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Root exudates compounds and microbial community composition regulates CH₄ dynamics in fire degraded tropical peatland

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Fires and drainage are common disturbance factors in tropical peatlands (TP) in Southeast Asia. These disturbances alter the hydrology, vegetation composition, and peat biogeochemistry; thereby affecting the microbiome where microbial communities reside. Studies from northern peatlands have well established the role of vegetation composition in regulating the labile C, in the form of plant root exudates, and microbial community composition affecting the peat decomposition; however, for tropics, it remains unexplored. Recent studies have also established how these fire-degraded TP areas become a hot spot of sedge-mediated CH₄ emission. To further our understanding of control mechanisms regulating CH₄ dynamics, we investigated the composition of plant root exudates (n=3 per plant species) from sedges (*Scleria sumatrensis*) and ferns (*Blechnum indicum*, *Nephrolepis hirsutula*), the most commonly occurring plant species at our fire-degraded tropical peatland site in Brunei, Northwest Borneo, as well as microbial community composition in plant (n=9 for *S. sumatrensis*, and *B. indicum*, and n=5 for *N. hirsutula*) rhizo-compartments (rhizosphere, rhizoplane, endosphere).

Using a targeted analysis, we found that the root exudates compounds secreted from sedge (*Scleria sumatrensis*) and one species of fern (*Blechnum indicum*) were significantly different (p<0.05) and showed a similar ratio of 2:1 for sugars (glucose, fructose) and organic acids (acetate, formate, lactate, malate, oxalate, succinate, tartrate), which is in contrast to that secreted from trees in intact tropical peatlands (1:2). Further, using 16S rRNA gene amplicon sequencing, we found that the microbial community composition in rhizo-compartments of plant species showed significant differences (p<0.001). Interestingly, the sedge species harboured a relatively higher abundance of methanogens (Thermoplasmata) and lesser methanotrophs (Alphaproteobacteria, Gammaproteobacteria) across all three compartments compared to fern species, which further supports the higher sedge-mediated CH₄ emissions from fire-degraded TP.

Our results provide fresh insights into the effects of post-fire vegetation composition in regulating the labile C and microbial community composition, and hence affecting CH₄ emissions

from fire-degraded TP. Further, our results can form an important basis for future CH₄ dynamics studies as the emissions might increase with the expansion of degraded TPs as a consequence of frequent fire episodes and flooding