



Ubiquitin-specific protease 22 promotes lipogenesis contributing to Hepatocellular Carcinoma pathogenesis



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Hepatocellular carcinoma (HCC), accounting for nearly 90% of liver malignancy, is the third most lethal human cancer worldwide with more than 830,000 deaths in 2020 [1]. Due to the complexity and heterogeneity of HCC, most patients are diagnosed at an advanced stage. While several phase III trials have been carried out for HCC therapy, limited clinical benefits were obtained because of quickly acquired drug resistance or considerable toxicity [2]. It remains an urgent need to explore effective novel strategies to combat HCC.

Dysregulation of cellular metabolism is a hallmark of cancers, especially HCC. Of note, the rapid increase of HCC cases is partly due to the epidemic of obesity, and the subsequent development and progression of metabolic-associated fatty liver disease (MAFLD), which made metabolic disorders a major risk factor for HCC [3–15]. Elevated de novo lipogenesis is a significant factor for the development of MAFLD and HCC [16]. Unfortunately, there is no substantial progress in therapeutic agents targeting lipid synthesis due to toxicity or complications [17]. It is of utmost importance to find more effective targets for fatty acid synthesis.

In the recent study published in *Nature Communications*, Ning et al. analyzed and observed that a variety of lipids including fatty acids, phospholipids and sphingomyelin were significantly enriched in HCC tissues compared with adjacent normal tissues using non-targeted metabolomics [18]. They found that the deubiquitinase ubiquitin-specific protease 22 (USP22), rather than the other USP members highly correlated with the abnormal upregulation of lipid synthesis [18]. Ning et al. performed Immunoprecipitation pulldown-Mass spectrometry (IP-MS) assay to reveal the regulatory mechanism and identified Peroxisome proliferator-activated receptor gamma (PPAR γ) as a reliable candidate substrate of USP22, which was

further confirmed by endogenous immunoprecipitation assay. PPAR γ is a key transcription factor that regulates lipid synthesis through the transcriptional activation of lipid synthesis enzymes such as acetyl-CoA carboxylase (ACC) and ATP citrate lyase (ACLY). PPAR γ is highly expressed in adipocytes, being implicated in lipid uptake, synthesis, and storage [19]. Ning et al. found that PPAR γ contributes to USP22-mediated ACC and ACLY upregulation and discovered a previously undescribed PPAR binding motif in the *ACACA* (the gene encoding ACC) promoter in HCC cells [18].

In addition, Ning et al. showed that USP22 deubiquitinated and stabilized PPAR γ by removing K48-linked ubiquitin chain which was catalyzed by E3 ligases CRL4BAH R and pVHL [18]. Genetic and pharmacological inhibition of PPAR γ abolished the regulatory effect on ACC and ACLY transcription, and significantly decreased lipogenesis and tumorigenesis caused by USP22 expression both in HCC cells and xenograft tissues. Furthermore, an *in silico* analysis revealed that HCC patients with upregulated USP22 and elevated levels of PPAR γ or ACC/ACLY exhibit poor prognosis and overall survival [18].

Taken together, in this study, Ning et al. demonstrated that USP22 promotes de novo synthesis of fatty acids and tumorigenesis by deubiquitinating PPAR γ in HCC (Fig. 1) [18]. Considering the limitation of clinical therapeutic options for HCC, emerging molecular targets are urgently needed. Findings from this research may provide a new option for targeting fatty acid synthesis that could yield therapeutic benefits to HCC patients with high USP22 expression.

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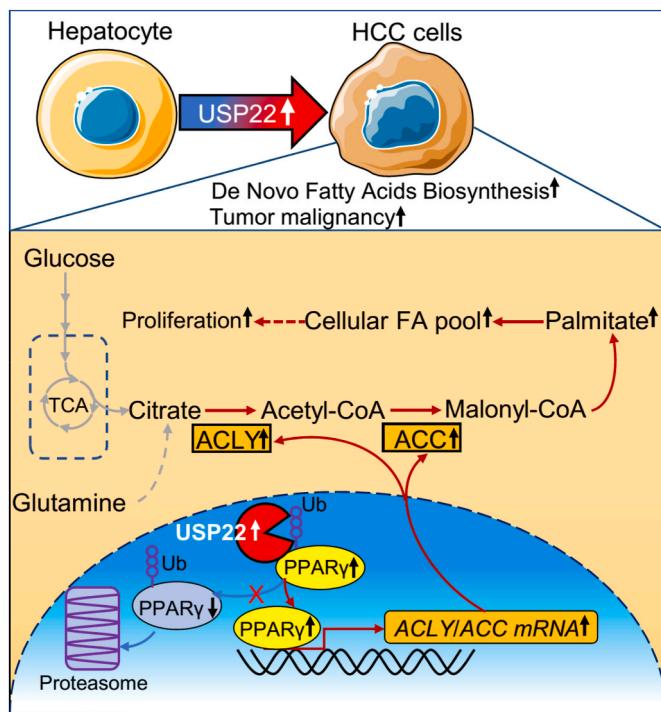


Fig. 1. USP22 regulates abnormal lipidome accumulation by lipogenetic mechanisms involving the PPAR γ -ACLY/ACC axis in HCC pathogenesis. This may provide a rationale for therapeutic targeting lipogenesis via USP22 inhibition.

