

Supplementary Materials

The rhizobial microbiome from the tropical savannah zones in Northern Côte d'Ivoire.

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Figure S1. Covering area of each sampling site and its corresponding features (not to scale).

Figure S2. Rarefaction curve of the seven samples, indicated by the number of ASVs depending on the size of the sequence sample.

Table S1. Physico-chemical properties of samples soils.

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Table S6. Measure of the richness and the alpha diversity per savannah zone.

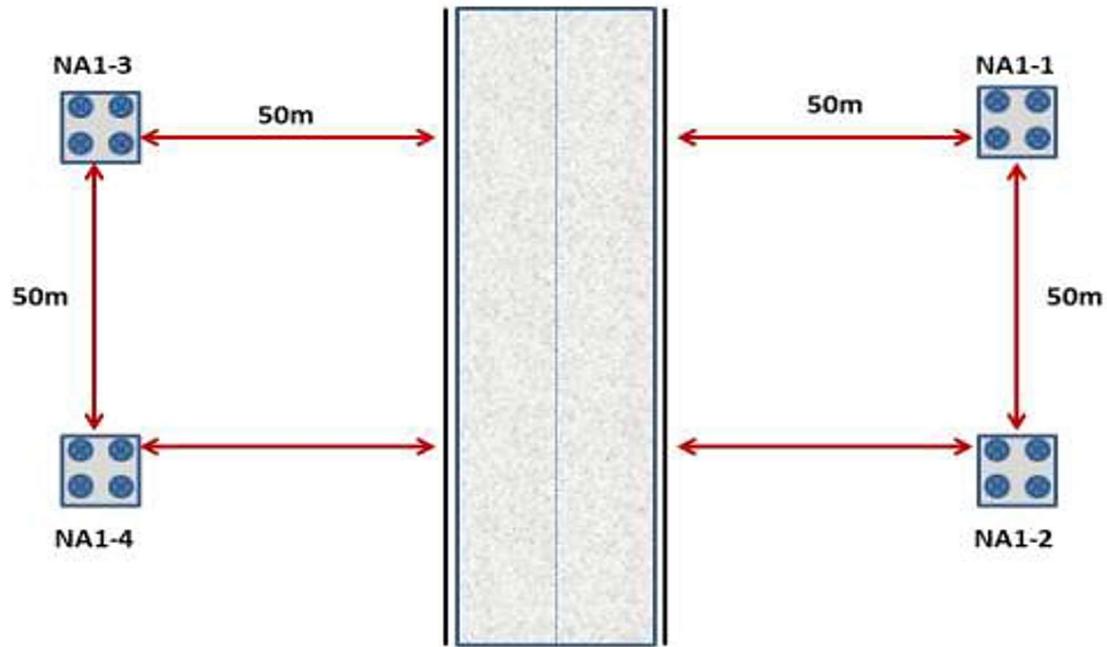


Figure S1. Covering area of each sampling site and its corresponding features (not to scale). Each sampling site was represented by an area of approximately. 100 m x 50 m with four independent sample locations (a virtual 1 m² quadrat) at the corners of the oblong. It is crossed in its length by a road of ca. 6 m wide. At each of the four independent sample locations, four topsoil cores (2 cm in diameter and 5 cm in depth) (pseudo-replicate samples) were collected, pooled together, and homogenized into a composite sample of approximately 25 g (replicate sample). Four independent replicate samples (4x25 g) obtained from four sample locations at each sampling site were kept in a labelled sterile plastic bag and formed an independent soil sample. A total of seven soil samples were collected following the same strategy for the seven sites studied in Northern Côte d'Ivoire.

