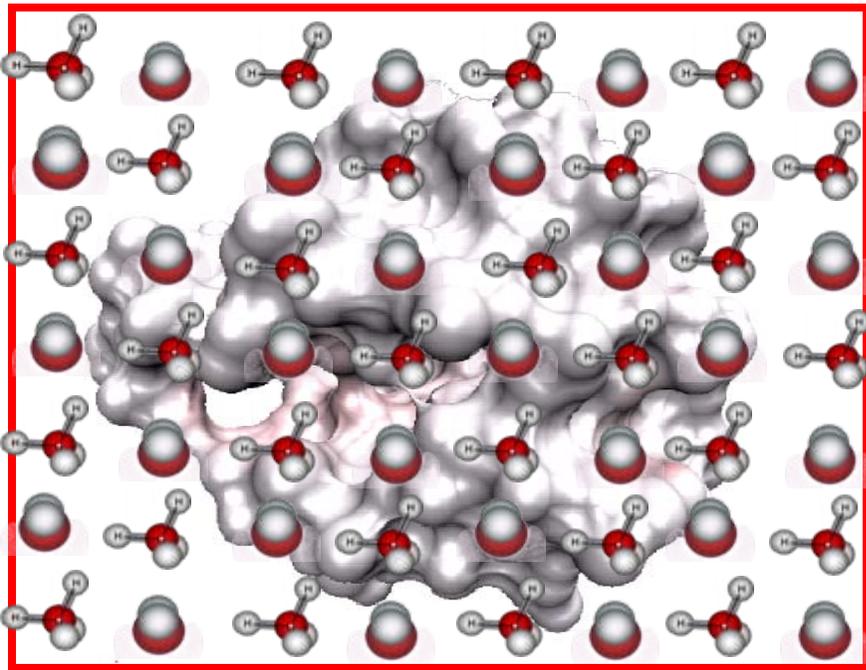
High dimension: ~ 10 million dimensions

Massive data sets: ~  $10^{18}$  data points

No viable physical/Mathematical models

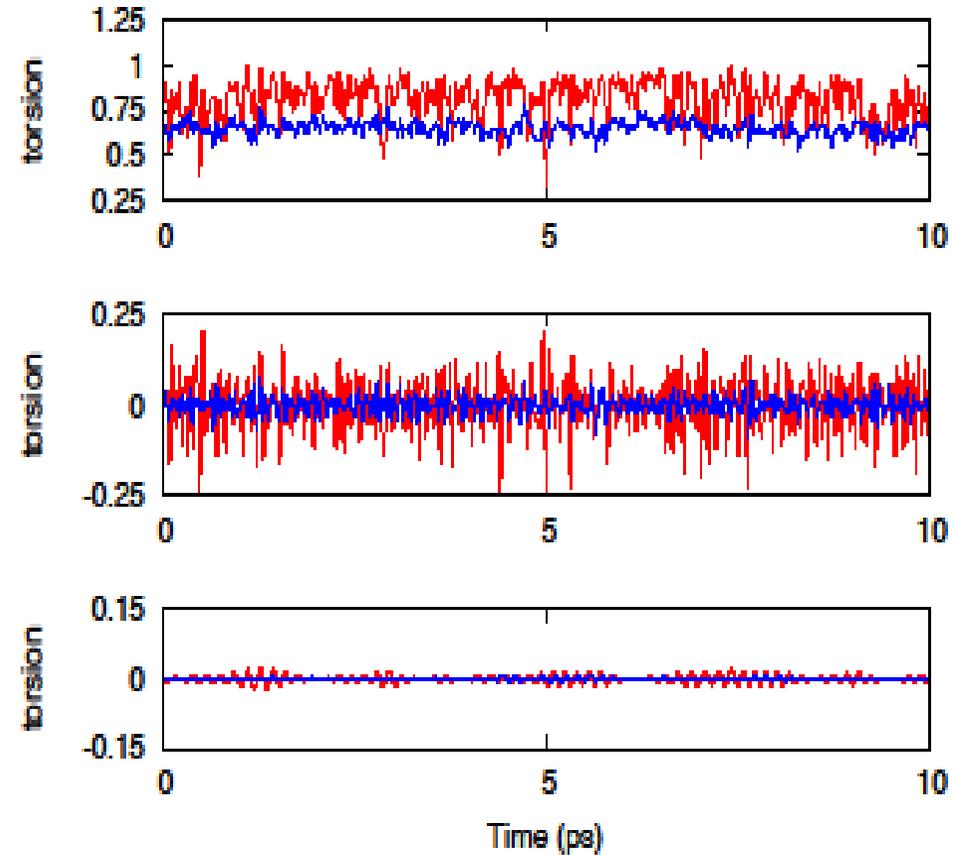
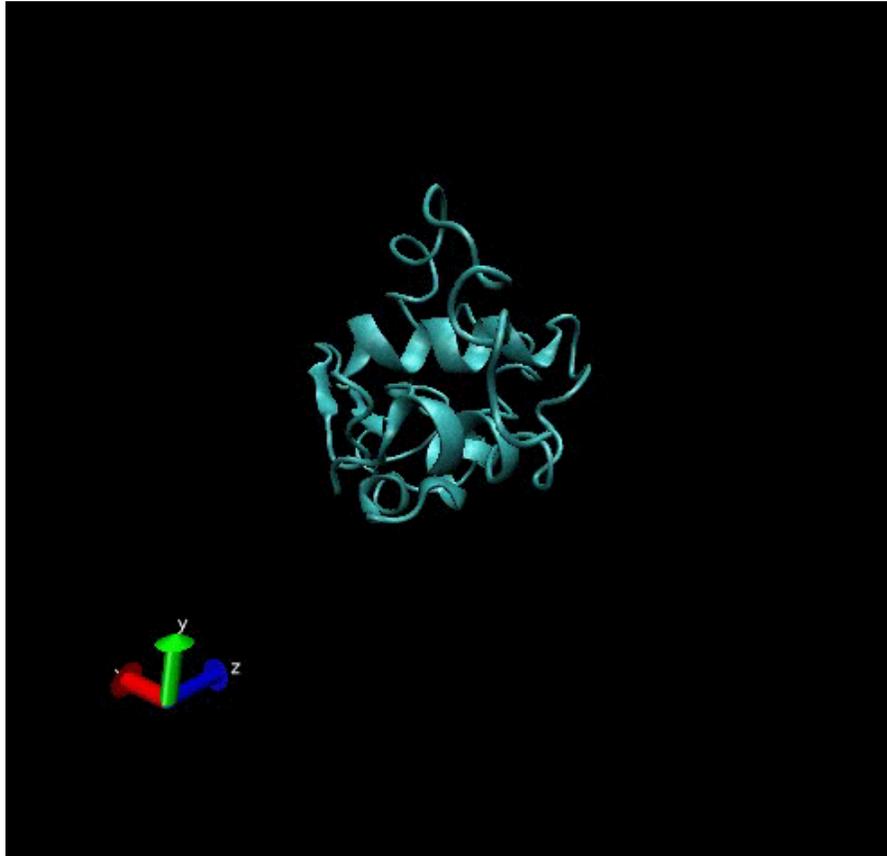
# Dimension reduction by multiscale analysis

- ⑩ Solvent is described by continuum models
- ⑩ Viruses are described by discrete models



The interface between the discrete and the continuum is described by differential geometry theory of surfaces  
(Reduce the dimension by about one order)

# Coarse-grained dynamic model based on persistently stable manifolds characterized by the time series of Frenet – Serret frames, torsion angles and curvatures



Machine learning approach to further reduce the dimension by 1 to 3 orders

(Tong, Wang, Wei, Zhou, 2010)

# Differential geometry based multiscale free energy functional for excessively large data size reduction of virus systems

$$\min G = \min \iiint \{ \textit{Geometric} + \textit{Electro} + \textit{Fluid} + \textit{MM} \} dx dz dt$$

$$G_{\textit{Geometric}} = \gamma |\nabla S| + Sp + (1-S)\rho_s u$$

$$G_{\textit{Electro}} = S \left[ \rho_m \phi - \frac{\epsilon_m}{2} |\nabla \phi|^2 \right] + (1-S) \left[ -\frac{\epsilon_s}{2} |\nabla \phi|^2 - k_B T \sum c_j \left( e^{-q_j \phi / K_B T} - 1 \right) \right]$$

$$G_{\textit{Fluid}} = -(1-S) \left[ \rho_s \frac{v^2}{2} - p + \frac{\mu}{8} \int^t \left( \frac{\partial v_i}{\partial x_j} + \frac{\partial v_j}{\partial x_i} \right)^2 dt' \right]$$

$$G_{\textit{MM}} = -S \sum \left[ \rho_j \frac{\dot{z}_j^2}{2} - U(z) \right]$$

(Wei, J Math Biol 2010)

