Controlling Foliar Blight of Wheat through Nutrient Management and Varietal Selection

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ABSTRACT

Helminthosporium leaf blight, a complex of spot blotch caused by Bipolaris sorokiniana and tan spot caused by Pyrenophora tritici-repentis, is one of the most important foliar diseases of wheat in Nepal. It appears in almost all wheat growing areas and causes severe yield loss every year. A study was conducted at Regional Agriculture Research Station (RARS), Tarahara, Sunsari during 2004-05 and 2005-06 wheat growing seasons to elucidate role of nitrogen in wheat genotypes for management of the disease. Field experiment was laid out on split plot design with three replications. Four doses of nitrogen in six different promising genotypes were tested. Nitrogen levels higher than 50 kg ha⁻¹ significantly reduced disease severity and increased grain yield in all genotypes but there was no significant differences in grain yield in the first year. In the second year, grain yield difference among the genotypes was significant. Area under disease progress curve (AUDPC) was not significant between two doses 100 and 150 kg ha⁻¹. The wheat genotypes showed different reactions to disease. Genotype BL 2047 had the lowest incidence of disease followed by BL 1887, whereas BL 2217 had the highest incidence of the disease. Genotype BL 2196 produced the highest grain yield (2172 kg ha⁻¹) and the lowest grain yield was obtained in Bhrikuti followed by BL 2089. These results suggested that fertilizer should be applied in soil at balanced dose 100:50:50 N:P₂O₅:K₂O kg ha⁻¹. Growing relatively resistant genotypes with the balance dose of fertilizers can reduce foliar blight severity in wheat.

Key words: AUDPC, Bipolaris sorokiniana, disease resistance, grain yield, Helminthosporium leaf blight, Pyrenophora tritici-repentis

INTRODUCTION

Wheat occupies the third position after rice and maize in Nepal representing 22% of total cultivated area. Major area of wheat production is covered by eastern terai region of Nepal. Successful production for wheat is constrained by several biotic and abiotic stresses. Among them, Helminthosporium leaf blight (HLB), a complex foliar disease of wheat caused by Bipolaris sorokiniana (Sacc.) Shoem. and Pyrenophora tritici-repentis (Died) is a serious constraint. It is considered a disease of major importance owing to its potential to cause yield loss. Globally an
estimation of 25 million ha of wheat land is affected by the disease (van Ginkel and Rajaram, 1998). In Nepal the disease is widely distributed throughout the wheat growing areas particularly in terai (NWDP 1976). The magnitude of yield loss due to HLB may vary among locations and years (Mahto 1995). In South Asia, yield loss due to HLB has been reported to 20-30% in farmers’ field and experiment stations (Dubin and Bimb 1991, Duveiller and Gilchrist 1994, Saari 1998, Duveiller et al 2005). In years when rain occurs late in the crop cycle, especially during grain filling, complete crop loss has been observed (Saunders 1988). Selection and breeding for resistant cultivars is the main disease management strategy for a sustainable agriculture. Duveiller and Gilchrist (1994) and Dubin and Rajaram (1996) reported several sources of spot blotch resistance in wheat but high yielding genotypes resistant to the disease is not yet available. Most of the improved and recommended wheat cultivars in Nepal are severely attacked by foliar blight (Shrestha et al 1998).

Crop management practices affect on the development of the disease epidemics. In the humid sub-tropics of South Asia, there is evidence of stress conditions, which favor foliar blight (Dubin and Bimb 1994). Factors such as minimum tillage or surface seeding, irrigation, late planting, or low soil fertility may be responsible for higher foliar blight severity in the wheat-based cropping systems of the Indo-Gangetic plains (Sharma and Duveiller 2003). Lower disease severity with higher nitrogen application was reported by Chaurasia and Duveiller (2006) and Sharma and Duveiller (2004). However, Singh et al (1998) reported more disease infection on higher nitrogen application.

The objective of the present study was to find out the influence of nitrogen in different wheat genotypes on the severity of foliar blight.

MATERIALS AND METHODS

A split plot experiment with three replications was conducted at Regional Agriculture Research station, Tarahara Sunsari, Nepal during wheat season of 2004-05 and 2005-06 where climatic and agro-ecological conditions favor rapid development of foliar blight pathogens. The main plot treatments were four levels of nitrogen: i) 0:50:50 N:P₂O₅:K₂O kg ha⁻¹, ii) 50:50:50 N:P₂O₅:K₂O kg ha⁻¹, iii) 100:50:50 N:P₂O₅:K₂O kg ha⁻¹, and iv) 150:50:50 N:P₂O₅:K₂O kg ha⁻¹. The sub plot treatments included most promising wheat genotypes: BL 2047, BL2089, BL2196, BL 2217, Gautam, and Bhrikuti. The individual plot size was 4×3-m and consisted of twelve rows with 25cm spacing. Seed was sown using standard seed rate of 120 kg ha⁻¹. Phosphorus and potassium were applied as basal dose. Half of N (for all three levels) was applied at basal dose and remaining half was top-dressed at tillering stage. Three irrigations were given. Plots were kept free from weeds by hand weeding. Daily rainfall, max and min temperature during wheat growing seasons were recorded from wheat emergence to harvest in the two test years.

