



Evaluation of some potential silkworm *Bombyx mori* L. genotypes during different seasons under temperate conditions

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Received: October 2, 2016; Revised received: February 19, 2017; Accepted: May 27, 2017

Abstract: In the present study twelve potential bivoltine silkworm *Bombyx mori* L. genotypes were evaluated for their performance at 25 ± 1 °C temperature and 75 ± 5 % relative humidity for twelve different traits during spring and summer seasons of 2012 and 2013, respectively. The data generated in respect of different traits was pooled separately, analyzed statistically and subjected to multiple trait evaluation indexes. The genotypes were ranked as per the cumulative score and the value of a particular trait in a particular genotype was compared with the ranking. Out of twelve genotypes, six genotypes viz., SKAU-R-1, SKAU-R-6, SKUAST-31, NB₄D₂, SH₆ and SKUAST-28 were shortlisted for spring season and eight genotypes viz., SKAU-R-1, SKAU-R-6, NB₄D₂, SH₆, SKUAST-31, CSR₁₈, DUN₆ and DUN₂₂ for summer season. These genotypes scored higher E I values (>50) and were identified as promising genotypes hence recommended for rearing under temperate climatic conditions to push up silk productivity in the valley. Furthermore, the genotypes viz., SKAU-R-1, SKAU-R-6, SKUAST-31, NB₄D₂, SH₆, and DUN₆ performed significantly better irrespective of the seasons and scored higher E I. values (>50). Hence, these genotypes can be recommended for both seasons to boost bivoltine silk production in temperate region.

Keywords: Evaluation, Performance, Promising, Silkworm, Seasons, Trait

INTRODUCTION

Sericulture has an important place in the economy of Jammu and Kashmir as more than 30,000 rural families which belong to economically backward sections of the society are generating their employment through this vocation (Economic Survey, J&K 2014-15). Being one of the traditional and eco-friendly agro-based labor intensive industries of the state, helps in improving the economic conditions of landless farmers by providing subsidiary employment and supplementing the income of rural farmers especially the economically weaker section of the society (Qadri *et al.*, 2010). The state presents an ideal and fertile land for the growth and development of bivoltine sericulture. Though, the state is known for producing bivoltine silk of international quality. However, production of quality bivoltine silk is still a challenge in J & K having enormous potential to produce bivoltine silk of international grade, which can help to reduce the import of bivoltine silk in the country (Malik, 2009). Traditional breeding methods employed during the last few decades has resulted in the development of many productive silkworm breeds which have contributed significantly in maximizing the

silk production in India in general and Jammu and Kashmir state in particular. Of late, major thrust has been given for quality rather than quantity of silk produced. Efforts made in this direction during the 90's have lead to the evolution of highly productive CSR bivoltine breeds which have the potential to produce international grade silk (Datta, 2000). However, these new breeds continue to suffer badly in adverse conditions of low/high temperature, humidity, poor leaf quality and low management practices prevalent with the small and marginal farmers in Kashmir. Unlike tropics, temperate sericulture being carried out under highly fluctuating environmental conditions and poor leaf quality urgently needs the development of broad based silkworm breeds with genetic plasticity to buffer the adverse situations.

Evaluation of genetic resources is an essential prerequisite for their effective utilization in order to gauge the extent of variability among genotypes. Silkworms have been evaluated in many environment and agro-climatic conditions in order to identify the season and region specific breeds for utilization (Malik *et al.*, 2002). The necessity for identification of season/region specific breeds/ hybrids arises due to variation in quan-

titative characters during different environmental conditions. In India, large number of silkworm *Bombyx mori* L. breeds/hybrids were evolved by various breeders so far suitable for different agro-climatic conditions (Lakashmanan and Suresh kumar,2012; Lakshmi *et al.*, 2012; Gowda *et al.*, 2013;Senapati and Hazarika 2014; Mandal and Moorthy2015 and Gowda *et al.*, 2016). A series of studies have also been conducted to identify suitable bivoltine silkworm breed for Kashmir valley particularly for spring and autumn seasons (Nisar *et al.*, 2005; Malik *et al.*, 2006; Nisar *et al.*, 2008; Malik *et al.*, 2009, Malik *et al.*, 2010; Malik *et al.*, 2010a and 2010b). In the present study, an attempt was made to evaluate bivoltine germplasm based on Evaluation index method developed by Mano *et al.* (1993) which is one such method that increases the precision of selection of breed among an array of breeds by a common index giving due weightage to all the yield component traits (Bhargava *et al.*, 1994). The information generated will be useful to identify most promising genotypes for future breeding of genotypes suitable for temperate regions during spring and summer seasons.

