

The Analysis and Application of the Emergent Electronic Properties of Self-Assembling Nucleopeptide Systems

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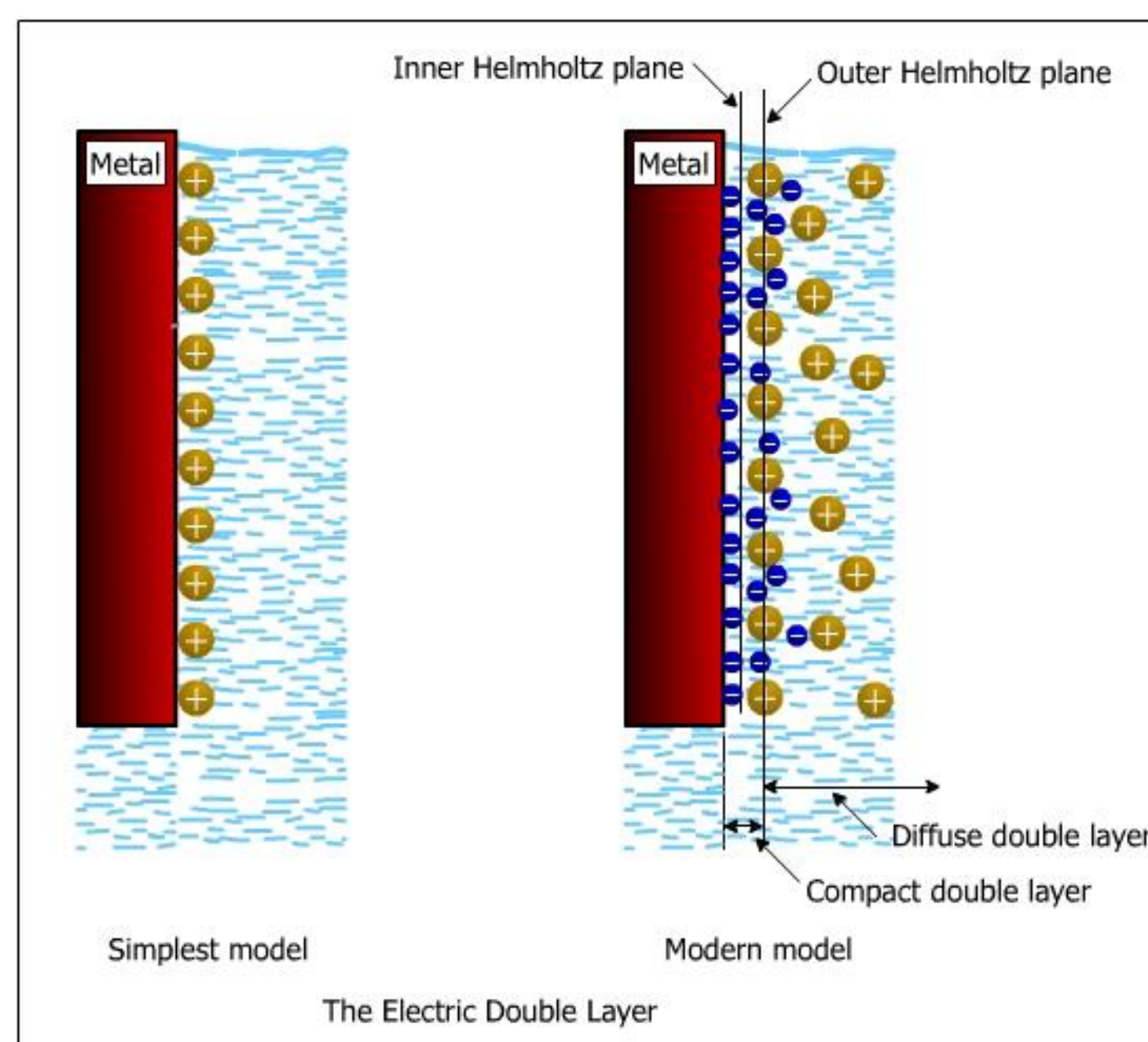
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Abstract

