

干旱及复水对油茶土壤养分及微生物的影响

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摘要:【目的】为探究土壤不同干旱程度对根际土壤养分与微生物的影响,指导油茶科学灌水。【方法】采用棚内盆栽试验,对3年生‘华金’进行不同土壤体积含水量处理(5%~10%、10%~15%、15%~20%、20%~25%),处理30天后均复水到体积含水量20%~25%,测定油茶生长量、土壤养分含量和土壤微生物多样性。【结果】1) 试验表明当土壤体积含水量降至较低水平(5%~15%)时,会导致油茶植株生长变缓。2) 不同处理下根际土壤土壤水分含量(SM)与pH、速效钾(AK)存在显著正相关关系,与全氮(TN)、硝态氮(NO₃-N)、有效磷(AP)和全钾(TK)存在极显著负相关关系;复水处理后各处理全磷(TP)、全钾(TK)及速效养分有所下降,pH提高。3) 高通量测序表明干旱复水能够改善油茶根际环境,提高微生物群落的丰富度和多样性;但复水降低了细菌酸杆菌纲(*Acidobacteria*)、嗜热油菌纲(*Thermoleophilia*)的相对丰度,提高了纤线杆菌纲(*Ktedonobacteria*)、*Saccharimonadia*和芽孢杆菌(*Bacilli*)的相对丰度。4) 冗余分析表明,油茶根际土壤理化因子SM、NO₃-N、TP对微生物群落结构存在显著影响。综合田间和盆栽试验结果,土壤含水量变化显著影响的细菌种类有 *Patescibacteria*、*Cyanobacteria*、拟杆菌属(*Bacteroidetes*),真菌种类有圆环菌纲(*Orbiliomycetes*)、*Leotiomycetes*、绿藻纲(*Chlorophyceae*)、根囊壶菌纲(*Rhizophydiomycetes*)和散囊菌纲(*Eurotiomycetes*)。【结论】控制土壤体积含水量在20%~25%时对土壤微生物群落多样性和丰富度最有利。

关键词: 干旱; 复水; 土壤养分; 土壤微生物

Effects of drought and re-water on nutrients and microorganisms in *Camellia oleifera* soil

Abstract: **【Objective】** To investigate the effects of different levels of soil drought on the growth of *Camellia oleifera* and the nutrient content and microbial community in the rhizosphere soil, provide guidance for scientific irrigation practices for *Camellia oleifera*. **【Methods】** A greenhouse pot experiment was conducted on 3-year-old 'Hujin' plants, subjected to different soil volumetric water content treatments (5%–10%, 10%–15%, 15%–20%, 20%–25%). After 30 days of treatment, all plants were rehydrated to a soil volumetric water content of 20%–25%. The growth parameters of *Camellia oleifera*, soil nutrient content, and soil microbial diversity were measured. **【Results】** 1) The experiment showed that when the soil volumetric water content decreased to lower levels (5%–15%), the growth of *Camellia oleifera* plants slowed down. 2) Under different treatments, the soil moisture content (SM) in the rhizosphere showed a significant positive correlation with pH and available potassium (AK), and a highly significant negative correlation with total nitrogen (TN), nitrate nitrogen (NO₃-N), available phosphorus (AP), and total potassium (TK). After rehydration, the total phosphorus (TP), total potassium (TK), and available nutrients decreased, while the pH increased. 3) High-throughput sequencing revealed that drought rehydration improved the rhizosphere environment of *Camellia oleifera*, enhancing the abundance and diversity of microbial communities. However, rehydration decreased the relative abundance of *Acidobacteria* and *Thermoleophilia*, while increasing the relative abundance of *Ktedonobacteria*, *Saccharimonadia*, and *Bacilli*. 4) Redundancy analysis indicated that soil physicochemical factors such as SM, NO₃-N, and TP had a significant impact on the microbial community structure. Based on the results of field and pot experiments, bacteria species such as *Patescibacteria*, *Cyanobacteria*, and *Bacteroidetes*, as well as fungal species such as *Orbiliomycetes*, *Leotioycetes*, *Chlorophyceae*, *Rhizophydiomycetes*, and *Eurotiomycetes*, were significantly influenced by soil moisture changes. **【Conclusion】** Maintaining a soil volumetric water content of 20%–25% is most favorable for the diversity and abundance of the soil microbial community.

Key words: Drought stress; Rewater; Soil nutrients; Soil microbial