

Measurement of N₂ fixation in 30 cowpea (*Vigna unguiculata* L. Walp.) genotypes under field conditions in Ghana, using the ¹⁵N natural abundance technique

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Abstract

In 2005 and 2006, 30 and 15 cowpea genotypes were respectively evaluated for plant growth and symbiotic performance at Manga in Northern Ghana, in order to identify N₂-fixing potential of these cowpea genotypes as source of N for cropping systems. The results showed differences in biomass production by the 30 or 15 cowpea genotypes. In 2005, cultivars Fahari, Mchanganyiko, IT97K-499-39, IT93K-2045-29 and IT84S-2246 produced the most shoot biomass, while Apagbaala, Brown Eye, ITH98-46, Vita 7 and Iron Grey produced the least. Of the 15 genotypes tested in 2006, cv. TVu11424 produced the largest amount of biomass, and ITH98-46, the least. Isotopic analysis of ¹⁵N in plant parts also revealed significant differences in δ¹⁵N of the cowpea genotypes studied. As a result, the percent N derived from fixation (% Ndfa) also differed among the cowpea genotypes tested in 2005, with only 5 out of the 30 cultivars obtaining over 50% of their N from symbiotic fixation. Whether expressed as mg N.plant⁻¹ or kg N.ha⁻¹, the levels of N₂ fixation by the cowpea genotypes varied considerably during 2005 and 2006, with values of N contribution ranging from 14.1 kg N.ha⁻¹ by cv. TVu1509 to 157.0 kg N.ha⁻¹ by IT84S-2246 in 2005. The amounts of N-fixed in 2006 ranged from 16.7 kg N.ha⁻¹ by cv. ITH98-46 to 171.2 kg N.ha⁻¹ by TVu11424, clearly indicating genotypic differences in symbiotic N yield. Re-evaluating 15 out of the 30 cowpea genotypes for N₂ fixation in 2006, revealed higher % Ndfa values (>50%) in all (15 cowpea genotypes) relative to those tested in 2005, indicating greater dependence on N₂ fixation for their N nutrition even though, the actual amounts of fixed-N were lower in 2006. This was due, in part, to reduced plant biomass as a result of very late sampling in 2006, close to physiological maturity (72 DAP in 2006 vs. 46 DAP in 2005) when considerable leaf matter was lost. The amount of N-fixed in 2006 can therefore be considered as being under-estimated.

Keywords: Cowpea genotypes, plant growth, δ¹⁵N, % Ndfa, N-fixed

1. Introduction

Cowpea (*Vigna unguiculata* L. Walp.) is the most important food grain legume in Africa. As a result, it has very high genetic diversity and wide adaptation to different soil ecologies through out the continent (Ehlers and Hall, 1997). Dietarily, the grain contains about 22–25% protein (Oyenuga, 1959; Platt, 1962), and 57% carbohydrate (Platt, 1962), while the green leaves which are eaten as vegetables in many African countries, contain up to about 27–34.3% protein (Jasper and Norman, 1983; Ahenkora et al., 1998).

Cowpea plays a major role in tropical cropping systems, as it has been estimated to fix as much as 201 kg N.ha⁻¹ per season and contribute up to about 42 kg N.ha⁻¹ to the N nutrition of a following maize crop (Dakora et al., 1987). Because of its ability to improve the N economy of cropping systems, cowpea is traditionally grown as an intercrop with millet, sorghum, maize, and more recently cassava (Ehlers and Hall, 1997).

Symbiotic N₂ fixation in legumes can be measured using the ¹⁵N natural abundance technique (Shearer and Kohl, 1986). However, the major problem limiting the use of this method is the choice of suitable reference plants that accurately measure the ¹⁵N natural abundance of soil N

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taken up by the test legume (Shearer and Kohl, 1986; Bremer et al., 1993; Pate et al., 1994; Unkovich et al., 1994; Nyemba and Dakora, 2005). Fortunately, a number of studies have successfully used the ^{15}N natural abundance method to estimate N_2 in a range of grain and pasture legumes (Peoples et al., 1997; Unkovich et al., 1994; Pate et al., 1994). In Africa, the ^{15}N natural abundance has been used to estimate N_2 fixation in food grain legumes such as cowpea, groundnut, Bambara groundnut and pigeon pea (Ayisi et al., 2000; Adu-Gyamfi et al., 2007; Ncube et al., 2007; Ojiem et al., 2007; Adjei-Nsiah et al. 2008). Other methods such as the N difference and the ^{15}N fertilizer techniques have also been used to quantify N_2 fixation in cowpea (Eaglesham et al., 1981; Dakora et al., 1987; Dakora and Keya, 1997; Carsky et al., 2001; Ofori et al., 1987; Peoples and Herridge, 1990; Bado et al., 2006).

The data so far obtained have been variable, suggesting the need for a broader program of selecting cowpea for enhanced N_2 fixation. Thus, there is a need to identify high N_2 -fixing cowpea genotypes that can contribute large amounts of biological N towards increased yields in traditional African cropping systems. This need for high N_2 -fixing cowpea genotypes in African agriculture is heightened not only by the low levels of N in many of Africa's agricultural soils, but also by the fact that cowpea plays a critical role in household food security in Africa, and thus in the dietary quality of plant protein intake.

This study, conducted over a 2-year period during 2005 and 2006, examined plant growth, symbiotic performance, and N yield in as many as 30 cowpea genotypes collected from Ghana, South Africa, Tanzania, and the International Institute for Tropical Agriculture (IITA) in Nigeria, with the hope of identifying high N_2 -fixing genotypes for use as biofertilizers in traditional cropping systems. To ensure high symbiotic performance, the cowpea collection had a good mix of both breeder-improved cultivars and farmer-selected varieties. The aim of the study was to screen and select cowpea genotypes that fix high levels of N_2 for use in enhancing N nutrition in the cropping systems of Africa.

