

Models for Simulation Based Selection of 3D Multilayered Graphene Biosensors

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Overview

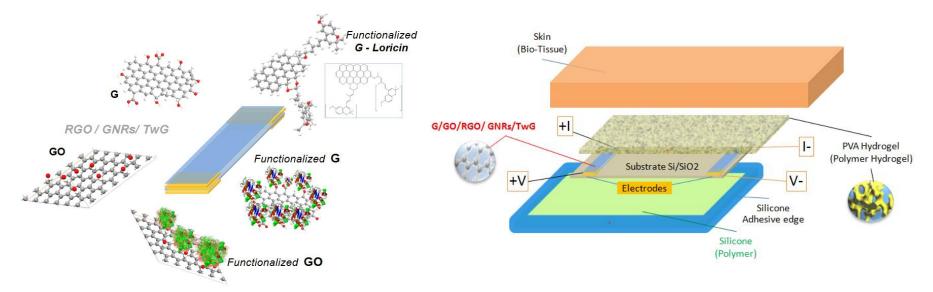
- Graphene biosensor models
- Density Functional Theory
- Graphene biosensor data
- Graphene models design & characterization (ChemBioOffice)
- Protein models design and characterization (ChemBioOffice)
- G/RGO/TwG Force Field characterization (MATLAB)
- Use of COMSOL Multiphysics
- COMSOL analysis: G-Biosensor structure model
- COMSOL analysis: Tween (TwG)- Biosensor structure model
- COMSOL Simulations (Acoustic Module): (TwG-E)/SiO₂; SiO₂ -environment
- COMSOL Simulations (Acoustic Module):G/GO SiO₂ interface
- Conclusions

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Graphene biosensor models



Graphene (G), graphene oxides (GO), reactive graphene oxides (RGO), graphene nanoribbons (GNRs) and other graphene based composite materials (TwG) are used on the biosensing area with the aim of defining new functionalized material solutions for personalized medical applications.



Density Functional Theory

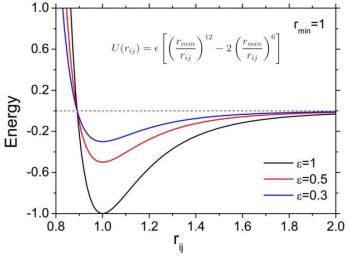
Lennard- Jones potential:

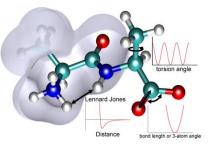
$$U_{(r_{ij})} = \epsilon \left[\left(\frac{r_{min}}{r_{ij}} \right)^{12} - 2 \left(\frac{r_{min}}{r_{ij}} \right)^{6} \right]$$

The force acting on the "*i*" atom of the system:

$$\vec{F}_i = -\frac{12 \epsilon}{r_{min}^2} \sum_{\substack{j=1\\j\neq i}}^{N} \left[\left(\frac{r_{min}}{r_{ij}} \right)^{14} - \left(\frac{r_{min}}{r_{ij}} \right)^8 \right] \vec{r}_{ij}$$

 ϵ — the depth of the potential energy well [eV] r_{min} —the distance at the minimum of the potential [A]





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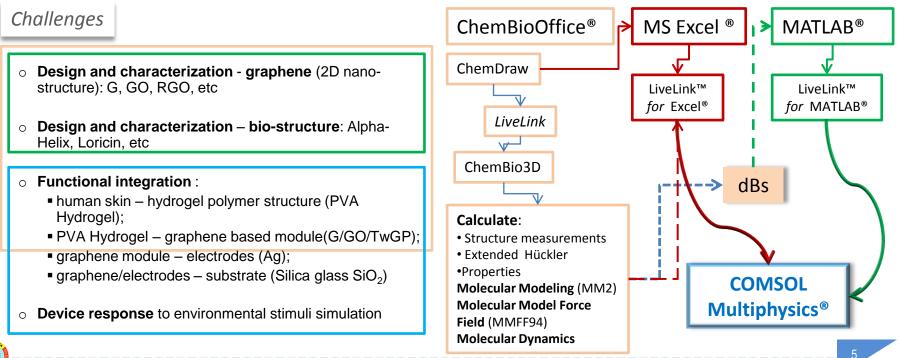
Parameters of van der Waals (vdW) interactions are specified through force field files [Lennard-Jones potential dBs]



