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3 Supplementary Figure 1 CHAC2 is aberrantly expressed in a variety of tumors. (A)

4 Pancancer analysis of the TCGA database indicated that the mRNA levels of CHAC2 were

5 abnormally expressed in most tumors. (B) Analysis of the Kaplan–Meier Plotter database

showed that the high or low expression of CHAC2 in most tumors affected the prognosis ofpatients.

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## 10 Supplementary Figure 2 CHAC2 promotes the proliferation of lung adenocarcinoma.

- 11 (A) Immunoblot assays verify the efficiency of CHAC2 overexpression in the SPCA1 cell
- 12 line. (B) CHAC2 enhanced the proliferation of lung adenocarcinoma SPCA1 cells in vitro.
- 13 (C), (D) Transwell assays were conducted to examine the migration and invasion abilities in
- 14 PC9 cells with CHAC2 overexpression (C) and in H1299 cells with CHAC2 knockout (D).
- 15 Scale bar: 200µm. (E) CHAC2 enhanced the proliferation of lung adenocarcinoma SPCA1
- 16 cells in vivo.



- 18 Supplementary Figure 3 Heatmap analysis of genes correlated with the mRNA
- 19 expression level of CHAC2. (A) Heatmap analysis of genes positively correlated with the
- 20 mRNA expression level of CHAC2. (B) Heatmap analysis of genes negatively correlated
- 21 with the mRNA expression level of CHAC2.



- 23 Supplementary Figure 4 CHAC2 promotes lung adenocarcinoma progression. (A) Flow
- 24 cytometry was conducted to detect the cell cycle of SPCA1 cells overexpressing CHAC2. (B)
- 25 Flow cytometry was conducted to detect the apoptosis of SPCA1 cells overexpressing
- 26 CHAC2.

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## 28 Supplementary Figure 5 CHAC2 decreased GSH content and increased ROS levels.

- 29 (A) The levels of GSH were examined in SPCA1 cells overexpressing CHAC2. (B) Flow
- 30 cytometry assays were conducted to examine the levels of ROS in SPCA1 cells
- 31 overexpressing CHAC2. (C) Levels of ROS were examined in SPCA1 cells overexpressing
- 32 CHAC2 or vector with NAC (1 mM) or without treatment. (D) Colony formation assays of
- 33 overexpressing vector and CHAC2 SPCA1 cells with NAC (1 mM) or without treatment.



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## 35 Supplementary Figure 6 CHAC2 activates the MAPK pathway. (A) Pathway enrichment

- 36 analysis suggested that overexpression of CHAC2 significantly regulated the MAPK
- 37 pathway. (B) Immunoblot assays were conducted to detect phospho-ERK in vector and
- 38 CHAC2 overexpression SPCA1 cells. (C) Immunoblot assays were conducted to detect
- 39 phospho-ERK in vector and CHAC2 overexpression SPCA1 cells with NAC (1 mM) or
- 40 without treatment. (D) Colony formation assays of overexpressing Vector and CHAC2
- 41 SPCA1 cells with PD98059 ( $5\mu M$ ) or without treatment.
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