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Supplemental Information

Long Noncoding RNA OIP5-AS1 Promotes the

Progression of Liver Hepatocellular Carcinoma

via Regulating the hsa-miR-26a-3p/EPHA2 Axis

Yu-Shui Ma, Kai-Jian Chu, Chang-Chun Ling, Ting-Miao Wu, Xu-Chao Zhu, Ji-Bin Liu, Fei Yu, Zhi-Zhen Li, Jing-Han Wang, Qing-Xiang Gao, Bin Yi, Hui-Min Wang, Li-Peng Gu, Liu Li, Lin-Lin Tian, Yi Shi, Xiao-Qing Jiang, Da Fu, and Xiong-Wen Zhang

1 Supplemental Information

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- 3 Long noncoding RNA OIP5-AS1 promotes the progression of liver

4 hepatocellular carcinoma via regulating hsa-miR-26a-3p/EPHA2 axis

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- 6 Supplemental information contains three supplemental figures and legends.
- 7



8 Supplementary figure and legend

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Supplementary Figure 1. Identification of significantly dysregulated IncRNAs in LIHC. (A) Hierarchical clustering of significantly dysregulated IncRNAs in LIHC was performed using the multiple experiment viewer 4.7.1 software programs. (B-D) The Kaplan-Meier method was used to evaluate the relationship between IncRNA CRNDE (B), OIP5-AS1 (C) and ZEB1-AS1 (D) expression and overall survival of 371 LIHC patients from TCGA datasets.



Supplementary Figure 2. The expression of OIP5-AS1 in LIHC samples. 18 (A-C) The expression level of CRNDE (A), OIP5-AS1 (B) and ZEB1-AS1 (C) in 19 371 LIHC tissues and 50 adjacent noncancerous tissues. (D-F) The correlation 20 21 of CRNDE (D), OIP5-AS1 (E) and ZEB1-AS1 (F) in 371 LIHC tissues and 50 adjacent noncancerous tissues. (G) OIP5-AS1 expression level was examined 22 in GSE104310 dataset. (H) OIP5-AS1 expression level was examined in 23 GSE84005 dataset. (I) The correlation of OIP5-AS1 expression level and 24 tumor stage in 371 LIHC patients. 25

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Supplementary Figure 3. The expression and prognostic value of 28 hsa-miR-26a-3p and in LIHC samples. (A) The expression level of 29 hsa-miR-26a-3p in 371 LIHC tissues and 50 adjacent noncancerous tissues. 30 (B, C) The expression level (B) and correlation (C) of hsa-miR-26a-3p in 50 31 pairs of LIHC tissues and adjacent noncancerous tissues. (D) The correlation 32 of hsa-miR-26a-3p expression level and tumor stage in 371 LIHC patients. (E) 33 The Kaplan-Meier method was used to evaluate the relationship between 34 hsa-miR-26a-3p expression and overall survival of 371 LIHC patients. (F) The 35 Kaplan-Meier method was used to evaluate the relationship between 36 OIP5-AS1/hsa-miR-26a-3p expression and overall survival of 371 LIHC 37

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patients. H/L, high OIP5-AS1 and low hsa-miR-26a-3p expression; L/H, low 38 39 OIP5-AS1 and high hsa-miR-26a-3p expression. (G, H) The correlation between hsa-miR-26a-3p and OIP5-AS1 in 50 adjacent noncancerous tissues 40 (H) and 371 LIHC tissues (G). (I) Hsa-miR-26a-3p expression level was 41 examined in GSE21362 dataset. (J) Hsa-miR-26a-3p expression level was 42 examined in GSE36915 dataset. (K) Hsa-miR-26a-3p expression level in 241 43 adjacent noncancerous tissues, 180 non-metastatic and 60 metastatic LIHC 44 45 tissues from GSE6857 dataset. (L) Hsa-miR-26a-3p expression level in 178 alive and 59 dead LIHC patients from GSE6857 dataset. 46