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Graphene biosensor data

Modeling approach



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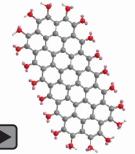
CONFERENCE 2015 GRENOBLE **ChemBioOffice® MATLAB®** COMSOL Graphene models design & Graphene Graphene G-OH 298.183 kcal/mol Total Energy of frame 524.036 kcal/mol 31.222 **RMS** gradient 16.969 characterization G-H₂ Molecular Dynamics Dynamics: 2.0 fs Step interval: 2.0 fs 10 fs 10 fs Frame interval: 306 steps 306 steps Terminate after: 1.000 kcal/atom/ps 1.000 kcal/atom/ps Heating/Cooling Rate: 300 Kelvin 300 Kelvin Target temperature: Properties: Exp →dB-Pi -BO_[G-H2] Pi bond order Exp →dB-Pi -BO_[G-OH] Exp \rightarrow dB-SEs [G-H2] Exp \rightarrow dB-SEs [G-OH] Steric energy summary Extended Hückel Calculate charges Calculate surfaces 209 Iteration 306 2.9399 Stretch 11.0341 1.7717 Bend 23.7161 Stretch-Bend 0.0463 0.5694 -82.6589 Torsion 28.8301 -8,4197 Non-1.4-vdW -16.8827 97.5110 59.1217 1.4 vdW 0.0504 Dipole/Dipole 84.8992 Total Energy -27.1485 kcal/mol 229.852 kcal/mol Non 1,4 vdW - energy for the through-space interaction between pairs of atoms that are separated by more than three atoms

1,4 vdW - energy for the through-space interaction atoms separated by two atoms

MM2 - Molecular Mechanics

MMFF94 - Merck Molecular Force Field



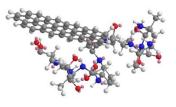


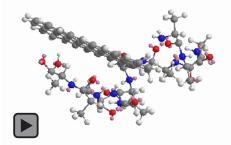
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Functionalized Graphene-Protein assembly-design & characterization



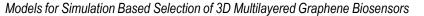


Protein models design &	 ChemBioOffice[®] 	MATLAB®	COMS
characterization	-4.488 kcal/mol	Total Energy of frame	524.036 kcal/mol
	6.4587	RMS gradient	16.969
		Molecular Dynamics	
1. 1		Dynamics:	
	2.0 fs	Step interval:	2.0 fs
X X X	10 fs	Frame interval:	10 fs
	1865 steps	Terminate after:	1494 steps
	1.000 kcal/atom/ps	Heating/Cooling Rate:	1.000 kcal/atom/ps
	300 Kelvin	Target temperature:	300 Kelvin
and a lot		· · · · · · · · · · · · · · · · · · ·	
- And and a second s		Properties:	
200	Exp →dB-Pi -BO_[αHelix] Exp →dB-SEs [αHelix]	Pi bond order	Exp \rightarrow dB-Pi -BO_[α Helix-G] Exp \rightarrow dB-SEs [α Helix-G]
T	Exp Hab-SES_[aHelix]	Steric energy summary	
		Extended Hückel	
		Calculate charges	
		Calculate surfaces	
~	1865	Iteration	1494
at the second	1.9981	Stretch	4.7891
	13.0319	Bend	13.2791
	0.9841	Stretch-Bend	0.8170
	6.0601	Torsion	-70.7286
T DATE A	-38.7740	Non-1,4-vdW	-68.6773
	25.8378	1,4 vdW	85.5095
	-13.6258	Dipole/Dipole	-7.3971
🖌 🖌 👘	-4.4876 kcal/mol	Total Energy	-42.4084 kcal/mol
	are separated	 energy for the through-space interaction between by more than three atoms argy for the through-space interaction atoms separal lar Mechanics 	