MATERIALS AND METHODS

Twelve potential bivoltine mulberry silkworm *B. mori* L. genotypes namely; SKAU-R-1, SKAU-R-6, SKUAST-28, SKUAST-31, CSR₂, CSR₄, CSR₁₈, CSR₁₉, NB₄D₂, SH₆, DUN₆ and DUN₂₂ formed basis for this study (Table 1). The disease free laying's (DFL's) of these selected silkworm races were obtained from the Germplasm Bank of Temperate Sericulture Research Institute (TSRI), Sher-e-Kashmir University of Agricultural Sciences and Technology of Kashmir (SKUAST-K) Mirgund, Central Sericultural Germplasm Resources Centre (CSGRC) Hosour, Tamilnadu, India and Central Sericultural Research and Training Institute (CSR&TI) CSB, Pampore, Srinagar Kashmir. The eggs were incubated under hygienic conditions at 25±1 °C temperature and 75-80 % relative humidity for about 10-12 days till their hatching. The silkworms were fed with mulberry leaves harvested from the popular mulberry varieties viz; Goshoraemi and Ichinose maintained in Mulberry Germplasm Bank of TSRI, SKUAST-Kashmir, Mirgund. The study was undertaken during the year 2012 and 2013. The spring rearing was conducted in April-May while as summer rearing was conducted during July-August by following the standard package of practices (Krishnaswami, 1978).

The experiment was laid out in Completely Randomized Block Design with three replications for each treatment. Each replication comprised of 250 silkworms of uniform age and size retained after third moult. Rearing was carried out under hygienic conditions. At the end of 5th instar, the mature larvae were collected manually and mounted in plastic collapsible mountages. During the rearing period, larvae and co-

Table 1. Characteristic features of different bivoltine silkworm *Bombyx mori* L. genotypes under study.

Genotype	Volitinism	Parental Source	Larval pattern	Cocoon colour	Cocoon shape	Origin/Evolution	Source
SKAU-R-1		Shunrei × Shogetsu	Marked	White	Constricted		
SKAU-R-6		Shogetsu × Hoshu	Plain	White	Slightly oval		
SKUAST-28		Evolved Under Broad Based Germplasm Complex, Comprising 10 Breeds With Marked Larvae (Kamili <i>et al.</i> , 2000)	Marked	White	Short dumbbell	TSRI, SKUAST-Kashmir- Mirgund	Silkworm Germplasm Bank, TSRI, SKUAST-K, Mirgund
SKUAST-31		Evolved Under Broad Based Germplasm Complex, Comprising 10 Breeds With Plain Larvae (Kamili <i>et al.</i> , 2000)	Marked	White	Oval		
CSR ₂	Bivoltine	Shunrei × Shogetsu	Plain bluish	Bright white	Oval		
CSR ₄		(BN18×BCS25) × NB ₄ D ₂	Plain bluish	Bright white	Dumbell		
CSR ₁₈		B201 × BCS12	Plain and marked	Creamish white	Oval	CSR&TI, Mysore-India	Silkworm Germplasm Bank, CSGRC, Hosur-Tamilnadu, India
CSR ₁₉		B201 × BCS12	Plain and marked	Creamish white	Dumbell		
NB ₄ D ₂		(Kokko × Seihaku) × (N124 × C124)	Plain faint bluish	White	Elongated, constricted		
SH ₆		Shogetsu × Hoshu	Moderately marked	White	Oval	RSRS, Majira, Dehradun- India	Silkworm Germplasm Bank, CSR&TI, CSB-Pampore
DUN ₆		CC1 × NN6D	Plain	White	Oval	CSR&TI, Pampore-Kashmir	
DUN ₂₂		(KS × NB ₄ D ₂) × (AT × NB ₄ D ₂)	Marked	White	Oval		

Table 2. Mean performance of twelve silkworm *B. mori* L. genotypes during spring (Data pooled over same seasons of 2012 and 2013).

