

## Responses of intestinalmicroflora diversity andlipid metabolism in humanflora associated C57BL/6J mice totea polyphenols diet

Xiaojing Zhang<sup>1\*</sup> Benhua Zeng<sup>2</sup> Zhenlin Liao<sup>1</sup> Zhiwei Liu<sup>1</sup> Hong Wei<sup>2</sup> Xiang Fang<sup>1</sup> (<sup>1</sup> College of Food Science, South China Agricultural University, Guangdong 510642; <sup>2</sup> Department of Laboratory Science, College of Basic Medicine Science, Third Military Medical University, Chongqing 400038)

Abstract: Diet is one of the mostimportantfactors to determine intestinal flora diversity and its functional genomics. Dietary regulation of the intestinal microbiotatopromotehuman health and it's mechanismare becoming abotresearch point. Diet polyphenols, an essential componentin the dietary structure, which has amost complicated interaction with the humanintestinal flora, has an important influence on the structure and function of the intestinal flora. Here we discussed the responses of colonflora diver-

sityandlipid metabolismto the tea polyphenols by using DGGE and RT-qPCR, body weight, fat tissue weight, glycemic index, blood lipid levels, insulin levels, sizeoffat cells. There has asignificantimpact when presents acertaindosetea polyphenols in the diet. These findings, which-preliminary revealed the relationship of polyphenols diet-colonic microflora-lipid metabolism, laid a foundation for further exploration of dietaryregulation of intestinal florato promote-human health.