

Review

Barnyard millet – a potential food and feed crop of future

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With 4 figures and 2 tables

Received May 28, 2014 / Accepted November 12, 2014

Communicated by M. Prasad

Abstract

The two species under genus *Echinochloa*, *E. frumentacea* (Indian barnyard millet) and *E. esculenta* (Japanese barnyard millet), are cultivated for food and fodder by hilly and tribal communities in Asia particularly in India, China and Japan. The crop has wide adaptability and occupies a special place in marginal rainfed areas because of its short life cycle. Although the area under the crop has come down drastically in last 50 years, the crop ability to survive under harsh conditions makes it a better choice during famine years. In the Indian Himalayan region, the crop was traditionally used as a substitute for rice. It has been identified as a suitable choice for climate-resilient agriculture. High nutrient content and antioxidant effects make it to be considered as a functional food crop. Recently, the demand of the crop has increased due to its highly nutritious grains. Thus, it has the potential to provide both food and nutritional security particularly in hills where nutritional deficiencies are in abundance. Despite enormous potential, the crop has not gained the popularity among masses and is still considered as poor man's food. This work therefore is an attempt to compile the meagre information available on crop history, evolution, crop breeding and present status to make the crop competitive and revamp its cultivation.

Key words: breeding — domestication — *Echinochloa* — germplasm — genomic resources — nutritional value — origin — postharvest processing

The genus *Echinochloa* belongs to the tribe Paniceae, subfamily Panicoideae of the family Poaceae (Clayton and Renvoize 2006). Barnyard millet (*Echinochloa* sp.) is one of the oldest domesticated millets in the semi-arid tropics of Asia and Africa. Of around 35 species (Table 1), two main species, *E. esculenta* (A. Braun) H. Scholz; syn. *E. utilis* Ohwi et Yabuno (Japanese barnyard millet) and *E. frumentacea* Link; syn. *E. colona* var *frumentacea* (Link) Ridl. (Indian barnyard millet), are cultivated as minor cereals in Japan, Korea, the north-eastern parts of China and India, Pakistan and Nepal, respectively (Yabuno 1987) (Fig. 1). Besides these two species viz., allohexaploid *E. crus-galli*, Lijiang millet and allotetraploid *E. oryzicola*, Mosou barnyard millet are under cultivation in China (Yabuno 1966, Yuichiro et al. 1999, Yamaguchi et al. 2005).

Barnyard millet is grown for human consumption as well as fodder. It is generally cultivated in areas where climatic and edaphic conditions are unsuitable for rice cultivation (Yabuno 1987). In India, barnyard millet is the second important small millet after finger millet having production and productivity 87 thousand tonnes and 857 kg/ha, respectively (Padulosi et al.

2009). In India, it is mainly cultivated in two different agro-ecologies, one in mid hills of Himalayan region of Uttarakhand in the North and another in Deccan plateau region of Tamil Nadu in the south. Wild barnyard millet (*Echinochloa colona*) is commonly found in rice fields as weed and consumed as food during drought years in many states of India (Padulosi et al. 2009).

Owing to fast growth and early maturity, barnyard millet has attracted some attention as a fodder in the United States and Japan and can produce as many as eight harvests per year (<http://www.fao.org/docrep/T0818E/T0818E01.htm>). The crop straw is considered superior fodder to rice, oat or Timothy straw in protein and calcium content (Obara 1936). Its grains are also used as a major feed for waterfowl and other birds in United States (Mitchell 1989).

History, Origin and Domestication

Echinochloa frumentacea showed parallel line of evolution both in India and Africa. It is an annual cultivated in India, Central African Republic, Tanzania and Malawi (Doggett 1989). Its wild progenitor is the tropical grass *E. colona* (L.) Link, popularly known as *Jungle rice*, but the exact date of domestication is uncertain. *Echinochloa esculenta* is annual in habit and is cultivated mostly in the temperate regions (De Wet et al. 1983) of Japan, Korea, China, Russia and Germany. Its wild ancestor is barnyard grass (*E. crus-galli* (L.) Beauv.) from which it was directly domesticated some 4000 years ago in Japan (Doggett 1989).

Archaeological evidence suggests that it was grown in Japan as early as Yayoi period, dating back some 4–5 millennia (Watanabe 1970). Another study puts the earliest records of domestication from Jomon period of Japan in 2000 B.C. (Nesbitt 2005). Nozawa et al. (2004) showed that *E. esculenta* was domesticated from a limited part of the *E. crus-galli* population. They used 13 SSR markers to study the genetic diversity of 170 *Echinochloa* accessions and grouped *E. esculenta* accessions into two classes, while ancestral species *E. crus-galli* was grouped into 11 classes.

The domestication syndrome, which refers to all modifications occurring in a crop plant during the course of evolution when it becomes cultivated from the wild form and is dependent on selection pressure (Hammer 1984, 2003), is not well studied in barnyard millet although both *E. frumentacea* and *E. esculenta* showed marked difference from their respective wild ancestors *E. colona* and *E. crus-galli* with respect to reduced vegetative