2. Methods and Materials

Field design and planting

Field experiments were conducted at Manga Station of the Savanna Agricultural Research Institute, Ghana, in 2005, using 30 cowpea genotypes collected from Ghana, South Africa, Tanzania, and the International Institute for Tropical Agriculture (IITA) in Nigeria. In 2006, 15 of the 30 cowpea genotypes were re-planted at Manga for further evaluation of N_2 fixation. A randomized complete block design was used with four replicate plots for each cowpea genotype.

Each plot measured 15 m^2 ($3 \text{ m} \times 5 \text{ m}$), with inter-row

spacing of 60 cm in 2005, and 75 cm in 2006. Cowpea seeds were planted 20 cm apart within each row. Weeds were manually controlled with hand hoes. Two sprays of lambda cyhalothrin (Karate 2.5 EC) insecticide were applied at flowering and at pod formation as minimum protection against insect pests.

Plant harvest and processing

In 2005, plants were sampled from the middle rows of each plot at 46 DAP corresponding to about 50% flowering, while in 2006, plants were similarly sampled from the middle rows at 72 DAP corresponding to early physiological maturity. Each plant was separated into shoots, roots and nodules, oven-dried (70°C) to constant weight, ground (0.85 mm sieve), and stored prior to ^{15}N analysis.

Determination of ^{15}N in plant samples, $\delta^{15}\text{N}$ and % Ndfa values

The ratio of $^{15}\text{N}/^{14}\text{N}$, the ^{15}N natural abundance (‰), and %N of both cowpea and reference plant material were analyzed using a Carlo Erba NA 1500 elemental analyzer (Fisons Instruments SpA, Strada Rivoltana, Italy) coupled to a Finnigan MAT 252 mass spectrometer (Finnigan MAT GmbH, Bremen, Germany) via a ConFlo II open-split device. Finely ground plant material was weighed (1.999 to 2.006 mg per shoot sample and 2.1 to 2.5 mg per root or non-fixing reference plant sample) into aluminum capsules, and loaded onto the Carlo-Erba system. *Nasturtium* spp. (4.62% N) was included as internal standard. All plant samples were combusted in evacuated quartz tubes in the presence of cupric oxide and metallic copper and the resultant gases cleaned on-line before entering the mass spectrometer. The $\delta^{15}\text{N}$ values of the *Nasturtium* spp. was used to correct machine errors.

As described by Unkovich et al. (1994), the ^{15}N abundance is usually expressed in a relative, δ (delta) notation, which is the ‰ deviation of the ^{15}N natural abundance of the sample from atmospheric N_2 (= 0.36637 atom % ^{15}N):

$$\delta^{15}\text{N} = \frac{\text{atom}\% \text{ } ^{15}\text{N sample} - \text{atom}\% \text{ } ^{15}\text{N air}}{\text{atom}\% \text{ } ^{15}\text{N air}} * 1000$$

The proportion of legume N derived from atmospheric N_2 fixation was estimated as (Shearer and Kohl 1986; Unkovich et al., 1993; Boddey et al 2000):

$$\% \text{ Ndfa} = [(\delta^{15}\text{N}_{\text{ref}} - \delta^{15}\text{N}_{\text{leg}}) / (\delta^{15}\text{N}_{\text{ref}} - B)] \times 100$$

where $\delta^{15}\text{N}_{\text{ref}}$ is the ^{15}N natural abundance of reference plant, $\delta^{15}\text{N}_{\text{leg}}$ is the ^{15}N natural abundance of legume and the B value is the ^{15}N natural abundance of cowpea

genotype which is dependent solely on symbiotic nitrogen fixation for its N nutrition. The *B* value used for cowpea was -1.759‰ and -0.940‰ for shoots and roots, respectively.

3. Results

Plant growth

Data collected on plant growth, symbiotic performance, and $\delta^{15}\text{N}$ are shown in Tables 1–6. The dry matter yield obtained in 2005 revealed significant differences among the 30 cowpea genotypes. At organ level, the genotype Fahari produced the largest amount of shoot dry matter, followed by the cvs. IT97K-499-39, IT84S-2246, and Mchanganyiko (Table 1). In contrast, two farmer varieties, Apagbaala and Bensogla, produced the least amount of shoot dry matter, generally about 3.4-fold less shoot biomass compared to cv. Fahari. The shoot dry matter yield of the remaining cowpea genotypes fell in between the two extremes (Table 1).

In 2005, root dry mass also differed among the 30 cowpea genotypes, with cvs. IT93K-2045-29, Brown Eye, Ngonji and Mchanganyiko showing greater root growth compared to cvs. Vuli-1, ITH98-46, TVu 1509, Iron Grey, ITH98-20, and Vita 7, which produced the least root dry mass (Table 1).

Because of the variation in organ biomass, whole-plant dry matter yield was also found to differ significantly among the 30 cowpea genotypes in 2005 (Table 1). For example, the cvs. Fahari, Mchanganyiko, IT97K-499-39, IT93K-2045-29 and IT84S-2246 produced the most biomass in 2005 (Table 1), while the genotypes Apagbaala, Brown Eye, Iron Grey, ITH98-46 and Vita 7 produced the least biomass on a per-plant basis (Table 1).

Similar to the 2005 study, there were large genotypic differences in the dry matter yield of the 15 cowpea genotypes evaluated in 2006. For example, cowpea cv. TVu11424 produced the greatest amount of shoot and plant total biomass in 2006, followed by the genotype Apagbaala, both of which were respectively 9- and 5-fold higher in biomass than ITH98-46, the cultivar with least dry matter yield (Table 2). Whole-plant (shoots + roots + nodules) dry matter yield of the remaining genotypes fell in between those of cvs. TVu11424 and ITH98-46.

