Seed Quality and Vigour Assessment of Fennel (Foeniculum vulgare L.) Genotypes

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A B S T R A C T

The present experiment was conducted during the 2015-16 and 2016-17 with sixty genotypes of fennel for the assessment of seed quality and vigour. Seeds of different genotypes were collected from different locations of India (Gujarat, Haryana, Rajasthan, Bihar etc.). These diverse genotypes were sown (13 November 2015 for first year and 11 November 2016 for second year) with Augmented block design and seed harvesting (20 May, 2015 for first year and 25 May 2017 for the second year) was done after full maturity. All seeds lots were analyzed in completely randomized design (CRD) with three replication for test weight (g), standard germination (%), seedling length (cm), fresh weight (cm), dry weight per seedling (mg), vigour index I and II. The results revealed significant variability among the sixty genotypes of fennel and HF-116, HF-124, HF-206, HF-207 found superior in terms of seed quality and vigour. Among all the genotypes most promising genotypes for various seed quality parameters were used for carryover seed and also can be used as a breeding material for further breeding of fennel genotypes.

Keywords: Fennel, Seed quality, Vigour, Viability, Variability.

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Introduction

Fennel (Foeniculum vulgare L.) is a biennial medicinal and aromatic plant belonging to the family Apiaceae (Umbelliferaeae). It is a hardy, perennial–umbelliferous herb with yellow flowers and feathery leaves. The flowers are produced in terminal compound umbels. The fruit is a dry seed 4–10 mm long. The seeds of fennel have an active substance, which is called essential oil and most important constituent is anethole that is used in pharmaceutical, food, perfumery and favoring industry (Miraldi, 1999; Sephidkon, 2001). Fennel essential oil possesses valuable antioxidant, and has antibacterial, anticancer and antifungal activity (Lucinewton et al., 2005; El-Alwadi and Esmat, 2010).

Mature fennel fruits are used as flavoring agents in food products such as pickles, liqueurs, bread, pastries and cheese (Zoubiri et al., 2014). Fennel fruits are used in diseases like cholera, nervous disorders, constipation, dysentery and colic pain. Fennel also contains minerals and vitamins like calcium, potassium, sodium, iron, phosphorus, thiamine, riboflavin and vitamin C. By virtue of its finest quality and high vitamin content, demand of export due to its high quality seed is increasing steadily. The quality seed is prerequisite to enhance the production and productivity. Use of quality seeds increased productivity of crop by 15-20% (Sidhawani, 1991). The quality of seed is mainly measured
by its genetic purity and capacity to develop into a healthy plant. Further, due to high value of seed spices, the quality of seeds becomes more important. This is mainly measured by its high genetical and physical purity, free from insect-pest and diseases, high vigour, germination percentage and uniformity in appearance. The advantages of high vigour seed are most often associated with rapid and high rate of emergence and stand establishment. There is also a need to have some more reliable parameters that evaluate the seed quality before it is sown in the field. Therefore, an attempt has been made to evaluate the fennel genotypes for the seed quality and vigour.

**Materials and Methods**

The present experiment was carried out during 2015-16 and 2016-17 at Chaudhary Charan Singh Haryana Agricultural University, Hisar. The field experimental site was located at between 29.15°N latitude 75.69°E longitude with a mean altitude of 215 m above msl. Sixty germplasms of fennel were studied for the experiment. The seeds are collected from different locations of India (Gujarat, Haryana, Rajasthan, Bihar etc.). These diverse genotypes were sown (13 November 2015 for first year and 11 November 2016 for second year) with Augmented block design having four blocks with fifteen entries in each block and having plot size of 3.0 m × 1.0 m with spacing of 50 cm × 20 cm. All recommended agronomic practices were followed timely for successful raising the crop. Seed harvesting (20 May, 2015 for first year and 25 May 2017 for the second year) was done after full maturity and seeds were sun dried for 3 to 4 days in the field. After proper drying, cleaning and attaining the optimum moisture content the seeds were collected and the completely randomized design (CRD) was followed to conduct laboratory testing for the seed quality parameters in the seed testing laboratory, Department of Seed Science and Technology, CCS Haryana Agricultural University, Hisar.

The observations recorded on seed quality parameters were test weight, standard germination, seedling length (cm) and seedling dry weight (mg), vigour index-I and Vigour index-II. Test weight: From each genotype seed lot, 1000 seeds were counted and weight was calculated in gram.

Standard germination (%): Hundred seeds per replication for individual genotypes were placed separately between two layers of moist germination paper (BP) and then kept in seed germinator at 20°C. The final count of normal seedlings was made on the 21st day and expressed as per cent germination.