Table 1: *Echinochloa* species along with their areas of adaptation

Sl No.	Species	Synonyms	Areas of geographical presence
1	<i>Echinochloa brevipedicellata</i>		Eastern Tropical Africa – Tanzania
2	<i>Echinochloa callopus</i>		Senegal, Mali, Ghana, Nigeria, Sudan, Congo, Tanzania
3	<i>Echinochloa chacoensis</i>		Bolivia to North Argentina
4	<i>Echinochloa colona</i>	<i>Echinochloa divaricata</i> ; <i>Echinochloa equitans</i> ; <i>Echinochloa subverticillata</i> ; <i>Echinochloa zonalis</i>	Tropical and subtropical old world
5	<i>Echinochloa crus-galli</i>	<i>Echinochloa caudata</i> ; <i>Echinochloa commutata</i> ; <i>Echinochloa crus-corvi</i> ; <i>Echinochloa disticha</i> ; <i>Echinochloa dubia</i> ; <i>Echinochloa echinata</i> ; <i>Echinochloa formosensis</i> ; <i>Echinochloa glabrescens</i> ; <i>Echinochloa hispida</i> ; <i>Echinochloa macrocarpa</i> var. <i>aristata</i> ; <i>Echinochloa macrocarpa</i> var. <i>mutica</i> ; <i>Echinochloa macrocorvi</i> ; <i>Echinochloa madagascariensis</i> ; <i>Echinochloa micans</i> ; <i>Echinochloa occidentalis</i> ; <i>Echinochloa paracorvi</i> ; <i>Echinochloa persistentia</i> ; <i>Echinochloa spiralis</i> ; <i>Echinochloa tzelevii</i> ; <i>Echinochloa zelayensis</i> ; <i>Echinochloa zenkowskii</i>	South and East Europe to Asia, West, East and South Tropical Africa to South Africa, Madagascar
6	<i>Echinochloa cruspavonis</i>	<i>Echinochloa aristata</i> ; <i>Echinochloa composite</i> ; <i>Echinochloa kimayalaensis</i> ; <i>Echinochloa sabulicola</i>	Tropical and South Africa, Assam to South China and Indo-China
7	<i>Echinochloa telmatophila</i>	<i>Echinochloa dietrichiana</i>	Queensland, Australia
8	<i>Echinochloa elliptica</i>		Queensland, Northern territory, Western Australia
9	<i>Echinochloa esculenta</i>	<i>Echinochloa frumentacea</i> var. <i>atherachne</i> ; <i>Echinochloa frumentacea</i> subsp. <i>utilis</i> ; <i>Echinochloa utilis</i>	Russian far east to Japan
10	<i>Echinochloa frumentacea</i>	<i>Echinochloa glabrescens</i> var. <i>barbata</i> ; <i>Echinochloa glabrescens</i> var. <i>glabra</i> ; <i>Echinochloa glabrescens</i> var. <i>pilosa</i>	Cultigen from India
11	<i>Echinochloa haploclada</i>	<i>Echinochloa aristifera</i> ; <i>Echinochloa haploclada</i> var. <i>stenostachya</i>	Africa-Ethiopia to Swaziland
12	<i>Echinochloa helodes</i>		Argentina
13	<i>Echinochloa holciformis</i>		Mexico, Guatemala
14	<i>Echinochloa inundata</i>		South Australia, Queensland and New South Wales
15	<i>Echinochloa jaliscana</i>		Mexico
16	<i>Echinochloa jubata</i>		Southern countries of Africa
17	<i>Echinochloa kimberleyensis</i>		Western Australia, Northern Territory and Queensland
18	<i>Echinochloa lacunaria</i>		South Australia, New South Wales
19	<i>Echinochloa macrandra</i>		Western Australia
20	<i>Echinochloa muricata</i>	<i>Echinochloa microstachya</i> ; <i>Echinochloa pungens</i> ; <i>Echinochloa wiegandii</i>	Canada, USA
21	<i>Echinochloa obtusiflora</i>		Senegal, North Nigeria to Sudan
22	<i>Echinochloa oplismenoides</i>		Arizona, Mexico, Guatemala
23	<i>Echinochloa oryzoides</i>	<i>Echinochloa coarctata</i> ; <i>Echinochloa erecta</i> ; <i>Echinochloa hispidula</i> ; <i>Echinochloa macrocarpa</i> ; <i>Echinochloa oryzicola</i> ; <i>Echinochloa pachychloa</i> ; <i>Echinochloa phyllopogon</i> ; <i>Echinochloa phylloryzoides</i>	Caucasus to Japan and Phillipines
24	<i>Echinochloa paludigena</i>		Florida
25	<i>Echinochloa picta</i>		Tropical Asia to West Pacific
26	<i>Echinochloa pithopus</i>		Tanzania
27	<i>Echinochloa polystachya</i>	<i>Echinochloa spectabilis</i>	Tropical and subtropical America
28	<i>Echinochloa praestans</i>		Papua New Guinea and Northern territory of Australia
29	<i>Echinochloa pyramidalis</i>	<i>Echinochloa frumentacea</i> var. <i>violacea</i> ; <i>Echinochloa guadeloupensis</i> ; <i>Echinochloa holubii</i> ; <i>Echinochloa kimpokoensis</i> ; <i>Echinochloa quadrifaria</i> ; <i>Echinochloa senegalensis</i> ; <i>Echinochloa verticillata</i>	Africa to Arabian Pen
30	<i>Echinochloa rotundiflora</i>		Nigeria to Eritrea
31	<i>Echinochloa stagnina</i>	<i>Echinochloa barbata</i> ; <i>Echinochloa hostii</i> ; <i>Echinochloa lelievrei</i> ; <i>Echinochloa malakuensis</i> ; <i>Echinochloa oryzetorum</i> ; <i>Echinochloa scabra</i>	Africa, tropical Asia
32	<i>Echinochloa telmatophila</i>		South West and East Australia
33	<i>Echinochloa turneriana</i>		Central and East Australia
34	<i>Echinochloa ugandensis</i>		Ethiopia to South Africa
35	<i>Echinochloa walteri</i>	<i>Echinochloa longearistata</i>	East Canada to Central America and Caribbean

Source – Kew Royal Botanical Gardens; <http://apps.kew.org/>.



Fig. 1: Panicles of two cultivated species of barnyard millet (a) *Echinochloa esculenta* (b) *Echinochloa frumentacea*

branching, more compact growth habit, larger inflorescence, reduced shattering and larger seed size. Yabuno (1975) considered low seed shattering, lack of seed dormancy, thick culms, wide leaves and round spikelets in *E. esculenta* were the main characters selected by man during the process of domestication. This suite of traits that constitutes 'domestication syndrome' for closely related foxtail millet (Defelice 2002, Doust et al. 2005, Li and Brutnell 2011) and pearl millet (Poncet et al. 1998, 2000) is likely for barnyard millet as well. Increase in seed size in Japanese barnyard millet during domestication is suggested by archaeological data. The mean size of *Echinochloa* caryopses from the Middle Jomon period (3470 B.C.E.–2420 B.C.E.) was about 20% larger than specimens from Early Jomon period (5000 B.C.E.–3470 B.C.E), indicating that selection for larger seed size was taking place over several millennia in Northern Japan (Crawford 1983, 2011, Takase 2009). The cross-compatibility between domesticated barnyard millet and their ancestral forms and the existence of naturally occurring intergrades between the two forms provide avenues to understand the mechanisms driving domestication and elucidate the genetics of domestication traits in this crop.