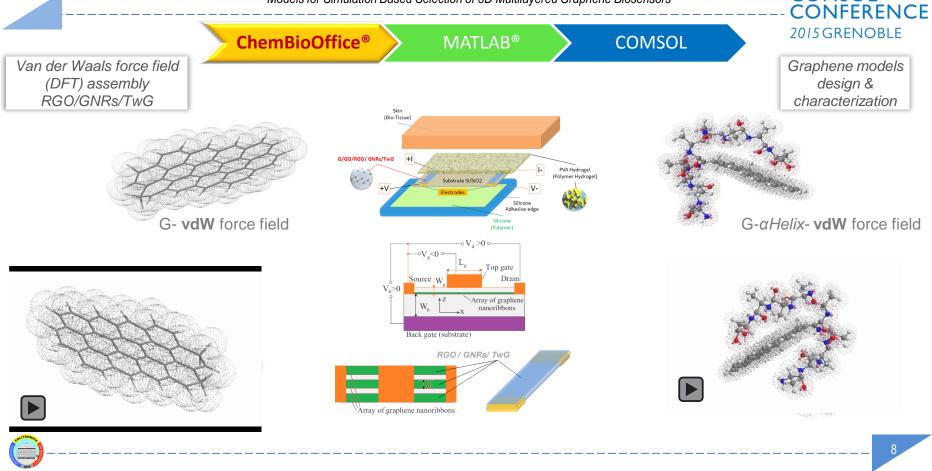
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MMFF94 - Merck Molecular Force Field

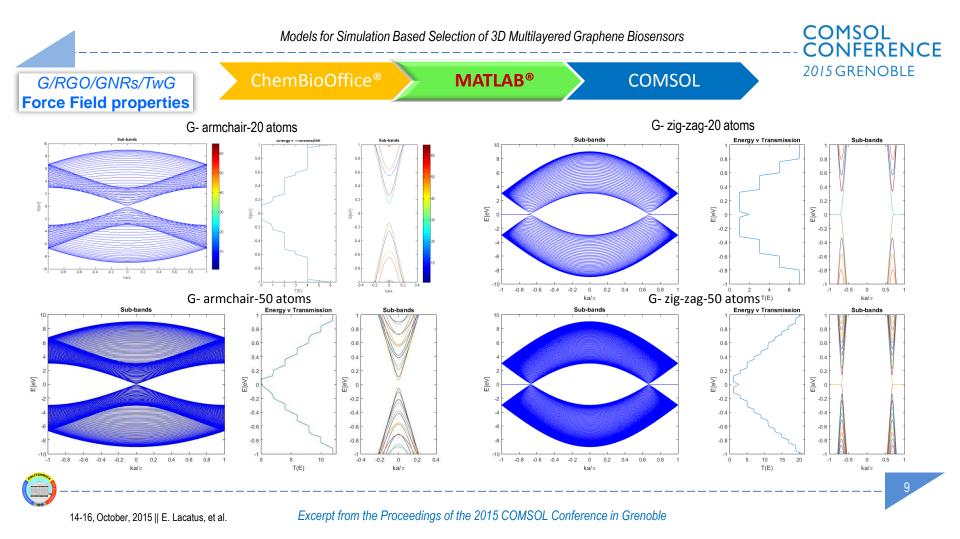
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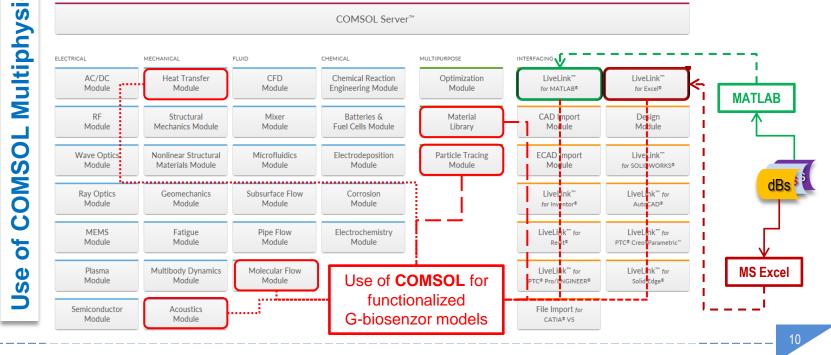
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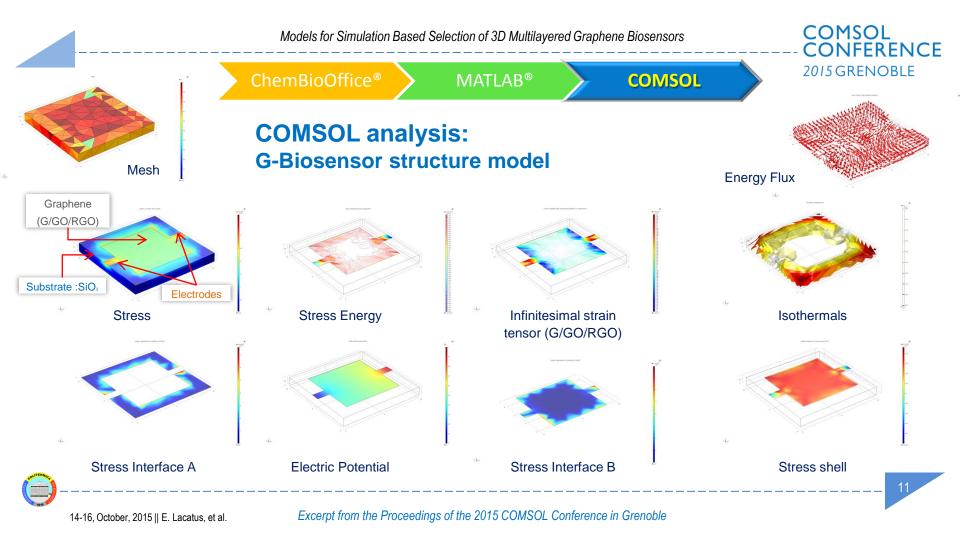


