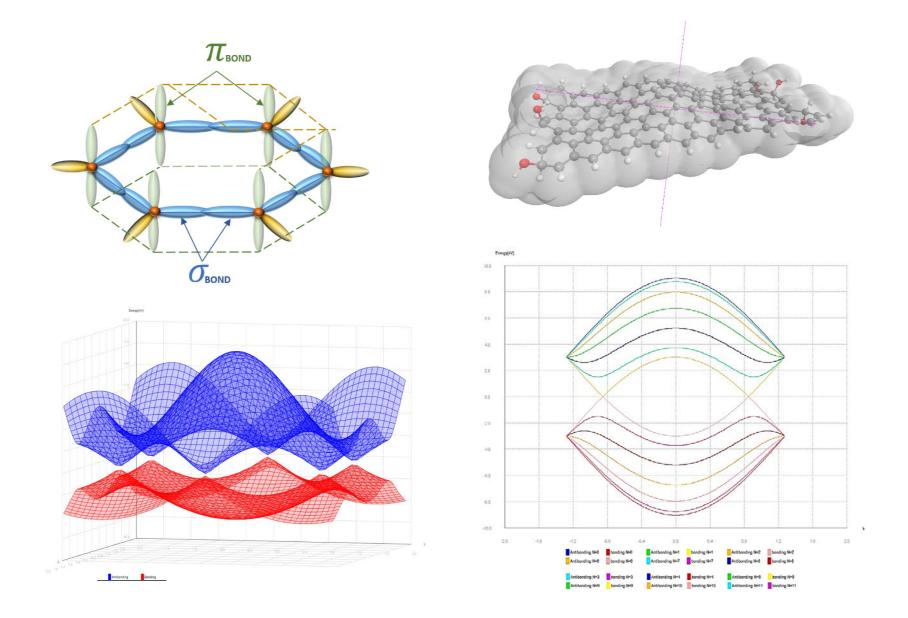
COMSOL CONFERENCE 2016 MUNICH

Functionalized Graphene Bio-Sensing **Building-Blocks under Environmental Stimuli**

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Introduction: This study continues to investigate the main properties of graphene (G/RGO/GO), and other C allotrope nanostructures for biosensors design, using COMSOL Multiphysics® modules.