Field nodulation in cowpea genotypes

There were strong differences in nodule numbers per plant in 2005. As shown in Table 1, Omondaw exhibited the highest nodulation, followed by Mchanganyiko, Fahari, Ngonji and Sanzie (all farmer-selected varieties) when compared to the other cowpea cultivars. In contrast, Apagbaala (a farmer variety) and four inbred cultivars (namely, CH14, IT82D-889, IT90K-76 and ITH98-20)

showed the least nodulation (Table 1). Nodulation levels of the remaining genotypes were interspersed between the low and highest nodulating genotypes. The nodule dry matter in 2005 followed a similar pattern as nodule number per plant. The cultivars Omondaw and Mchanganyiko, which produced the highest number of nodules per plant, also had the largest nodule dry mass per plant (Table 1). In contrast, cowpea genotypes TVu11424 and Apagbaala, which were poorly nodulated, showed the lowest nodule dry matter per plant (Table 1).

In 2006, nodulation in the 15 genotypes tested was highest in cvs. Brown Eye and CH14 compared to the rest, with cv. Glenda showing the least nodule number per plant (Table 2). Because of the high nodulation of cv. Brown Eye, its nodule dry matter weight was significantly greater in 2006 (Table 2). Cultivar IT93K-2045-29 showed the lowest nodule dry weight in 2006 (Table 2).

Amounts of N in organs and whole plants of cowpea genotypes

There were genotypic differences in the N content of shoots, roots and whole plants in 2005 (Table 1). For example, the N content of shoots and roots were highest in cultivars Fahari, Mchanganyiko, and IT93K-2045-29 and thus resulted in markedly greater total biomass of the three genotypes (Table 1). This was in contrast to cultivars Apagbaala, Bensogla, ITH98-46 and Omondaw, which generally had reduced shoot and root dry matter yield, resulting in much smaller total plant biomass (Table 1). The remaining genotypes produced intermediate plant biomass.

The N content of shoots and whole plants were markedly greater in cowpea cv. TVu11424, followed by Apagbaala in 2006 (Table 2). In contrast, the N content of shoots and whole plants of cv. ITH98-46 were significantly lower than those of all the other cowpea varieties (Table 2).

$\delta^{15}\text{N}$ values (‰) of reference plants

In 2005, maize, sorghum and the weed *Hyptis specifigera* were the reference plant species sampled from the experimental site. The mean shoot and root ^{15}N natural abundance values ($\delta^{15}\text{N}$) of these reference plants were +5.123‰ and +3.57‰, respectively. The reference plant species sampled in 2006 included *Metracarpus villosus*, *Cynodon dactylon*, *Pennisetum pedicellatum*, *Eleusine indica*, *Cenchrus ciliaris* and *Hyptis specifigera*; and their mean shoot and root $\delta^{15}\text{N}$ values were +4.31‰ and +2.44‰, respectively. These averaged mean $\delta^{15}\text{N}$ values of the reference plants sampled for each year were used to estimate % Ndfa in the different cowpea genotypes, as earlier studies have shown that the combined mean $\delta^{15}\text{N}$ value of herbaceous weeds from a site can provide an accurate estimate of N-fixed (Pate et al., 1994; Nyemba and Dakora, 2005).

Table 1. Dry matter and grain yield, and N content of field-grown cowpea plants sampled at 46 DAP at Manga, Ghana in 2005. Mean values followed by dissimilar letters in a column are significant at $P \leq 0.05$. *ps = photosensitive genotypes, therefore not podded at physiological maturity of the other genotypes.

Genotype	Nodule no. per plant	Dry matter				Grain yield (kg.ha ⁻¹)	N content		
		Nodule dry mass (mg.plant ⁻¹)	Shoots (g.plant ⁻¹)	Roots (g.plant ⁻¹)	Shoots+ roots+ nodules (g.plant ⁻¹)		Shoots (mg.plant ⁻¹)	Roots (mg.plant ⁻¹)	Shoots+ roots (mg.plant ⁻¹)
Apagbaala	6.4n	300.0o	14.1d	6.3de	20.9d	458de	639.3c	65.5def	704.8cd
Bengsogla	26.3h	1500.0ef	14.9d	5.6de	55.2abc	680bc	633.7c	166.4cde	800.2bcd
Botswana White	54.8d	1333.3fg	17.0cd	15.0bc	33.9cd	903ab	571.5c	71.5def	642.9d
Brown Eye	19.9i	2466.7cd	29.8abcd	22.7ab	22.0cd	637bcd	1306.7abc	317.9a	1624.6ab
CH14	10.9m	1300.0fgh	24.2cd	10.3cd	36.1cd	473cde	954.1abc	118.1def	1072.3bcd
Fahari	65.5c	1633.3e	47.8a	18.4abc	68.2a	444def	1823.9a	231.5abc	2055.4a
Glenda	33.4g	1133.3ghi	34.5abc	12.9cd	48.8abc	622bcd	1389.8abc	161.7cde	1551.4abc
Iron Grey	17.4ij	500.0n	19.2cd	5.9de	25.8cd	221g	715.4bc	70.1cde	785.4cd
IT82D-889	12.8lm	1500.0ef	29.4bcd	16.3bc	47.4abcd	533cd	1277.5abc	173.4cde	1450.9abc
IT84S-2246	17.2ij	2500.0c	43.3ab	14.5bc	60.5abc	444def	1598.0ab	170.4cde	1768.4ab
IT90K-59	32.3g	2300.0d	30.5abcd	17.2bc	50.3abc	436def	1115.1abc	206.9bcd	1322.0abcd
IT90K-76	13.1klm	866.7jkl	31.2abcd	15.2bc	47.5abcd	488def	1311.6abc	178.7cde	1490.3abc
IT93K-2045-29	36.4f	1000.0jk	36.0abc	23.6a	60.9abc	324efg	1412.2abc	290.4ab	1702.6ab
IT93K-452-1	43.2e	700.0lm	34.2abc	10.9cd	46.1abcd	429ef	1258.9abc	123.4def	1382.3abc
IT94D-437-1	16.0jk	500.0n	27.1bcd	16.1bc	43.8abcd	*ps	1053.3abc	204.4bcde	1257.6abc
IT97K-499-39	19.2i	1200.0ghi	44.9ab	15.5bc	62.1abc	429ef	1678.1ab	181.1cde	1859.2ab
ITH98-20	13.2klm	566.7mn	21.3cd	5.4de	27.4cd	518cde	971.1abc	60.0ef	1031.0bcd
ITH98-46	36.7f	800.0kl	16.3cd	4.9de	22.2cd	1036a	668.0c	60.4ef	728.4cd
Line 2020	37.3f	1066.7ij	20.4cd	13.5cd	35.2cd	*ps	888.9bc	166.1cde	1055.1bcd
Mamlaka	24.6h	800.0kl	18.3cd	13.3cd	32.6cd	470cde	790.8bc	186.5cde	977.2bcd
Mchanganyiko	73.6b	3133.3b	42.9ab	20.0abc	66.2ab	*ps	1773.0ab	252.4abc	2025.4a
Ngonji	52.2d	1066.7ij	27.9bcd	21.0abc	50.3abc	429ef	972.5abc	236.1abc	1208.6abcd
Omondaw	85.0a	4133.3a	15.9cd	8.5de	28.6cd	673bc	599.4c	101.7def	701.1cd
Pan 311	15.6jkl	800.0kl	31.2abcd	8.5de	40.6bcd	666bc	1330.5abc	102.5def	1433.0abcd
Sanzie	52.1d	1100.0hi	34.3abc	12.8cd	48.4abc	888abc	1201.5abc	171.2cde	1372.8abcd
TVu11424	16.9ij	100.0p	28.9bcd	6.5de	35.8cd	251fg	1207.7abc	79.8def	1287.5bcd
TVu1509	39.1f	2300.0d	17.9cd	5.9de	26.3cd	251fg	785.6bc	82.0def	867.6cd
TVX3236	37.7f	1333.3fg	29.9abcd	9.1de	40.5bcd	547cd	1218.3abc	102.1def	1320.3abcd
Vita 7	25.1h	833.3kl	17.8cd	5.0de	23.9cd	429ef	812.6bc	61.1ef	873.7cd
Vuli-1	25.4h	1633.3e	23.1cd	4.3e	29.2cd	606bcd	997.2abc	51.1f	1048.2bcd
LSD _{0.05}	398.5***	177.4***	2.2**	3.2***	3.2***	6.5**	1.8**	5.9***	2.8***
C.V. (%)	5	9	4	32	30	11	13	27	12