Biomolecular structures are held together by a complex network of molecular interactions that direct assembly and stabilize structures. In order to translate the fundamental molecular interactions of biomolecules into the design of functional biomaterials, we have developed a model system that integrates nucleic acids and self-assembling peptides (shown below). These nucleopeptides serve as a small-model system for the study of the non-covalent molecular interactions involved in biomolecule self-assembly measured through impedance. The peptides are in this case our system of electronic circuit. The emergent electronic properties found in most of our 21 samples, which were measured by Electrical Impedance Spectroscopy (EIS) and analyzed by ZMAN software shows promise. This is because one of the most frequent applications to show up was Battery and Supercapacitor which can be used in Biosensors.

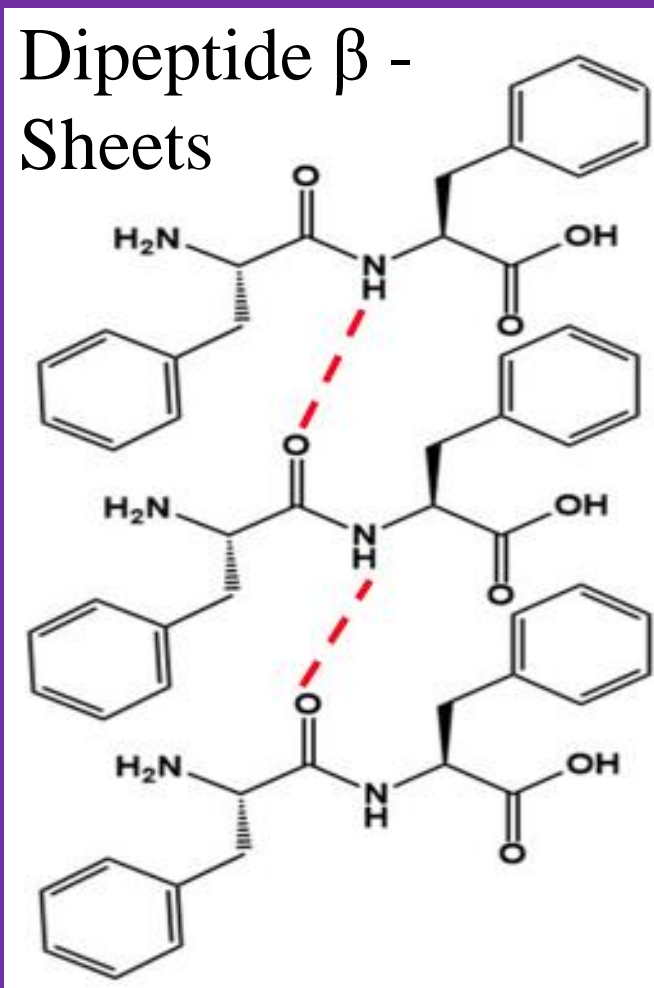
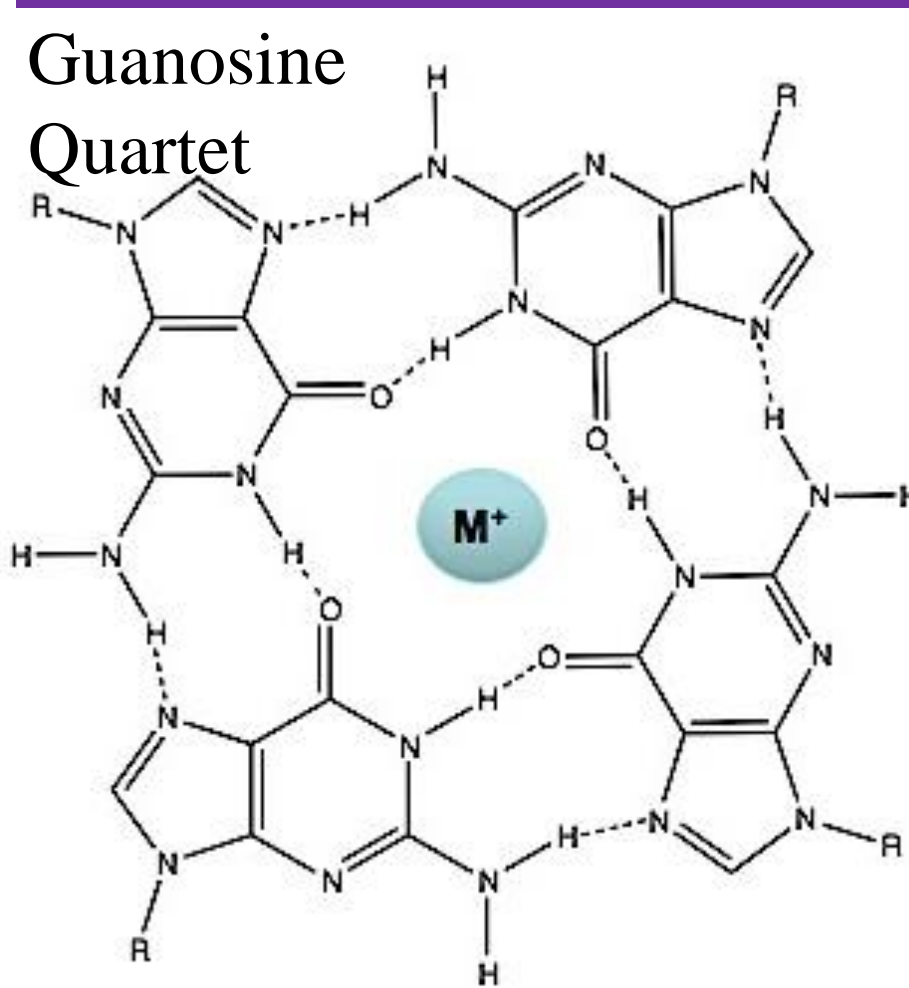
Electrochemical Impedance Spectroscopy (EIS)