Abbreviations: ACC: acetyl-CoA carboxylase; ACLY: ATP citrate lyase; HCC: Hepatocellular Carcinoma; PPAR γ : Peroxisome proliferator-activated receptor gamma; USP22: ubiquitin-specific protease 22.

Declaration of competing interest

None.

References

- [1] Sung H, Ferlay J, Siegel RL, Laversanne M, Soerjomataram I, Jemal A, Bray F. Global Cancer Statistics 2020: GLOBOCAN estimates of incidence and mortality worldwide for 36 cancers in 185 countries. CA A Cancer J Clin 2021;71(3):209–49.
- [2] Qin S, Bai Y, Lim HY, Thongprasert S, Chao Y, Fan J, Yang TS, Bhudhisawasdi V, Kang WK, Zhou Y, Lee JH, Sun Y. Randomized, multicenter, open-label study of oxaliplatin plus fluorouracil/leucovorin versus doxorubicin as palliative chemotherapy in patients with advanced hepatocellular carcinoma from Asia. J Clin Oncol 2013;31(28):3501–8.
- [3] Liu J, Dalamaga M. Emerging roles for stress kinase p38 and stress hormone fibroblast growth factor 21 in NAFLD development. Metabol Open 2021;12:100153.
- [4] Fotis D, Liu J, Dalamaga M. Could gut mycobiome play a role in NAFLD pathogenesis? Insights and therapeutic perspectives. Metabol Open 2022;14:100178.
- [5] Dalamaga M, Zheng L, Liu J. Gut mycobiome as a promising preventive and therapeutic target for metabolic disorders. Metabol Open 2022;13:100168.
- [6] Dalamaga M, Liu J. DRAK2-SRSF6-regulated RNA alternative splicing is a promising therapeutic target in NAFLD/NASH. Metabol Open 2021;13:100157.
- [7] Dalamaga M, Christodoulatos GS, Karampela I, Vallianou N, Apovian CM. Understanding the Co-epidemic of obesity and COVID-19: current evidence, comparison with previous epidemics, mechanisms, and preventive and therapeutic perspectives. Curr Obes Rep 2021;10(3):214–43.
- [8] Marouga A, Dalamaga M, Kastania AN, Antonakos G, Thrasyvoulides A, Kontelia G, Dimas C, Vlahakos DV. Correlates of serum resistin in elderly, non-diabetic patients with chronic kidney disease. Clin Lab 2013;59(9–10):121–128.
- [9] Hroussalas G, Kassi E, Dalamaga M, Delimaris I, Zachari A, Dionyssiou-Asteriou A. Leptin, soluble leptin receptor, adiponectin and resistin in relation to OGTT in overweight/obese postmenopausal women. Maturitas 2008;59(4):339–49.
- [10] Dalamaga M, Karmaniolas K, Arsenis G, Pantelaki M, Daskalopoulou K, Papadavid E, Migdalis I. Cedecea lapagei bacteremia following cement-related chemical burn injury. Burns 2008;34(8):1205–7.
- [11] Papadavid E, Gazi S, Dalamaga M, Stavrianeas N, Ntelis V. Palmoplantar and scalp psoriasis occurring during anti-tumour necrosis factor-alpha therapy: a case series of four patients and guidelines for management. J Eur Acad Dermatol Venereol 2008;22(3):380–2.
- [12] Marouga A, Dalamaga M, Kastania AN, Kroupis C, Lagiou M, Saounatsou K, Dimas K, Vlahakos DV. Circulating resistin is a significant predictor of mortality independently from cardiovascular comorbidities in elderly, non-diabetic subjects with chronic kidney disease. Biomarkers 2016;21(1):73–9.
- [13] Karampela I, Chrysanthopoulou E, Christodoulatos GS, Dalamaga M. Is there an obesity paradox in critical illness? Epidemiologic and metabolic considerations. Curr Obes Rep 2020;9(3):231–44.
- [14] Lempesis IG, Tsilingiris D, Liu J, Dalamaga M. Of mice and men: considerations on adipose tissue physiology in animal models of obesity and human studies. Metabol Open 2022;15:100208.
- [15] Dalamaga M, Liu J. A chromatin remodeling checkpoint of diet-induced macrophage activation in adipose tissue. Metabol Open 2022;15:100204.
- [16] Rohrig F, Schulze A. The multifaceted roles of fatty acid synthesis in cancer. Nat Rev Cancer 2016;16(11):732–49.
- [17] Vander Heiden MG. Targeting cancer metabolism: a therapeutic window opens. Nat Rev Drug Discov 2011;10(9):671–845.
- [18] Ning Z, Guo X, Liu X, Lu C, Wang A, Wang X, Wang W, et al. USP22 regulates lipidome accumulation by stabilizing PPAR γ in hepatocellular carcinoma. Nat Commun 2022;13(1):2187.
- [19] Desvergne B, Michalik L, Wahli W. Transcriptional regulation of metabolism. Physiol Rev 2006;86(2):465–514.

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