# **Right to information Act, 2005**

# 4 THE GAZETTER OF INDIA EXTRAORDINARY [PART II-4]

# **CHAPTER II**

# Right to information and obligation of public authorities

3. Subject to the provisions of this Act, all citizens shall have the right to information.

- 4. (1) Every public authority shall
  - a) maintain all its records duly catalogued and indexed in a manner and the form which facilitates the right to information under this Act and ensure that all records that are appropriate to be computerized are, within a reasonable time and subject to availability of resources, computerized and connected through a network all over the country on different systems so that access to such records is facilitated;
  - b) publish within one hundred and twenty days from the enactment of this Act

# (i) The particulars of its organization, functions and duties;

# **Organization:**

SERI-BIOTECH RESEARCH LABORATORY CENTRAL SILK BOARD, MINISTRY OF TEXTILES GOVERNMENT OF INDIA, KODATHI, CARMELRAM P.O. BENGALURU – 560 035, KARNATAKA

### Mandate:

The Seri-Biotech Research Laboratory [SBRL] was established during 1993 under the World Bank aided National Sericulture Project as per the advice of a high level committee headed by Prof. Lynn Riddiford, University of Washington, USA to carry out research in the frontier areas of biology for the development of the sericulture industry. The following are the broad mandates of the laboratory:

> To conduct research in frontier areas of modern biology and to seek potential applications of these work towards improving silk productivity.

> To interact with other institutions doing basic or applied research in areas related to sericulture and other allied areas.

> To disseminate technology developed to the target groups through the other R & D constituents of CSB.

### **Functions:**

Presently the laboratory is implementing externally funded and in-house research projects as indicated below:

# Host-Plant Biotechnology

Identification, characterization and utilization of genes/markers involved in disease resistance and abiotic stress tolerance, work was carried out related to molecular characterization and identification of various mulberry species and other host plants, development of microsatellites for mulberry, characterization of drought resistant genes, study on microspore embryogenesis etc.

### Silkworm Biotechnology

The focus is on identification of silkworm genes and their functions associated with resistance to viral and fungal pathogens, development of NPV tolerant silkworm lines based on marker assisted selection and RNAi technique, regulation of yolk proteins, characterization of RNA dependent RNA polymerase gene, characterization of oviposition stimulating factors in silkworms and Identification of molecular markers associated with filament characters and its use in improvement of multivoltine breeds, whole genome sequencing and transcriptome analysis of golden silk moth etc.

The laboratory is also focusing on identification and molecular characterization of various pathogens like virus, bacteria, microsporidia etc. infecting silkworms and development of diagnostic tools for their easy and early detection. Development of Densovirus resistant marker assisted selection productive breeds through and Transkingdom RNA interference (tkRNAi) approach for resistance against BmNPV infection in silkworm. Research is also carried out on Identification of immune response proteins and their interactions, under stress from pests like uzi fly, under microsporidia infection etc. for improvement of silkworm breed.

### Silk Biomaterials

Programmes on Characterization of Sericin for Cosmetic Applications in collaboration with CSTRI, Bangalore, M/s Hindustan Lever and work on development of fibroin fusion silk with antimicrobial, antioxidant and UV protective properties is on-going.

### 1. Significant achievements

SBRL has been implemented 54 research projects supported by CSB, DBT, DST and international funding. The brief achievements since inception are as indicated below:

### 1.1. <u>Research highlights</u>

- Recombinant silk Fibroin- Cecropin B, a fusion protein was expressed in a heterologous expression system. This fusion protein was effective against gram-positive and gram negative bacteria. It has shown enhanced wound healing activity in rats and human dermal cells. The fusion protein has also shown strong anti- oxidant activity.
- Transgenic CSR2 races over-expressing immune gene Relish 1 and antifungal Drosomycin B are developed. These transgenic silkworms showed enhanced resistance/tolerance against pathogenic bacteria and microsporidian infection. Utilization of the transgenic silkworms enhance silk production by reducing the crop loss due to pathogens.
- Bacteria-based dsRNA is developed for RNA interference (RNAi) of transcripts of NPV genes ie-1 and lef-1 involved in viral DNA multiplication. Feeding bacterially expressed dsRNA led to restriction of the viral multiplication in the silkworm. The results obtained indicates successful use of non-pathogenic bacteria as a vector for delivering dsRNA to elicit RNAi against BmNPV infection in silkworm and suggest as an alternative tool for insect pest management.
- Screening for molecular markers associated with silk filament traits in *B. mori* larva showed influence of genes encoding ion transport (ITP), VATPase, CPR genes and SG metabolism pathway genes on efficient processing and transportation of silk proteins from posterior to anterior silk gland. Further a deletion region is linked to filament length, cocoon weight and shell weight. The deletion can be utilized as marker for screening germplasm to identify multivoltine hybrids with better post cocoon traits and its improvement.
- The genome of viral pathogen (Accession: GI: 1371952746) infecting oak tasar silkworm Antheraea proylei has six conserved sequences