EIS is used to measure the impedance or total resistance in an AC electrical circuit. For an electric circuit to function it has 3 basic components: a resistor, a conductor, and an inductor. Each component has its own impedance, represented by the symbol Z . Through the LCR Reactance Phasor Diagram, one can eventually find Z_{total} . The first step in analyzing the peptides is by applying a small voltage perturbation and measuring the current response.

This is done by taking the bulk solution or peptides and introducing an electrode or cable. In the diagram above, the cable is shown as a metal that attracts the anions and forms an Electric Double Layer. This creates a region in the vicinity of the charged surface up to a distance called the Debye length, with the inner and outer Helmholtz regions forming the first few layers of ions near the charged surface. Once the data from the sample is collected, it is then plotted in two ways: Bode plots and Nyquist plots.

Supramolecular Secondary Structures



The Self-Assembling Nucleopeptides System

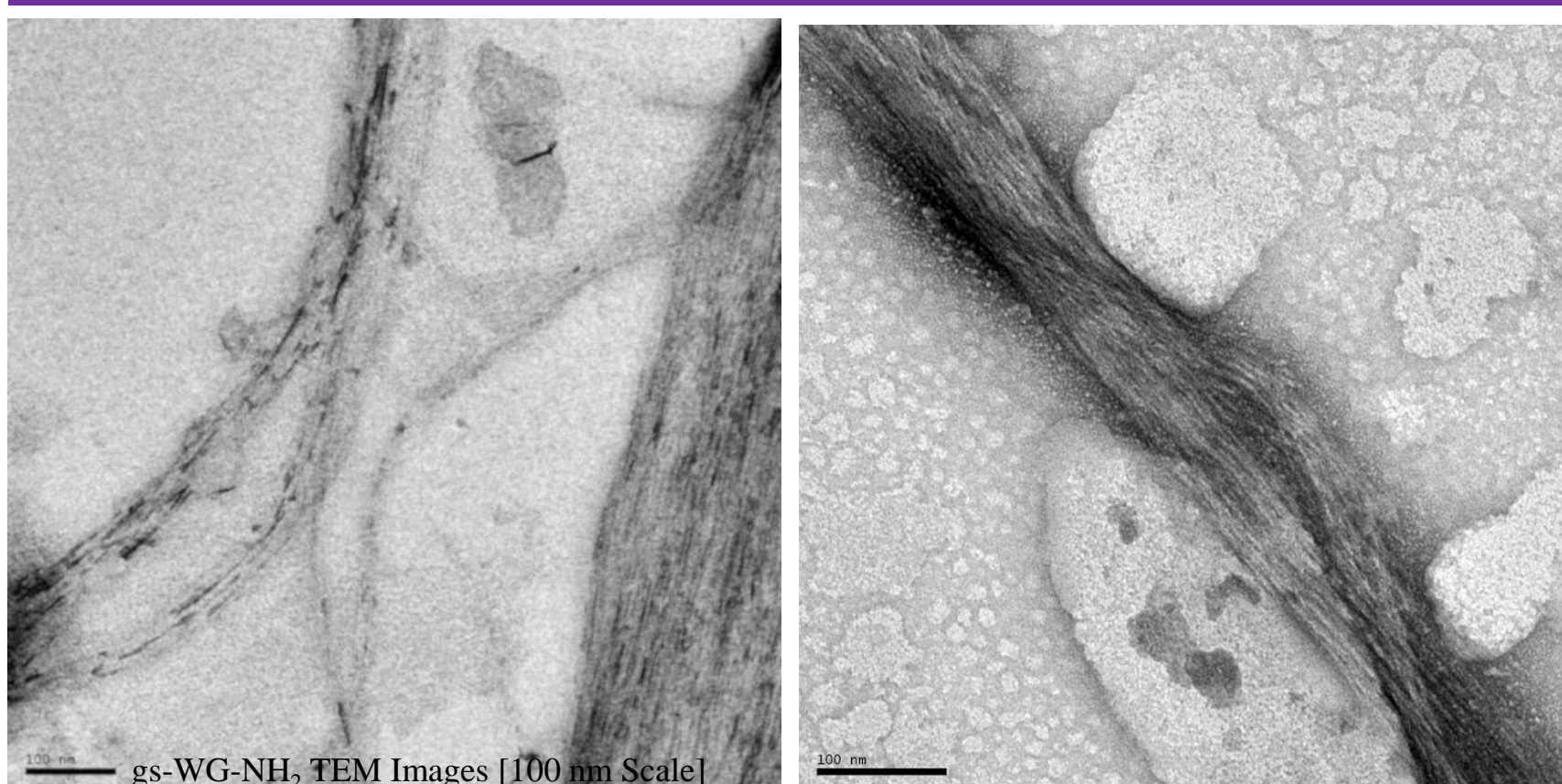


Assembly Conditions:

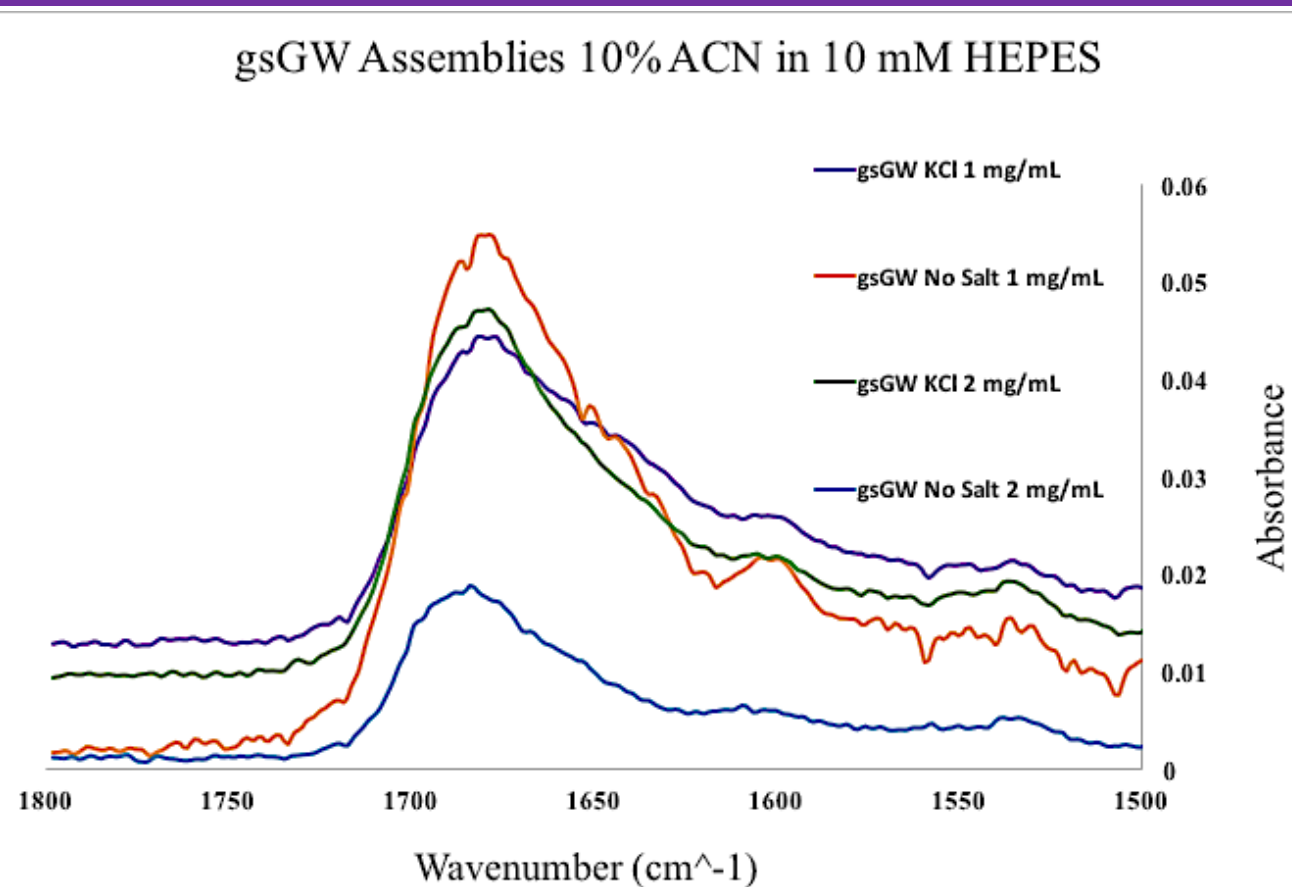
1 mg/mL and 2 mg/mL gsGW concentrations assembled in 10% ACN in 10 mM HEPES