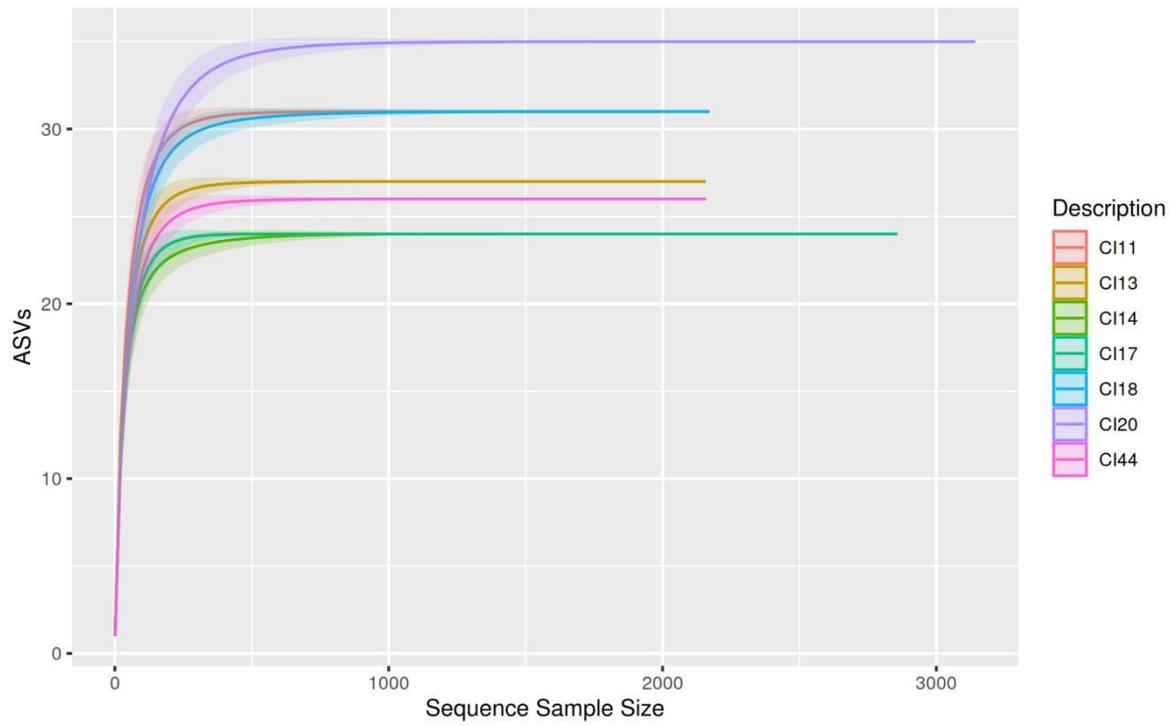


Figure S2. Rarefaction curve of the seven samples, indicated by the number of ASVs depending on the size of the sequence sample. Most rarefaction curves were able to reach the asymptote with a smaller number of sequences (<1000 sequences) than the subsampling size, suggesting that the sequencing effort and the subsampling size were appropriate.

Table S1. Physico-chemical properties of soil samples.

Soil ID	Soil texture (%)			Mineral elements (mg/kg)					Other parameters				
	Clay	Sand	Silt	Al	Ca	Fe	K	Mg	Mn	Na	P	C (%)	pH (H ₂ O)
CI11	13	77	10	413.0	958.9	57.0	56.6	220.9	42.5	23.4	8.1	1.3	6.6
CI13	8	84	8	216.3	646.0	42.8	32.3	103.9	101.9	17.8	11.5	0.7	6.6
CI14	12	81	7	391.1	1230.2	66.4	59.5	175.3	63.2	22.1	14.6	1.4	6.4
CI17	12	78	10	461.0	254.8	93.3	24.5	88.9	183.1	20.3	5.3	0.6	6.0
CI18	12	80	7	495.7	819.7	48.4	84.2	155.1	38.5	21.1	16.4	1.2	5.8
CI20	14	72	14	414.7	359.7	83.5	30.3	104.7	37.2	19.8	7.3	0.7	6.0
CI44	12	85	3	241.9	338.0	43.4	22.8	85.4	49.2	12.5	22.7	0.6	5.7

Table S5. ASVs richness, cumulative relative abundance of sequences and prevalence of ASVs per soil. *Microvirga*, *Paraburkholderia* and *Bradyrhizobium* which are the most important genera in terms of ASVs richness, relative abundance and ubiquity are highlighted in bold.

Genus (Final affiliation)	Number of ASVs per ge- nus	Relative abun- dance of ASVs per genus	Cumulative relative abundance of se- quences	Soil # CI11	Prevalence of ASVs per soil					
					# CI13	# CI14	# CI17	# CI18	# CI20	# CI44
<i>Microvirga</i>	24	31.17%	21.43%	12	13	9	4	9	5	5
<i>Paraburkholderia</i>	11	14.29%	9.03%	3	2	1	5	5	4	5
<i>Bradyrhizobium</i>	9	11.69%	49.05%	5	4	4	5	5	8	5
<i>Ensifer</i>	7	9.09%	1.23%	1	1	1	0	2	1	1
<i>Cupriavidus</i>	5	6.49%	5.14%	1	1	1	5	4	3	3
<i>Trinickia</i>	5	6.49%	2.57%	0	0	0	2	1	4	0
<i>Methylobacterium</i>	4	5.19%	1.72%	4	2	3	1	0	3	2
<i>Mesorhizobium</i>	3	3.90%	4.69%	2	2	1	1	2	3	3
<i>Devosia</i>	3	3.90%	2.68%	1	1	2	0	1	1	1
<i>Rhizobium</i>	3	3.90%	1.94%	1	1	1	1	2	2	1
<i>Shinella</i>	2	2.60%	0.31%	0	0	1	0	0	1	0
<i>Neorhizobium</i>	1	1.30%	0.19%	1	0	0	0	0	0	0
Total	77	100%	100%	31	27	24	24	31	35	26

Table S6. Measure of the richness and the alpha diversity per savannah zone. The richness is reported as the mean Chao1 index per soil and the alpha diversity is reported as the mean Shannon, Simpson and Fisher index per soil, with standard deviations shown in parentheses. All the richness and alpha diversity values were found to be statistically identical at the level =5 %.

Savannah zone	Chao1	Shannon	Simpson	Fisher
Sudanian savannah	27.33 (\pm 3.51)	2.91 \pm (0.12)	0.92 \pm (0.007)	4.55 (\pm 0.58)
Sub-sudanian savannah	29 (\pm 4.96)	2.77 (\pm 0.04)	0.89 (\pm 0.004)	4.59 (\pm 0.87)