Effectsof Pre-Emergence Herbicides, *Rhizobium* Strains and their Integration on Black Root Rot and Faba Bean Productivity

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የባቂላ የስር አበስብስ በሽታ በባቂላ ምርታማነት ላይ ከፍተኛ ጉዳት የሚያስክትል በሽታ ነው። የባቂላ የስር አበስብስ በሽታን ለመቆጣጠር የሚያስችል ምርምር በአምቦ የዕዕዎት ጥበቃ ምርምር ማዕክል እና በሆሉታ ግብርና ምርምር ማዕክል ስታስቲክስ ዲዛይንን በመጠቀም ተሰርቷል። የምርምሩ ውጤት እንደሚያሳየው በአምቦ አዕዋት ጥበቃ ምርምር ማዕክል እና በሆሉታ ግብርና ምርምር ማዕክል ከፍተኛ ምርት (በሄክታር2.26 እና3.64 ተን) በራይዞብየም ባክቴርያዝርያ 1035 (በሄክታር 500 ግራም) ከታከመው ማሳ ላይ ማግኘት ተችሏል። ከዚህም በተጨማሪ በራይዞብየም ባክቴርያ ዝርያ 1035 ከታከመው ማሳ ላይ ዝቅተኛ የባቂላ የስር አበስብስ በሽታ ክስተት (34.94 እና 13.68) እና የበሽታው የማጥታት ደረጃ (25.38 እና 14.39) ተመዝግባል። እንዲሁም ከፍተኛ የበሽታው ክስተት (61.77 እና 29.03) እና የበሽታው የማጥታት ደረጃ (66.23 እና 32.69) ካልታከመው ማሳ ላይ በአምቦ አዕዋት ጥበቃ ምርምር ማዕክል እና በሆሉታ ግብርና ምርምር ማዕክል ተመዝግባል። ምጣኔ ሀብታዊ ጥቅሙን ስናይ ከፍተኛ ጥቅም በራይዞብየም ባክቴርያ ዝርያ 1035 ከታከመው ጥሳ አይባለት ጥሰቃ ምርምር ለማዕክል እና በሆሉታ ግብርና ምርምር ማዕክል ተመዝግባል። ምጣኔ ሀብታዊ ጥቅሙን ስናይ ከፍተኛ ጥቅም በራይዞብየም ባክቴርያ በርያ 1035 ከታከሙት ማሳይች ላይ በሁለቱም ማዕክላት ተመዝግባል።

Abstract

Black root rot disease (Fusarium solani) is considered as a major problem in faba bean crops causing great losses in seed yield and low productivity. A randomized complete block design with nine treatments and three replications were employed at Ambo Plant Protection Research Center (APPRC) and Holetta Agricultural Research Center fields. The analysis of variance revealed that highly significant ($P \leq 0.001$) variation among the herbicides and Rhizobium strains treatments for grain yield, thousand seeds weight and Black root rot disease incidence and severity on faba bean. Thus, higher grain yields of 2.26 and 3.64t ha⁻¹were recorded from plots treated with Rhizobium strain FB1035. However, plots treated with herbicides plus Rhizobium strains gave lower yield than plots treated with Rhizobium strain alone. Plots treated with Rhizobium strain FB1035 showed less disease incidence (34.943 and 13.6767%) and severity (25.377 and 14.391%) respectively. However, the highest disease incidence (61.77 and 29.026 %) and severity (66.23 and 32.687 %) were recorded from the control plot. The disease severity was found negatively correlated with yield and yield components at both locations. Considering economic benefits, Partial budget analysis revealed that the highest total gross yield benefit (birr 35030 ha^{-1} and birr 56420 ha^{-1}) were obtained from plots treated with Rhizobium strain FB1035, proving that the application of Rhizobium strains alone were profitable. Thus, it is recommended to use these Rhizobium strains for the best protection against faba bean black root rot and improve the productivity of faba bean.

Introduction

Black root rot disease caused by *Fusarium solani* is one of the main constraints contributing to the low productivity of faba bean resulting in yield losses of up to 78 % in Ethiopia (Bogale *et al.*, 2009). In the highlands of Ethiopia, faba bean black root rot is among the most important disease in black clay soils, where water- logging is a big problem (Eshetu *et al.*, 2013).

Many species of *Rhizobium* are reported to inhibit significantly the growth of pathogenic fungi (Estevez de Jensen *et al.*, 2002). *Rhizobia* are reported as effective bio-control agent for the inhibition of soil-borne plant pathogens (Mazen *et al.* 2008).

In addition, in recent years farmers have begun to apply several herbicides to soil, not only for weed control but also because of an unconfirmed belief in a secondary controlling effect on root rot (Johnston, *et al.*, 1980). However, the presence of herbicide residues in soil could have direct impacts on soil microorganisms is matter of great concern. The effects of herbicides on soil fungi varied amongst herbicides depending on their application rates and the type of herbicide used (Sebiomo *et al.*, 2011). When herbicides applied frequently, they are accumulated in to the soil and at elevated levels impair the metabolic activities resulting in reduced growth of *rhizobia*, legumes or both (Khan *et al.*, 2004).

In view of the above, the present study was carried out with the objectives to determine the effect of pre-emergence herbicides and *Rhizobium* strains on black root rot and faba bean productivity and to assess the cost-benefit of herbicides and *Rhizobium* strains for faba bean production.