and it displayed homology with alpha-baculovirus (AnprNPV) infecting saturniid silkworms. Presence of the virus on the surfaces of *A. proylei* eggs was confirmed through PCR showing vertical transmission from the infected eggs as the source of infection.

- The RNA virus known as Iflavirus causing flacherie in tropical tasar Antheraea mylitta was characterized through whole genome sequencing. The multiplication of virus was detected in the various tissues such as fat body, midgut, Malpighian tubule & Ovary. Transmission of the virus from mother moth to the offspring is confirmed.
- Under the DBT sponsored project a new hybrid (MASN) was developed using marker assisted selection against NPV infection. The rearing performances of newly developed bivoltines and cross breeds were compared with the ruling varities at different sericultural areas under CSRTI of Mysore, Berhampore and Pampore. The MASN4 X CSR4 hybrids yielded 53.5 -63 Kg /100DFLs against the yield of 51% in the ruling variety at observed farmers under CSRTI Berhampore. Similarly, the cross breeds Nistari x MASN4 yielded 52kgs/100DFLs whereas the ruling variety yielded 48.5kg/100DFLs. The performance of newly developed hybrids was significantly higher than the ruling variety.
- Mildew resistance locus O (MLO) proteins from mulberry were analyzed for motifs and domains in comparison with functionally characterized MLOs from other dicot species and phylogeny was established for identification of clade-V MLO genes from mulberry. This analysis led to identification of candidate MLO genes involved in powdery mildew resistance in mulberry.
- Whole genome of the wild and reared Indian muga silkworm, Antheraea assamensis is sequenced and transcriptome analysis to identify differentially expressed genes in response to pathogen and silk gland specific genes are in progress.
- > Three lines of transgenic silkworms with construct for

overexpression of immune genes relish1, drosomycin and cecropinB were synthesized from pure Mysore and CSR2 for enhanced immunity.

- Causal organism of tiger band diseases in oak tasar silkworm, Antheraea proyeli identified as an Antheraea proyeli nucleopolyhedrovirus (AnprNPV).
- Developed NPV tolerant MASN4, MASN6 and MASN7 breeds through marker assisted backcrossing using Sarupat as donor and CSR2 as a recurrent parent.
- Developed DNV-2 resistant improved CSR6, CSR26, MASN6, MASN7 and J2P breeds.
- Effect of dsRNA against NPV multiplication shown that 30-40% increased survivability of NPV infected larvae in comparison to without dsRNA treatment.
- Utilizing RNAi technique NPV resistance gene from NPV resistant transgenic Nistari multivoltine lines has been introgressed to NPV susceptible CSR4 and CSR27 bivoltine lines which revealed 30% enhanced resistance to NPV.
- The immune response genes viz. Gloverin, SERPIN, Hsp 23.7 and Hsp 40 revealed significant up-regulation in NPV resistant genotype [Sarupat] suggesting that these genes could play an important role in baculovirus resistance in *B. mori*.
- DNV-2 infection in *B.mori* destructively damaged the midgut tissue suggesting the necessity to carry out systematic survey for DNV-2 in flacherie prone areas to devise appropriate control measures.
- B.mori germplasm possessing the nsd-2 gene for resistance to DNV-2 was identified that can be utilized to strengthen silkworm breeding programmes.
- SSU-rRNA gene sequence was found to be highly useful to decipher the correct taxonomic status of the 14 different microsporidians identified from the silkworms and different insect pests harboring various mulberry fields, the latter possibly being the main source

for cross infecting silkworms.