Phylogeny

The members of *Echinochloa* include minor cereal crops and major weeds in fields. The interspecific relationship is poorly understood due to the high morphological variation. Several attempts aimed at solving the difficulties have led to an ambiguous understanding of phylogenetic relationships among taxa (Hilu 1994, Michael 1994, 2001). The best understanding of this genus has come from series of studies by Yabuno (1962, 1984, 1996, 2001). Two cultivated species, Japanese barnyard millet (*E. esculenta*) and Indian barnyard millet (*E. frumentacea*), have a close relationship with their wild counterparts, *E. crus-galli* and *E. co-*

lona, respectively. Both of the cultivated species as well as their progenitors are hexaploid with $2n = 6x = 54$ where $x = 9$ (Yabuno 1962, 1966). Interspecific hybrids obtained by crosses between *E. crus-galli* × *E. esculenta* and *E. colona* × *E. frumentacea* were found to have normal meiotic division with 27 bivalents. The crosses between these two groups, that is, between two cultivated species (*E. esculenta* × *E. frumentacea*) and their ancestors (*E. crus-galli* × *E. colona*), resulted in meiotic irregularities, univalents, laggards and micronuclei. These cytogenetic evidences suggested that the hexaploid wild species *E. colona* and *E. crus-galli* are possible progenitors of *E. frumentacea* and *E. esculenta*, respectively, and the two cultivated species have different genomic composition (Yabuno 1966). The crosses between *E. crus-galli* and *E. oryzoides* showed 18 bivalents and nine univalents suggesting that two of the three genomes of *E. crus-galli* are homologous to *E. oryzoides* and *E. crus-galli* is an allohexaploid produced through natural hybridization between the tetraploid *E. oryzoides* with a not-yet-discovered diploid species of *Echinochloa* (Yabuno 1966, 1984).

On the basis of these crossing experiments and observation of meiotic chromosomal behaviour of interspecific hybrids, *Echinochloa* species were grouped into several cross-compatible groups, and polyploidization followed with amphidiploidization had taken place in the evolutionary process of species diversification in this genus (Yabuno 1966, 1984, 2001). However, the morphological, reproductive and ecologic traits relied upon by conventional taxonomists for the identification or discrimination of taxa might present ambiguous results due to domestication and adaptive syndromes (Yabuno 1966, 1984, 2001, Yamaguchi et al. 1996).

Later on to understand, genetic relationships among Eurasian annual *Echinochloa* species molecular markers were used by several workers (Yasuda et al. 2002, Yamaguchi et al. 2005, Ruiz-Santaella et al. 2006). The molecular analysis also confirmed that the Asian annual weedy *Echinochloa* species including their domesticated counterparts fall into three cross-compatible groups, the *Echinochloa oryzicola* group, *Echinochloa crus-galli* complex (group) and *Echinochloa colona-frumentacea* group (Yamaguchi et al. 2005). Species divergence is well illustrated by the nucleotide sequencing of the particular regions of the cpDNA, and the three groups have shown different cytoplasmic lineages (Yamaguchi et al. 2005). The tetraploid *E. oryzicola* includes the wild representatives, *E. oryzicola* (Vasing.) and *Echinochloa phyllopogon* (Stapf) Koss. and Mosuo barnyard millet. The hexaploid *E. crus-galli* group consists of four wild *E. crus-galli* (L.) P. Beauv. varieties – var. *crus-galli*, var. *praticola* Ohwi, var. *formosensis* Ohwi and var. *oryzoides* (Ard.) Lindm. (syn: *Echinochloa oryzoides* [Ard.] Fritsch) – and one cultivated species, *Echinochloa esculenta* (A. Braun) Scholz (Yabuno 1984, 1996). The hexaploid *E. colona-frumentacea* group consists of wild *E. colona* (L.) Link and cultivated *Echinochloa frumentacea* (Roxb.) Link. The hypothesis that *E. crus-galli* was derived from a hybrid between *E. oryzicola* and unknown diploid species (Yabuno 1966) was confirmed using internal transcribed spacer and chloroplast DNA sequences by Aoki and Yamaguchi (2008). They found same nuclear lineage between *E. oryzicola* and *E. crus-galli* indicating that *E. oryzicola* is the paternal donor of *E. crus-galli*. Further, phylogenetic analysis of the *sh4* gene (control spikelets shattering) of *Oryza* detected homoeologue copies in *E. oryzicola*, *E. crus-galli*, *E. stagnina*, *E. colona* and *E. crus-pavonis*. This indicated a genomic relationship between the Asian *Echinochloa* species

and supported that the allohexaploid *E. crus-galli* shares two genomes with its parental donor, *E. oryzicola*. The Asian perennial tetraploid species, *E. stagnina*, shares one genome with *E. oryzicola* and possesses an unknown genome, *E. crus-pavonis*, from the New World, shows a close affinity of two genomes with *E. crus-galli* and *E. oryzicola*, while *E. colona* showed distant affinities in all homoeologous copies (Aoki and Yamaguchi 2009).

Botanical Description

Barnyard millet has a wide adaptation capacity and can grow up to an altitude of 2000 m above mean sea level during summer season (Gupta *et al.* 2009a). It is variable in flowering time, inflorescence shape, morphological features, pigmentation of spikelets, plant type and other plant traits (Obara 1938). The crop plant is a tall, robust annual and grows up to 220 cm high. It has a short generation time, fastest growth among all small millets and completes the life cycle from seed to seed in 45–60 days (depending upon accession and growth environment) (Denton 1987, Padulosi *et al.* 2009), however, may take longer time under northern hill ecosystem. Hulse *et al.* (1980) reported that *Echinochloa* millets grow well in different seasons but at high elevations may require 3–4 months to mature. Leaf blades are flat and wide with no ligules. The inflorescence is a terminal panicle 10–25 cm long with dense racemes of 3- to 4-mm-long spikelets (Napper 1965). The inflorescence is usually erect, rarely drooping with shapes varying from cylindrical, pyramidal and globose to elliptic. Racemes are few to numerous, densely crowded with spikelets arranged in four irregular rows on the triquetrous rachis and are loosely or tightly appended on the rachis. The spikelets are green, brown to purple in colour and crowded on one side of the rachis. Spikelets are two flowered, awnless or awned, with red or green awns and placed on short rough pedicels subtended by two glumes (Mitich 1990). Lower floret is neuter (sterile) with lemma and small palea, and upper floret is bisexual (Gupta *et al.* 2010a). The sterile lemma is 5-veined. Fertile lemma is plano-convex, elliptic, smooth and shiny, abruptly sharp-pointed or cuspidate, and margins are inrolled below over palea with apex of palea not enclosed. The palea is flat, and surface texture is similar to fertile lemma (Napper 1965). Stamens are three in number, and ovary superior contains two distinct styles with plumose stigma (Sundararaj and Thulasidas 1976). Grain is 2–3 mm long and 1–2 mm wide and enclosed in white shining hardened lemma and palea.