Materials and Methods

Planting and experimental materials

Experimental materials used for this field experiments were: Two *Rhizobium* strains, *Rhizobium* strain 1017 and *Rhizobium* strain 1035 at the rate of 500 gram per hectare, (carrier based powder inoculants) identified and released from HARC soil microbiology department were inoculated on the seed by using seed dressing method; Two pre- emergence herbicides, Dual Gold 960 EC at rate of 1.5 l/ha and Codal Gold 412.5 DC at the rate of 2.5 l/ha were sprayed immediately after planting. The faba bean Wolki cultivar released for Vertisols soil area from HARC was used as testing cultivar. The causal pathogen of black root rot on faba bean *Fusarium solani* was inoculated in the sick plot soil at Ambo. At Holetta, field experiment was conducted without inoculation of black root rot pathogen however; the presence of natural infestation of the pathogen was checked. These herbicides and *Rhizobium* strain 1017 alone, *Rhizobium* strain 1035 alone, Dual Gold 960 EC + *Rhizobium* strain 1017, Dual Gold 960 EC + *Rhizobium* strain 1035, Codal gold 412.5 DC + *Rhizobium* strain 1017, Codal Gold 412.5 DC + *Rhizobium* strain 1035 and *Fusarium solani* alone at APPRC and Un inoculated at HARC) were used as a treatments.

Experimental design and management

The treatment combinations were tested using faba bean cultivar Wolki in a randomized complete block design with three replications at both locations. The net plot size was 3.2 m x 4 m. Spacing of 0.4 m between rows and 0.1m between plants with 8 rows per plot. Diammonium phosphate (DAP) fertilizer was applied at 100kg ha⁻¹ with 18 kgha⁻¹ Urea at planting. The trials were kept weed-free by hand weeding.

Disease assessment

Disease severity

Randomly selected 10 Plants from 2^{nd} and 7^{th} rows were removed from the soil and roots were washed with tap water and disease severity was estimated visually by assessing necrotic lesions on the roots and hypocotyls using a rating scale of 0-4 described by Ondrej *et al.*, (2008) where 0:hypocotyls and roots white and firm, no root pruning, 1:slightly brown or discolored hypocotyls and roots, slight root pruning, 2:moderately discolored hypocotyls and roots, extensive root pruning, 3: darkly discolored hypocotyls and roots, hypocotyls completely collapse or, severe root pruning and 4:Dead or dying.

Disease severity = $\frac{\Sigma(ab)}{AK} * 100$ Bekriwala*et al.*, (2016)

a= number of diseased plants having the same degree of infection, b= degree of infection,

A= total number of examined plants, K= highest degree of infection

Disease incidence

It was measured as counting the dead plants out of the total plants.

Disease incidence(%) =
$$\frac{\text{Number of infected plant}}{\text{Total number of plant}} \times 100$$

Phenological and growth parameters

Days to 50% emergence, Day to 50% flowering, Days to 90% maturity, Plant height (cm), Stand count at seedling emergency and Stand count at harvest were recorded from the middle four rows for each plot.

Yield and yield components data

Number of pods per plant, Number of seeds per pod, 1000 seed weight (g), Total grain yield (t/ha), Viable nodule count and Non-viable nodule count were measured from five randomly taken plants from the middle four rows for each experimental unit and the means were recorded.

Relative yield loss

Percent relative grain yield loss was calculated as follows:

$$RYL (\%) = \frac{(Yp - Yt)}{Yp} \ge 100$$

Where, RYL = relative yield loss in percent, Yp = yield from the maximum protected plots and Yt = yield from other plots.

Correlation between yield and disease parameters and regression analysis

Correlations among the disease parameters and with the all yield component were tested at 5% probability level. The reliable yield loss was estimated based on the severity level by employing regression equations.

Cost - benefit analysis

The price of Faba bean seeds (birr/kg) was assessed from the local market and the total price of the yield of faba bean obtained from each treatment was computed on hectare basis. Input cost of herbicides and labor was converted into plot based. Since there was significant difference between mean yields of treatments, the obtained data was analyzed using the partial budget analysis method (CIMMYT, 1998). Then the treatment with the highest rate of return was recommended for use by the farmers. Marginal rate return was calculated using the following formula:

$$MRR = \frac{DNI}{DIC}$$

Where, MRR is marginal rate of returns, DNI is difference in net income compared with control, DIC is difference in input cost compared with control

Data analysis

All the data collected from field experiments were used and the analysis of variance (ANOVA) was performed using the PROC GLM in SAS (SAS Institute, 2012). The data on disease incidence was subject to square root transformation before analysis. The correlation coefficients of selected yield components were calculated using the SAS procedure PROC CORR (SAS Institute, 2012).

Results and Discussion

Disease assessment in the field

Days to disease onset

The analysis of variance (ANOVA) exhibited that there was highly significant (P< 0.001) difference among treatments in days to disease onset both at APPRC and at HARC. The longest (33.00 and 37.667 days) period to disease onset was recorded on the plot treated with the *Rhizobium* strain FB1035 and *Rhizobium* strain FB1017 at APPRC and plot treated with the *Rhizobium* strain FB1035 at HARC, while the shortest, 26.667 and 34.667 days period to disease onset were observed on the control plots at Ambo and Holetta, respectively. Similarly, the longest period to disease onset (30.667 and 37.333 days) was recorded on the plot treated with Codal gold + *Rhizobium* strain FB1035 and plot treated with *Rhizobium* strain FB1017 at Ambo and Holetta, respectively.

