Graphene and immune cells: a box of opportunities

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Abstract

Despite significant progress of nanotechnology also in medicine, the development of new nanomaterials able to treat and prevents diseases such as cancer is still one of the biggest challenges [1]. In this challenge, a critical step is represented by the assessment of their impact on the complexity of the immune system [2]. Graphene oxide (GO), has attracted tremendous interest being explored for many potential applications in medicine [3]. To clarify its action on the immune cells, here, we propose an integrative analytical pipeline encompassing genomic and cellular characterization of the impact of graphene on primary immune cells. We used whole-transcriptomic analysis (Illumina BeadArray) for functional and molecular characterization of GO and GO functionalizated through the addition of amino groups (GONH₂) on human T-cells and monocytes. We then employed single cell mass cytometry. We identify 15 cellular main populations corresponding to 200 of distinct logically nodes but interconnected cell sub-populations. Notably, thanks to several analytical tools (i.e.SPADE and viSNE) we found that only the functionalized GONH₂ was able to

induce a specific dendritic cell and monocyte activation skewed toward a Th1/M1 response, as demonstrated by the production of increased classic M1 cytokines (TNFa, IL6, and CCL4) [Figure 1]. This effect was proved also by the overexpression of pathways critical for the development of an effective anti-tumor immune response (i.e. interferon signalling, CCR5 and CXCR3 ligands) [Figure 2]. Our findings report a new pipeline able to immunological carefully analyse the impact of new nanoscale platforms for a successful application in medicine.

References

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Figure 1: Heat map analysis of the immune cell behavior using CyTOF.