The flowering starts from top of the inflorescence and moves downward completing in 10–15 days. Flowers open from 5 to 10 am with maximum number of flower opens between 6 and 7 am (Sundararaj and Thulasidas 1976, Jayaraman *et al.* 1997). In the individual raceme, the flowering first starts at marginal ends and then proceeds to the middle of the raceme. The flowers are hermaphrodite (have both male and female organs). Before the anthers dehiscence, the stigmatic branches spread and flower opens (Seetharam *et al.* 2003). Late season florets are cleistogamous (not opening) (Maun and Barrett 1986). It is primarily self-pollinating (Maun and Barrett 1986, Potvin 1986, 1991) and self-compatible (Maun and Barrett 1986). Some degree of out-crossing recorded which was facilitated by wind pollination (Maun and Barrett 1986). Hot water treatment of inflorescence at 48°C for 4–5 min (personal observation) was effective in inducing male sterility under hill condition in both the cultivated species.



Fig. 2: Barnyard crop trial at VPKAS, ICAR, Almora, Uttarakhand, India

The two cultivated millet species namely Indian and Japanese barnyard millet can easily be distinguished on the basis of panicle morphology, texture of the glumes and lower lemma (Yabuno 1971, 1987, De Wet *et al.* 1983). *Echinochloa frumentacea* has smaller awnless spikelets, with membraneous glumes in comparison with large usually awned spikelets and chartaceous upper glumes and lower lemma in *E. esculenta*. Based on inflorescence morphology, the species *E. frumentacea* was classified into four races namely *Stolonifera*, *Intermedia*, *Robusta* and *Laxa*. Similarly, *E. utilis* (Syn. *E. esculenta*) was classified into two races namely *utilis* and *intermedia*.

The mean diploid 2C DNA content of barnyard millet is reported to be 2.65–2.7 pg *bot agr* (Schlegel 2010).

Growth and Development

Barnyard millet grown under natural precipitation is a fast-growing annual summer crop for both food and fodder. Rapid growth, drought tolerance and ability to grow in marginal environments make barnyard millet an important crop for famine areas (De Wet *et al.* 1983, Kono *et al.* 1987, Majid *et al.* 1989) (Fig. 2). The crop seeds germinate within approximately 24 h of water imbibition (Rahn *et al.* 1968). After 5 days of emergence, plants produce adventitious roots (Rahn *et al.* 1968) and the first tillers appear 10 days after emergence (Holm *et al.* 1991). Temperature plays a major role in the growth of both the species (Holm *et al.* 1991), and low temperature conditions (15/10°C day/night) cease the growth and affect the plant development particularly leaf area and dry weight. For proper growth of *E. frumentacea*, the optimum temperature range is 27–33°C and 15–22°C day and night, respectively (Muldoon *et al.* 1982). In high hills of Uttarakhand, *E. frumentacea* and *E. esculenta* showed significant variation in seedling emergence and other characters in April sowing, whereas the differences were non-significant in May–June sowing. This is because of *E. esculenta* had better tolerance to low temperature and showed little reduction in relative growth rate in comparison with *E. frumentacea* (Muldoon *et al.* 1982,

Bandyopadhyay 1999). If conditions are favourable, plants show profuse growth for 3 weeks and the transition from vegetative growth into inflorescence takes place rapidly within approximately 40 days after emergence (Maun and Barrett 1986). Although both the cultivated species have been reported to be quantitative short-day plants (Muldoon 1985), most *Echinochloa* spp. can grow and reproduce in a range of photoperiods, short days (8–13 h) and long days (16 h) (Maun and Barrett 1986, Mitich 1990). Under short-day conditions, plants are small and flowering is quick, yet abundant; under long-day conditions, the plants are more robust with a large seed output (Maun and Barrett 1986, Manidool 1992). Late plantings shorten the vegetative phase and adversely affect the crop yield. Seeds mature in about 20–40 days after the reproductive phase begins. The optimum time of harvesting in Japanese barnyard millet is 30–35 days after heading (Kumagai et al. 2011), when the moisture content of panicles is 16–18%. Inmate seed dormancy in freshly harvested seeds varies from 4 to 48 months in *Echinochloa crus-galli* (Maun and Barrett 1986, Sung et al. 1987, Manidool 1992). All *Echinochloa* species have a C₄ photosynthetic pathway, show a great competitive advantage when they grow together with C₃ crops (Bouhache and Bayer 1993) and perform better than rice under water stress conditions due to high water-use efficiency and 60% higher carbon efficiency ratio (Migo et al. 1991).

Germplasm and Its Impact

Diversity in barnyard millet is being fast eroded due to considerable reduction in acreage and changing sociocultural and economic dimensions of the farming community in India (Maikhuri et al. 2001). Considerable efforts have been made to preserve the crop diversity *ex situ*, but the information regarding on-farm, *in situ* conservation is very scarce (Padulosi et al. 2009). The largest *ex situ* collection at international level is maintained by the Consultative Group on International Agricultural Research (CGIAR) with 2365 collections. A few of them (44 accessions) have been duplicated with the Global Crop Diversity Trust for safe storage at the Svalbard Gene Bank in Norway (Padulosi et al. 2009). India holds the largest barnyard millet collection at National Bureau of Plant Genetic Resources (1718 accessions) followed by All India Coordinated Small Millet Improvement Project at the University of Agricultural Sciences, Bangalore, Karnataka (985 accessions). Vivekananda Parvatiya Krishi Anusandhan Sansthan (VPKAS), Almora, is mainly maintaining more than 300 local collections of barnyard millet from Uttarakhand. ICRISAT has a total of 743 active collections and 487 base collections from nine countries for research and distribution (Upadhyaya et al. 2008). In addition, the US GRIN database contains 306 accessions of 18 *Echinochloa* species from 33 countries housed at the National Centre for Genetic Resources Conservation (Fort Collins, Colorado) and a smaller collection of 67 accessions at Australian Plant Genetic Resource Information Service, Biloela (Dwivedi et al. 2012). Notwithstanding the impressive size of global barnyard millet collection at both National and International organizations, it is imperative to analyse and fill germplasm gaps to make the collection more comprehensive. For example, accessions belonging to *laxa* race endemic to Sikkim state of India hitherto unrepresented in the *ex situ* collections need to be collected before it becomes extinct (Dwivedi et al. 2012).