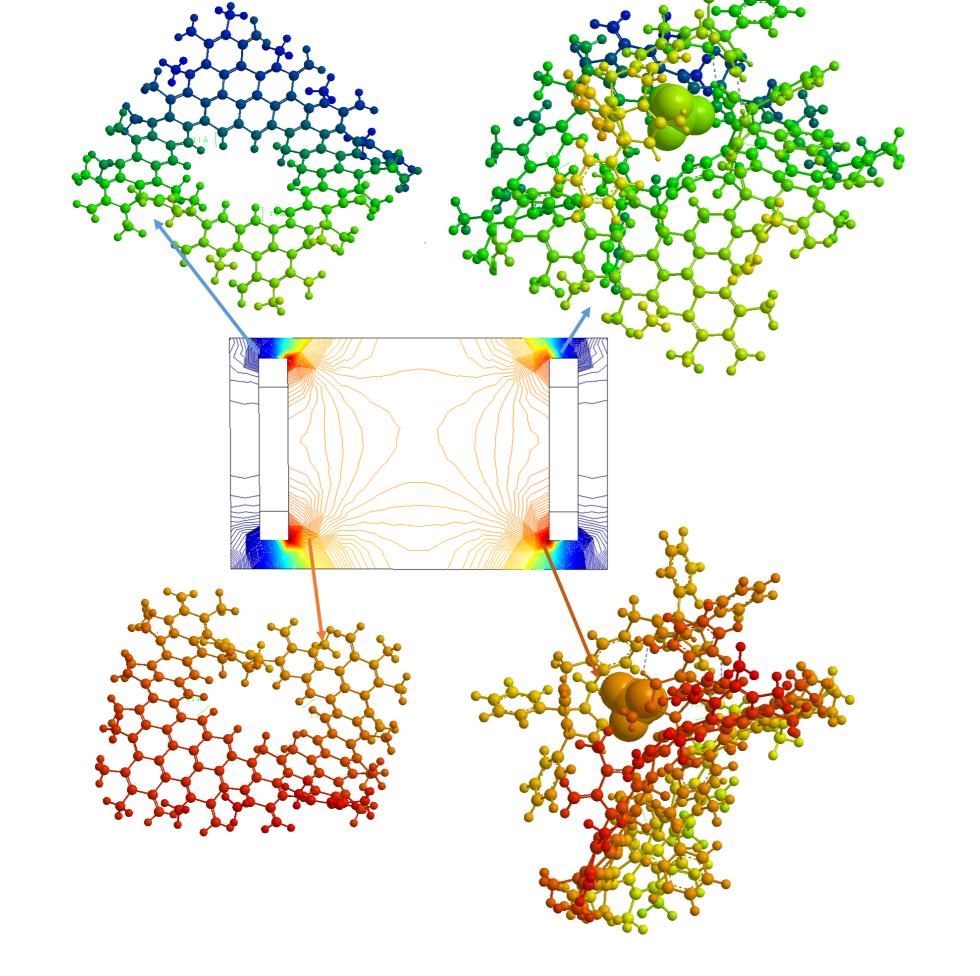


Figure 1. Functionalized G/GO/RGO bio-sensing modules

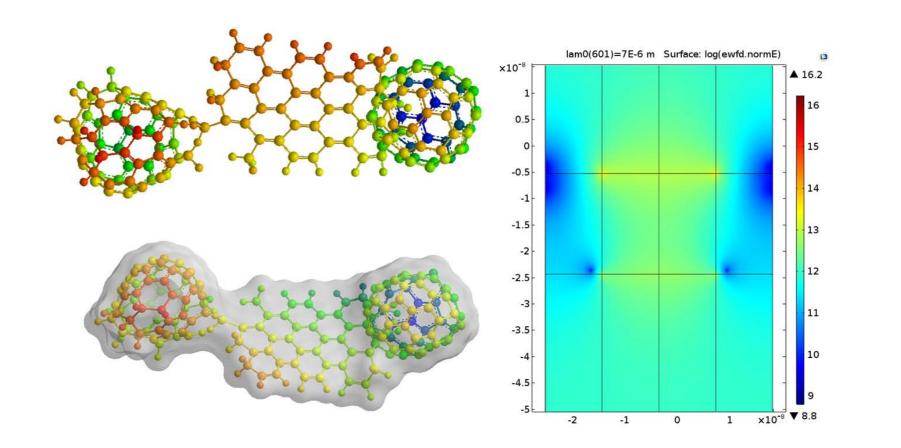
Methods: Computational Electrical Potential Profile is a function of distance (x) from graphene surface. The shape of potential function is determined by Modified Poisson-Boltzmann equation (Eq.1):

Figure 2. Carbon allotropes studies

(a) G unit cell; (b)G -T.Char.Density;

(b) MATLAB computed properties (c) G (10,10); (d) CNT (6,6,10)

Results: With COMSOL Multiphysics® modules, importing geometry and properties from ChemBio3D on Electrochemistry module, and the electrical and the thermal properties from MATLAB through LiveLinkTM on Multiphysics were performed different functionalization simulations (Fig. 3-5)



$$\frac{d^2 V}{dx^2} = \sum_i \frac{z_i e c_{0i}}{\varepsilon} \cdot \frac{2 \sinh\left(\frac{z_i e V}{kT}\right)}{1 + 2v_i \sinh^2\left(\frac{z_i e V}{2kT}\right)}$$

In MATLAB code (P- potential, used instead V from Eq.1) all simulation parameters are (Eq.2): $[x, P, R] = potential_{1D}(P_0, Z_i, C_i, E_R, EFF, MPB)$ One of the simulation output parameters is ion concentration as a function of the distance from graphene surface (Eq.3):

Figure 3. RGO- 2xC60 self-assembly : (a) model; (b) Total Charge Density; (c) Electrical Potential Profile