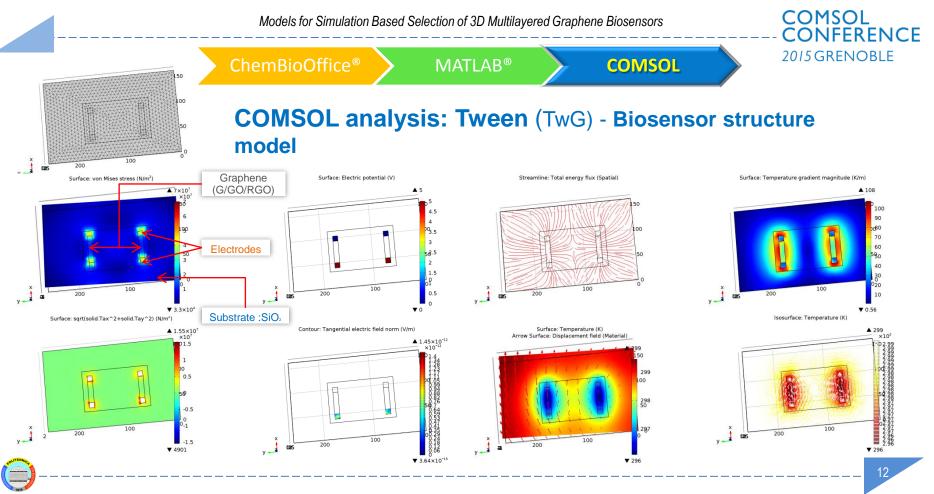
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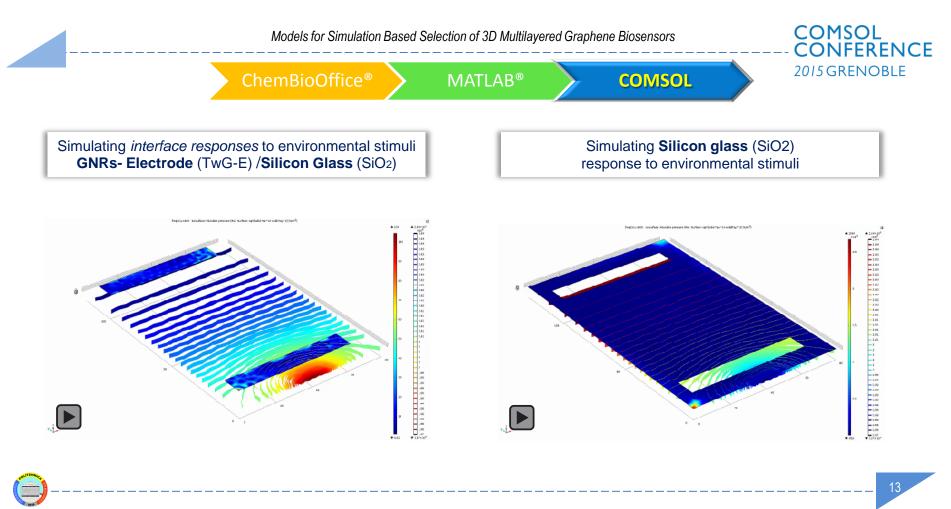
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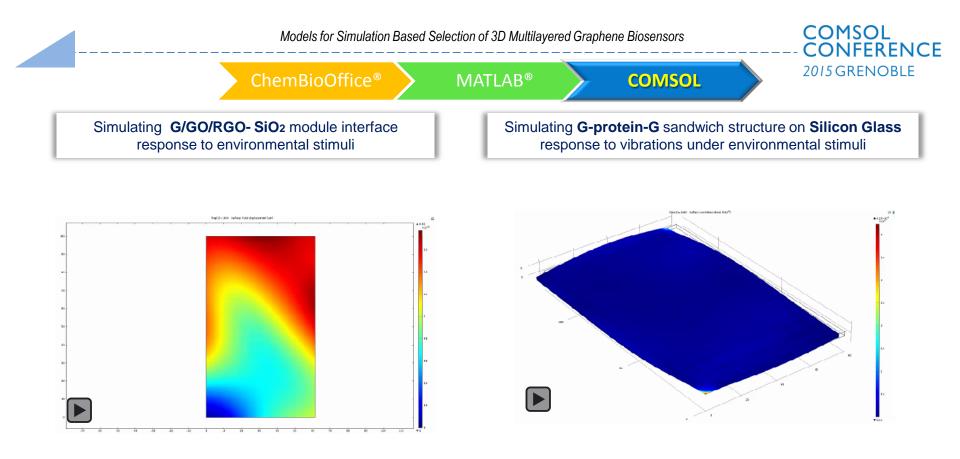