- Six different microsporidians identified from tasar silkworms had clear homology with *Nosema* species.
- RNAi silencing of yolk protein receptors disrupted ovary growth, yolk deposition and developments indicating their role in egg development. VgR gene expression levels were high in Pure Mysore, Sarupat, CSR2, CSR26 and NB4D2.
- The Indian isolate of DNV-2 was identified and found to be severely widespread under field conditions causing flacherie disease.
- Twenty four host-response and immune proteins from uzi infected B. mori and 3 from S. ricini were identified, while, 3 proteins were observed to interact with Hsp70 a major stress protein induced after uzi fly infection in the integument and hemocytes most of which were activated in the early stages of infection and based on this outcome, immunocompetent B. mori strains can be identified for breeding programs.
- The Eri silkworm larvae exhibited innate defence mechanism by forming cysts around the infected uzi maggot killing it by inhibiting growth.
- Identified and characterized several Simple Sequence Repeats (SSR) and Inter-simple sequence repeats (ISSR) for the first time from tropical strains of the silkworm *B.mori*.
- Identified markers viz. RAPD, RFLP, SSR and ISSR were utilized for characterizing 60 multivoltine and 90 bivoltine *B.mori* using specific primers.
- One hundred and fifty mulberry germplasm were characterized using various DNA marker systems to determine their taxonomic status and association of the markers with important characters.
- Twenty eight SSR markers were developed for characterization of diapausing and non diapausing silkworms for higher & lower cocoon characters.
- Silkworm DNA markers for yield attributes and larval duration (one)

with potential use in Marker Assisted Breeding program were identified.

- Studies were attempted for the first time to understand the genetic relationship of the ecoraces of muga and tasar silkworms using well defined marker systems like SSRs.
- Fifteen Eri silkworm populations studied showed genetic homogeneity and less variability with positive correlation of genetic distance between populations with geographic distance and negative with altitude.
- Localized differentiation of Eri silkworm subpopulations noticed and recommended Eri population increase by widening area of localized populations.
- Wild tasar ecorace, Jatta Daba was identified as a primitive strain from which other strains have originated by adapting to different environment in Jharkhand and the muga Selsela population from the West Garo Hills as heterogenous population, both of which need to be protected from natural disasters.
- ISSR marker studies revealed taxonomic complexities of mulberry germplasm identifying highly divergent cultivars based on genetic distance for use as ideal material for breeding program
- First mulberry genetic linkage map was constructed with S36 x V1 mapping populations based on which dense genetic linkage map can be developed and association as well as linkage mapping studies can be taken up.
- Fifty five ISSR and 179 RAPD putative/species specific markers were identified for existence of microsporidians in addition to *Nosema bombycis*.
- Twenty five RAPD putative/species specific markers were identified to detect various microsporidians.
- A PCR based multiplex detection system has been developed for the simultaneous detection of microspridia, nucleopolyhedrosis virus, and densovirus. The technology has been filed for patenting with

NRDC, New Delhi. No. 1746/CHE/2008 and published in Patent Journal by NRDC: 29/01/2012. The technique is being used in NSSO Basic Seed Farm Mysore to diagnose pebrine disease in the silkworm seed.

- Silkworm brain-specific lipophorin receptor (a key molecule functioning in reproduction and development) variant gene (LpR4) identified to have a role in signal transduction apart from uptake of lipids which is a new finding in insect lipoprotein research aiding in future related studies.
- Diapause specific genes have been identified in diapause induced multivoltine silkworms through semi-quantitative polymerase chain reaction (PCR), Suppressive Subtractive Hybridization and Microarrary analysis. The multivoltine silkworm germplasm can be screened based on expression of these genes and subjected to diapause induction for cost effective conservation.
- DNA markers viz. Nag 34, Nag 65, Nag 84 and Nag 88 were identified to be associated with NPV resistance and were mapped on chromosome numbers 12, 5, 7 and 3. Introgression of these markers in NPV susceptible CSR2 lines through Marker Assisted Selection to develop NPV tolerant bivoltine lines and their field trials are in progress so that, they can be utilized as parents for making commercial F1 hybrids.
- Hsp70 was identified as a major stress protein induced after uzi fly infection in the integument and hemocytes of *B. mori* larvae.