Disease incidence

There was highly significant (P \leq 0.001) difference among treatments on the black root rot incidence at APPRC and HARC. The highest disease incidences (61.77 and 29.026 %) were recorded on the control plots at APPRC and HARC, respectively and the lower disease incidences (34.943 and 13.6767 %) were recorded from plots treated with *Rhizobium* strain

process of *Phizobia* were found

FB1035 at APPRC and HARC, respectively (Table 1). Many species of *Rhizobia* were found to promote plant growth and to inhibit the growth of various soil-borne pathogens including *Macrophomina phaseolina, Rhizoctonia solani,* and *Fusarium solani* in both leguminous and non-leguminous plants (El-Batanony *et al.* 2007). In this line Buonassisi *et al.* (1986) was observed a significant reduction in root rot of bean caused by *Fusarium solani f.sp. Phaseoli* by inoculating the seeds with strains of *Rhizobium leguminosarum pv. Phaseoli*.

Thus, treatment of faba bean with effective strains of *Rhizobium leguminosarum* alone or in combination with other beneficial microorganisms may be preferred over the fungicides, because of their multiple potentials to fix nitrogen, control disease, improve of soil fertility, and increase crop productivity besides reducing the negative environmental impact associated with chemical use (Huang and Erickson, 2007).

Disease severity

The analysis of variance (ANOVA) indicated that there was highly significant (P< 0.001) difference among treatments in disease severity at both locations, APPRC and HARC. The highest disease severities of 66.23 and 32.69% were recorded on the control plots, while the lowest 25.377 and 14.391% were recorded from the plots treated by the *Rhizobium* strain FB1035 at APPRC and HARC, respectively (Table 1). Several previous studies reported that *Rhizobia* increase seed germination significantly and improve plant growth and yields through a reduction of soil-borne pathogens (Mazen *et al.* 2008). In this line, Rakib Al- ani *et al.* (2012) stated that, root rot severity caused by *Fusarium solani* was reduced by the addition of *Rhizobia* to the contaminated soil with the pathogen. Correspondingly, Samavat *et al.* (2011) also found that Common bean seeds treated with *Rhizobia* and *Pseudomonas fluorescens* isolates reduced root rot and damping-off severity.

| DDO | | | DI (%) | | DS (%) | |
|--------------|---------|---------|---------|---------|--------|---------|
| Treatment | APPRC | HARC | APPRC | HARC | APPRC | HARC |
| FB1017 | 33.00a | 37.33ab | 37.55fg | 15.31e | 32.01g | 16.31f |
| FB1035 | 33.00a | 37.66a | 34.94g | 13.68e | 25.38h | 14.39f |
| CG+ FB1035 | 30.67b | 35.00ef | 39.34f | 18.1d | 46.05e | 22.64cd |
| DG+ FB1035 | 30.00bc | 35.67de | 42.24e | 18.56cd | 38.96f | 17.14ef |
| DG | 30.00bc | 35.67de | 56.95b | 20.4c | 59.87b | 31.42a |
| CG | 29.67c | 36.67bc | 54.05c | 19.97cd | 55.39c | 27.09b |
| DG+ FB1017 | 28.33d | 36.00cd | 51.99c | 18.72cd | 51.98d | 19.77de |
| CG+ FB1017 | 28.33d | 35.00ef | 47.29d | 22.49b | 47.81e | 24.01c |
| FS (control) | 26.67e | 34.67f | 61.77a | 29.03a | 66.23a | 32.69a |
| Mean | 29.96 | 35.96 | 47.35 | 19.58 | 47.08 | 22.83 |
| LSD | 0.726** | 0.726** | 2.78** | 2.04** | 2.19** | 2.96** |
| CV (%) | 1.399 | 1.17 | 3.39 | 6.02 | 2.69 | 7.49 |

Table 1. Black root rot disease onset, incidence, and severity

Where: DDO: Days to disease onset, DI: Disease incidence, DS: Disease severity, FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1017, FB1035: Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobium strain FB1017, FB1017: Rhizobium strain FB1017, **- significant at $p \le 0.001$.

Phenological data

The analysis of variance (ANOVA) revealed that highly significant differences (P< 0.0001) among the treatments on days to 50% emergence and days to 90% maturity at APPRC. Similarly, the ANOVA indicated that highly significant differences (P<0.0001) on days to 90% maturity and significant difference (P<0.005) on days to 50% emergence at HARC (Table 2). The results showed that days to emergence were varying over locations. This difference can be explained by due to variation in sowing depth, land preparation and environment influences, which affected the germination and emergence of the seedlings (Pal, 2004). The plots treated with *Rhizobium* strain FB 1035 and *Rhizobium* strainFB1017 were found to decrease days to emergence while, the plot treated with the treatment Dual gold + *Rhizobium* strain FB1035 took longer days to seedling emergence at APPRC and HARC, respectively.

Statistically, significant (P<0.05) difference was observed for days to 90% maturity among the treatments at both locations APPRC and HARC (Table 2). The longest days to 90% maturity i.e. 136 and 145 days were recorded from plots treated with Codal Gold alone, Codal Gold with *Rhizobium* strain FB 1017 and control at APPRC, and Dual Gold alone and Codal Gold with *Rhizobium* strain FB1017at HARC, respectively. However, the plot treated with *Rhizobium* strain FB1017 and Dual gold + *Rhizobium* strain FB1017 reached to 90% maturity relatively in a shorter period of 134 and 143 days at APPRC and HARC, respectively. The reduction in days to maturity at APPRC might be due to the variation in the environmental conditions and disease pressure in the growing season. Thus, the variability in attaining maturity for the treatments might be attributed to environmental condition and the effect of the disease (Wulita, 2015).