Genotype	Fecundity (no.)	Hatching (%)	V th age larval Duration (d:h)	Total larval Duration (d:h)	Weight of Ten Mature Larvae (g)	Single cocoon weight (g)	Single shell weight (g)	Pupal weight (g)	Shell ratio (%)	Filament Length (m)	Cocoon yield/10,000 larvae (kg)	
											by number	by weight
SKAU-R-1	577	95.24	7.06	27.09	50.19	2.09	0.45	1.63	21.45	1197.00	9367	19.23
SKAU-R-6	617	96.05	7.03	27.06	49.47	2.04	0.43	1.60	21.14	1168.00	9128	18.79
SKUAST-28	621	94.00	7.03	27.03	48.27	1.84	0.38	1.45	20.74	1060.50	8986	16.40
SKUAST-31	635	94.38	7.09	27.09	49.17	1.91	0.40	1.50	20.85	1148.00	9217	17.30
CSR ₂	519	92.52	7.03	27.02	40.53	1.71	0.34	1.37	19.94	898.17	8994	15.42
CSR ₄	514	93.73	7.06	27.08	43.48	1.74	0.35	1.38	20.07	917.48	8918	15.61
CSR ₁₈	516	94.58	6.22	25.10	40.78	1.69	0.34	1.34	20.17	928.23	8713	14.54
CSR ₁₉	533	95.35	6.19	25.08	39.70	1.67	0.32	1.34	19.24	920.17	8754	14.23
NB ₄ D ₂	582	94.87	7.03	26.09	48.87	1.89	0.39	1.49	20.70	1121.00	9128	16.95
SH ₆	613	95.89	7.01	26.08	42.17	1.84	0.38	1.45	20.60	1098.00	9069	16.19
DUN ₆	596	95.15	7.08	27.11	41.18	1.78	0.37	1.40	20.69	1019.50	9312	16.72
DUN ₂₂	588	94.53	7.06	27.09	40.13	1.76	0.36	1.39	20.40	1013.67	9187	16.29
Mean	575.92	94.69	6.91	26.58	44.5	1.83	0.38	1.45	20.5	1040.81	906.42	16.47
S.D	44.39	0.98	0.33	0.79	4.28	0.13	0.04	0.10	0.59	107.08	202.63	1.5
CD p≤0.05	6.51	0.13	0.87	1.25	0.32	0.60	0.61	0.45	0.15	23.91	9.72	0.58

Table 3. Mean performance of twelve silkworm *B. mori* L. genotypes during summer (Data pooled over same seasons of 2012 and 2013).

Genotype	Fecundity (no.)	Hatching (%)	V th age larval duration (d:h)	Total larval duration (d:h)	Weight of Ten mature Larvae (g)	Single cocoon weight (g)	Single shell weight (g)	Pupal weight (g)	Shell ratio (%)	Filament length (m)	Cocoon yield/10,000 larvae (kg)	
											by number	by weight
SKAU-R-1	582	92.43	7.01	26.10	43.65	1.74	0.36	1.37	20.60	998.23	9296	15.98
SKAU-R-6	623	93.03	7.01	26.09	45.17	1.76	0.38	1.38	21.53	1012.60	9279	16.67
SKUAST-28	629	91.02	7.03	27.01	42.23	1.70	0.32	1.37	18.76	934.62	8890	14.94
SKUAST-31	616	93.90	6.12	26.03	42.70	1.72	0.33	1.39	19.13	987.43	9001	15.13
CSR ₂	512	91.73	6.17	26.08	36.20	1.59	0.29	1.29	18.30	752.00	8598	14.11
CSR ₄	533	93.46	6.14	26.03	38.82	1.62	0.31	1.30	19.07	789.00	8617	14.26
CSR ₁₈	521	94.10	6.09	25.06	34.70	1.64	0.33	1.30	20.08	929.12	9288	15.56
CSR ₁₉	540	94.53	6.06	25.01	33.80	1.49	0.28	1.20	18.91	878.16	8907	13.49
NB ₄ D ₂	590	92.70	6.13	26.02	43.36	1.74	0.34	1.39	19.60	993.36	9023	15.43
SH ₆	609	95.02	6.11	26.01	39.87	1.73	0.33	1.39	19.12	975.97	9012	15.22
DUN ₆	603	94.98	6.21	26.08	38.48	1.67	0.32	1.34	19.22	896.61	9181	14.97
DUN ₂₂	594	94.93	6.19	26.06	37.40	1.64	0.32	1.33	19.45	883.17	9213	14.86
Mean	579.33	93.49	6.36	25.97	39.70	1.67	0.33	1.34	19.48	919.19	9025.42	15.05
S.D	41.70	1.33	0.40	0.51	3.74	0.08	0.03	0.06	0.88	83.71	242.8	0.85
CD p≤0.05	4.42	0.10	0.61	0.09	0.71	0.90	0.33	0.90	1.43	7.76	6.27	0.28

Table 4. Multiple trait evaluation index in respect of twelve silkworm *B. mori* L. genotypes during spring*.