$\delta^{15}\text{N}$ values (‰) and % Ndfa of cowpea plants

The $\delta^{15}\text{N}$ natural abundance ($\delta^{15}\text{N}$ values) of shoots, roots, and whole plants differed significantly for the 30 cowpea cultivars in 2005. Farmer varieties such as Bengsogla, Sanzie, Ngonji, Mamlaka, and the improved cv. IT84S-2246, showed the lowest $\delta^{15}\text{N}$ values in shoots in contrast to cvs. Apagbaala, Botswana White, TVu1509, TVX3236, IT90K-76, ITH98-20, IT94D-437-1, IT97K-499-39 and Fahari which generally exhibited the highest $\delta^{15}\text{N}$ in shoots (Table 3). Root $\delta^{15}\text{N}$ values also varied significantly among the 30 cowpea genotypes. Estimates of N derived from fixation (% Ndfa) in the 2005 trial revealed large genotypic differences in plant dependence on N_2

fixation for N nutrition. Only 5 out of the 30 cowpea genotypes derived over 50% of their N from symbiotic N_2 fixation (Table 3).

In 2006, there were significant differences in the $\delta^{15}\text{N}$ of shoots, roots and whole plants of field-grown cowpea at Manga. The $\delta^{15}\text{N}$ values of shoots and whole plants were generally very low compared to 2005 data (Table 4). Data on estimates of N derived from symbiotic N_2 fixation showed that genotypes with low $\delta^{15}\text{N}$ at organ level (e.g. TVu11424 and Botswana White) also had the highest % Ndfa values. In contrast, genotypes with high $\delta^{15}\text{N}$ values (e.g. Soronko, IT90K-59 and Glenda) exhibited the lowest % Ndfa values (Table 4).

Table 2. Dry matter yield and N content of cowpea varieties sampled at 72 DAP at Manga, Ghana, in 2006. Mean values with different letters in a column are significant at $P \leq 0.05$.

Genotype	Nodule no. per plant	Dry matter				Grain yield (kg.ha ⁻¹)	N content		
		Nodule dry mass (mg.plant ⁻¹)	Shoots (g.plant ⁻¹)	Roots (g.plant ⁻¹)	Shoots+ roots+ nodules (g.plant ⁻¹)		Shoots (mg.plant ⁻¹)	Roots (mg.plant ⁻¹)	Shoots+ roots (mg.plant ⁻¹)
Apagbaala	7.8e	73.3cd	17.4b	2.6c	20.5b	772.6cd	564.8b	61.0bcde	625.8b
Botswana white	8.3d	97.3bc	12.3de	2.5c	15.4d	688.2d	382.6cd	57.6cdef	440.2cd
Brown Eye	11.9a	137.3a	13.0cde	3.7b	17.2bcd	1007.1abcd	389.7bcd	68.2abcd	457.9bcd
CH14	11.3b	74.3cd	15.4bcd	1.59d	17.9bcd	1494.7a	525.4bc	33.9fghi	559.3bc
Glenda	3.7j	42.3f	15.3bcd	2.67c	18.4bcd	1147.8abcd	559.8b	53.2defg	613.0bc
IT82D-889	8.7c	103.7b	14.3bcd	4.59a	19.5bcd	1264.6abcd	431.0bcd	84.6ab	515.6bc
IT84S-2246	4.5hi	60.7def	14.5bcd	1.3de	16.2cd	915.4bcd	509.5bc	24.6hi	534.1bc
IT90K-59	5.6g	47.7ef	9.9ef	1.3de	11.7e	1360.5ab	312.7d	23.5hi	336.1d
IT93K-2045-29	4.7h	16.3g	14.2bcd	4.6a	19.4bcd	1332.9abc	516.4bc	88.8a	605.2bc
IT97K-499-39	7.5e	54.7def	15.7bc	2.5c	18.7bcd	734.1d	542.1bc	55.0def	597.1bc
ITH98-46	6.5f	62.3def	3.1g	0.8e	4.3f	749.9d	111.0e	14.8i	125.9e
Sanzie	6.7f	65.3def	13.6cd	2.0cd	16.1cd	880.4cd	412.8bcd	43.0efgh	455.7bcd
Soronko	5.5g	104.0b	13.7cd	3.5b	17.6bcd	771.6cd	427.9bcd	81.3abc	509.2c
TVu11424	5.6g	85.3bcd	33.0a	2.0cd	35.5a	839.5bcd	1177.1a	40.6efgh	1217.7a
Vuli-1	4.3i	91.7bc	7.6f	1.5de	9.5e	993.3abcd	265.1d	28.7ghi	293.8d
LSD _{0.05}	308.2***	14.1***	29.7***	21.0***	30.3***	2.3**	19.4***	9.1***	20.6***
C.V. (%)	8	10	14	18	13	21	20	28	18