Germplasm is the basic material for crop improvement programme in any crop. Descriptor list has been developed and

used to characterize barnyard millet germplasm for sets of morphologic and agronomic traits (http://www.ecpgr.cgiar.org/fileadmin/bioiversity/publications/pdfs/394_Echinochloa_millet_descriptors.pdf?cache=1372427061). Phenotypic characterization of the collected materials has indicated that barnyard millet germplasm is highly diverse (Halaswamy et al. 2001, Gowda et al. 2008, Gupta et al. 2009a, Nirmalakumari and Vetriventhan 2009). Statewise grouping of morphological characters of Indian collections of seven states did not reveal concentration of any group to a specific state (Halaswamy et al. 2001). In a similar study, different genotypes clustered together irrespective of their place of origin suggesting that geographical isolation was not the only factor causing genetic diversity (Mehta et al. 2005, 2007). Similar results were also observed by Prabha et al. (2010) where the accessions of two different cultivated species of barnyard millet clustered together although, with the help of isozyme markers, they also found accessions within each species forming two different clusters, which is consistent with the morphological evidence on existence of intergrades and overlaps between the two species. On the basis of principal component analysis of quantitative traits, Gupta et al. (2009a) classified 194 barnyard accessions of Indian origin into three races *robusta*, *intermedia* and *stolonifera* on the basis of resemblance of morphological traits of the accessions with these races. Race *laxa*, however, was absent in the group.

For optimum and precise utilization of diversity for agronomic and nutritional improvement, a core collection representing 50 and 89 accessions has been developed by Gowda et al. (2009) and Upadhyaya et al. (2014), respectively. Halaswamy et al. (2001) characterized the national collection and identified promising accessions for higher plant height (seven accessions), higher number of basal tillers (nine accessions), longer inflorescence (10 accessions) and early flowering (27 accessions). New lines, IEC 566 and IEC 566/2, which differ in several aspects from cultivated varieties and produce abundant pollen grains and have flowers that open for a long time with stigma protruding enough to facilitate emasculation and pollination, have also been identified (Nirmalakumari and Vetriventhan 2009). Similarly, an easy dehulling accession B29 was identified in accessions from Uttarakhand hills. The dehulled grain recovery of B29 was 40% and 140% higher than the check varieties VL 172 and PRJ 1, respectively. The accession is registered with NBPGR vide number INGR09023 (Gupta et al. 2009b).

Characterization and evaluation of germplasm for important agronomic traits constitutes the most critical component driving utilization of germplasm by crop breeders. Utilization of barnyard millet genetic resources for crop improvement falls far short of the desired. The release of variety PRJ 1, which was a direct selection from ICRISAT germplasm, in 2003 for Uttarakhand state is among few notable examples of effective utilization of barnyard millet germplasm. The variety yielded 45.4% higher than the check variety VL 29 (Upadhyaya et al. 2008). PRJ 1 belongs to *E. esculenta*, whereas all the existing adapted material in Uttarakhand hills was of species *E. frumentacea*. Much, therefore, needs to be carried out to enhance utilization of barnyard millet germplasm for genetic improvement of the crop.

Breeding Objectives and Achievements

Echinochloa species show a high degree of autogamy, but the rate of cross-pollination is sufficient to assure gene exchange among their populations (Maun and Barrett 1986). Emasculation and artificial hybridization is difficult due to small flower size,

early hours of flowering, short viability of pollen, non-availability of pollen grain and slight opening of flowers that too for a short period (Nirmalakumari and Vetriventhan 2009). The crop is still considered as a minor food and feed crop of poor tribal people, has not attracted research efforts like other major crop plants and very limited work has been carried out for its improvement. In India, barnyard millet breeding is carried out mainly in the states of Uttarakhand and Tamil Nadu. Thus far, more than 20 improved cultivars have been developed and released for different barnyard millet growing regions of the country. Mass selection and pure line selection have remained the major breeding strategies for the improvement of the crop; however, some varieties have been developed through hybridization followed by pedigree method of selection.

Although it is difficult to induce a mutant phenotype in a polyploid species like *Echinochloa*, full waxy stable mutant lines have been developed through γ irradiation of low amylase landrace 'Nogehie' (Hoshino *et al.* 2010). Gamma irradiation also increased genetic variance for tiller number, plant height, head length and grain yield (Mehra *et al.* 1985). Mehta *et al.* (2005) emphasized interspecific hybridization programme involving early maturing *E. frumentacea* and high-yielding *E. esulenta* to develop early maturing high-yielding segregants. However, hybrids between *E. frumentacea* and *E. esculenta* are sterile both ways, whereas those with its wild progenitors are fertile. These wild progenitors can provide valuable genetic resources for the improvement of cultivated species (Mandelbaum *et al.* 1995). Large variation in protein and calcium values within accessions of a species suggests selective breeding for nutritive values (Mandelbaum *et al.* 1995).

Barnyard millet can grow up to 2 m or more under high moisture conditions prevalent during rainy season in India, making it prone to lodging. Reducing plant height to about 120–130 cm could minimize the problem. The resultant reduction in fodder yield may be compensated by developing genotypes with higher number of basal tillers, adding to grain yield as well. In barnyard millet, the length and number of spikes is positively correlated with grain yield. Developing genotypes with more and longer spikes should be the principal approach for enhancing grain yield. Breeding for easy dehulling is another important breeding objective to reduce the drudgery involved in postharvest processing of barnyard millet. Availability of easy dehulling types has been reported in the local germplasm (Anonymous 2010), which can be used to transfer the trait to high-yielding adapted cultivars. As barnyard millet is traditionally consumed just like rice, increasing grain size is likely to enhance its appeal among the consumers. This, however, remains a challenge due to narrow range of variability for the trait.

Among biotic stresses, grain smut caused by *Ustilago* spp. is a major yield constraint in barnyard millet causing yield losses of up to 60.8% (Jain *et al.* 1997). Source of resistance to grain smut is not reported in Indian barnyard millet, while Japanese barnyard millet is near immune to the disease. The transfer of resistance from Japanese to Indian barnyard millet, however, is hampered by sterility in the hybrids between the two species (Sood *et al.* 2014). The wild progenitor of Indian barnyard millet, *E. colona*, which is resistant to grain smut as well as crossable with cultivated barnyard millet may be an alternate source for the trait. Breeding efforts should also be directed towards resistance to *Helminthosporium* leaf blight which is a serious problem in specific areas.

Improvement in grain yield remains the most important breeding objective in barnyard millet. A wide production gap

still exists between yields realized at farmer's fields because of prevalence of local cultivation practices (Gupta *et al.* 2006). Most of barnyard-millet-growing areas are still under local cultivars and land races with low grain yields of 1.0–1.5 tons per hectare, whereas the crop has the potential of >2.0 tons/ha (Harinarayana 1989) and yield levels of 3.0 tons/ha have been reported in the *E. esulenta* lines (Bandyopadhyay 2001). Improved varieties coupled with modern agronomic practices can bring about significant improvement in barnyard millet productivity and bridge the production gap. Barnyard millet straw is preferred over rice straw. Therefore, selection of variety which could give high grain as well as high fodder is the best alternative.