After anthesis, the peak stage of disease development, disease severity was assessed using the double-digit scale (00-99) developed as a modification of Saari and Prescott’s scale for assessing severity of foliar diseases of wheat (Saari and Prescott 1975, Eyal et al 1987). The first digit (D1) indicates disease progress in canopy height from the ground level and second digit (D2) refers to severity measured based on diseased leaf area. Both D1 and D2 are scored on a scale of 1 to 9. Three individual disease scores per plot were recorded at every 7-days interval. For each evaluation, percent disease severity was estimated using the formula: % severity = (D1/9) × (D2/9) × 100. The area under disease progress curve (AUDPC) was calculated using the percentage disease severity estimations corresponding to the three ratings as outlined by Shaner and Finney (1977).

Yield (kg/ha) and 1000-grain weight were measured after harvesting all plots. The data were analyzed using computer software MSTATC and MINITAB 11 for windows. Statistical analysis
includes ANOVA test, mean separation based on Duncan’s multiple range test, regression analysis for AUDPC and the grain yield over different levels of nitrogen.

RESULTS AND DISCUSSION

Weather conditions and disease severity
There was high pressure of disease in both years. However, the disease severity was significantly higher (p < 0.05) in the second year (Table 1). The difference in the disease might be due to climatic variations in these two years (Figure 1). Higher temperature in early stage of the crop cycle was observed in the second year. This result suggested the influence of temperature on disease development. Naitao and Yousan (1998) reported that Bipolaris sorokiniana resistance traits in wheat are polygenic and greatly affected by climatic factors such as rainfall, humidity and temperature. Similarly, Gilbert et al (1998) reported that humid environment resulting from high temperature and high rainfall promoted the development of the disease.

Influence of nitrogen (N) levels on disease severity and wheat yield
N levels had a significant effect (p < 0.01) on AUDPC in both years (Table 1). Among four levels of N tested, the mean AUDPC value was higher for N levels 0 and 50 kg ha\(^{-1}\) as compared to 100 kg ha\(^{-1}\) and 150 kg ha\(^{-1}\). Such variation in mean values for disease severity among the four levels of N confirmed its role in reducing foliar blight infection in wheat. Regression analysis showed a significant negative linear relation of AUDPC, \(Y = 657.458 - 6.7583X\), R-Sq = 0.048** over dose of nitrogen application (0-150 kg/ha; Figure 2A). This result indicated that foliar blight severity was lowered with increased dose of N.

Mean values for grain yield differed in the test years. Average grain yield was lower in 2005 than in 2004. The lower yield with more disease in the second year suggested the effect of the disease in wheat yield reductions. The effect of N level was significant for grain yield and thousand kernel weight (TKW) in both the years. The mean grain yield was high for N levels of 100 and 150 kg ha\(^{-1}\) as compared with 0 and 50 kg ha\(^{-1}\) but the difference between the first two levels was insignificant (Table 1). Similarly AUDPC value was also not significantly different between these two N levels (100 and 150 kg ha\(^{-1}\)). Regression analysis showed that grain yield was increased as a function of nitrogen level, \(Y = 521.168 + 716.125X\), R-Sq = 0.667** (Figure 2). There was sharp\(100\) and 150 kg ha\(^{-1}\) increment of grain yield over N application up to the dose 100 kg ha\(^{-1}\) in all the genotypes.
but after the level, the rate of increment decreased (Figure 3). This result suggested that nitrogen level more than 100 kg ha\(^{-1}\) was not significantly effective to lower the disease severity and to increase the grain yield. There was no specific trend for TKW over N levels but average value was the lowest in 150 kg ha\(^{-1}\).

The results suggested that N levels had a significant influence on foliar blight development, grain yield and TKW. It was observed that low and imbalanced use of nitrogenous fertilizer increased severity of the disease. Krupinsky et al (1998) noted more leaf necrosis and chlorosis caused by *Pyrenophora tritici-repentis* in the plots with no additional nitrogen and/or low level of nitrogen than higher nitrogen levels. Similarly higher tan spot severity in N deficient condition was observed by Fernandez et al (1998). In agreement with the results of Sharma and Duveiller (2004) and Chaurasia and Duveiller (2006) our results clearly suggested that higher dose of nitrogen had significant effect on reducing the disease and thereby increase the grain yield. This finding differs from the observation by Singh et al (1998) who reported higher disease infection with higher level of nitrogen.

