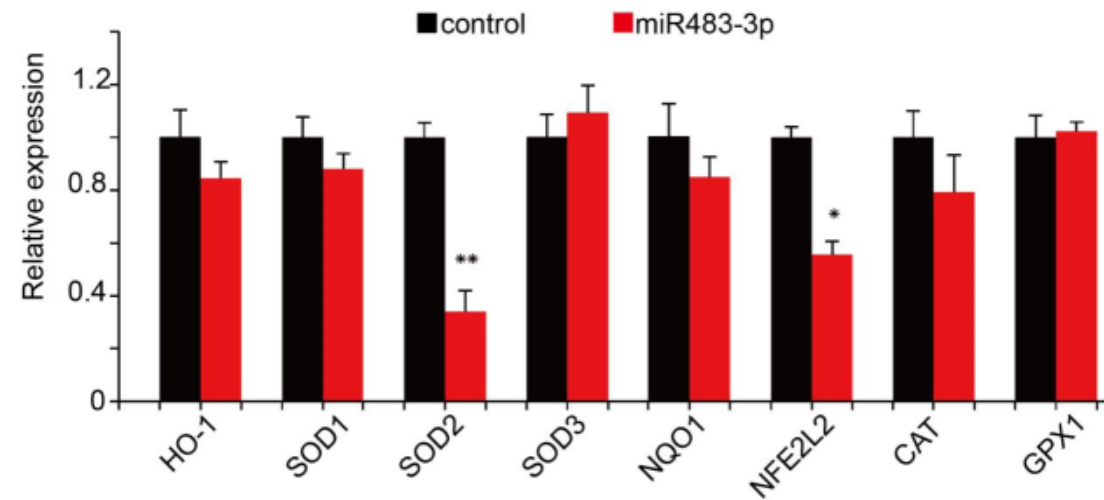
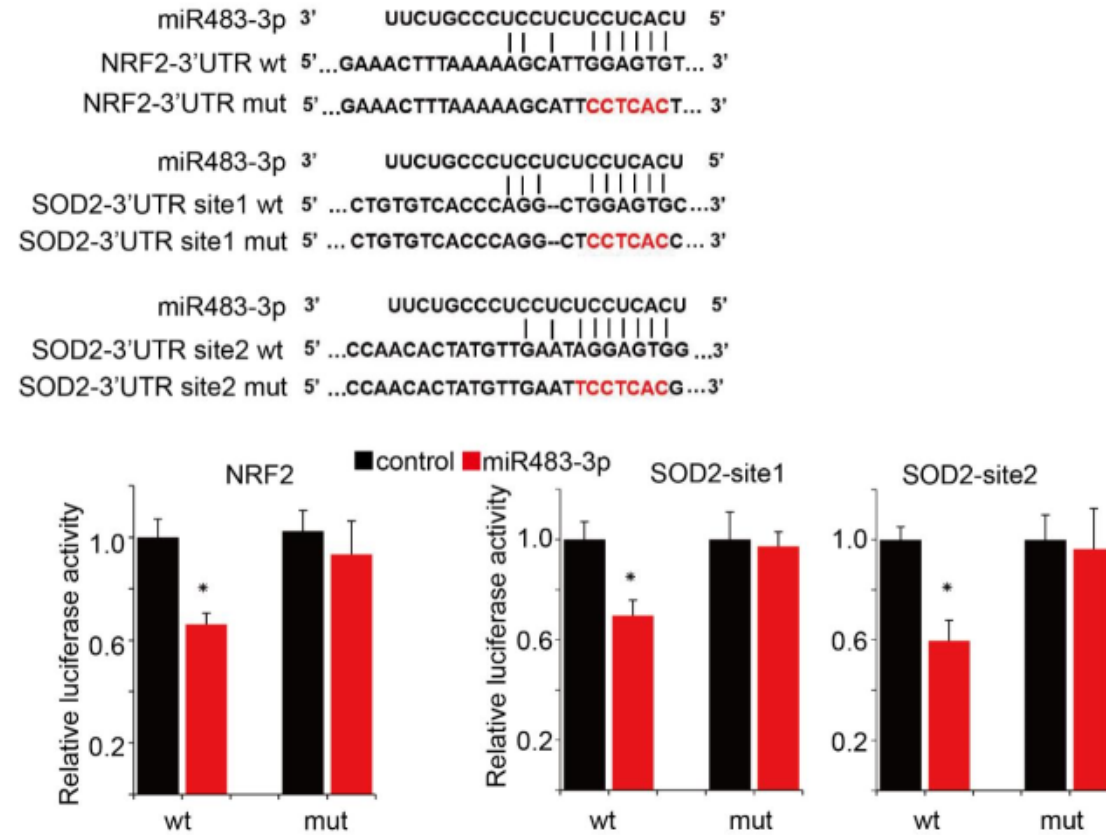