Salt and No Salt Conditions: +/- 10 mM KCl

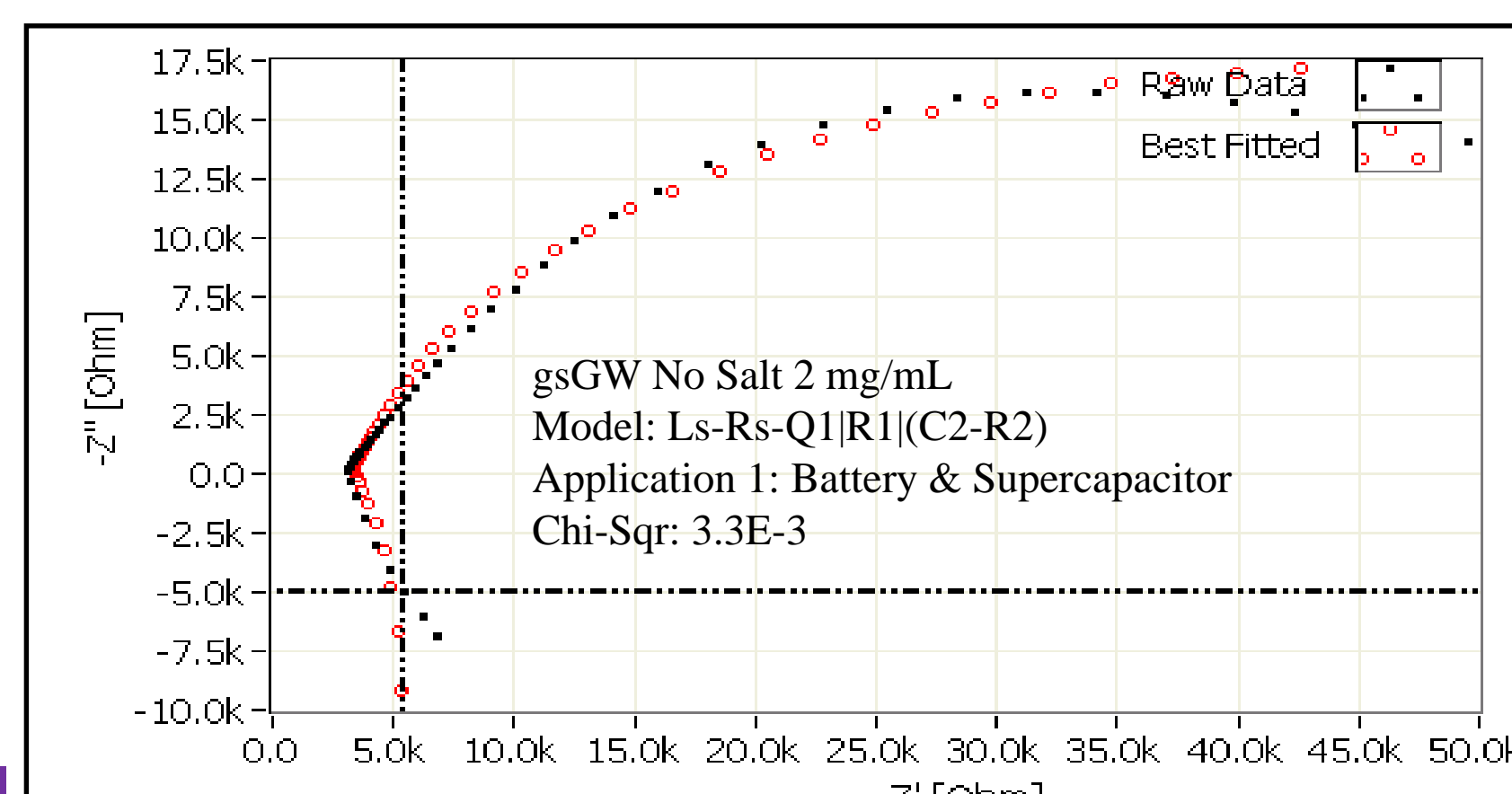
