

10.1071/SR15069_AC
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Supplementary Material: *Soil Research*, 2016, 54(3), 265–275.

Biochemical properties of highly mineralised and infertile soil modified by acacia and spinifex plants in northwest Queensland, Australia

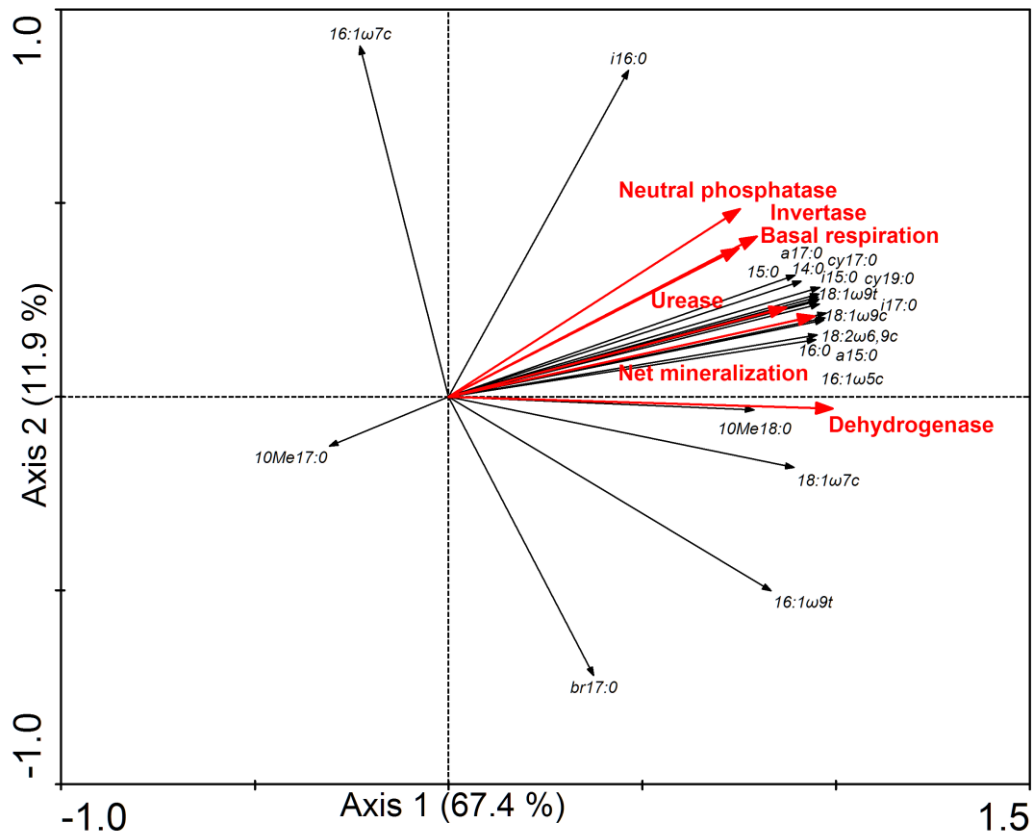
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Supplementary Fig. 1. RDA-function ordination biplot of individual phospholipid fatty acids (PLFAs) and biochemical processes.

Supplementary Table 1. Summary of representative fatty acids for typical microbial community

Microbial community		Representative PLFAs
Actinomycetes		10Me 18:0
AMF		16:1 ω 5c
Fungi		18:2 ω 6,9c, 18:1 ω 9c and 18:1 ω 9t.
Bacteria	G+ bacteria	i15:0, a15:0, i16:0, a17:0, i17:0 and br17:0
	G- bacteria	16:1 ω 7c, cy17:0, 18:1 ω 7 and cy19:0.
	Other bacteria	14:0, 15:0, 16:0,
Others		11:0, 18:0, and 10Me 19:0

Supplementary Table 2. Correlation between soil properties and microbial properties

	Basal respiration rate	Net mineralization rate	Dehydro genase	Invertase	Urease	Neutral phosphatase	Total PLFAs
WH C	0.61	0.83*	0.77	0.70	0.87*	0.74	0.83*
pH	-0.96**	-0.93**	-0.80	-0.80	-0.93*	-0.74	-0.86*
CEC	0.86*	0.84*	0.69	0.82*	0.90*	0.79	0.77
EC	-0.94**	-0.98**	-0.87*	-0.95*	-0.97**	-0.76	-0.93**
TOC	0.92**	0.97*	0.85*	0.92*	0.99*	0.76	0.91*
TN	0.91*	1.00**	0.93*	0.92**	0.98**	0.71	0.97**
MBC	0.91	0.99**	0.95**	0.90*	0.95*	0.66	0.97**

** $P < 0.01$; * $P < 0.05$; $n = 6$.