| Treatment | Days to 50% | | Days to 50 | Days to 50% flowering | | % maturity |
|--------------|-------------|---------|------------|-----------------------|-----------|------------|
| | emergence | | | | | |
| | APPRC | HARC | APPRC | HARC | APPRC | HARC |
| FB1017 | 6.00bc | 8.33c | 49.67ab | 56ab | 134.33d | 143.67bc |
| FB1035 | 5.66c | 9abc | 49.00b | 55.67b | 134.67cd | 143.67bc |
| CG+ FB1035 | 5.67c | 8.67bc | 49.00b | 56.33ab | 134.67cd | 144.67ab |
| DG+ FB1035 | 6.33bc | 10a | 49.33ab | 56ab | 135.33abc | 144abc |
| DG | 6.67a | 10a | 49.00b | 57a | 135.00bcd | 145a |
| CG | 6.00bc | 8.35c | 49.33ab | 57a | 136.00a | 144abc |
| DG+ FB1017 | 8.67a | 9.33abc | 50.00a | 56.33ab | 135.67ab | 143.33c |
| CG+ FB1017 | 6.00bc | 8.67bc | 50.00a2 | 56.33ab | 136.00a | 145a |
| FS (control) | 6.33bc | 9.67ab | 49.33ab | 56.67ab | 136.00a | 144.33abc |
| Mean | 6.37 | 9.11 | 49.407 | 56.37 | 135.3 | 144.19 |
| LSD | 0.86** | 1.21* | 0.91 ns | 1.13 ns | 0.97** | 1.09** |
| CV (%) | 7.78 | 7.65 | 1.06 | 1.16 | 0.41 | 0.44 |

Table 2. Number of days to 50% emergence, days to 50% flowering and days to 90% maturity

Where: FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1017, FB1035: Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobiumstrain FB1017, FB1017: Rhizobium strain FB1017, ns: non-significant, *- significant at 5%, **Significant at p≤0.001.

Growth parameters

The analysis of variance (ANOVA) revealed significant plant height differences among treatments ($P \le 0.001$) at APPRC and ($P \le 0.05$) at HARC (Table 3). At Ambo the tallest plant height measurement was recorded from the plots treated with *Rhizobium* strain FB1035, CG + *Rhizobium* strain FB1035 and *Rhizobium* strain FB1017. This result was similar to the report of Sameh *et al.*, (2017) who found significantly higher plant height (127–134 cm) from plants inoculated with *Rhizobium* strains NGB-FR62, 126 and 128, than un-inoculated control plants. Sharma *et al.* (2000) also reported that the significant effect of seed inoculation on plant height and biomass dry mater compared to control. The shortest (60.93 cm) plant height was recorded from the control plot. This might be due to the presence of severe *Fusarium* root rot on the control plot. At Holetta, the tallest (118 cm) plant height was recorded from the control plot (Table 3). The mean plant height recorded at HARC (111.11cm) was greater than the mean plant height recorded at APPRC (76.822 cm). This might be due to the presence of high inoculum of *Fusarium solani* at APPRC field.

The data of stand count at seedling growth and at harvest stage from both locations (Table 3) showed highly significant (P<0.001) difference among treatments. The highest number of stand count at seedlingi.e. 158.7 and at harvest 117.3 were recorded from the plots treated with the treatments *Rhizobium* strains FB1035 at APPRC and 154.7 and 119.7 were recorded from plots treated with *Rhizobium* strains FB1017 and *Rhizobium* strains FB1035 at HARC for stand count at seedling stage and at harvest stage respectively. The lowest number of seedlings i.e. 147.3 and 144.0 were recorded from the control plot, plots treated with the herbicide Dual gold at APPRC and HARC, respectively, and smallest stand number at harvest (86.00 and 102.00) was recorded from the control treatment at the two locations APPRC and HARC, respectively.

| Treatment | PH (cm) | | S | SCS | | бCH |
|--------------|---------|----------|----------|-----------|----------|---------|
| | APPRC | HARC | APPRC | HARC | APPRC | HARC |
| FB1017 | 86.467a | 104.33d | 156ab | 154.67a | 107bc | 116b |
| FB1035 | 86.533a | 113.33ab | 158.67a | 150bc | 117.33a | 119.67a |
| CG+ FB 1035 | 86.53a | 118a | 153.33bc | 148.33cd | 113ab | 114bc |
| DG+ FB 1035 | 74.4b | 114.67ab | 156.33ab | 147.33cde | 112.33ab | 112.67c |
| DG | 66.47c | 111bc | 151.33c | 144e | 106bc | 112.67c |
| CG | 76.73b | 115.33ab | 156.33ab | 152.33ab | 103.33cd | 115bc |
| DG+ FB 1017 | 68.53c | 107cd | 155.67ab | 146.33cde | 96.33d | 105.33e |
| CG+ FB 1017 | 73.73b | 113ab | 155.33bc | 148.33cd | 98.67cd | 108.67d |
| FS (control) | 60.93c | 103.33d | 147.33d | 144.67de | 86e | 102f |
| Mean | 75.64 | 111.11 | 154.48 | 148.44 | 104.44 | 111.78 |
| LSD | 4.74** | 5.06* | 3.6** | 3.82** | 8.45** | 3.13** |
| CV (%) | 3.62 | 2.63 | 1.35 | 1.49 | 4.67 | 1.62 |

Table 3. Effects of treatments on plant height stand count at seedling and stand count at harvest

PH: plant height, SCS: stand count at seedling, SCH: stand count at harvest, FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1017, FB1035: Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobium strain FB1017, FB1017: Rhizobium strain FB1017, *-significant at 5%, **- Significant at $p \le 0.001$.