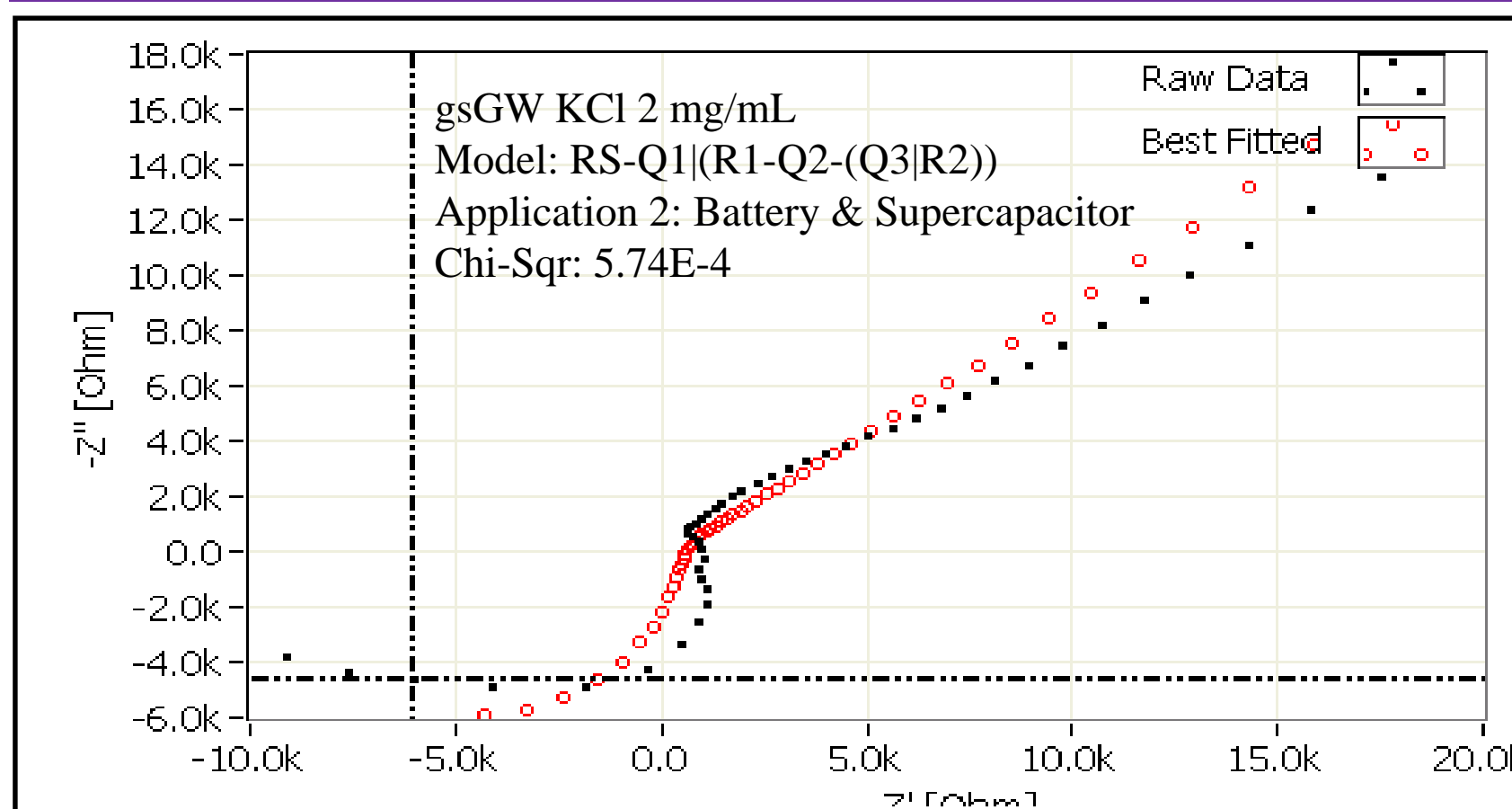
Transmission Electron Microscopy