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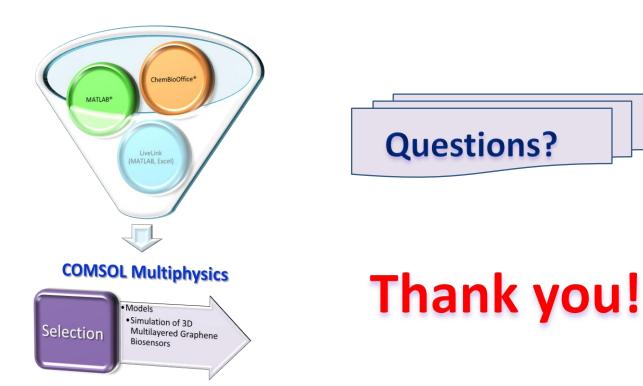
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Conclusions

- Graphene (G/GO/RGO/GNRs/TwG) models, characterizations and field properties imported through LiveLink[™] (for Excel,and for MATLAB) can be used for simulating graphene-biosensors' responses to environmental stimuli
- Van der Waals field forces potential *(based on: DFT, Lennard-Jones, Kirchhoff, and Frölich continuum)* applied for molecular and quantum level interactions *(electron-electron, phonon-phonon, electron-phonon)* can explain and simulate the G/GO/RGO-biosensing device continuum
- The functionalized G-protein-G structures, simulated as self-assembled structures, evolved during iterations towards stabile configurations (*DFT, vdW*)
- Biological and environmental stimuli *(thermal, electric, acoustic, chemical, etc)* are harvested by the functionalized graphene structures in continuum like models through the use of COMSOL Multiphysics modules





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