Yield and yield components

Rhizobium strains application significantly affect the number of pods per plant compared to non-inoculated treatments at the two locations APPRC and HARC. Significant ($P \le 0.05$) variations were observed among the treatments on number of pods per plant and on number of seeds per pod at both locations (Table 4). In this study, the highest number of pods per plant was recorded from plot treated with *Rhizobium* strain FB1035 at APPRC and from plot treated with *Rhizobium* strain FB1035 at HARC. The highest number of seeds per pod was recorded from plot treated with *Rhizobium* strain FB1035 at HARC. The highest number of seeds per pod was recorded from plot treated with *Rhizobium* strain strain FB1035 at HARC. The highest number of seeds per pod was recorded from plot treated with *Rhizobium* strains alone, plot treated with Codal gold and Dual gold with *Rhizobium* strain FB1017 at HARC. This is in agreement with the study by Rodriguez-Navaro *et al.* (2000) observed significantly increase in pod numbers per plant among different bean cultivars inoculated by *Rhizobium* strain. Argaw and Tsigie (2015) also reported that the inoculation of *Rhizobium* increased the seed number of common bean. The lowest number of pods per plant and number of seeds per pod (4.7 and 1.83) at APPRC and (5.0 and 2) at HARC were obtained from the control plot, respectively (Table 4).

Analysis of variance also showed highly significant (P<0.001) statistical difference among the treatments at both study fields, on viable nodule count and non-viable nodule count. The highest number of viable nodules (449 and 467.7) was obtained from plots treated with *Rhizobium* strain FB1035 and the lowest number of viable nodules (242.67 and 270.67) was observed from the control plots both at APPRC and HARC, respectively (Table 4). These finding is in agreement with Habtemichial *et al.* (2007) who found that *Rhizobium* inoculation increased the nodule number and fresh weight of faba bean in north Ethiopia by 53% and 95% over un-inoculated control plots, respectively. The current result also in line with the study conducted on lentil by Beshur *et al.*, (2015) observed that *Rhizobial* inoculation increase in nodule number and dry weight. Similarly, Mehrpouyan (2011) reported significant increase in nodule number and dry weight in common bean cultivars when inoculated with *Rhizobium leguminosarum* strain Rb117.

The lowest (286 and 310) viable nodule count next to the control plot was recorded on plots treated with herbicide Codal gold at both APPRC and HARC, respectively. This is in line with the finding of Khan *et al.*, (2006) who stated that the common use of herbicides may negatively affect nodulation either directly by affecting *Rhizobium*, or indirectly by reducing photosynthate allocation to the nodules for N fixation, or by restricting root growth and hence the number of root sites available for infection. Also other studies showed that pendimethalin when applied under field conditions significantly decreased nodulation on soybean (Vaziritabar *et al.*2014).

The largest number of non-viable nodule (40 and 39.7) was recorded from the plot treated with *Rhizobium* strain FB1017. Sheng (1993) reported that besides fixation of atmospheric nitrogen, the nodulation effect of rhizobial isolates is due to the production of plant growth regulators such as Auxins and cytokinins like substance. The smallest number of non viable nodule (20.7 and 25.7) was recorded from the plots treated with Codal gold + *Rhizobium* strain FB1017 at the two locations, APPRC and HARC, respectively.

| [65] |
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|------|

| Treatment | N | P | NS | SP | VNC | | NNC | |
|--------------|--------|---------|---------|-------|---------|----------|---------|---------|
| | APPRC | HARC | APPRC | HARC | APPRC | HARC | APPRC | HARC |
| FB1017 | 10.67b | 7.67ab | 2.67ab | 3a | 445.67a | 457.33ab | 40a | 39.67a |
| FB1035 | 14.67a | 8.33a | 3a | 3a | 449a | 467.67a | 26d | 31.33ab |
| CG + FB1035 | 9.33b | 7abc | 2.33abc | 2.3b | 443.33b | 447.67b | 31c | 27.33b |
| DG+ FB 1035 | 7.67c | 8.33a | 2.33abc | 2.67a | 405c | 417.33c | 30.33c | 30.67ab |
| DG | 10.33b | 6.33bcd | 2.33abc | 2.67a | 304.33e | 327.33e | 22.67ef | 32.67ab |
| CG | 9.33b | 7abc | 1.92c | 3a | 286f | 310f | 22.67ef | 27.33b |
| DG+ FB 1017 | 6.67cd | 6cd | 2bc | 3a | 360.67d | 368d | 33.67b | 29b |
| CG+ FB 1017 | 5.67de | 7.33abc | 2bc | 2.67a | 310e | 320ef | 20.67f | 25.67b |
| FS (control) | 4.67e | 5d | 1.83c | 2b | 242.67g | 270.67g | 24.33de | 26.67b |
| Mean | 8.78 | 7 | 2.27 | 2.67 | 359.63 | 376.22 | 27.93 | 30.04 |
| LSD | 1.567* | 1.61* | 0.69* | 0.58* | 7.36** | 16.36** | 2.40** | 9.99** |
| CV (%) | 10.31 | 13.26 | 17.79 | 12.5 | 1.18 | 2.51 | 4.97 | 19.22 |

Table 4. Yield and yield components of faba bean on different treatmentsduring the main cropping season, 2016

Where: NPP: number of pods per plant, NSP: number of seeds per pod, VNC: viable nodule count, NNC: non-viable nodule count, FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold, + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1035; Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobium strain FB1017, FB1017: Rhizobium strain FB1035, CG + Significant at $p \le 0.001$.