Infrared Spectroscopy



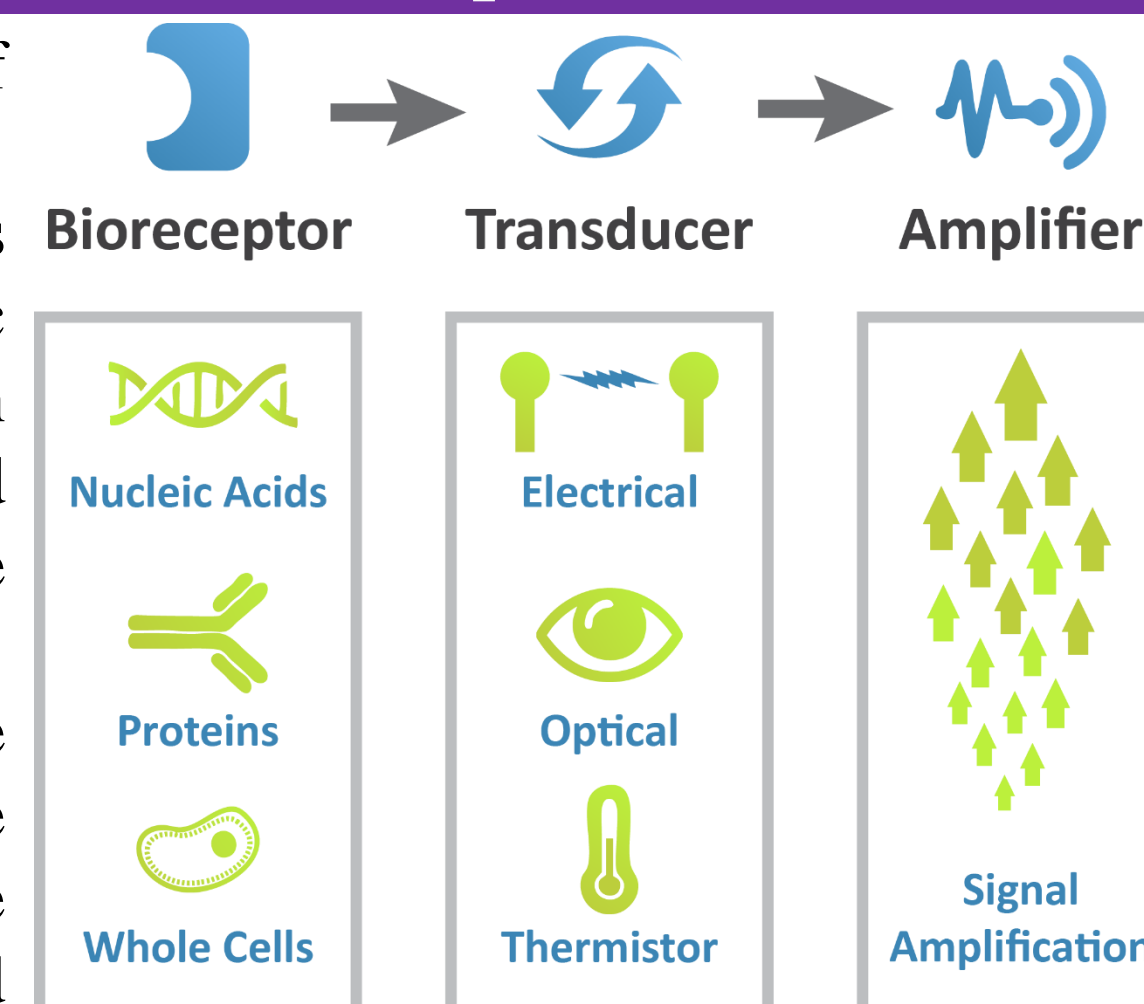
Nyquist Plots



The Nyquist plot is read right to left, and the variables are Z_{real} and $Z_{imaginary}$. From the 21 samples tested, a representative Nyquist plot is shown on the left, including the sample name, model, application, and chi-squared value. The first 3 applications shown by the software were chosen and analyzed. The model shows the different components of the system, where the lower the Chi-Sqr, the more reliable the equation.

Preliminary Successes and Future Experiments

- Successful synthesis and purification of guanosine modified nucleopeptides.
- Characterization of nucleopeptide supramolecular structures by Atomic Force Microscopy, Transmission Electron Microscopy, Infrared Spectroscopy, and Electrochemical Impedance Spectroscopy.
- In the future, further analysis on more samples with different variables will be done. The samples above show promise to be used as Batteries and Supercapacitors in biosensors. More trials will be needed to perfect the solutions.



References

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