Fodder Quality

Barnyard millet grows rapidly and produces voluminous fodder. Its fodder is highly palatable and can be used for making hay or silage. It is considered superior to rice and oat straw because of high protein and calcium content (Yabuno 1987). Barnyard millet straw contains up to 61% total digestible nutrients and good amount of protein and digestible fibre (National Research Council 1996). The average dry fodder yield of barnyard under Indian conditions is 5 tons/ha (AIC-SMIP 2014). In Uttarakhand hills of India, barnyard millet fodder contributes 11.5% of the total fodder consumption of the State (Singh and Singh 2005), where even grains of Indian barnyard millet are also fed to animals. Recent studies revealed that Japanese barnyard millet has higher fodder production potential than Indian barnyard millet, and recommended fertilizer application significantly enhances the fodder yield in barnyard millet (Yadav and Yadav 2013). Japanese barnyard millet has also been found superior than Indian barnyard millet under double fodder cutting management for dual purpose use (Bandyopadhyay 2009). However, non-significant differences were observed between the two species for dry matter digestibility (Unpublished data).

In Australia, China and the United States of America, Japanese barnyard millet is grown for grazing or hay. In United States of America, it is reported to produce eight harvests per year (Kajuna 2001). It has the highest protein content of all the millet species (<http://www.fao.org/docrep/008/y5831e/y5831e06.htm>). In Australia, it has been recognized as a valuable short-term rotation crop for spring-early summer grazing. The rapid early growth of the crop can fill feed shortfalls in early summer after floods or drought. The crop gives two good grazings in northern New South Wales and repeated grazing in the cooler southern areas. The major weakness of the crop is quick heading under hot and dry conditions (Dairy Link-Establishing pastures-Pasture Species http://www.dpi.nsw.gov.au/__data/assets/pdf_file/0018/163116/establishing-pastures-1-5.pdf).

It has superior feed quality when fed young to the animals (Metabolizable energy 8.5–9.5 MJ/kg); however, protein content declines from 25% to 6% at maturity. In Kyabram, Australia, *Echinochloa* millet cv. 'Shirohie' produced dry matter (DM) and digestible dry matter (DDM) (16.3 t/ha, 10.8 t/ha) equivalent to sorghum \times sudan grass hybrid cv. 'Sudax' (17.1 t DM/ha and 10.7 t/ha) and showed higher digestibility (65.9%) and nitrogen concentration (1.9%) than sorghum \times sudan hybrids (63.3% dry matter digestibility, 1.5% N). Similar studies in Bangladesh also showed higher productivity and better nutritive value of *Echinochloa crus-galli* L. (Kanak *et al.* 2013).

Genomic Resources and Molecular Breeding Advancements

The use of molecular markers in barnyard millet closely followed their use in other small millets such as finger millet and foxtail millet. Hilu (1994) using RAPD markers observed high degree of variability in Indian barnyard millet (*E. frumentacea*) in consistency with high degree of morphological variability observed in this species. Danquah et al. (2002) developed five primer pairs of microsatellite loci for studying genetic diversity and interspecific classification of three agronomically important *Echinochloa* spp viz., *E. crus-galli* (L.) Link., *E. colona* (L.) Beauv. and *E. crus-pavonis* (Kunth.). They reported that microsatellites were useful in discriminating the three species and could aid in classification of species within this complex genus. SSR markers were used by Nozawa et al. (2006) to study population structure and diversity of 155 accessions of barnyard millet including 49 from var. *esculenta*, 94 from var. *crus-galli* and 12 from var. *formosensis*. The SSR markers clustered the *esculenta* accessions into two groups, *crus-galli* accessions into 12 groups and *formosensis* accessions into six groups. The results also revealed that accessions belonging to var. *esculenta* were less diverse than those of var. *crus-galli* or var. *formosensis*. The information on genomic resources in barnyard millet is meagre; however, foxtail millet has emerged as a model crop for studying the system biology of other millets due to the availability of genomewide sequence resources. In foxtail millet, large-scale development of molecular markers such as SSRs (Pandey et al. 2013), EST-derived SSRs (Kumari et al. 2013) and ILPs (Muthamilarasan et al. 2014) was reported along with the demonstration of these markers in germplasm characterization, transferability, phylogenetics and comparative mapping studies in millets and bioenergy grasses (Muthamilarasan et al. 2014). In addition, a unique web-based Foxtail millet Marker Database (FmMDb) had also been constructed (Venkata Suresh et al. 2013) for access of breeders and researchers to these marker resources. The high levels of transferability in these studies substantiate the applicability of sequence-based markers in comparative genome mapping and evolutionary studies in other grass species including barnyard millet where the cross-transferability was more than 90% (Table 2) and was placed in same group with foxtail varieties and species (Yadav et al. 2014). The development and large-scale validation of such genomic microsatellite markers in a genomewide scale could be useful for barnyard millet also, for which very little or no genomic information is available (Lata et al. 2013). Molecular breeding studies such as genetic diversity, linkage mapping and identification of QTLs require a large number of molecular markers. As there is no genome sequence information available in barnyard millet, comparative genomics plays an important

Table 2: Percentage transferability of foxtail sequence-based genetic markers in barnyard millet

Sl No.	Marker type and number	Percentage transferability	References
1	SSR, 58	91.3	Pandey et al. (2013)
2	EST SSR, 106	90.6	Kumari et al. (2013)
3	ILP, 100	94.1	Muthamilarasan et al. (2014)
4	miRNA, 66	89	Yadav et al. (2014)

SSR, simple sequence repeat; EST, expressed sequence tag; ILP, intron length polymorphism; miRNA, micro-ribonucleic acid.

role. In this concern, the primary studies conducted earlier in foxtail millet (Kumari et al. 2013, Pandey et al. 2013, Muthamilarasan et al. 2014, Yadav et al. 2014) will help in barnyard crop improvement through molecular approaches. The identified SSRs can be utilized in the genetic diversity studies, linkage map construction and their further use in identification of markers linked to the QTLs of important agro-morphological characters in barnyard millet. The identified QTLs can be effectively introgressed in the locally adapted genotypes of barnyard millet for yield improvement and stress amelioration through marker-assisted selection.