Amounts of N-fixed in cowpea genotypes

Estimates of N-fixed in whole cowpea plants differed significantly, and ranged from 84.5 to 941.7 mg.plant⁻¹ in 2005 (Table 5). Cultivars IT84S-2246, Mchanganyiko and Sanzie fixed significantly more N on per-plant basis in 2005 compared to the other genotypes (Table 5). The lowest amount of N-fixed was observed in cowpea cv. TVu1509 at both organ and whole-plant level (Table 5). In 2006, the amounts of N-fixed in the 15 cowpea genotypes ranged from 111.0 to 1177.1 mg.plant⁻¹, with cv. TVu11424 showing significantly more fixed-N in shoots and whole plants compared to the other 14 genotypes. Cowpea cv. ITH98-46 indicated the least amount of N-fixed (Table 6). Genotypes intermediate between low and high levels of N₂ fixation included cvs. Apagbaala, CH14, IT84S-2246, IT93K-2045-29 and IT97K-499-39 (Table 6).

Whether expressed on a per-hectare or per-plant basis, the amount of N-fixed in 2005 remained significantly greater in cowpea genotypes IT84S-2246, Mchanganyiko and Sanzie, with cv. TVu1509 being the lowest, followed by Apagbaala, Bensogla and IT90K-76 (Table 5). As with the per-plant data, cowpea cv. TVu11424 fixed more N on a per-hectare basis in 2006 relative to the other genotypes (Table 6), and produced 10.3-fold more fixed-N per hectare than ITH98-46, the lowest N₂-fixing genotype in 2006 (Table 6).

4. Discussion

In this study, there were strong variations in organ development and overall plant growth of the 30 or 15 cowpea genotypes tested in 2005 and 2006, respectively (Tables 1 and 2). In 2005, the cowpea genotypes Fahari, Mchaganyiko, IT97K-499-39, IT93K-2045-29 and IT84S-2246 produced significantly more biomass of the 30 cultivars (about 3.3, 3.2, 2.97, 2.91 and 2.89-fold respectively greater than cv. Apagbaala, the least in dry matter production). However, the data also showed that, in 2005, the same genotypes (i.e. Fahari, Mchaganyiko, IT97K-499-39, IT84S-2246 and IT93K-2045-29) exhibited the highest N content per plant, which were respectively 2.9, 2.87, 2.63, 2.51 and 2.41-fold greater than that of Apagbaala, the cultivar with least N yield (Table 1). Similar findings were obtained in 2006, with genotypes TVu11424, Apagbaala and IT82D-889 being the highest in both dry matter production and whole-plant N content compared to cv. ITH98-46, which was least (Table 1). More specifically, the genotypes TVu11424, Apagbaala and IT93K-2045-29 produced 8.26, 4.77 and 4.51-fold more biomass, and had 9.7, 4.97 and 4.80-fold more N than cv. ITH98-46. As symbiotic parameters are reported to correlate with plant biomass (Herridge et al., 1990), the level of growth of a legume crop is therefore a function of its symbiotic N nutrition. In that regard, our data on plant growth and plant N yield of the best performing genotypes probably reflect their symbiotic performance. The higher nodulation (nodule

Table 3. $\delta^{15}\text{N}$ values and % Ndfa and B values of field-grown cowpea genotypes sampled at 46 DAP at Manga, Ghana, in 2005. Mean values followed by dissimilar letters in a column are significant at $P \leq 0.05$. Mean $\delta^{15}\text{N}$ values of shoots (5.12‰) and roots (3.57‰) of three plant species (i.e. maize, grain sorghum and the weed *Hyptis specifigera*) were used as reference plant values for estimating % Ndfa in cowpea.

