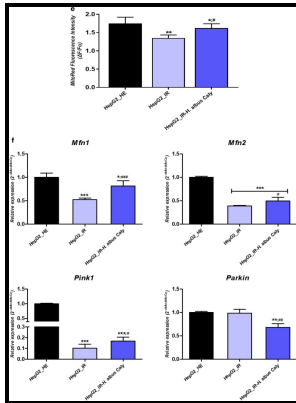# Characterisation of altered gene expression in response to Oxidative stree in HepG2 cells

University of Birmingham - A comparative proteomic analysis for capsaicin

Description: -
-Characterisation of altered gene expression in response to Oxidative stree in HepG2 cells
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Tags: #Characterization #of #hepcidin #response #to #holotransferrin #in #novel #recombinant #TfR1 #HepG2 #cells

Investigation into the effects of antioxidant

From the ELISA results, the level of 4-HNE adducts were found to be reduced after the leaf treatment as compared to the untreated HepG2 cells. You can through your profile settings.

Differential response of antioxidant defense in HepG2 cells on exposure of Livotrit®, in a concentration dependent manner

Padgett LE, Broniowska KA, Hansen PA, Corbett JA, Tse HM. Networks of interacting genes identified using STRING from the RNA-seq results from Huh7 cells at 3 DPI.

Altered global gene expression profiles in human gastrointestinal epithelial Caco2 cells exposed to nanosilver

International journal of infectious diseases: IJID: official publication of the International Society for Infectious Diseases. Therefore, the oxidatively-mediated damage may be from a combination of hyperglycemia and elevated FFA concentrations.

Glucose 6

Besides antioxidant enzymes, Nrf2 also regulates expression of HO-1.

CR

Fold change values were calculated by comparative C t analysis after normalizing with GAPDH mRNA.

CR

The higher the ROS amount, the higher the fluorescent intensity.

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