Besides the use of molecular markers, literature abounds in reports of successful callus and plant regeneration in *E. frumentacea* (Talwar and Rashid 1989, Sankhla et al. 1992, Bobkov 2005), *E. crus-galli* (Gupta et al. 2001) and *E. colona* (Tyagi et al. 1985, Samantaray et al. 1995, 1996, 2001, Rout et al. 1997). Bobkov (2005) optimized the media to maintain viable callus of *E. frumentacea* for many years. He cultured the seeds in 2KC media which contained salts according to the MS medium and vitamins according to the B5 medium, 100 mg/l myo-inositol, 4 g/l sucrose, 2 mg/l glycine, 6 g/l agar and 2 mg/l 2,4-D. There is only one preliminary study of transformation in barnyard millet (Gupta et al. 2001) which was carried out along with finger millet for assessment of the efficiency of five gene promoters (CaMV35, Act1, uql, rice ribulose 1, 5-biphosphate carboxylase small subunit (RbcS) and *Flaveria trinervia* (Ft)). In barnyard millet, only the uql promoter was effective for the expression of GUS gene; other four promoters (CaMV35S, Act1, RbcS and Ft) were ineffective based on the histochemical analysis.

Nutritional Importance and Antinutritional Compounds

Both the cultivated species have higher protein content than their wild ancestors. The protein content in barnyard millet ranged from 11.1% to 13.9% (Monteiro et al. 1987). However, the reverse is true for mineral content which suggests that wild species might consist of greater proportion of embryo/endosperm because of selection of larger seeds that increases endosperm size (Mandelbaum et al. 1995). The barnyard millet grain contains about 65% carbohydrate, majority of which is in the form of non-starchy polysaccharide and dietary fibre. This helps in the prevention of constipation, lowering of blood cholesterol and slow release of glucose into the blood stream during digestion. Barnyard millet has been found to be most effective in reducing blood glucose and lipid levels compared to other millets (Proso, Foxtail, Kodo and Little millet) and rice (Krishna Kumari and Thayumanavan 1998). There are several other reports suggesting effectiveness of barnyard and its products for diabetics (Arora and Srivastava 2002, Surekha 2004, Ugare et al. 2014). Polishing barnyard millet results in loss of fibre and other nutritional components, and maximum loss occurs at 14% moisture level (Lohani et al. 2012). Barnyard has non-glutinous type of endosperm (Tomita et al. 1981). Starch of barnyard millets was more digestible than maize starch in terms of *in vitro* amylolysis by pancreatic amylase (<http://www.fao.org/docrep/T0818E/T0818E0c.htm#Carbohydrate>). However, high fibre content and poor digestibility of nutrients adversely affects consumer acceptability of all small millets. The nutritional potential of millets is limited by the presence of phytates, phenols and tannins. Kulkarni et al. (1992) assessed the tannin content of five minor millets viz., proso, kodo, Italian, little and barnyard millet and recorded

lowest level in barnyard millet (102.96 mg). It has been shown that dehulling of the seeds reduces phytate and tannin levels (Lorenz 1983, Kulkarni *et al.* 1992).

Recently, the demand of barnyard millet has increased due to its highly nutritious grains and presence of strong antioxidative compounds (Watanabe 1999). He isolated three antioxidative phenolic compounds, one serotonin derivative and two flavonoids, from Japanese barnyard millet (cv. 'Kurohie') grains. High nutrient content and antioxidant effects make barnyard millet to be considered as a functional food crop (Kim *et al.* 2011), and the crop has potential to be included in normal and therapeutic diet formulations (Veena *et al.* 2005). Gluten presence in main cereal crops such as wheat make them allergic to some people, but barnyard millet grains are gluten free and, therefore, offer good opportunity for their use as health foods also (Hoshino *et al.* 2010).

Millet as a group contain several antinutritional compounds such as polyphenols, tannins, phytic acid and phytate, goitrogens and oxalic acid. The presence of antinutrients such as phytate, polyphenols, oxalates and tannins affect the mineral bioavailability. These antinutrients form complexes with dietary minerals, such as calcium, zinc and iron, leading to a marked reduction in its bioavailability and make them biologically unavailable to human organism (Arora *et al.* 2003). Polyphenols and tannin compounds are concentrated in the bran and affect *in vitro* protein digestibility. Oxalic acid forms an insoluble complex with calcium and thereby reducing its biological availability. The work on antinutritional compounds in barnyard millet is very limited, and there are no reports of any kind of antinutritional compounds in barnyard millet (Dwivedi *et al.* 2012). A study by Gupta *et al.* (2013) also showed that the total tannin content (0.301), oxalate content (0.02) and trypsin inhibitor activity (31.95) were well below the threshold level and cyanide content and haemagglutinin activity were absent in two National varieties VL 29 and VL 172 of Indian barnyard millet. However, we have to wait for much more reports to draw any valid conclusion on the presence of antinutritional compounds in barnyard millet.

Diseases and Insect Pests

Grain smut, caused by *Ustilago panici-frumentacei* Brefeld, is the major disease in barnyard millet (Fig. 3). In the infected panicles, the flower ovaries are round and hairy and may enlarge 2–3 times than normal size. Sometime, gall-like swellings are also observed on the nodes, axils of the older leaves and the

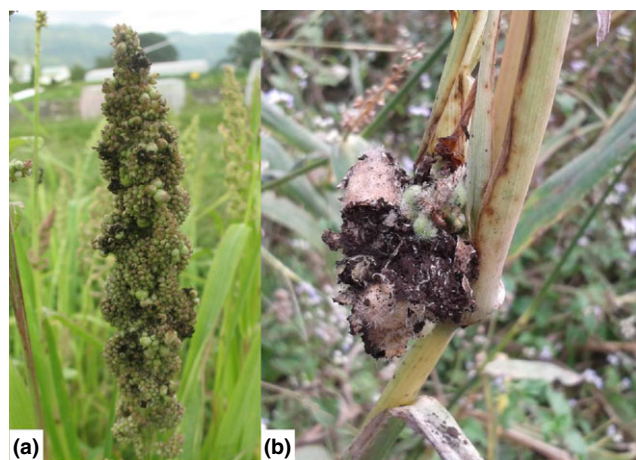


Fig. 3: Smut in barnyard millet (a) grain smut (b) head smut

stem. This disease is called head smut and is caused by *Ustilago crus-galli* Tracy and Earle (Nagaraja *et al.* 2007). This occurs late in the season when the crop is about to mature.