Genotype	$\delta^{15}\text{N}$ (‰)		B values		% Ndfa	
	Shoots	Roots	Shoots	Roots	Shoots	Roots
Apagbaala	4.3bc	3.8ab	-1.759	-0.940	19.3jk	16.1ghi
Bengsogla	1.3l	2.0bcd	-1.759	-0.940	19.7jk	6.9i
Botswana White	4.2bc	3.9ab	-1.759	-0.940	59.6a	36.4cd
Brown Eye	3.8cd	3.6ab	-1.759	-0.940	25.3hijk	13.6i
CH14	3.5ef	3.2ab	-1.759	-0.940	30.2ghi	24.2fgh
Fahari	4.5b	2.3bcd	-1.759	-0.940	25.3hijk	25.8ef
Glenda	3.4ef	3.7ab	-1.759	-0.940	30.6ghi	15.3hi
Iron Grey	3.8de	2.4bcd	-1.759	-0.940	26.1hijk	15.9ghi
IT82D-889	3.1fg	3.1ab	-1.759	-0.940	34.3gh	32.6cdef
IT84S-2246	1.5kl	2.5bcd	-1.759	-0.940	56.0abc	27.2def
IT90K-59	2.1ij	2.6bcd	-1.759	-0.940	48.0bcd	31.6cdef
IT90K-76	5.1a	3.8ab	-1.759	-0.940	8.5l	16.2ghi
IT93K-2045-29	3.4ef	2.6bcd	-1.759	-0.940	30.8ghi	26.4ef
IT93K-452-1	2.3ij	2.8bc	-1.759	-0.940	45.6cd	24.6fg
IT94D-437-1	4.5b	3.5ab	-1.759	-0.940	16.3kl	25.1efg
IT97K-499-39	4.2bc	3.4ab	-1.759	-0.940	20.5ijk	26.2ef
ITH98-20	4.5b	3.2ab	-1.759	-0.940	16.3kl	15.1hi
ITH98-46	3.8cd	1.2d	-1.759	-0.940	25.3hijk	76.1a
Line 2020	3.6ef	2.8bc	-1.759	-0.940	27.7hij	25.6ef
Mamlaka	1.9jk	2.7bc	-1.759	-0.940	51.1abcd	24.2fgh
Mchanganyiko	2.4ij	2.1cd	-1.759	-0.940	44.0cde	26.0ef
Ngonji	1.5kl	2.8bc	-1.759	-0.940	56.0abc	38.6c
Omondaw	2.0ij	2.2cd	-1.759	-0.940	49.5bcde	29.3cdef
Pan 311	4.3bc	4.3a	-1.759	-0.940	19.1jk	8.7i
Sanzie	1.4l	2.5bcd	-1.759	-0.940	58.2ab	34.5cde
TVU11424	3.3ef	2.5bcd	-1.759	-0.940	32.6gh	33.8cdef
TVU1509	5.1a	2.2cd	-1.759	-0.940	7.9l	47.6b
TVX3236	5.1a	2.2cd	-1.759	-0.940	8.6l	53.4b
Vita 7	2.8hi	2.4bcd	-1.759	-0.940	39.6efg	50.4b
Vuli-1	3.1gh	2.6bcd	-1.759	-0.940	35.0fgh	32.4cdef
LSD _{0.05}	39.6***	1.7**	nd	nd	39.6***	1.7**
C.V. (%)	10	28	nd	nd	16	17

number and weight) observed in genotypes with better plant growth in 2005 (e.g. Mchanganyiko, Fahari and IT84S-2246, see Table 1), and cowpea cultivars that showed improved growth in 2006 (e.g. TVu11424 and IT93K-2045 in 2006, see Table 6) would seem to suggest that plant growth benefits from nodule functioning in symbiotic legumes (Table 6).

Evaluating N_2 fixation in legumes using the ^{15}N natural abundance technique usually requires isotopic analysis of both legume and reference plants (Shearer and Kohl, 1986; Unkovich et al., 1993; Ramos et al., 2001; Teixeira et al., 2006). In this study, different plant species were sampled as non- N_2 -fixing reference plants for measuring soil N uptake by the cowpea genotypes during both 2005 and 2006. Due to the large volume of data collected in this study, the mean $\delta^{15}\text{N}$ values of reference plant (shoots and roots) in 2005 and 2006 were used for estimating N_2 fixation in the

cowpea genotypes. Because there were large differences in $\delta^{15}\text{N}$ values of organs and whole cowpea plants, estimates of N derived from fixation in 2005 and 2006 were also found to vary considerably at both organ and whole-plant level (Tables 3 and 4). As predicted above, the cowpea genotypes with greater biomass accumulation were generally among those with the lowest whole-plant $\delta^{15}\text{N}$ values (see cvs. IT84S-2246 and IT93K-2045-29 in Table 3, or TVu11424 and IT93K-2045-29 in Table 4) and also showed significantly greater Ndfa values (whether measured as a percentage; see Tables 3 and 4, or in mg $\text{N} \cdot \text{plant}^{-1}$, see Tables 5 and 6). Interestingly, in 2005, only 5 out of the 30 cowpea genotypes obtained about 50% or more of their N from symbiotic fixation of atmospheric N_2 (Table 3), in contrast to 2006 where all 15 genotypes obtained over 50% of their N nutrition from symbiosis (Table 4). This difference in symbiotic functioning of the

Table 4. $\delta^{15}\text{N}$ (‰) and % Ndfa and B values of cowpea varieties sampled at 72 DAP at Manga, Ghana, in 2006. Mean values with dissimilar letters in a column are significant at $P \leq 0.05$. Mean $\delta^{15}\text{N}$ values of shoots (4.31‰) and roots (2.44‰) of six plant species (namely, *Metracarpus villosus*, *Cynodon dactylon*, *Pennisetum pedicellatum*, *Eleusine indica*, *Cenchrus ciliaris* and *Hyptis specifigera*) were used as reference plant values for estimating % Ndfa in cowpea genotypes.