## Generalized Navier-Stokes Equation for fluid flow

$$\rho_s \left( \frac{\partial v}{\partial t} + v \cdot \nabla v \right) = -\nabla p + \frac{1}{1-S} \nabla \cdot (1-S)T + F$$

$$F = \frac{S}{1-S} \left( -\nabla p - \frac{1-S}{S} \nabla(\rho_s u) + \frac{\rho_m}{S} \nabla(S\phi) \right)$$

$$\nabla \cdot v = 0$$

## Generalized Newton equation for molecular dynamics

$$\rho_j \frac{d^2 z_j}{dt^2} = f_{SSI}^j + f_{RF}^j + f_{PI}^j$$

$$f_{SSI}^j = -\frac{1-S}{S} \nabla_j(\rho_s u)$$

$$f_{RF}^j = \frac{\rho_m}{S} \nabla_j(S\phi)$$

$$f_{PI}^j = -\nabla_j U(z)$$

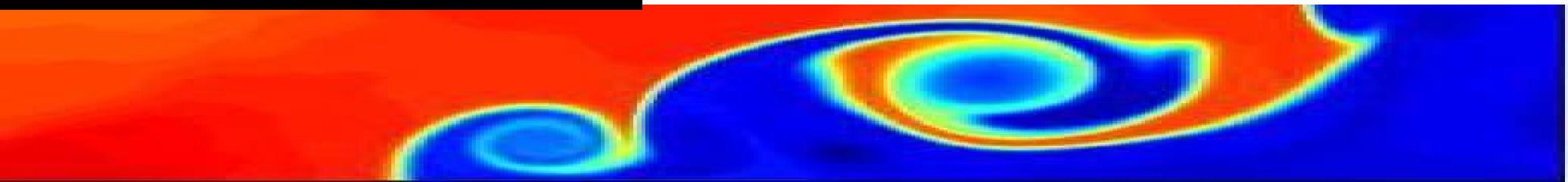
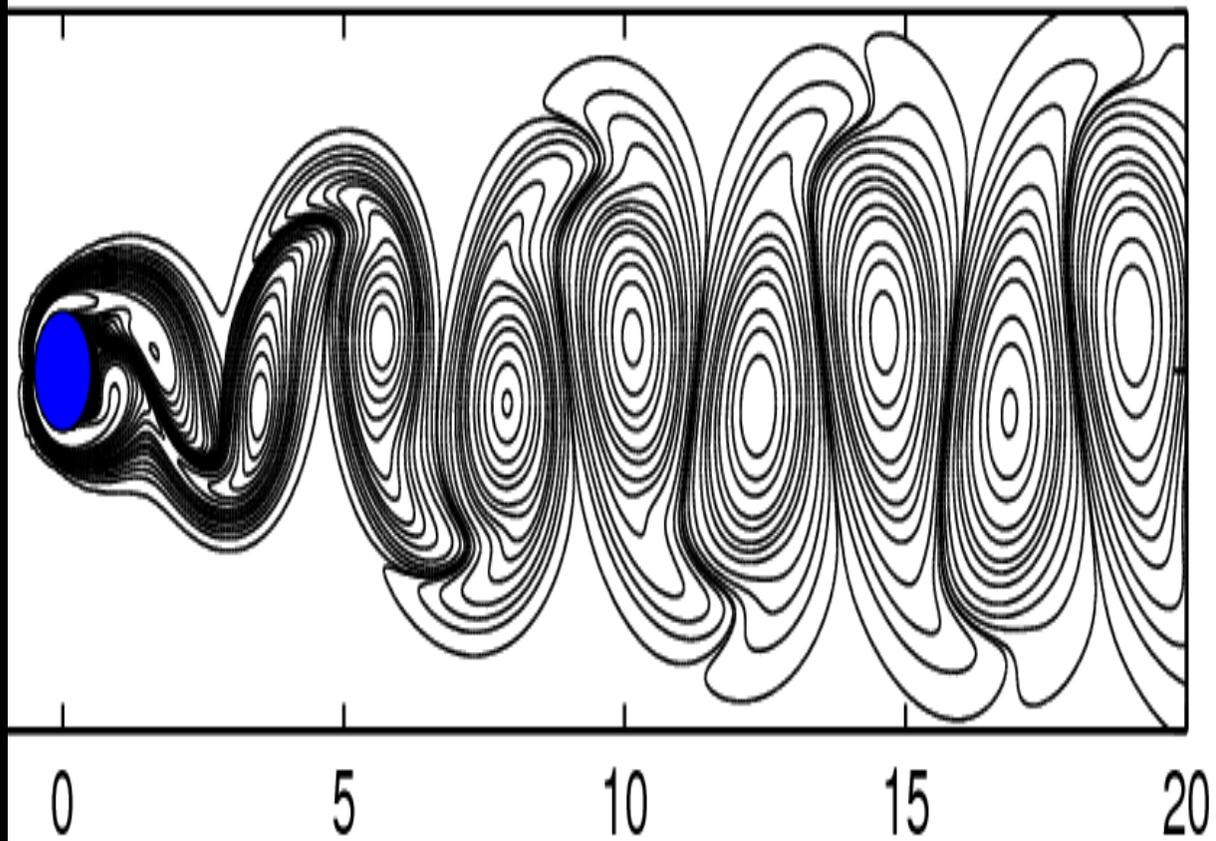
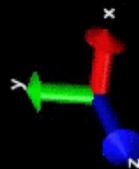
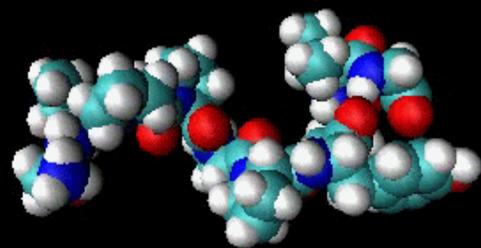
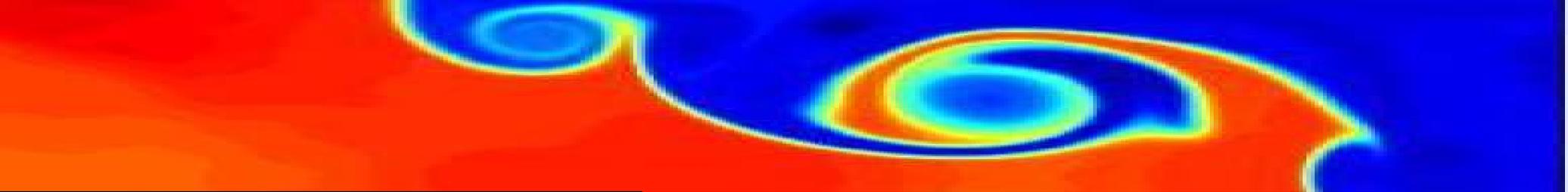
## Generalized Poisson-Boltzmann Equation for electrostatics

$$-\nabla \cdot \varepsilon(S) \nabla \phi = S \rho_m + (1-S) \sum q_j c_j e^{-q_j \phi / k_B T}$$

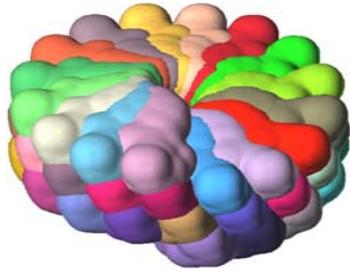
$$\varepsilon(S) = S \varepsilon_m + (1-S) \varepsilon_s$$