Genotype	Fecundity (no.)	Hatching (%)	V th age larval duration (d:h)	Total larval duration (d:h)	Weight of Ten mature Larvae (g)	Single cocoon weight (g)	Single shell weight (g)	Pupal weight (g)	Shell ratio (%)	Filament length (m)	Cocoon yield/10,000 larvae		Average EI (%)	Rank
											By number	By weight (Kg)		
SKAU-R-1	50.24	55.62	45.37	43.50	63.31	69.43	69.51	69.37	66.21	64.59	64.93	68.41	60.87	I
SKAU-R-6	59.25	63.90	46.28	43.88	61.63	65.69	64.25	66.23	60.93	61.88	53.14	65.47	59.38	II
SKUAST-28	60.16	42.93	46.28	44.26	58.82	50.75	51.50	50.52	54.11	51.84	46.13	49.52	50.57	VI
SKUAST-31	63.31	46.82	44.45	43.50	60.93	55.98	56.36	55.76	55.98	60.01	57.53	55.52	54.68	III
CSR ₂	37.18	27.79	46.28	44.39	40.73	41.03	40.57	42.15	40.47	36.68	46.52	42.97	40.56	XII
CSR ₄	36.05	40.17	45.37	43.63	47.63	43.27	43.20	43.19	42.68	38.48	42.77	44.24	42.56	X
CSR ₁₈	36.50	48.87	70.90	68.69	41.32	39.54	40.57	39.00	44.39	39.49	32.66	37.10	44.92	IX
CSR ₁₉	40.33	56.74	71.81	68.94	38.79	38.04	35.31	39.00	28.53	38.73	34.68	35.03	43.83	XI
NB ₄ D ₂	51.37	51.83	46.28	56.16	60.22	54.48	53.73	54.71	53.42	57.49	53.14	53.19	53.83	IV
SH ₆	58.35	62.27	46.88	56.29	44.57	50.75	51.10	50.52	51.72	55.34	50.23	48.11	52.18	V
DUN ₆	54.52	54.70	44.76	43.25	42.25	46.26	48.47	45.29	53.25	48.01	62.22	51.65	49.55	VII
DUN ₂₂	52.72	48.35	45.37	43.50	39.80	44.77	45.83	44.24	48.31	47.47	56.05	48.78	47.10	VIII

* Data pooled over same seasons of 2012 and 2013

Table 5. Multiple trait evaluation index in respect of twelve silkworm *B. mori* L. genotypes during summer*.

Genotype	Fecundity (no.)	Hatching (%)	V th age larval duration (d:h)	Total larval duration (d:h)	Weight of Ten mature larvae(g)	Single cocoon weight (g)	Single shell weight (g)	Pupal weight (g)	Shell ratio (%)	Filament length (m)	Cocoon yield/10,000 larvae		Average EI (%)	Rank
											By number	By weight (kg)		
SKAU-R-1	50.64	42.05	33.67	47.37	60.56	58.90	62.60	55.66	62.69	59.44	61.14	60.95	54.64	II
SKAU-R-6	60.47	46.57	33.67	47.57	64.63	61.45	69.97	57.41	73.24	61.16	60.44	69.08	58.80	I
SKUAST-28	61.91	31.44	33.17	29.67	56.77	53.82	47.85	55.66	41.83	51.84	44.42	48.69	46.42	IX
SKUAST-31	58.79	53.12	55.89	48.74	58.02	56.36	51.54	59.15	46.02	58.15	48.99	50.81	53.80	V
CSR ₂	33.85	36.78	54.64	47.76	40.65	39.83	36.79	41.72	36.61	30.03	32.40	38.91	39.16	XII
CSR ₄	38.89	49.81	55.39	48.74	47.65	43.64	44.16	43.47	45.34	34.45	33.18	40.68	43.78	X
CSR ₁₈	36.01	54.62	56.64	67.61	36.64	46.18	51.54	43.47	56.79	51.19	60.81	56.00	51.46	VI
CSR ₁₉	40.57	57.86	57.38	68.58	34.23	27.11	33.10	26.04	43.53	45.10	45.12	31.61	42.52	XI
NB ₄ D ₂	52.56	44.08	55.64	48.93	59.79	58.90	55.22	59.15	51.35	58.86	49.90	54.47	54.07	III
SH ₆	57.11	61.55	56.14	49.12	50.46	57.63	51.54	59.15	45.91	56.78	49.45	51.99	53.90	IV
DUN ₆	55.68	61.25	53.64	47.76	46.74	50.00	47.85	50.44	47.04	47.30	56.41	49.05	51.10	VII
DUN ₂₂	53.52	60.87	54.14	48.15	43.86	46.18	47.85	48.69	49.65	45.70	57.73	47.75	50.34	VIII

