

论著

蛇床子素通过激活过氧化物酶体增殖物激活受体 α 调节肝细胞内脂肪酸代谢

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摘要 目的 探讨蛇床子素是否通过激活过氧化物酶体增殖物激活受体 (PPAR) α 调节肝细胞内的脂肪酸代谢。方法 大鼠肝细胞在用蛇床子素 $12.5\sim 100\ \mu\text{mol}\cdot\text{L}^{-1}$ 作用24 h后,用比色法测定细胞内甘油三酯(TG)和游离脂肪酸(FFA)含量,用逆转录聚合酶链反应法测定PPAR α mRNA表达的变化。PPAR α 抑制剂MK886 $1\ \mu\text{mol}\cdot\text{L}^{-1}$ 预处理肝细胞24 h后,观察蛇床子素 $100\ \mu\text{mol}\cdot\text{L}^{-1}$ 作用24 h后对细胞内TG和FFA含量以及PPAR α 调控的靶基因包括固醇调节元件结合蛋白(SREBP)-1/2、脂肪酸合酶(FAS)、二脂酰甘油酰基转移酶(DGAT)、肉碱软脂酰转移酶(CPT)-1a、脂肪酸转运蛋白(FATP)4和肝脂肪酸结合蛋白(L-FABP)mRNA表达的影响。结果 蛇床子素 $12.5\sim 100\ \mu\text{mol}\cdot\text{L}^{-1}$ 可明显降低肝细胞内 TG和FFA的含量($P<0.01$),同时也能明显增加肝细胞内PPAR α mRNA的表达($P<0.01$)。在用PPAR α 抑制剂MK886预处理后,蛇床子素降低肝细胞内TG和FFA的作用则明显被减弱($P<0.01$),同时抑制SREBP-1/2, FAS和DGAT mRNA表达的作用明显减弱或完全消失($P<0.01$),增加CPT-1a, FATP4和L-FABP mRNA表达的作用也明显减弱或完全消失($P<0.01$)。结论 蛇床子素通过激活肝细胞中PPAR α 后可降低细胞中的TG和FFA含量,其机制与激活PPAR α 后随后抑制SREBP-1/2, FAS和DGAT的基因表达以及增加CPT-1a, FATP4和L-FABP的基因表达有关。

关键词 [蛇床子素](#) [脂肪酸](#) [过氧化物酶体增殖物激活受体 \$\alpha\$](#) [肝细胞](#)

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Osthole regulates fatty acid metabolism in hepatocytes by activation of PPAR α

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Abstract

OBJECTIVE To determine whether osthole regulates fatty acid metabolism in hepatocytes of rats by activating peroxisome proliferator-activated receptor (PPAR) α . **METHODS** Rat hepatocytes were cultured and treated with osthole $12.5\sim 100\ \mu\text{mol}\cdot\text{L}^{-1}$ for 24 h, before triglycerides (TG) and free fatty acid (FFA) contents in hepatocytes were determined by colorimetric method. PPAR α mRNA expression was determined by reverse transcription polymerase chain reaction. In order to determine whether the lipid-regulating effect of osthole was associated with activation of PPAR α , hepatocytes were pretreated with PPAR α inhibitor MK886 $1\ \mu\text{mol}\cdot\text{L}^{-1}$ for 2 h before incubation with osthole $100\ \mu\text{mol}\cdot\text{L}^{-1}$ for 24 h. TG and FFA contents, PPAR α -regulated target genes including sterol regulatory element-binding protein (SREBP)-1/2, fatty acid synthase (FAS), diacylglycerol acyltransferase (DGAT), carnitine palmitoyltransferase (CPT)-1a, fatty acid transporter protein (FATP)4, and liver fatty acid binding protein (L-FABP) mRNA expressions in hepatocytes were examined. **RESULTS** Osthole $12.5\sim 100\ \mu\text{mol}\cdot\text{L}^{-1}$ could significantly reduce TG and FFA contents and enhance the PPAR α mRNA expression in hepatocytes of rats($P<0.01$), but reduction of TG and FFA contents were significantly alleviated after pretreatment with PPAR α inhibitor MK886($P<0.01$). Similarly, the reduction of SREBP-1/2, FAS and DGAT mRNA expressions as well as the increment of CPT-1a, FATP4 and L-FABP mRNA expressions in hepatocytes of rats by osthole were also alleviated or abrogated after pretreatment with PPAR α inhibitor MK886($P<0.01$). **CONCLUSION** Osthole can decrease TG and FFA contents in hepatocytes. The mechanisms might be associated with activation of PPAR α , subsequent reduction of SREBP-1/2, FAS and DGAT gene expressions and increment of CPT-1a, FATP4 and L-FABP gene expressions.

Key words [osthole](#) [fatty acids](#) [peroxisome proliferator-activated receptor \$\alpha\$](#) [hepatocytes](#)

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