## Generalized Laplace-Beltrami Equation for surface dynamics

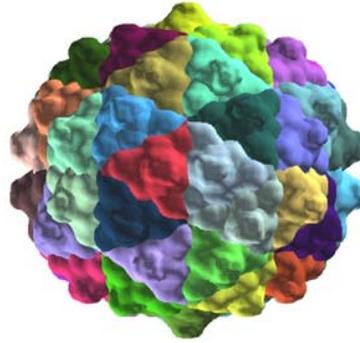
$$\frac{\partial S}{\partial t} = |\nabla S| \left\{ \begin{array}{l} \nabla \cdot \frac{\gamma \nabla S}{|\nabla S|} - p + \rho_s u - \rho_m \phi + \frac{\varepsilon_m}{2} |\nabla \phi|^2 \\ - \frac{\varepsilon_s}{2} |\nabla \phi|^2 - k_B T \sum c_j \left( e^{-q_j \phi / K_B T} - 1 \right) \\ - \left[ \rho_s \frac{v^2}{2} - p + \frac{\mu}{8} \int^t \left( \frac{\partial v_i}{\partial x_j} + \frac{\partial v_j}{\partial x_i} \right)^2 dt' \right] \\ + \sum \left[ \rho_j \frac{\dot{z}_j^2}{2} - U(z) \right] \end{array} \right\}$$



1CGM



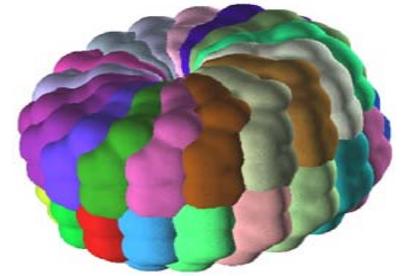
1NOV



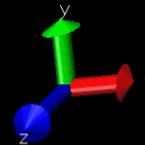
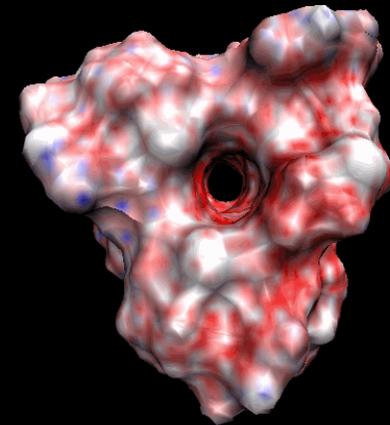
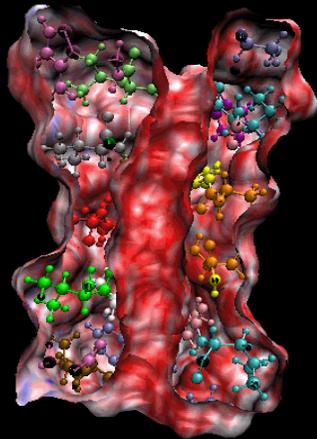
2BK1

