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Protein network module-based identification of key pharmacological pathways of *Curcuma* phaeocaulis Val. acting on hepatitis

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Abstract:

Ethnopharmacological relevance: Curcuma phaeocaulis Val. (CP), as the vital medicines for blood-breaking and disorder-eliminating, has been widely used for hepatitis with good curative effects. Owing to the complexity of traditional Chinese medicine, the pharmacological mechanism of CP remains unclear. To solve this problem, a protein network module-based approach was proposed in this study.

Materials and methods: Firstly, the content of active components of CP was detected based on HPLC-DAD. Then the liver protection of CP on Con A-induced hepatitis was validated via the analysis of serum levels of ALT, AST and LDH and histological findings. Next, the targets of CP components obtained from TCMD database were predicted by STITCH and ChEMBL retrieval. In addition, the protein interaction network (PIN) of CP was constructed by Cytoscape based on protein-protein interaction of targets obtained from STRING database. Following the topological analysis of CP PIN, it showed to exhibit the properties of scale-free, small world, and modularity matched with the property of complex biological networks. Finally, the functional modules were

Abbreviations: CP, Curcuma phaeocaulis Val.; ALT, Alanine aminotransferase; AST, Aspartate transaminase; LDH, Lactate dehydrogenase; PIN, Protein interaction network; HBV, Hepatitis B virus; HCV, Hepatitis C virus; TCM, Traditional Chinese medicine; PPIs, Protein-protein interactions; NFKB1, Nuclear factor kappa B subunit 1, p50; RELA, Transcription factor p65, p65; NFKBIA, NF-κB inhibitor alpha; NFKBIB, NF-κB inhibitor beta; IKBKB, inhibitor of nuclear factor kappa-B kinase subunit beta; CHUK, Conserved helix-loop-helix ubiquitous kinase; TGFB1, Transforming growth factor beta 1; TGFBR1, TGF beta type I receptor; TGFBR2, TGF beta type II receptor; SMAD3, Mothers against decapentaplegic homolog 3; SMAD4, Mothers against decapentaplegic homolog 4; ZFYVE9, SMAD anchor for receptor activation

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