# Graphene oxide lateral dimensions can mediate different molecular response of human immune cells.

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## Abstract

In the last few years, there has been enormous interest in graphene oxide (GO) for its wide variety of applications [1, 2]. However, for any medical application, the immune system-impact of GO still remain to be fully understood [3]. Here we focused on the molecular effects of two GOs, different for lateral size dimensions, on human primary immune cell populations: peripheral blood mononuclear cells (PBMCs). GOs were fully characterized, then we performed a wide range of standard assays looking at cell viability, cell activation and multiple cytokines secretion. We characterized the molecular impact of GOs on 84 genes immune-response-related. Additionally, a whole genome analysis was conducted on T cells and monocytes as representative of the innate and adaptive immune responses. In Fig. 1 TEM and AFM characterization of GO-Small (140 nm) and GO-Large (4µm). We did not detect any toxicity in GO PBMCs treated samples. The 84 gene expression analysis evidenced a clear dimension-dependent impact of GOs on cell activation (Fig. 2). In particular, GO-Small modulated 16 genes (Fold Regulation >4) compared to only 5 of GO-Large (in red in Fig.2 C). Action confirmed also by cytokine analysis (Fig. 2 D). Further evidences were given by microarray analysis on T and monocytes cell lines. GO-Small impact the immune cell activation, underlined by the over expression of many pathways such as leukocyte chemotaxis pathway (Fig.3), genes such as CXCL10 and its receptor CXCR3 (Fig 3, red box). These genes are commonly activated during acute inflammatory processes as those associated with immune-mediated tumor rejection and pathogen clearance [4].

Moreover, we found a strong action on cell metabolism with a down-regulation on energetic pathways such as oxidative-phosphorylation pathway in both cell types (data not shown). Our work represents a comprehensive molecular-characterization of different sized GOs on immune cells giving crucial information for the chemical and physical design of graphene for biomedical applications i.e. as a new possible drug delivery systems and nanoimmunotherapy tools.

## References

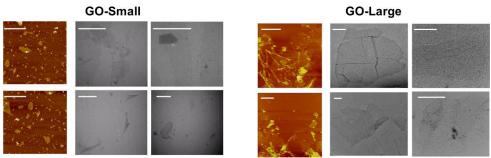
[1] Sechi G, Bedognetti D, Sgarrella F, Van Eperen L, Marincola FM, Bianco A, Delogu LG. Nanomed. (Lond) 9 (2014) 1475-86.

## [2] Goldberg MS Cell 161 (2015) 201-4

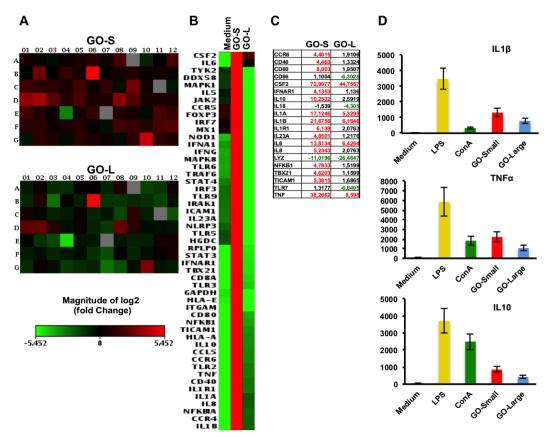
[3] Orecchioni M, Bedognetti D, Sgarrella F, Marincola FM, Bianco A, Delogu LG, Journal of translational medicine 21; (2014) 12:138.

[4] Pescatori M, Bedognetti D, Venturelli E, Menard- Moyon C, Bernardini C, Muresu E, Piana A, Maida G, Manetti R, Sgarrella F, Bianco A, Delogu LG. **Biomaterials 34 (2013) 4395-403.** 

## Figures



**Figure 1. Characterizations of GO-Small and GO-Large.** Atomic force microscopy (AFM) and transmission elettronic microscopy (TEM) images of GO-Small and GO-Large respectively. All scale bars are 1µm.



**Figure 2. Immune gene expression array**. A) Heatmap comparison of 84 genes after exposure to GO-Small or GO-Large. B) Heat map detail showing the immune transcript upregulated by GO-Small in PBMCs. C) Table of modulate genes in GO-Small and GO-Large versus control. Red show genes with a fold change greater than 4, green show genes with a fold regulation less than 4. D) Multiplex cytokine secretion analysis of GO-small and GO-large samples, Interleukin 1 $\beta$  (IL1 $\beta$ ), Tumor Necrosis factor  $\alpha$  (TNF $\alpha$ ) and Interleukin 10 (IL10) are showed.

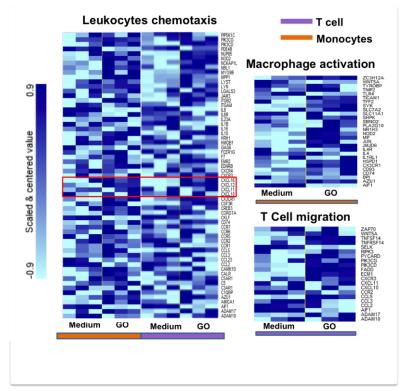


Figure 3. Whole genome expression analysis. Heatmap representation of GO-Small treatment for relevant Gene Ontology categories.