* Data pooled over same seasons of 2012 and 2013

Table 6. Overall evaluation index of two seasons (spring and summer -2012 and 2013) for different traits in twelve silkworm *B. mori* L. genotypes.

Genotype	Fecundity (no.)	Hatching (%)	V th age larval duration (d:h)	Total larval duration (d:h)	Weight of mature larvae(g)	Single cocoon weight (g)	Single shell weight (g)	Pupal weight (g)	Shell ratio (%)	Filament length (m)	Cocoon yield/10,000 larvae		Average EI (%)	Rank
											By number	By weight (kg)		
SKAU-R-1	50.44	47.46	36.27	44.74	62.23	66.21	67.14	65.30	65.71	62.73	65.51	66.63	58.36	II
SKAU-R-6	59.90	54.52	36.95	45.06	63.24	64.74	67.14	63.89	70.42	61.94	58.59	67.76	59.51	I
SKUAST-28	61.07	34.20	36.61	37.95	57.99	51.97	49.74	52.64	46.36	51.90	44.21	49.17	47.82	VI
SKUAST-31	61.18	50.52	50.82	45.38	59.73	56.39	54.48	57.56	50.00	59.49	53.47	54.05	54.42	III
CSR ₂	35.49	30.34	51.16	45.54	40.54	40.17	38.66	41.39	36.79	33.23	36.52	41.01	39.24	XI
CSR ₄	37.35	45.06	51.16	45.38	47.60	43.12	43.41	42.79	43.63	36.28	34.98	42.54	42.77	IX
CSR ₁₈	36.19	52.52	66.04	69.29	38.96	41.65	44.99	39.98	52.05	44.45	47.60	43.58	48.11	X
CSR ₁₉	40.39	58.53	67.06	69.77	36.45	33.30	33.92	32.95	36.11	41.25	38.39	32.84	43.41	XII
NB ₄ D ₂	51.96	46.96	51.83	53.46	60.19	56.39	54.48	56.86	52.43	58.35	51.66	53.87	54.04	IV
SH ₆	57.80	63.68	52.51	53.62	47.27	53.44	51.32	54.04	48.03	56.17	49.76	49.49	53.09	V
DUN ₆	55.11	59.78	49.46	44.74	44.25	47.54	48.15	47.01	49.47	47.63	60.91	50.75	50.40	VII
DUN ₂₂	53.52	56.42	50.14	45.06	41.55	45.09	46.57	45.60	49.01	46.58	58.40	48.32	48.86	VIII

* pooled mean of spring and summer seasons of 2012 and 2013

coons were assessed for different parameters viz; fecundity, hatching percentage, larval weight, larval durations (fifth age and total), cocoon yield by weight and by number, single cocoon weight, single shell weight, pupal weight, shell ratio and filament length, during spring and summer seasons (2012 and 2103). The data generated in respect of different traits (Table 2 and 3) was pooled separately, analyzed statistically and subjected further to multiple trait evaluation index method as per the procedure outlined by Mano *et al.* (1993).

$$EvaluationIndex(EI) = \frac{(A - B)}{C} \times 10 + 50$$

Where, as evaluation index for Vth age and total larval durations was computed separately by using the modified formula given by Talebi and Subramanaya (2009).

$$EvaluationIndex(EI) = \frac{(B - A)}{C} \times 10 + 50$$

Where, A = Value of a particular breed for particular trait,

B = Mean value for a particular trait of all the breeds,

C = Standard Deviation of a particular trait for all the breeds,

10 = Standard unit,

50 = Fixed value. Minimum/average EI value fixed for selection of a breed is >50. The evaluation index value for negative traits viz., Vth age larval duration and total larval duration was calculated separately. The index value obtained for the entire negative traits was combined and the average EI values were obtained. The EI value fixed for the selection of breed / genotype is 50 or >50 for positive traits and 50 or <50 for negative traits. The genotypes, which scored above the limit, were considered to possess greater economic value.