Phenotypic screening for grain smut allowed identification of disease-resistant germplasm. Natural intensity of 0–75% at maturity has been reported by Pawar *et al.* (1982). Two accessions (PRB 9402 and PRB 9602) belonging to *E. esculenta* showed immune reaction to grain smut at Almora (AICSMIP 2001); however, two separate studies at two locations in India (Almora and Bangalore) failed to identify a single accession of *E. frumentacea* immune to disease (Nagaraja and Mantur 2008, Gupta *et al.* 2010b). Nevertheless, availability of variation for grain smut resistance, particularly low grain smut incidence in accessions of different origin groups (Bihar and Karnataka), provides opportunity to significantly enhance the resistance level through hybridization and selection (Gupta *et al.* 2010b). More research input is needed to identify sources of resistance to grain smut in Indian barnyard millet.

Other diseases of minor importance in the crop are anthracnose, *Cercospora* leaf spot (*Cercospora fujimaculans*), leaf blight (*Exserohilum monoceras*) and sheath blight (*Rhizoctonia solani*) (<http://www.nilgs.affrc.go.jp/db/diseases/contents/edisease.htm>). The causal organism of the anthracnose disease is *Colletotrichum echinoclhoe* (Moriwaki and Tsukiboshi 2009) and is distinct from *C. graminicola* which was earlier considered as the causal organism for the disease in the crop.

The major insect pests of barnyard millet are pink stem borer (*Sesamia inferens*) (Fig. 4) and shootfly (*Altherigona falcata*) (Jagadish *et al.* 2008). The pink stem borer tunnels the stem resulting in dead heart/white head formation, while the shootfly causes tunnelling in the nursery stage. The crop has long storage life and keeping quality and is practically free of stored grain pests.

Postharvest Processing

Barnyard millet grain requires dehulling prior to making it suitable for human consumption (Lohani *et al.* 2012). The dehulling is conventionally performed by repeated pounding in mortar, which is time consuming and also labour intensive as the grains



Fig. 4: Stem borer and its damage in barnyard millet

are firmly encased in the lemma and palea. The drudgery involved in manual processing is an important factor contributing to reduction in consumption of millets. The small seed size also makes processing of these crops difficult. To reduce the drudgery, Singh et al. (2003) developed Vivek Millet Thresher 1, which can thresh 40–60 kg barnyard millet grains per hour. This machine is suitable for marginal farmers in hilly and tribal areas due to its low cost, small size, light weight and ability to do both threshing and dehusking. The machine can dehusk 5–6 kg grains per hour by changing the sieves but require 3–4 passes. Another similar easy-to-use affordable mill has been developed through collaborative efforts by Indian and Canadian researchers, which is 98% efficient in dehusking with <2% broken grains. This machine produces about 2 kg clean seed in an hour. Higher capacity machines/mills, which can process 100 kg/h, are also available for large farms. These machines with higher capacities have been designed for dehusking only. One such machine developed by Central Institute of Agricultural Engineering (CIAE, ICAR), Bhopal, India, has a capacity of dehusking and grinding 100 kg in an hour at 10–12% moisture content. It operates with one-horsepower single-phase electric motor and can even process one kg of grains also. These machines can significantly reduce the work load and time for postharvest processing of small millets.

Uses

In the Indian Himalayan region, barnyard millet is traditionally used a substitute for rice. The grains are dehulled, cooked and consumed like rice. Barnyard millet porridge (locally called *madi-ira ki kheer*) is a popular sweet dish in Uttarakhand. In southern India, barnyard millet is used in traditional preparations such as idli, dosa and chakli. The millet has also been used to develop products such as biscuits, sweets, noodles, rusk, ready mix, popped products and some other speciality foods (Arora and Srivastava 2002, Poongodi et al. 2003, Veena et al. 2004, Ugare 2008), but large-scale production needs industry involvement to commercialize the products globally.

Barnyard millet is highly suitable for commercial foods for diabetics, infants and pregnant women because of high iron content. However, the non-availability of ready-to-use processed products has limited the usage and acceptability of barnyard millet, despite its nutritional superiority. There is a need to develop millet-based food products in the form of ready-to-use products and functional foods to meet the demands of the present-day consumers. Value addition to minor millets not only offers variety, convenience and quality food to consumers, but is important for revival of barnyard millet cultivation as well.

Barnyard millet is also an important source of fodder in the Himalayan region. Barnyard millet leaves are broad, and the plant picks up good growth in short time and thus produces voluminous fodder. Barnyard fodder is highly palatable and can be used for making hay or silage.

Future Scope of Work and Prospects

The genetic base of cultivated barnyard millet is relatively narrow particularly with respect to traits such as disease resistance and seed size. The genetic base of the crop can be broadened by exploiting the wild relative gene pool, particularly the progenitors. The introgression of traits from the wild relatives can be facilitated by developing closely linked markers for these traits.

Interspecific hybridization between Indian and Japanese barnyard millet offers a promising avenue for mutual genetic improvement of the two species. The fertility barriers between the two species need to be analysed, and modern biotechnological tools employed to overcome them. There is also a need to systematically analyse the available genetic diversity to identify sources of agronomically important traits such as higher yield, disease resistance, nutritional quality of the grain and others. Molecular breeding efforts in barnyard millet lag considerably behind those in other small millets such as foxtail millet (genome sequenced) and finger millet (genetic map developed). Development of a genetic map will greatly boost the breeding efforts towards targeted improvement of barnyard millet. Availability of foxtail millet genome sequence will be enormously advantageous for comparative genomics, genome mapping, marker development and molecular breeding of barnyard millet. These genomic microsatellite markers of foxtail millet showing cross-transferability in barnyard millet would be of enormous and immense use for various large-scale genotyping applications, including germplasm characterization, cultivar identification and QTL discovery. High levels of transferability of ILP markers also demonstrate the usability as anchor markers for comparative genome mapping and understanding phylogenetics across diverse crop species, which eventually will be helpful for map-based isolation of genes in crop plants.

On a worldwide scale, barnyard millet like all other small millets is losing its importance as a food crop in competition with major cereals such as wheat, rice, maize and sorghum. The decline in production has resulted in reduced consumption, which could also be attributed to changing lifestyle and government policies. Drudgery associated with processing of the crop has also contributed to decline in area and production of the crop. However, because of its short crop cycle and ability to grow on a wide range of soil types, it may continue to remain a useful crop in Asia on poor agricultural land in regions with low rainfall or a short growing season. The prospects for barnyard millet in India seem limited, but the crop may gain importance as a niche crop in dry regions at medium to high altitudes under changing climate conditions. Millets have been referred as climate-resilient crops because of their greater ability to resist biotic and abiotic stresses. This offers an additional advantage for this crop, and the potential can be harnessed by concerted research efforts. Collective actions including public awareness on nutritional value, enhanced research on issues associated with crop production, processing and utilization, value addition and government support for marketing could save the crop for future.

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