Supplementary Fig. 1



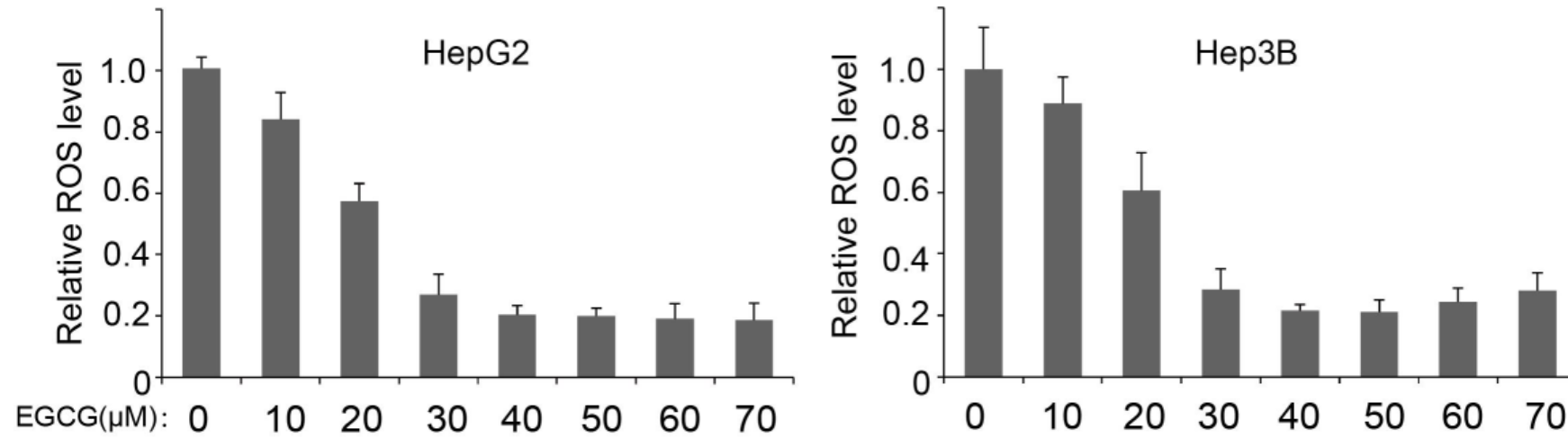
Supplementary Fig. 1. Identification of antioxidant genes targeted by miR483-3p.