Genotype	$\delta^{15}\text{N}$ (‰)		B values		% Ndfa	
	Shoots	Roots	Shoots	Roots	Shoots	Roots
Apagbaala	-0.2d	1.1de	-1.759	-0.940	74.0d	38.7ab
Botswana white	-0.8ef	1.1de	-1.759	-0.940	83.7bc	39.4ab
Brown Eye	-0.4d	1.6bc	-1.759	-0.940	76.8d	26.2bcd
CH14	-0.4d	0.8e	-1.759	-0.940	76.8d	48.3a
Glenda	0.9a	1.4cd	-1.759	-0.940	56.7g	31.6b
IT82D-889	0.1c	2.1a	-1.759	-0.940	69.3e	10.7e
IT84S-2246	-0.9f	1.5cd	-1.759	-0.940	85.0b	28.4bc
IT90K-59	0.4b	2.0a	-1.759	-0.940	65.0f	13.7de
IT93K-2045-29	-0.6e	1.3cd	-1.759	-0.940	81.0c	32.4b
IT97K-499-39	0.2bc	2.1a	-1.759	-0.940	67.4ef	9.4e
ITH98-46	-1.1g	2.2a	-1.759	-0.940	89.0a	7.8e
Sanzie	-0.8f	1.1de	-1.759	-0.940	84.4bc	39.1ab
Soronko	0.9a	1.9ab	-1.759	-0.940	55.7g	15.1de
TVu11424	-0.9fg	1.2cde	-1.759	-0.940	85.8ab	37.6ab
Vuli-1	-0.9f	1.5cd	-1.759	-0.940	85.0b	28.1bc
LSD _{0.05}	80.6***	9.1***	nd	nd	12.0***	111.8***
C.V. (%)	5	16	nd	nd	3	27

cowpea genotypes between 2005 and 2006 at Manga was due to an earlier history of very high N fertilization of cereal crops in that same field prior to its use in this experiment. Thus, while soil N may not have limited nodule formation in this study (Herridge and Betts, 1988; Streeter, 1988), it probably decreased nodule functioning in some genotypes (Dakora et al., 1992; Dakora and Keya, 1997; Ayisi et al., 2000), hence the low levels of N-fixed.

Whether expressed as mg N.plant⁻¹ or kg N.ha⁻¹, the levels of N₂ fixation by the cowpea genotypes varied considerably during 2005 and 2006 (Tables 5, and 6), with values of symbiotic N contribution ranging from 14.1 kg N.ha⁻¹ by cv. TVu1509 to 157.0 kg N.ha⁻¹ by IT84S-2246 in 2005. The amounts of N-fixed in 2006 also ranged from 16.7 kg N.ha⁻¹ by cv. ITH98-46 to 171.2 kg N.ha⁻¹ by TVu11424, clearly indicating genotypic differences in symbiotic N yield. When 15 out of the 30 cowpea genotypes were re-evaluated for N₂ fixation in 2006, all 15 genotypes showed higher shoot % Ndfa values relative to those tested in 2005 (Tables 3 and 4), indicating greater dependency on N₂ fixation for N nutrition (i.e. greater % Ndfa), even though the actual amounts of N-fixed per hectare were generally lower in 2006. This was due to the fact that the late sampling of plants in 2006 (at 72 DAP), resulted in loss of nodules and leaves from senescence, leading to reduced biomass and N accumulation (even though % Ndfa values and grain yield were much greater in 2006; see Tables 1–4). The data on N-fixed in 2006 can therefore be considered as being under-estimated. However, despite these differences in the age of cowpea plants (46

DAP in 2005 vs. 72 DAP in 2006), the amounts of N-fixed per shoot were generally similar for genotypes such as Botswana White, Brown Eye, IT82D-889, IT93K-2045-29, Glenda, and ITH98-46 during both 2005 and 2006.

In 2005, cowpea genotypes Fahari, Mchanganyiko, IT84S-2246 and Ngonji showed better plant growth and N-fixed in contrast to cvs. Apagbaala, ITH98-46, TVu1509 and Iron Grey, which exhibited the least plant growth and N₂ fixation (Tables 1 and 5). Interestingly, cowpea cv. Sanzie was not only the highest N₂-fixing genotype, it also produced the highest amount of grain yield compared to the other cowpea genotypes (Table 1). A similar pattern was observed in 2006, where cvs. TVu11424, Apagbaala, IT93K-2045-29 and CH4 showed greater plant growth and N-fixed, which led to significantly more grain yield by cvs. CH4 and IT93K-2045-29 compared to genotypes that produced low biomass and N-fixed (Tables 2 and 6). For example, cowpea cvs. ITH98-46, Vuli-1, IT90K-59 and Botswana White showed the least plant growth and N-fixed, which resulted in ITH98-46 and Botswana White producing the least grain yield (Tables 2 and 6). In a number of instances, there was thus a direct relationship between plant growth, N₂ fixation and grain yield, with the best performing genotypes showing greater plant growth, higher N₂ fixation and increased grain yield in contrast to low performing genotypes which showed poor plant growth, low N₂ fixation and low grain yields (Tables 1, 2, 5 and 6).

Compared to this study, cowpea has been shown to fix 24–29 kg N.ha⁻¹ in Kenya (Ssali and Keya, 1984), 122.0 kg

Table 5. Amount of N derived from fixation in field-grown cowpea genotypes sampled at 46 DAP at Manga, Ghana, in 2005. Mean values followed by dissimilar letters in a column are significant at $P \leq 0.05$.