RESULTS AND DISCUSSION

Multiple trait evaluation indices (EI) assessment is the multiple performance of a population for selection/short-listing of the genotypes/hybrid combinations by taking into consideration all the economic traits. The data recorded from twelve bivoltine silkworm *B. mori* L. genotypes in respect of different traits viz., fecundity, hatching, larval duration, cocoon weight, shell weight, pupal weight, shell ratio, filament length and cocoon yield by number and by weight was subjected to multiple trait evaluation as per the procedure outlined by Mano *et al.* (1993). Based on the performance of the 12 silkworm genotypes, individual indices were calculated for each of the 12 parameters. Evaluation index values were calculated for each of the genotype in all the 12 parameters for both spring and summer seasons. The data of which is presented in Tables 4-5. The indices obtained from all the traits in respect of each genotype were combined and the average EI values were assessed. The criteria for selection of the genotype was based on the average EI value >50. The gen-

otypes which scored above the limit of 50 in many of the traits were considered to possess greater economic value.

During spring, out of 12 genotypes, 6 genotypes scored the average EI value >50. SKAU-R-1 occupied the top position with average EI score value of 60.87 while as in summer, 8 genotypes out of 12 scored average EI value >50. SKAU-R-6 recorded the highest average E. I. value of 58.80 among the 12 genotypes evaluated during summer. However, the analysis of data indicates that all the genotypes utilized in the study vary significantly with respect to most of the parameters studied during spring and summer seasons. During spring, NB₄D₂ scored E.I values >50 for the maximum of 11 traits viz., fecundity, hatching percentage, total larval duration, larval weight, single cocoon weight, single shell weight, pupal weight, shell ratio, filament length, cocoon yield by weight and number. However, this genotype occupied fourth position in the average evaluation index score with an average E.I of 53.83. SKAU-R-1 and SKAU-R-6 scored E.I indices >50 for 10 characters. SKUAST-31 and SH₆ obtained E.I values >50 for 9 characters each (Table-4). While in summer, SKAU-R-1, SKAU-R-6, SKUAST-31, NB₄D₂, and SH₆ recorded E.I values >50 for the maximum of 9 characters each. Whereas, SKUAST-31, NB₄D₂, and SH₆ recorded E.I values >50 for other 9 traits followed by CSR₁₈, DUN₆ and DUN₂₂ which obtained E.I values >50 for 8, 6 and 4 traits respectively (Table-5). Furthermore, evaluation indices obtained irrespective of the seasons revealed that out of 12 genotypes evaluated, 7 genotypes scored average E.I values >50. SKAU-R-6 scored average E. I value of 59.51. Out of 12 characters evaluated, SKUAST-31 and NB₄D₂ recorded E.I values >50 for the maximum of 11 traits. However, these genotypes occupied third and fourth position in the average evaluation index score with an average E.I of 54.42 and 54.04 respectively. SKAU-R-6 recorded E.I values >50 for 10 characters. SKAU-R-1 and SH₆ which obtained E.I values >50 for 9 and 8 characters respectively (Table-6).

In the recent past evaluation index method developed by Mano *et al.* (1993) has been utilized for short listing some promising silkworm genotypes/hybrids for commercial exploitation in different seasons under Kashmir climatic conditions (Malik *et al.*, 2009; Malik *et al.*, 2010; Malik *et al.*, 2010a; 2010b; Maqbool *et al.*, 2015 and Nooruldin *et al.*, 2015) and the same has been utilized in the present study as well for evaluating 12 bivoltine silkworm *Bombyx mori* L. genotypes in respect of different traits viz., fecundity, hatching, larval duration, cocoon weight, shell weight, pupal weight, shell ratio, filament length and cocoon yield by number and by weight. All the top ranking genotypes recorded > 95 % hatching which is well supported by the earlier findings (Nisar *et al.*, 2013). The