Supplementary Fig. 2



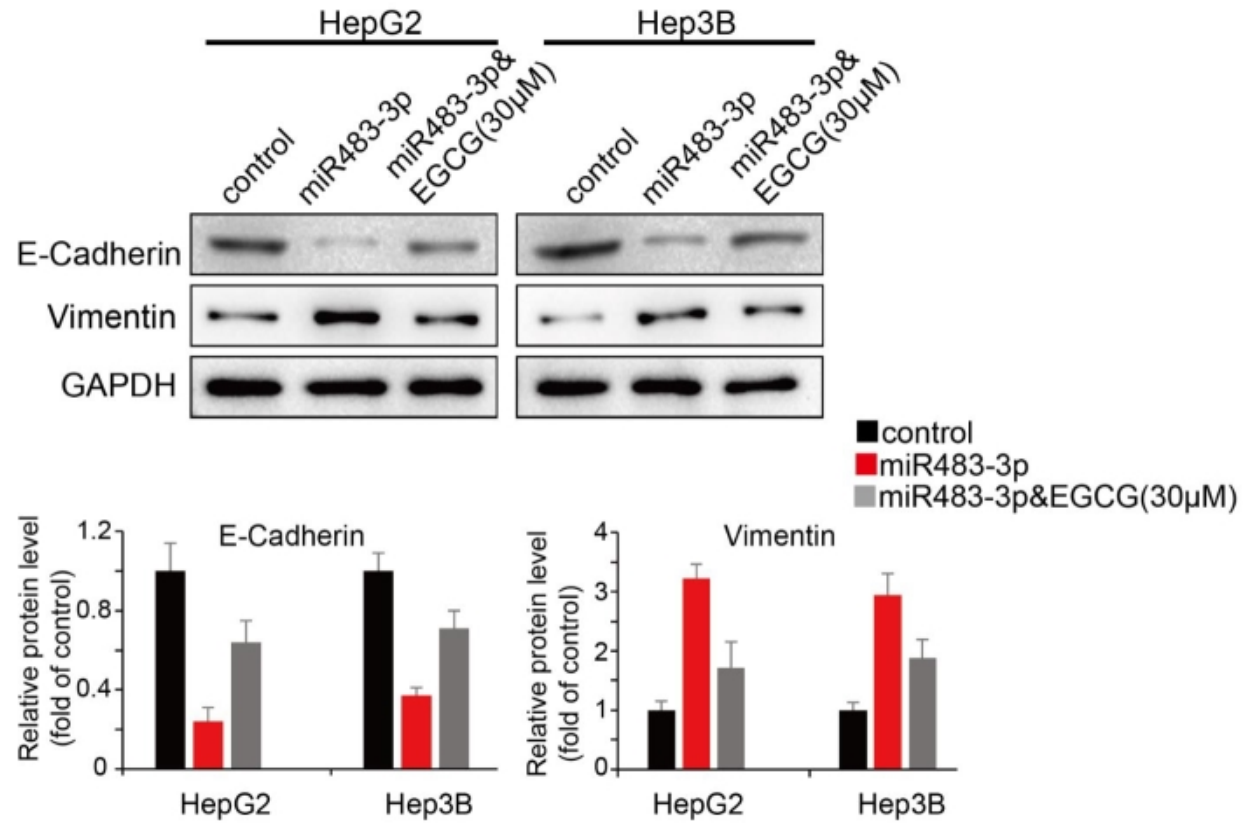
Supplementary Fig. 2. Luciferase activity assay of antioxidant genes targeted by miR483-3p.

Supplementary Fig. 3



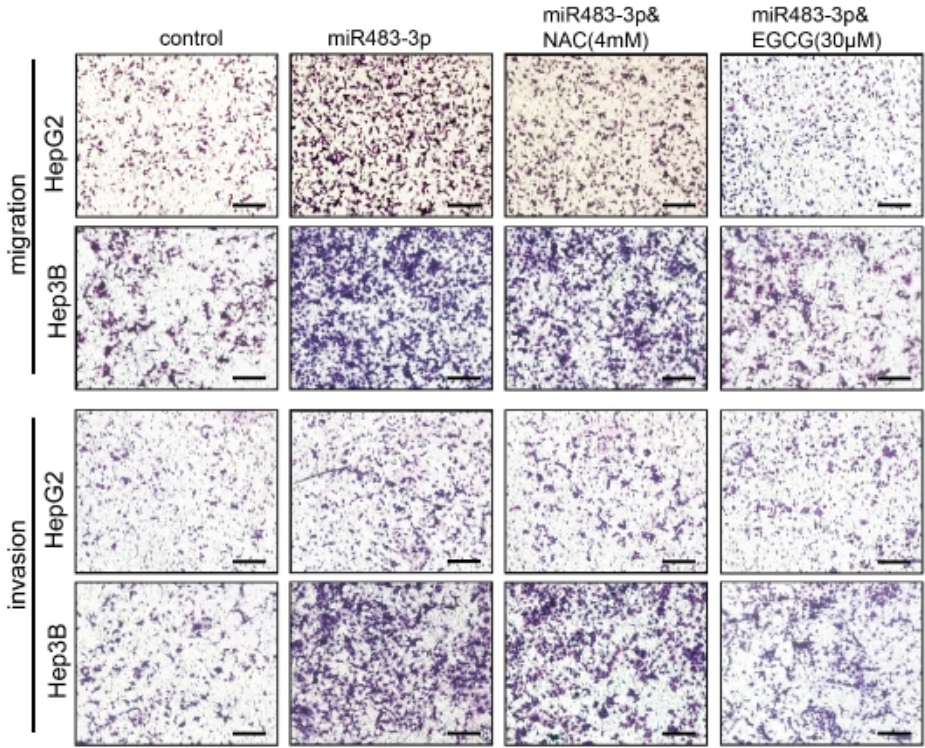
Supplementary Fig. 3. Effect of EGCG on ROS levels in miR483-3p-overexpressed HCC cells.