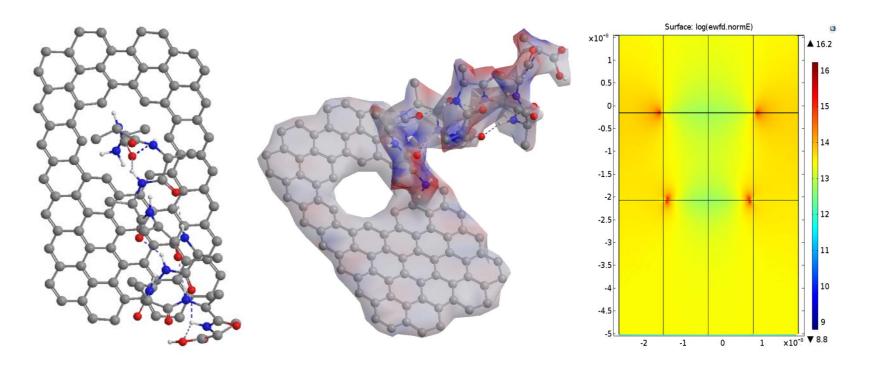
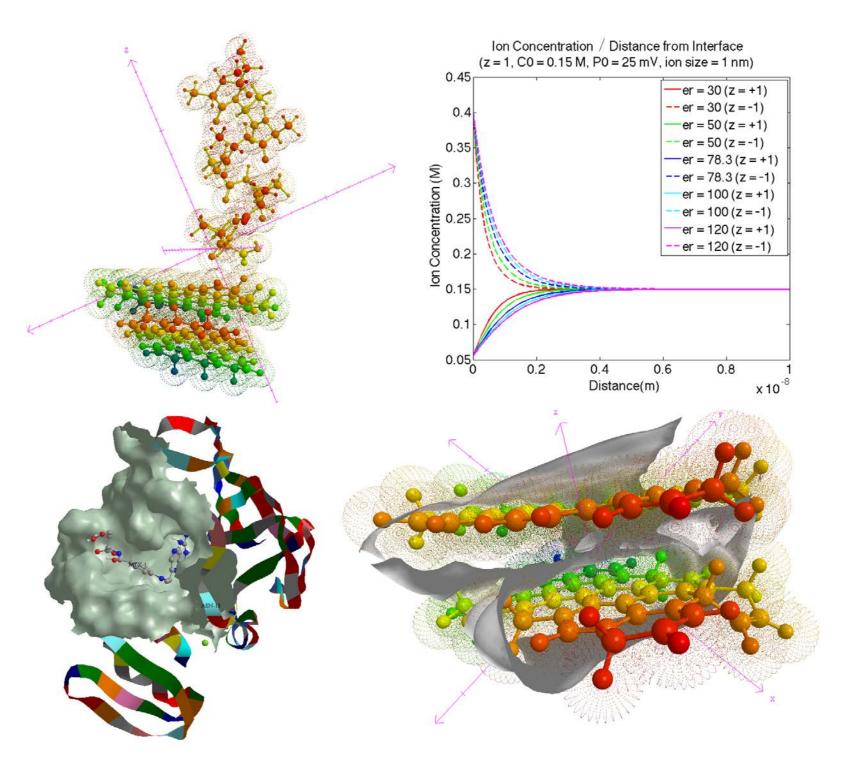


Figure 4. GO- Alpha-Helix (protein) self-assembly : (a) model; (b) Total Charge Density; (c) Electrical Potential Profile



 $C_i = C_{i0} e^{\frac{-z_i eV(x)}{kT}}$

MATLAB application of graphene properties (Fig.2a,b,c) and CNTs (Fig.2c) in specific models and simulations were exported through LiveLink in COMSOL Multiphysics® (Fig.3c,4c,5b).

Figure 5. 2x RGO(1-pore)- Alpha-Helix- self-assembly : (a) Self-Assembly model; (b) Ion concentration; (b) (c) Local Charge Density – Protein-RGO pore ; (c) (d) 2xRGO – Total Charge Density

Excerpt from the Proceedings of the 2016 COMSOL Conference in Munich

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