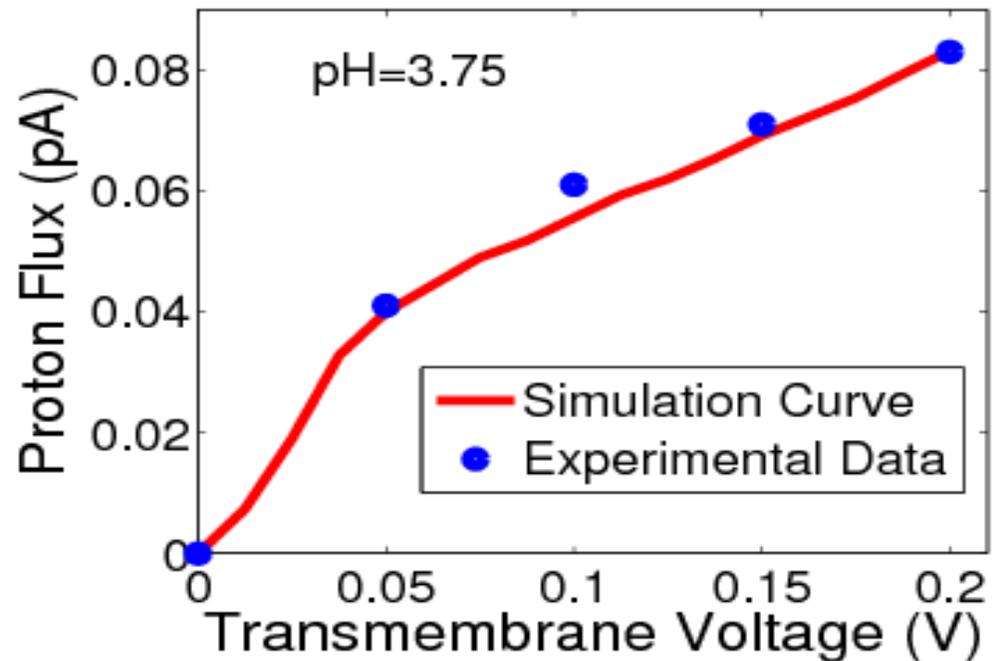
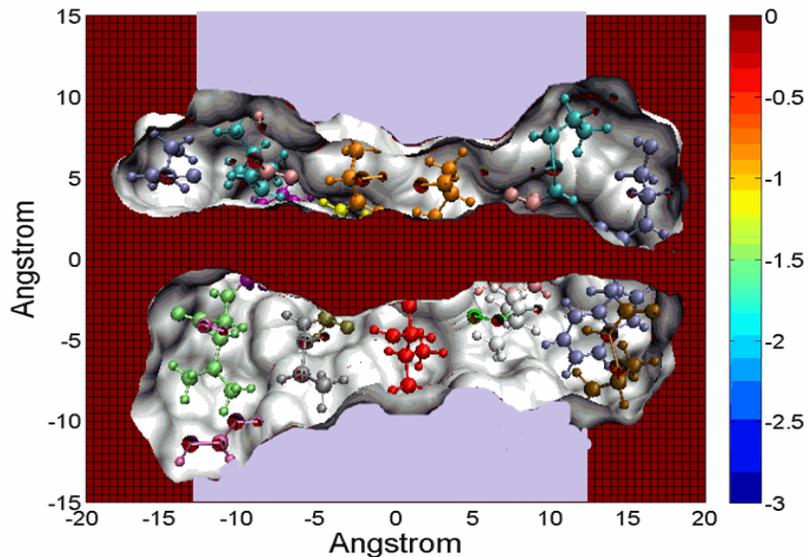


