

Table S1. Putative proteins (≈ 35 kDa) co-precipitated with LC3

| Protein | NCBI Reference Sequence | Score |
|-------------------------|--------------------------------|--------------|
| histone H1.2 | NP_005310.1 | 391 |
| histone H1.3 | NP_005311.1 | 391 |
| histone H1.4 | NP_005312.1 | 391 |
| unnamed protein product | BAG36431.1 | 391 |
| liver histone H1e | ABN72276.1 | 391 |
| Histone cluster 1, H1e | AAH96170.1 | 391 |
| histone H1.5 | NP_005313.1 | 330 |

Table S2 Putative proteins (≈ 11 kDa) co-precipitated with LC3

| Protein | NCBI Reference Sequence | Score |
|---------------------------|--------------------------------|--------------|
| unnamed protein product | CAA24918.1 | 306 |
| HIST2H4B protein, partial | AAI28106.1 | 306 |
| histone 1, H4c | EAW55528.1 | 306 |
| histone 1, H4e | EAW55544.1 | 306 |

The nuclear fractions were isolated from GO/CDDP-treated Skov-3 cells and same amount of proteins were subjected to immunoprecipitation using anti-LC3 MAb and Dynabeads[®] Protein A. The immunocomplex associated with LC3 was separated by 12% SDS-PAGE and the bands (≈ 11 kDa and ≈ 35 kDa) that were apparently upregulated after the GO/CDDP treatment were cut for LC/MS/MS analysis.

Table S1 shows putative proteins (≈ 35 kDa) with the highest scores (>300) after LC/MS/MS analysis and suggests that the protein was presumably histone H1.

Table S2 shows putative proteins (≈ 11 kDa) with the highest scores (>300) after LC/MS/MS analysis and suggests that the protein was presumably histone H4.

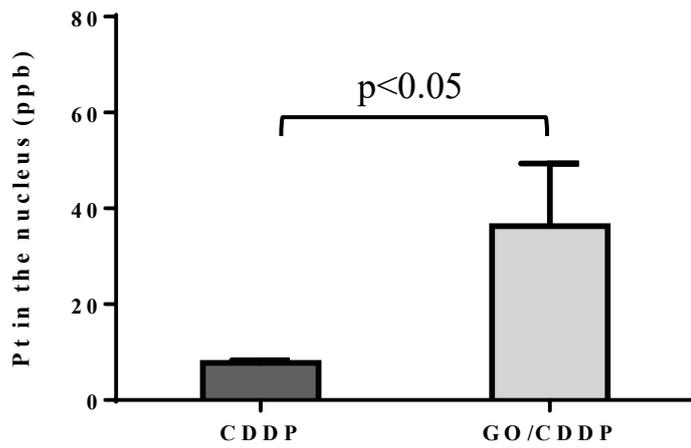


Fig. S1. To assess whether CDDP was concurrently transported into nucleus, we treated the Skov-3 cells with GO, CDDP or GO/CDDP as in Fig. 2A and separated the nuclear fractions, followed by ICP-MS analysis that detects the platinum in CDDP. As depicted in the figure, GO/CDDP triggered the nuclear import of significantly more CDDP than the CDDP group in Skov-3 cells.

Supplementary Materials and Methods

GO preparation and characterization

GO nanosheets (thickness < 2 nm) with characteristic functional groups and a lateral size of ≈ 450 nm in mean diameter (size range ≈ 100 -800 nm) were prepared as described [1] and dispersed in deionized water. Briefly, GO with a size of ≈ 2.4 μm was first prepared from natural graphite by the modified Hummers method and dispersed in water. The solution was centrifuged ($7,200 \times g$ for 5 min) to remove unexfoliated GO and byproducts and centrifuged again ($400 \times g$ for 15 min) to remove broken fragments and debris. The pellet was dried under vacuum overnight, weighed on an ultra-micro balance with 0.1 μg resolution and dissolved in deionized water (≈ 250 $\mu\text{g ml}^{-1}$). The GO solution was tip sonicated (Misonix Sonicator 3000) in an ice bath at a power of 30 W for 1 h, filtered through a 0.45 μm syringe filter and dried under vacuum overnight. The GO was characterized by atomic force microscopy, High-resolution X-ray photoelectron spectroscopy and Fourier transform infrared spectroscopy as described [1]. The GO nanosheet was weighed, reconstituted and well-dispersed in deionized water to ≈ 250 $\mu\text{g ml}^{-1}$. The size distribution of GO was characterized using Dynamic Light Scattering (380 ZLS, Nicomp, USA) as described [1].

1. Chen G-Y, Yang H-J, Lu C-H, Chao Y-C, Hwang S-M, Chen C-L, et al. Simultaneous induction of autophagy and toll-like receptor signaling pathways by graphene oxide. *Biomaterials*. 2012; 33: 6559-69.