Seedling length (cm): The seedling length was measured for ten randomly selected normal seedlings taken from three replications of standard germination test and recorded in centimeter. At last, average of ten seedlings was taken for final calculation.

Seedling dry weight (mg): Ten normal seedlings selected for measuring seedling length were further kept in hot air oven for taking dry weight. These are dried at 80°C for 48 h and then seedling dry weight was recorded in milligram. The average weight of ten seedlings was taken for further calculations.

Vigour indices: The vigour indices were calculated according to the following formulae suggested by Abdul Baki and Anderson (1973).

Vigour Index – I: Standard germination (%) x Average seedling length (cm)

Vigour Index – II: Standard germination (%) x Average seedling dry weight (mg)
Results and Discussion

The weight of 1000 seeds denotes the extent of development of seed and is an important yield attribute besides contributing towards yield and quality of the seed. All the genotypes test weight and overall mean of sixty genotypes for the 2015-16 and 2016-17 were presented in fig-1. It is revealed that overall mean during 2015-16(4.77 g) is slightly higher than 2016-17 (4.58). Genotype HF-120 (3.03g) recorded with lightest weight and HF-116 (6.37 g) was found to be heavier. Similarly, Sengupta (2011) in fenugreek, Yadav (2016) in coriander and Tanuj (2014) in fennel assess the different genotypes on the basis of test weight and found variability in test weight of different genotypes. Test weight is an important seed quality test as it represents the sample of crop and will give an indication to compares the seed quality standards. Standard germination test is an acceptable measure of seed quality and it provides information about emergence capacity of lots under favorable conditions (ISTA, 2003). Standard germination percentage demonstrate the significant variations (fig-2) among all the genotypes. The maximum germination (92.01 %) was observed in seeds harvested from the genotype HF-124 followed by HF-202(92.0%) HF-115 (91.55 %), HF-210 (91.33%), HF-125 (91.01 %), and HF-109 (90.0 %) while the minimum germination (84.6 %) was observed in seeds harvested from the genotypes HF-105. The overall mean during 2015-16(88.80 %) is slightly lower than 2016-17 (89.0 %). The results indicating that most of genotypes with more test weight and size of seed bear higher seedling length and seedling dry weight. Large and heavy seeds have a competitive advantage over smaller and light seeds by having higher germination rates and greater nutrient reserves for the young seedlings, which enables the seedlings to grow larger and vigorous to tap resources earlier than their small-seeded counterparts (Flenner, 1983; Milberg et al., 1996 and Soltani et al., 2002). However, a negative relationship between seed size and germination percentage, root and shoot length was also reported by Kaya et al., (2008). Kapoor et al., 2010 reported that seedling growth is the morphogenetic expression of genetic programming. Significant variations were recorded in all the genotypes with respect to seed vigour index-I and II. The maximum value for vigour index-I (1742.99) was registered in seeds extracted from HF-207 and the minimum value for vigour index- I (1090) was recorded in seeds extracted from the genotypes PF-35 (fig-5).
Fig. 1 Test Weight of genotypes (A, B, C) and overall mean of genotype (D)
Fig. 2 Standard germination percentage of genotypes (A, B, C) and overall mean of genotypes (D)
**Fig. 3** Seedling length (cm) of genotypes (A, B, C) and overall mean of genotypes (D)
Fig. 4 Dry weight (mg) of genotypes (A, B, C) and overall mean of genotypes (D)
**Fig. 5** Vigour index-I of genotypes (A, B, C) and overall mean of genotypes (D)
Fig. 6 Vigour index-II of genotypes (A, B, C) and overall mean of genotypes (D)
Similarly, seed vigour index-II was maximum in HF-206(368) and the minimum seed vigour index-II (185.67) was recorded in HF-120 (fig-6). Wani et al., (2013) revealed that seed vigour index is strongly under genetic control and highest heritability accompanied with genetic advance was observed in seed vigour index.

The use of genotype with vigorous seed is pre-requisite for successful adoption, production and improvement of any crop, because the variety with quality seed determines the beneficial effects of other production inputs. The differences in the seed viability and vigour are a function of complex interaction of genetic constitution and environmental factors. Overall, it is concluded that variation in genotypes in respect of different vigour parameters resulted due to physiological, biochemical and genetical differences in genotypes of fennel. As physiological and biochemical differences controlled by genetic constitution of plant, therefore, ultimately plants genetic constitution is responsible for all such differences in genotypes. However, best (HF-116, HF-124, HF-206, HF-207) genotypes can be undertaken as a carryover seed and also seed material for further breeding programme.

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