Thousand seed weight and total grain yield

Analysis of variance results revealed the presence of highly significant ($P \le 0.001$) differences at APPRC and significant ($P \le 0.05$) variation at HARC among the treatments for thousand seed weights and highly significant ($P \le 0.001$) variation among the treatments for grain yield at both locations (Table 5). At APPRC and HARC, the highest thousand seed weight of 541.23 and 545.1 g were obtained from the plots treated with *Rhizobium* strain FB1035 and the lowest thousand seed weight of 400.63 and 477.27 g were recorded from the control plots, respectively. Elsheikh and Elzidany (1997) reported that *Rhizobium* inoculation significantly increased hundred seed weight of faba bean. Similarly, Kazemi *et al.* (2005) reported that seed inoculation by *Rhizobial* bacteria significantly increased the number of pods and seed per plant, thousand grain weights and finally the yield of soybean.

The highest (2.26 and 3.64 t/ha) grain yield were also obtained from the plots treated with *Rhizobium* strain FB1035. In this line, Abbasi *et al.* (2010) reported that soybean seed yield quadratically increased by *Rhizobium* inoculation and phosphorous application. Elsheikh and Elzidany (1997) also found that *Rhizobium* inoculation and 40 kg N ha⁻¹ application of inorganic N gave a comparable amount of grain yield increase. However, the lowest grain yields of 1.63and 2.99 t/ha was observed from the control plots at APPRC and HARC, respectively. The production of faba bean is constrained by several yield-limiting factors, of which diseases are the main factors (Eshetu *et al.*, 2013). The production is threatened by black root rot caused by *Fusarium solani* (Bogale *et al.*, 2009).

| [66] |
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|------|

| Treatment | TSV | V (g) | TGY (t | ha-1) |
|--------------|----------|----------|--------|--------|
| | APPRC | HARC | APPRC | HARC |
| FB1017 | 514.17ab | 529.33ab | 2.25b | 3.59b |
| FB1035 | 541.23a | 545.1a | 2.26a | 3.64a |
| CG+ FB1035 | 491.53bc | 506.17bc | 2.01e | 3.3e |
| DG+ FB1035 | 500.3bc | 514.57ab | 1.95f | 3.28f |
| DG | 479.47c | 497.63bc | 2.11d | 3.43d |
| CG | 486.3bc | 503.73bc | 2.16c | 3.46c |
| DG+ FB1017 | 437.23d | 479.17c | 1.85g | 3.22g |
| CG+ FB1017 | 424.43de | 498.37bc | 1.78h | 3.18h |
| FS (control) | 400.63e | 477.27c | 1.63i | 2.99i |
| Mean | 475.03 | 505.70 | 1.62 | 2.72 |
| LSD | 2.83** | 3.41* | 0.23** | 0.45** |
| CV (%) | 3.45 | 3.33 | 0.52 | 0.95 |

Table 5.Yield and thousand seed weights of faba bean tested at APPRC and HARC during the main cropping season, 2016

Where: TSW: thousand seed weight, TGY: total grain yield, FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1017, FB1035: Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobium strain FB1017, FB1017: Rhizobium strain FB1017, *- significant at 5%, **- Significant at $P \le 0.001$.

Relative yield loss

Grain yield losses and thousand seeds weight losses were computed relative to the average grain yield and thousand seeds weight of plots with the maximum protection against the disease (Plots with lowest disease severity in the treatments). The highest relative grain yield losses (39.04% and 21.6%) and thousand seeds weight losses(35.1% and 14.21%) were occurred on control plots.Complete crop losses could occur in severe infection conditions and when favorable conditions prevail for the pathogen. In farmers' fields, a loss of about 45% was estimated due to this disease (Nigussie *et al.*, 2008). The lowest yield losses(0.75% and 1.3%) and thousand seed weight losses(5.26% and 2.98%) were recorded on plots treated with *Rhizobium* strain 1017 at APPRC and HARC, respectively (Table 6). This indicated thatmany species of *Rhizobium* are inhibiting significantly the growth of pathogenic fungi and reduce yield loss due to diseases (Estevez de Jensen *et al.*, 2002).

| Table 6.Yield loss of faba bean due to black root rot disease at APPRC and HARC on different |
|--|
| treatments during the main cropping season, 2016. |

A) APPRC

| / ///////////////////////////////////// | | | | | | | |
|---|-------------------------------|-----------|---------|----------|----------|----------|--|
| Treatment | Yield and relative yield loss | | | | | | |
| | Yield (t/ha) | RYL (ton) | RYL (%) | TSW | loss (g) | Loss (%) | |
| FB1035 | 2.26a | 0 | 0 | 541.23a | 0 | 0 | |
| FB1017 | 2.25b | 0.017 | 0.75 | 514.17ab | 27.06 | 5.26 | |
| CG | 2.16c | 0.095 | 4.4 | 486.3bc | 54.93 | 11.3 | |
| DG | 2.11d | 0.015 | 7.18 | 479.47c | 61.76 | 12.88 | |
| CG+ FB1035 | 2.01e | 0.025 | 12.61 | 491.53bc | 49.7 | 10.11 | |
| DG+ FB1035 | 1.95f | 0.032 | 16.28 | 500.3bc | 40.93 | 8.18 | |
| DG+ FB1017 | 1.85g | 0.042 | 22.53 | 437.23d | 104 | 23.79 | |
| CG+ FB1017 | 1.78h | 0.048 | 27.19 | 424.43de | 116.8 | 27.52 | |
| FS (control) | 1.63i | 0.064 | 39.04 | 400.63e | 140.6 | 35.1 | |