1EI7

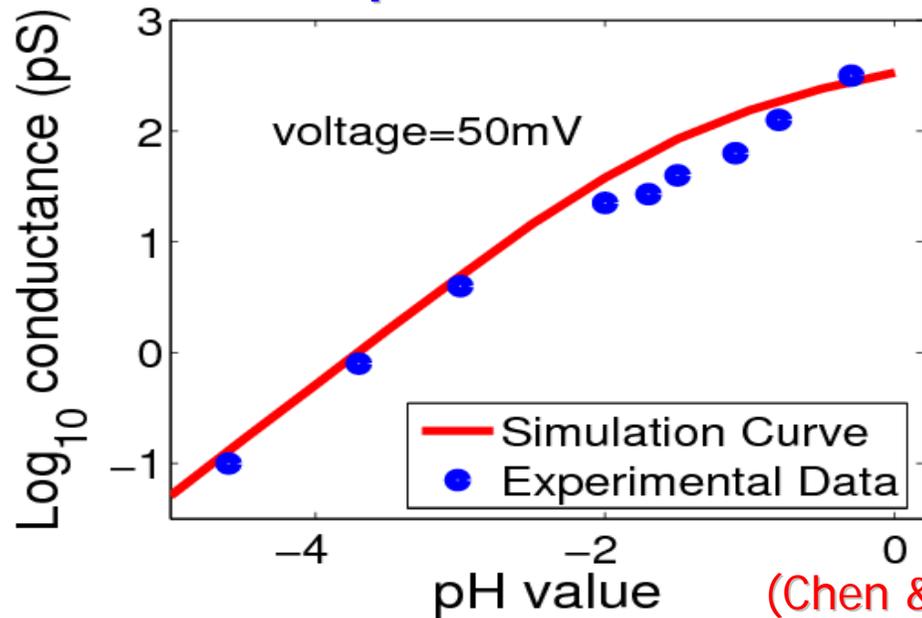


## Virus morphology and virus ion channel



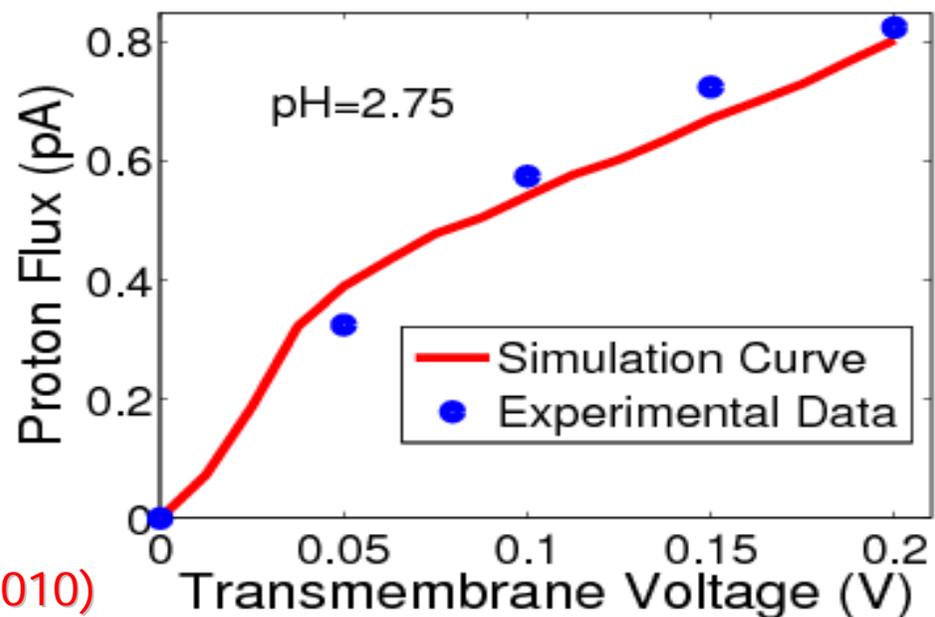