Supplementary Fig. 4



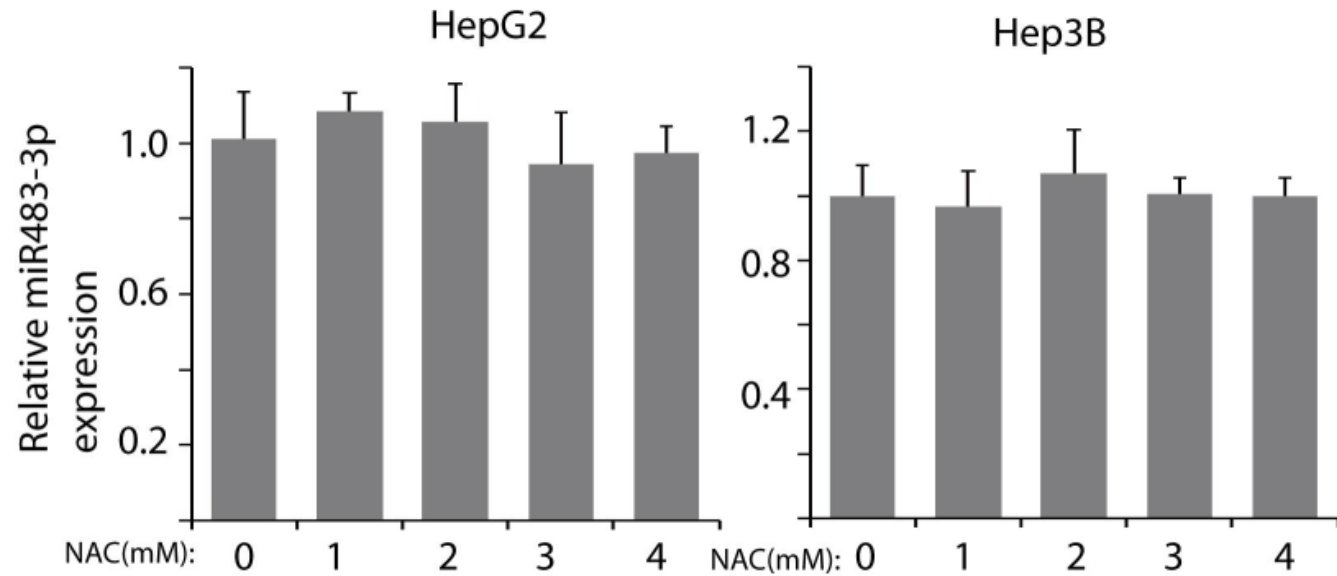
Supplementary Fig. 4. Levels of EMT markers (E-cadherin and vimentin) in HCC cells subjected to different treatments.