healthiness of larvae is a very important character from the point of view of silkworm rearers and as such stabilization of cocoon crop is very important for the sericulture industry. The genotypes that have shorter larval duration have less chance to get infected with diseases (Basavaraja *et al.*, 2005). In the present study, out of 12 genotypes two genotypes viz., CSR₁₈ and CSR₁₉ recorded shorter larval duration below 26 days irrespective of seasons (Table 2-3). The single cocoon weight in the top ranking genotypes ranged from 1.84 g in SKUAST-28 to 2.09 in SKA-U-R-1 during spring (Table-2) while it ranged from 1.64 g in DUN₂₂ to 1.76 g in SKA-U-R-6 during summer (Table-3). High cocoon shell weight is an important trait for high productivity. The cocoon shell weight shows variability in different environments. According to Mano *et al.* (1993), if the breed is showing cocoon shell weight of 0.45 g and above, it becomes weak and not suitable for summer rearing. The identified genotypes for spring and summer climatic conditions recorded shell weight in the range of 0.38 g (SKUAST-28) to 0.45 g (SKA-U-R-1) in spring while in summer it ranged from 0.32 g (DUN₂₂) to 0.38 g (SKA-U-R-6). The length of cocoon filament is one of the important attributes of the silkworm breed/hybrid. Longer the filament better, it is to the filature and textile industry. In Japan, almost all the commercial races have been synthesized to satisfy this. The present study identified spring and summer specific genotypes having filament length in the range of 1060 m to 1197 m and 883 m to 1012 m respectively. Cocoon yield is another very important character connected with the production of cocoons by Sericulturists. It is necessary to have high yielding silkworm genotypes to raise the farmer's income. During spring, the top ranking genotypes recorded cocoon yield/10,000 larvae by number in the range of 8986 to 9367 and yield by weight in the range of 16.40 kg to 19.23 kg. While in summer it ranged from 9213 to 9279 by number and 14.86 kg to 16.67 kg by weight, respectively (Table. 2-3).

In silkworm *Bombyx mori* L. large numbers of breeds were tested and promising ones were selected based on the economic traits (Lakshmi *et al.*, 2012; Gowda *et al.*, 2013; Mandal *et al.*, 2013; 2015; 2016 and Gowda *et al.*, 2016). Evaluation index is one such method that increases the precision of selection of breed among an array of breeds by a common index giving due weightage to all the yield component traits (Bhargava *et al.*, 1994). The silk yield is contributed by more than 21 traits (Thiagarajan *et al.*, 1993) and there exists an interrelationship between multiple traits in silkworm. Any effort to improve the yield requires consideration of cumulative effect of the major traits, which influences the silk yield impartially. To judge the superiority of the silkworm breeds, a common index method is required (Bhargava *et al.*, 1994, Mano *et al.* 1993). A selection index makes it possible to select for a charac-

ter by selecting simultaneously for two or more characters related to it. Obviously, the present investigation on performance of some potential bivoltine silkworm genotypes with reference to important economic traits has yielded rich information, to identify promising breeds which can be recommended for commercial exploitation in the interest of the temperate bivoltine silk industry for pushing up the quality silk production.

Conclusion

Based on the performance and evaluation indices of 12 silkworm *B. mori* L. genotypes during spring and summer seasons, six genotypes viz., SKAU-R-1, SKAU-R-6, SKUAST-31, NB₄D₂, SH₆ and SKUAST-28 were shortlisted for spring season and eight genotypes viz., SKAU-R-1, SKAU-R-6, NB₄D₂, SH₆, SKUAST-31, CSR₁₈, DUN₆ and DUN₂₂ for summer season. These genotypes scored higher E I. values (>50) and have been identified as promising genotypes hence recommended for rearing under spring and summer climatic conditions of temperate region to push up silk productivity in the valley. Based on the overall performance of genotypes during both seasons (spring and summer) SKAU-R-1, SKAU-R-6, SKUAST-31, NB₄D₂, SH₆, and DUN₆ performed significantly better irrespective of the season and scored higher E I. values (>50). Hence, these genotypes can be reared in both seasons to boost bivoltine silk production in the temperate region.

ACKNOWLEDGEMENTS

The first author wishes to thank Directorate of Research, Sher-e- Kashmir University of Agricultural Sciences and Technology of Kashmir, Srinagar-India for financial support. The author sincerely acknowledges Director, CSGRC Hosur, Director, CSR and TI Pampore (Central Silk Board, Ministry of Science and Technology, Govt. of India) and Head, TSRI, SKUAST-K, Mirgund for providing silkworm germplasm laboratory facilities to conduct silk worm rearing.

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