B) HARC

| / | | | | | | | | |
|-------------|-------------------------------|-----------|---------|----------|----------|----------|--|--|
| Treatment | Yield and relative yield loss | | | | | | | |
| | Yield (t/ha) | RYL (ton) | RYL (%) | TSW | loss (g) | Loss (%) | | |
| FB1035 | 3.64a | 0 | 0 | 545.1a | 0 | 0 | | |
| FB1017 | 3.59b | 0.047 | 1.3 | 529.33ab | 15.77 | 2.98 | | |
| CG | 3.46c | 0.177 | 5.11 | 503.73bc | 41.37 | 8.21 | | |
| DG | 3.43d | 0.212 | 6.2 | 497.63bc | 47.47 | 9.54 | | |
| CG+ FB1035 | 3.3e | 0.339 | 10.28 | 506.17bc | 38.93 | 7.69 | | |
| DG+ FB1035 | 3.28f | 0.361 | 11.01 | 514.57ab | 30.53 | 5.93 | | |
| DG+ FB1017 | 3.22g | 0.415 | 12.87 | 479.17c | 65.93 | 13.76 | | |
| CG+ FB1017 | 3.18h | 0.455 | 14.31 | 498.37bc | 46.73 | 9.38 | | |
| FS(control) | 2.99i | 0.646 | 21.6 | 477.27c | 67.83 | 14.21 | | |

RYL: Relative yield loss, TSW: thousand seed weight, FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1017, FB1035: Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobium strain FB1017, FB1017: Rhizobium strain FB1017, FB1017; Rhizobium strain FB1017; Rhizobium stra

Correlation between yield and disease parameters Correlation

Correlation analysis showed that black root rot severity had strongly negative correlation (r = -0.801 and r = -0.811) with grain yields at APPRC and HARC, respectively (Table 7). The correlation coefficient between disease severity and number of pods per plant (r = -0.617 and r = -0.77) were not significant (P> 0.05) at APPRC and HARC, respectively. A strong and negative association with correlation coefficient of r = -0.805 and r = -0.614 were observed between the number of seeds per pod and disease severity values at APPRC and HARC, respectively.

Ahighly and positive correlation coefficient was recorded between yield and number of seeds per pod were r = 0.806 and r = 0.583 at APPRC and HARC, respectively (Table 7). Similarly, the thousand seed weight showed highly significant (P< 0.01) correlation with yield and disease severity at HARC and significant (p< 0.05) with disease severity at APPRC. A strong and negative association with correlation coefficient of r = -0.759 and r = -0.801 values were observed between thousand seed weight and disease severity (Table 9).

| Table 7. Correlation coefficients (r) among the traits of black root rot disease yield and yield |
|--|
| components of faba bean evaluated at both locations in 2016 during the main |
| cropping season |

| A) APPR | С |
|---------|---|
|---------|---|

| Parameter | NPP | NSP | TSW (g) | TGY (kg/ha) | DS (%) |
|------------|----------|----------|---------|-------------|--------|
| NPP | 1 | | | | |
| NSP | 0.879** | 1 | | | |
| TSW (g) | 0.909** | 0.861** | 1 | | |
| TGY (t/ha) | 0.955** | 0.806** | 0.899** | 1 | |
| DS (%) | -0.617ns | -0.805** | -0.759* | -0.801** | 1 |

B) HARC

| Parameter | NPP | NSP | TSW (g) | TGY (kg/ha) | DS (%) |
|------------|---------|----------|----------|-------------|--------|
| NPP | 1 | | | | |
| NSP | 0.468ns | 1 | | | |
| TSW (g) | 0.871** | 0.422ns | 1 | | |
| TGY (t/ha) | 0.582ns | 0.583ns | 0.833** | 1 | |
| DS (%) | -0.77ns | -0.614ns | -0.801** | -0.811** | 1 |

Where: NPP: number of Pod per plant, NSP: number of Seed per pod, TSW: thousand seed weight, TGY: total grain yield, DS: disease severity,*: refers to mean square values significant at α =0.05, **: refers to mean square values not significant at α =0.05

Regression

Linear regression model fitted to disease severity at APPRC and HARC as predictor and grain yield as dependent variable were regressed to estimate the yield loss due to the disease. Regression analysis of disease severity with yield revealed significant ($P \le 0.001$) relationship. Thus, the linear regression indicated for every one unit increases in severity (%), there is a corresponding 0.02and 0.043 t/ha loss in yield of faba bean at APPRC and HARC, respectively (Figure 1). The regression equation: thousand seed weight (g) = -2.696X + 601.97, ($R^2 = 57.7\%$, P = 0.000) and thousand seed weight (g) = -2.377X + 559.97, ($R^2 = 44.8\%$, P = 0.000) demonstrated reduction of about 2.696 and 2.337 gram in thousand seed weight with the increase of 1% severity at APPRC and HARC, respectively (Figure 2).

As the severity increased there was a decreasing trend in yield and thousand seed weight, and declined towards zero asymptote, which indicated the reverse relation between yield and severity. In addition, the R^2 value of 36.4 and 12.4% indicated that these amounts of the variation in yields were explained by disease severity at APPRC and HARC, respectively.