## Proton transport of Gramicidin A



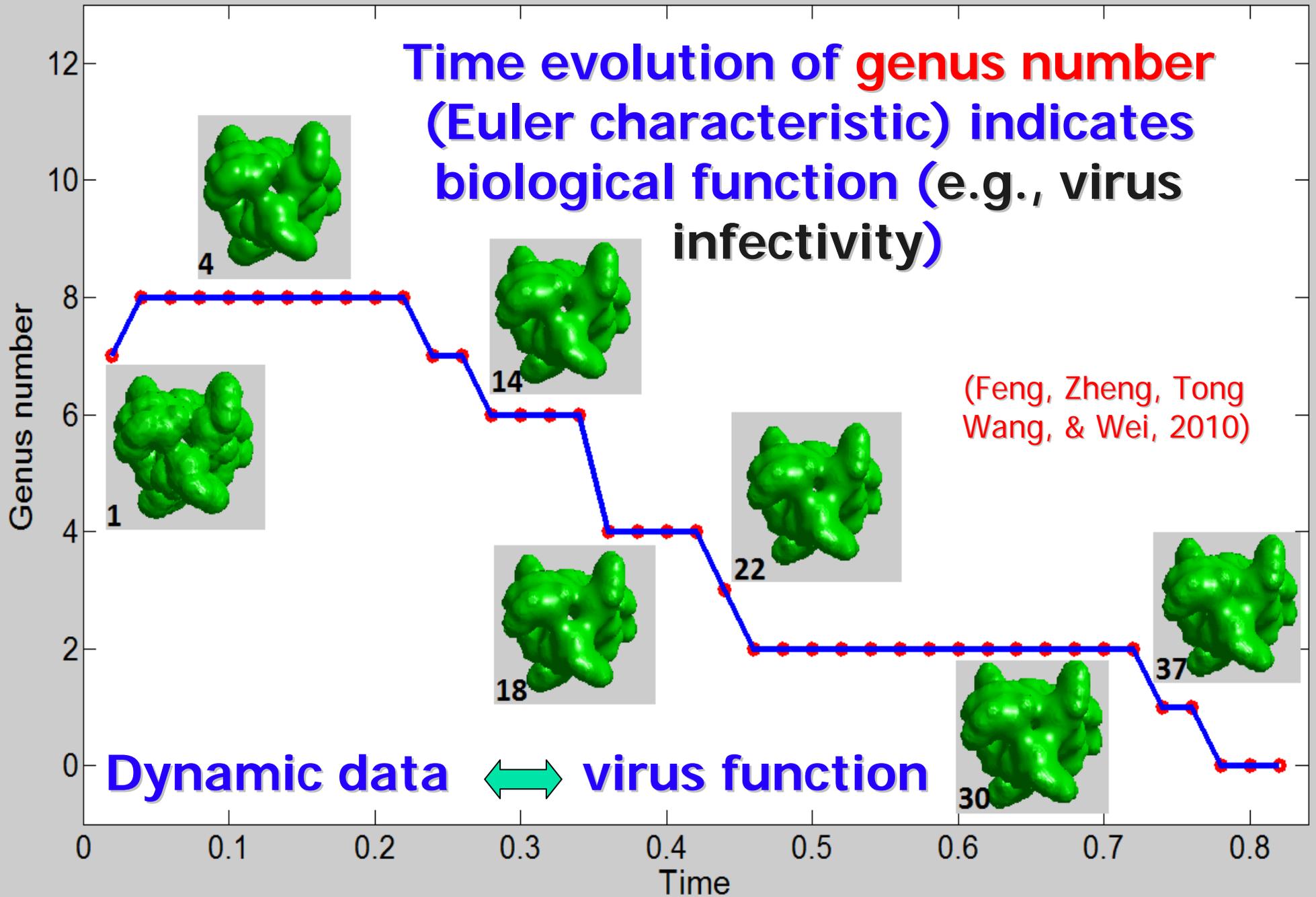
(Chen & Wei, 2010)