Table 1. Means values for disease and agronomic variables at four level of nitrogen application and of six genotypes recorded in 2004 and 2005 wheat growing seasons at Tarahara, Sunsari, Nepal

<table>
<thead>
<tr>
<th>Effect of nitrogen levels</th>
<th>AUDPC</th>
<th>Grain yield, kg/ha</th>
<th>TKW, g</th>
</tr>
</thead>
<tbody>
<tr>
<td>N0</td>
<td>446 a</td>
<td>774 a 610 a</td>
<td>1550 c</td>
</tr>
<tr>
<td>N50</td>
<td>452 a</td>
<td>770 a 611 a</td>
<td>2606 b</td>
</tr>
<tr>
<td>N100</td>
<td>378 b</td>
<td>671 b 525 b</td>
<td>3259 a</td>
</tr>
<tr>
<td>N150</td>
<td>368 b</td>
<td>665 b 517 b</td>
<td>3474 a</td>
</tr>
<tr>
<td>Mean</td>
<td>441</td>
<td>719</td>
<td>2722</td>
</tr>
<tr>
<td>F-test</td>
<td>**</td>
<td>**</td>
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</table>

<table>
<thead>
<tr>
<th>Effect of genotypes</th>
<th>AUDPC</th>
<th>Grain yield, kg/ha</th>
<th>TKW, g</th>
</tr>
</thead>
<tbody>
<tr>
<td>BL 2217</td>
<td>76 c</td>
<td>473 ab 840 a 657 a</td>
<td>2649</td>
</tr>
<tr>
<td>BL 2089</td>
<td>77 c</td>
<td>483 a 774 b 629 b</td>
<td>2682</td>
</tr>
<tr>
<td>Bhrikuti</td>
<td>81 b</td>
<td>438 b 742 b 590 c</td>
<td>2871</td>
</tr>
<tr>
<td>BL 2196</td>
<td>77 c</td>
<td>396 c 777 b 586 c</td>
<td>2581</td>
</tr>
<tr>
<td>BL 1887</td>
<td>80 b</td>
<td>332 d 632 c 482 d</td>
<td>2792</td>
</tr>
<tr>
<td>BL 2047</td>
<td>86 a</td>
<td>346 d 555 d 450 c</td>
<td>2761</td>
</tr>
<tr>
<td>Mean</td>
<td>441</td>
<td>719</td>
<td>2722</td>
</tr>
<tr>
<td>F-test</td>
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AUDPC, Area under disease progress curve, TKW= Thousand kernel weight. Means within a column followed by the same letter do not differ significantly based on least significant difference at P = 0.05.
Figure 2. Regression lines indicating linear negative relation of area under disease progress curve (AUDPC; A) and linear positive relation of grain yield (kg/ha; B) over level of N in wheat grown at Sunsari, Nepal during 2004-05 and 2005-06 wheat growing seasons.

Varietal performance
Analysis of variance explicitly showed highly significant differences among genotypes (P < 0.01) for AUDPC in both the years (Table 1). Interaction between the fertilizer dose and genotypes was insignificant and negative linear trend for AUDPC across nitrogen levels was observed in all the genotypes (Figure 3). Among the tested genotypes no resistance reaction to the disease was observed conforming previous observations of van Ginkel and Rajaram (1998). The six genotypes showed variations in level of resistance to the disease and agronomic traits. Such variation in disease level confirmed genetic differences among the genotypes. BL 2047 showed relatively higher disease resistance characterized by lower AUDPC value than others, and was followed by Gautam. Similarly, BL 2217 had the highest AUDPC value demonstrating the most susceptible to the disease among the tested genotypes (Table 1). The genotypes were significantly different in days to heading (P < 0.01). Among the tested genotypes BL 2047 was relatively late and BL 2217 was early for days to heading (that foliar Table 1). This result suggested blight resistance in wheat genotypes was associated with maturity and confirmed the previous observations by Dubin et al (1998). They reported the best foliar blight resistant wheat genotypes in South Asia were late. A significant negative correlation between the days to heading and AUDPC was found by Mahto (1999).
Figure 3. Changes in grain yield, thousand kernel weight (TKW), and area under disease progress curve (AUDPC) in six wheat genotypes across four level of nitrogen in the 2004 and 2005 wheat seasons at Tarahara, Sunsari, Nepal.

The wheat genotypes differed significantly only in the second year for grain yield (Table 1). In this year, genotype BL 2196 produced the highest yield (2172 kg ha\(^{-1}\)) which was followed by BL 2217 (2049 kg ha\(^{-1}\)). Despite having higher AUDPC value these genotypes produced higher grain yield suggesting they are tolerant to the disease. Among the tested genotypes thousand kernel weight was relatively high in BL 2089 followed by BL 2196.
CONCLUSION

The findings of the study have implications for integrated management of foliar blight (HLB) disease of wheat. Nutrient management in field is very critical to minimize the disease severity. Crop without or with low dose of Nitrogenous fertilizer are vulnerable to severe foliar blight attack. Genotypes BL 2047 and BL 1887 showing relatively resistance, and BL 2196 having tolerance to the disease should be promoted. These three genotypes are potential and be crossed to combine resistance and tolerance to foliar blight along with high yielding progenies. By growing the selected cultivars with balanced dose of fertilizer @ 100:50:50 N:P₂O₅:K₂O kg ha⁻¹, yield loss due to the disease in warm wheat growing climate could be minimized.

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