Supplementary Fig. 5



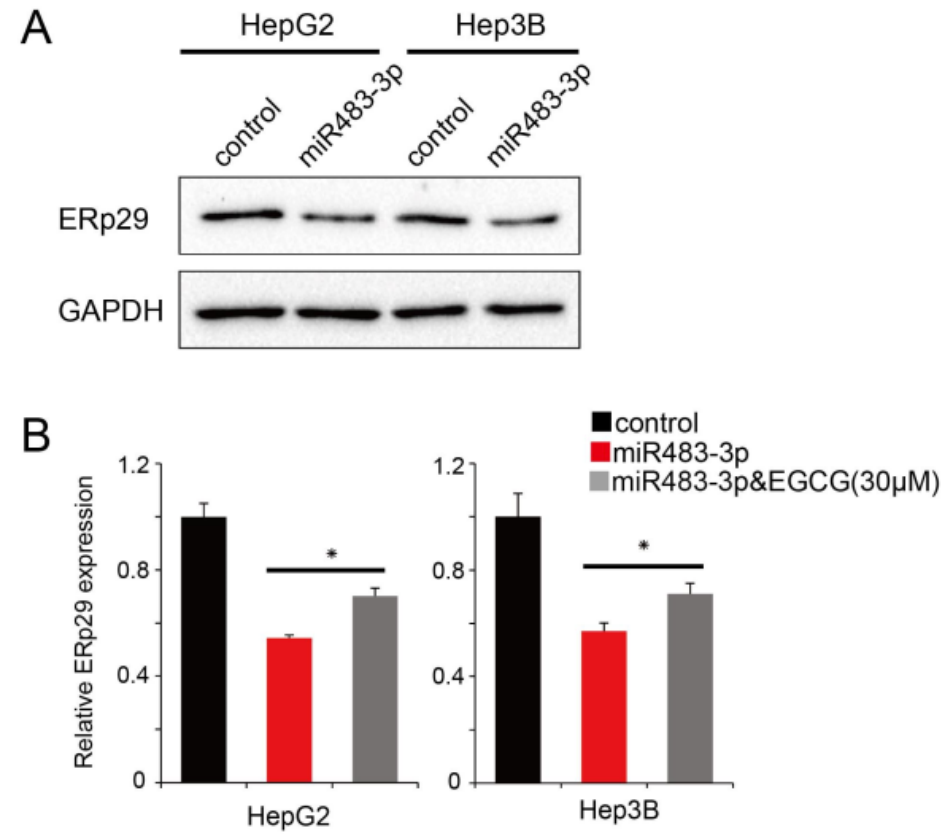
Supplementary Fig. 5. Migration and invasion assays of HCC cells subjected to different treatments. scale bar = 100µm.

Supplementary Fig. 6



Supplementary Fig. 6. Effect of different concentrations of NAC on the expression of miR483-3p in HCC cells.

Supplementary Fig. 7



Supplementary Fig. 7. Non-cytotoxic dose of EGCG partly restores the miR483-3p induced downregulation of ERp29 in HCC cells.

qPCR primers

| Forward | Reverse |
|--------------------------------|-------------------------|
| SOD2: GGAAGCCATCAAACGTGACTT | CCCGTTCCTTATTGAAACCAAGC |
| NRF2: TCAGCGACGGAAAGAGTATGA | CCACTGGTTTCTGACTGGATGT |
| SOD1: GGTGGGCCAAAGGATGAAGAG | CCACAAGCCAAACGACTTCC |
| SOD3: ATGCTGGCGCTACTGTGTTT | CTCCGCCGAGTCAGAGTTG |
| HO-1: AAGACTGCGTTCCTGCTCAAC | AAAGCCCTACAGCAACTGTCG |
| GAPDH: AATCCCATCACCATCTTCCA | TGGACTCCACGACGACTCA |
| CAT : TGGAGCTGGTAACCCAGTAGG | CCTTTGCCTTGGAGTATTTGGTA |
| GPX1: CAGTCGGTGTATGCCTTCTCG | GAGGGACGCCACATTCTCG |
| NQO1: GAAGAGCACTGATCGTACTGGC | GGATACTGAAAGTTCGCAGGG |
| miR483-3p: GCGAGCACTCACTCCTCTC | TGGTGTCGTGGAGTCGGC |

Stem loop primers for miR483-3p reverse transcription:

GCGTCTCAACTGGTGTCGTGGAGTCGGCAATTCAGTTGAGACGCAAGACGGGA

sgRNA targeting miR483-3p promoter

sgRNA-1:TCCAGCTGAGCATTGCTGTG

sgRNA-2:CTTGGGGGACCCCCGTGATG

Primers for BSP

forward: TTTTTTGAGAGGAGGGGAGGG reverse: CACCACCCCTAAAACACTAAAACA