(Expl: Eisenman et al., 1980)



**Time evolution of genus number (Euler characteristic) indicates biological function (e.g., virus infectivity)**

(Feng, Zheng, Tong Wang, & Wei, 2010)



1. Qiong Zheng and Guo-Wei Wei, Implicit Poisson Nernst-Planck model for ion channels. (2010)
2. Yang Wang, Guo-Wei Wei and Siyang Yang, Model decomposing evolution equations, (38 pages) (2010).
3. Zhan Chen and G. W. Wei, Differential geometry based solvation model III: Quantum formulation (35 pages). (2010).
4. Kelin Xia, Meng Zhan and Guo-Wei Wei, The matched interface and boundary (MIB) method for multi-domain elliptic interface problems. (2010)
5. Yang Wang, Guo-Wei Wei and Siyang Yang, Empirical Model Decomposition using Local Spectral Evolution Kernels, (38 pages) (2010).
6. Qiong Zheng, Duan Chen and Guo-Wei Wei, Second-order convergent Poisson Nernst-Planck solver for ion channels. (2010)
7. Duan Chen and Guo-Wei Wei, Quantum dynamics in continuum model for proton channel transport. (2010)
8. Sigal Gottlieb, Guo-Wei Wei, and Shan Zhao, A unified discontinuous Galerkin framework for time integration (46 pages), (2010)
9. Zhan Chen, Nathan Baker and G.W. Wei, Differential geometry based solvation model II: Lagrangian formulation (53 pages) submitted, (2010)
10. Weihua Geng, and G. W. Wei, Multiscale molecular dynamics via the matched interface and boundary (MIB) method, *Journal of Computational Physics*, 230, 435-457 (2010).
11. Duan Chen, Zhan Chen, Changjun Chen, Weihua Geng and Guo-Wei Wei, MIBPB: A software package for electrostatic analysis, *Journal of Computational Chemistry*, published online: 15 SEP (2010).
12. Zhan Chen, Nathan Baker and G. W. Wei, Differential geometry based solvation model I: Eulerian formulation, *Journal of Computational Physics*, 229, 8231-8258 (2010).
13. Duan Chen and Guo-Wei Wei, Modeling and simulation of nano-electronic devices, *Journal of Computational Physics*, 229, 4431-4460, (2010).
14. Changjun Chen, Rishu Saxena, and Guo-Wei Wei, Differential geometry based multiscale model for virus capsid dynamics, *Int. J. Biomed Imaging*, Volume 2010, Article ID 308627, 9 pages (2010)
15. Guo-Wei Wei, Differential geometry based multiscale models, *Bulletin of Mathematical Biology*, volume 72, 1562-1622, (2010).

# Summary

- *Differential geometry based multiscale models for viruses*
- *Topological invariants for virus function characterization*
- *Stable manifolds for identifying coarse-grain clusters*
- *Machine learning methods for dimension reduction*
- *Theoretical prediction agrees with experimental data*