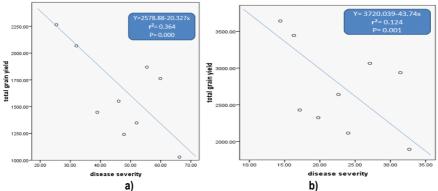


Figure 1: Linear regression of grain yield and disease severity at a) APPRC and b) HARC

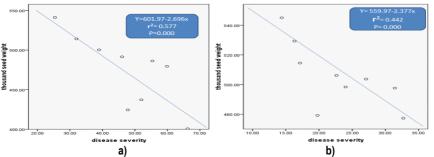


Figure 2: Linear regression of thousand seed weight and disease severity at a) APPRC and b) HARC

Cost - benefit analysis

Net benefit variations were obtained among treatments (herbicides and *Rhizobium* strains) at APPRC and HARC. Partial budget analysis was calculated based on cost of variable inputs during the main cropping season of the year 2016 and net benefit was estimated based on mean values of local market price when most of farmers supplied the produce to market.

Partial budget analysis revealed that the highest (birr 35030per hectare and birr56420per hectare) total gross yield benefit were obtained from plots treated with *Rhizobium* strain FB1035 at APPRC and HARC, respectively (Table 8). On the other hand, the lowest total gross yield (birr25265and birr46345per hectare) were obtained from the control plots at APPRC and HARC, respectively.

The highest (birr34870 and 56260per hectare)net profit was obtained from plots treated with *Rhizobium* strain FB1035, followed by birr34715 and 55485 per hectareobtained from plots treated with *Rhizobium* strain FB1017, whereas the lowest birr25265 and 46345 per hectare net benefit was obtained from the control plots at APPRC and HARC, respectively.

Generally, the highest values of marginal rate return of 60.03125 and 61.96875% were obtained from plots treated with *Rhizobium* strain FB1035, followed by 59.0625 and 57.125% obtained from plots treated with *Rhizobium* strain FB1017 at APPRC and HARC, respectively.

In this study, the application of *Rhizobium* strain FB1035 and *Rhizobium* strain FB1017 are found best to reduce black root rot disease of faba bean and to increase the yield. The economic analysis on these strains also showed more profit than the other treatments at current market prices.

Table 8. Partial budget analysis for herbicides and *Rhizobium* strains treated for the management of the faba bean black root root disease at APPRC and HARC.

| A) | APPRC |
|----|-------|
|----|-------|

| Treatment | General cost benefit | | | | | | |
|--------------|----------------------|-----------|----------|-----------|---------------|----------------|----------|
| | (A) adj. | (B) price | (C) sale | (D) | (E) net | (F) | MRR |
| | yield(t/ha | (birr/ t | revenue | marginal | profit (birr) | marginal | (F/D)(%) |
| |) | | (A*B) | cost | (C-D) | benefit | |
| | | | | (birr/ha) | | (birr) | |
| FB1035 | 2.26 | 15500 | 35030 | 160 | 34870 | 9605 | 60.03125 |
| FB1017 | 2.25 | 15500 | 34875 | 160 | 34715 | 9450 | 59.0625 |
| CG | 2.17 | 15500 | 33635 | 750 | 32885 | 7620 | 10.16 |
| DG | 2.11 | 15500 | 32705 | 600 | 32105 | 6840 | 11.4 |
| CG+ FB1035 | 2.01 | 15500 | 31155 | 910 | 30245 | 4980 | 5.472527 |
| DG+ FB1035 | 1.94 | 15500 | 30070 | 760 | 29310 | 4045 | 5.322368 |
| DG+ FB1017 | 1.85 | 15500 | 28675 | 760 | 27915 | 2650 | 3.486842 |
| CG+ FB1017 | 1.78 | 15500 | 27590 | 910 | 26680 | 1415 | 1.554945 |
| FS (control) | 1.63 | 15500 | 25265 | 0 | 25265 | 0 | 0 |
| B) HARC | | | | | | | |
| Treatment | General cost benefit | | | | | | |
| | (A) adj. | (B) price | (C) | (D) | (E) net | (F) | MRR |
| | yield(t/ha | (birr/ t | sale | marginal | profit (birr) | marginal | (F/D)(%) |
| |) | | revenue | cost | (C-D) | benefit (birr) | |
| | | | (A*B) | (birr/ha) | | | |
| FB1035 | 3.64 | 15500 | 56420 | 160 | 56260 | 9915 | 61.96875 |
| FB1017 | 3.59 | 15500 | 55645 | 160 | 55485 | 9140 | 57.125 |
| CG | 3.46 | 15500 | 53630 | 750 | 52880 | 6535 | 8.713333 |
| DG | 3.43 | 15500 | 53165 | 600 | 52565 | 6220 | 10.36667 |
| CG+ FB1035 | 3.3 | 15500 | 51150 | 910 | 50240 | 3895 | 4.28022 |
| DG+ FB1035 | 3.28 | 15500 | 50840 | 760 | 50080 | 3735 | 4.914474 |
| DG+ FB1017 | 3.23 | 15500 | 50065 | 760 | 49305 | 2960 | 3.894737 |
| CG+ FB1017 | 3.19 | 15500 | 49445 | 910 | 48535 | 2190 | 2.406593 |
| FS (control) | 2.99 | 15500 | 46345 | 0 | 46345 | 0 | 0 |

MRR: marginal rate return, FS: Fusariumsolani, DG + FB1035: Dual gold + Rhizobium strain FB1035, CG + FB1035: Codal gold + Rhizobium strain FB1035, DG: Dual gold, CG: Codal gold, DG + FB1017: Dual gold + Rhizobium strain FB1017, FB1035: Rhizobium strain FB1035, CG + FB1017: Codal gold + Rhizobium strain FB1017, FB1017: Rhizobium strain FB1017,

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