Genotype	N-fixed					
	Shoots	Roots (mg.plant ⁻¹)	Shoots+Roots	Shoots	Roots (kg N.ha ⁻¹)	Shoots+Roots
Apagbaala	122.1cd	10.7f	132.8d	20.3cd	1.8f	22.1d
Bengsogla	123.5cd	11.1f	134.6d	20.6cd	1.8f	22.4d
Botswana White	337.8bcd	26.8def	364.6bcd	56.3bcd	4.5def	60.8bcd
Brown Eye	330.7bcd	42.9cde	373.6bcd	55.1bcd	7.1cde	62.3bcd
CH14	283.7bcd	28.7def	312.3bcd	47.3bcd	4.8def	52.1bcd
Fahari	640.1abc	59.0bcd	699.1abc	106.7abc	9.8bcd	116.5abc
Glenda	432.7bcd	24.3def	457.0bcd	72.1bcd	4.0def	76.2bcd
Iron Grey	186.2cd	11.4f	197.7cd	31.0cd	1.9f	32.9cd
IT82D-889	434.3bcd	56.0bcd	490.3bcd	72.4bcd	9.3bcd	81.7bcd
IT84S-2246	895.5a	46.2bcd	941.7a	149.3a	7.7bcd	157.0a
IT90K-59	531.3abc	65.4abc	596.7abc	88.6abcd	10.9abc	99.5abc
IT90K-76	112.2cd	29.0def	141.2d	18.7cd	4.8def	23.5d
IT93K-2045-29	445.8bcd	75.5ab	521.3bcd	74.3bcd	12.6ab	86.9bcd
IT93K-452-1	572.9abc	29.4def	602.3abc	95.5abcd	4.9def	100.4abc
IT94D-437-1	184.8cd	51.2bcd	236.0cd	30.8cd	8.5bcd	39.3cd
IT97K-499-39	345.4bcd	47.3bcd	392.8bcd	57.6bcd	7.9bcd	65.5bcd
ITH98-20	158.5cd	9.1f	167.6cd	26.4cd	1.5f	27.9cd
ITH98-46	168.6cd	43.4cde	212.0bcd	28.1cd	7.2cde	35.3cd
Line 2020	245.7bcd	42.6cde	288.3bcd	40.9bcd	7.1cde	48.0bcd
Mamlaka	422.0bcd	43.7cde	465.7bcd	70.3cd	7.3cde	77.6bcd
Mchanganyiko	767.8ab	65.5abc	833.3ab	128.0ab	10.9abc	138.9a
Ngonji	544.2abc	91.2a	635.4abc	90.7abcd	15.2a	105.9ab
Omondaw	297.0bcd	28.3def	325.3bcd	49.5bcd	4.7def	54.2bcd
Pan 311	244.8bcd	9.0f	253.8bcd	40.8bcd	1.5f	42.3bcd
Sanzie	701.2abc	59.3bcd	760.6abc	116.9abc	9.9bcd	126.8ab
TVu11424	396.4bcd	27.1def	423.5bcd	66.1bcd	4.5def	70.6bcd
TVu1509	48.2d	36.4def	84.5de	8.0d	6.1def	14.1d
TVX3236	105.3cd	54.4bcd	159.7cd	17.6cd	9.1bcd	26.6cd
Vita 7	322.0bcd	31.0def	353.1bcd	53.7bcd	5.2def	58.8bcd
Vuli-1	365.6bcd	16.3ef	381.9bcd	60.9bcd	2.7ef	63.7bcd
LSD _{0.05}	3.2***	19.2***	2.0**	3.2***	19.2***	2.0**
C.V. (%)	19	12	18	6	4	6

N.ha⁻¹ in Nigeria (Eaglesham et al., 1981), and 201.0 kg N.ha⁻¹ in Ghana (Dakora et al., 1987). It has also been shown in Australia that cowpea can derive about 53–69% of its N supply from N₂ fixation, and contribute about 87 kg N.ha⁻¹ in monoculture (Ofori et al., 1987). Because a large number of cowpea genotypes were evaluated in this study, the proportions and amounts of N derived from symbiotic fixation were within the range of values obtained in similar studies in Southern Ghana, Western Kenya and Southeastern Zimbabwe where the ¹⁵N natural abundance method was used to measure N₂ fixation in field-grown cowpea plants (Ncube et al., 2007; Ojiem et al., 2007; Adjei-Nsiah et al., 2008).

In conclusion, our results show that cowpea genotypes differ in their levels of N₂ fixation. This indicates that the choice of a high N₂-fixing genotype is important where biofertilization is the major objective for legume inclusion in the cropping system. The data further show that the low-

N status of African soils can easily be overcome if high N₂-fixing cowpea genotypes such as IT84S-2246, Mchanganyiko, Sanzie, IT93K-2045-29, IT93K-452-1, Ngonji and TVu11424 are used in cropping systems to provide symbiotic N, which can range from 61.0 to 171.2 kg N.ha⁻¹ (Tables 5 and 6).

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Table 6. Amount of N-fixed in field-grown cowpea varieties sampled at 72 DAP at Manga, Ghana, in 2006. Mean values followed by dissimilar letters in a column are significant at P≤0.05.

Genotype	N-fixed					
	Shoots	Roots (mg.plant ⁻¹)	Shoots+Roots	Shoots	Roots (kg N.ha ⁻¹)	Shoots+Roots
Apagbaala	564.8b	61.0bcde	625.8b	69.7b	3.9ab	73.7b
Botswana white	382.6cd	57.6cdef	440.2cd	53.2bcd	3.8ab	57.0bcd
Brown Eye	389.7bcd	68.2abcd	457.9bcd	50.0bcd	3.0abc	53.0bcd
CH14	525.4bc	33.9fghi	559.3bc	67.2b	2.7bcd	70.0b
Glenda	559.8b	53.2defg	613.0bc	53.0bcd	2.8bc	55.7bcd
IT82D-889	431.0bcd	84.6ab	515.6bc	50.0bcd	1.5cde	51.5bcd
IT84S-2246	509.5bc	24.6hi	534.1bc	72.1b	1.1cde	73.2b
IT90K-59	312.7d	23.5hi	336.1d	33.8de	0.5e	34.3de
IT93K-2045-29	516.4bc	88.8a	605.2bc	69.7b	4.7a	74.4b
IT97K-499-39	542.1bc	55.0def	597.1bc	60.9cd	0.8de	61.7bc
ITH98-46	111.0e	14.8i	125.9e	16.5e	0.2e	16.7e
Sanzie	412.8bcd	43.0efgh	455.7bcd	58.0bc	3.0abc	61.0cd
Soronko	427.9bcd	81.3abc	509.2bc	39.7cd	2.0bcde	41.7cd
TVu11424	1177.1a	40.6efgh	1217.7a	168.5a	2.7bcd	171.2a
Vuli-1	265.1d	28.7ghi	293.8d	37.4cde	1.4cde	38.8cd
LSD _{0.05}	9.1***	20.6***	6.89**	21.7***	14.0***	6.9**
C.V. (%)	27	20	22	20	5	19

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