# 1.2. <u>Publications</u>

SBRL has published more than 100 research papers in peer reviewed international and national journals (Attached as Annexure 1), copies of which can be obtained on request.

# 2. <u>LIST OF R & D PROJECTS (1994 – July 2022)</u>

SN	Project Code	Project title	Project
	_		duration
1	ARP-08007MI	Biological and molecular characterization of	March
	CSB	virosis in Muga silkworm, (Antheraea	2022-
		assamensis Helfer)	February
			2025
2	AIB-08008MI	Development and Evaluation of Eri silkworm	March
	CSB	(Samia ricini Donovan) breeds/ hybrids with	2022-
		improved productivity	February
			2025
3	AIC08009CN	Profiling of lipid, protein and carbohydrate of	April 2022-
	CSB	mulberry mealybug Maconellicoccus hirsutus	March
		(Green)	2025
4	AIT08010M	Transcriptome studies for susceptibility of	April 2022-
	CSB	Muga Silkworm (Antheraea assamensis) to	March
		diseases.	2026
5	AIT 08006 EF	Development of lateral flow assay (LFA) kit for	March
	(Indo-Swedish	diagnosis of pebrine disease in silkworms	2021-
	project)	January	
	100000000		2023
6	AIT08003CN	Gene Expression Profiling for the	August
	DBT	Identification of Resistant/Tolerant Genes to	2019- July
		Microsporidian Infection in Lamerin Breed of	2022
7	AIT08005MI	Silkworm, Bombyx mori L.	
1	CSB	Development and Evaluation of Bidensovirus March	
	COD	resistant silkworm hybrids developed from 2020-Feb marker assisted breeding lines -Phase II 2023	
8	PIT08004MI	Study on Epigenetic and autophagy modifiers	March
0	CSB		2020- Feb
	COD	oninductionofhaploidmicrospore2020- Febembryogenesis in mulberry2023	
9	PRP08002MI	Identification of powdery mildew resistant	May 2019-
	CSB	genes and validation of CAPS marker for	April-2022
		Chalcone synthase	
10	ARP- 08001 CI	Studies on the genetic characterization,	April 2018
	(Indo-Swedish	transmission and tissue distribution of	to March
	project)	Iflavirus infecting the Indian tropical tasar 2021	
	· · · · ·	silkworm, Antheraea mylitta	

11	ARP 3606	Development of diagnostic tool for early	Feb. 2017-
11	DBT	detection of baculovirus causing tiger band	Feb. 2017-
	DDI	disease in Antheraea Proylei	
10		Validation of the DNA markers in silkworm Feb. 2017-	
12	ARP 3605		
	DBT	breed developed by introgression of DNA	Feb. 2020
		markers associated with NPV resistance using	
		marker Assisted Selection Breeding and large	
		scale field trial of the breed	
13	AIT-3582	Development of Densovirus Resistant	Sept. 16 –
	CSB	productive Bivoltine Silkworm breeds through	Aug. 18
		Marker Assisted Selection	
14	AIT-3584	Identification of molecular markers associated	Sept. 16 –
	CSB	with filament characters and its use in	Aug. 19
		improvement of multivoltine breeds (In	U
		collaboration with CSTRI, Bangalore	
15	AIT-3583	Transkingdom RNA interference ( <i>tk</i> RNAi)	Sept. 16 –
	CSB	approach for resistance against BmNPV	Aug. 19
	0.02	infection in silkworm Bombyx mori L (in	
		collaboration with University of Delhi)	
16	CFC-7064	Characterization of Sericin for Cosmetic	Apr.15 –
10	CSB	Applications (CSTRI, Bangalore, SBRL, M/s	Mar.16
	CSD	Hindustan Lever)	Mai.10
17	ARP-3522	Isolation, Cloning and Characterization of Apr.15 –	
17	CSB		Mar.18
	CSB	Antibacterial Protein (s) from Silkworm,	Mar.18
10		Bombyx mori (CSR&TI, Berhampore & SBRL)	A 1 L
18	AIT 3540	Development of transgenic silkworms for the	Apr.15 –
	CSB	over expression of disease-resistant genes for	Mar.18
		enhanced immunity [in collaboration with	
10		IISc, Bangalore]	
19	AIT-3538	Development of fibroin fusion silk with	Apr.15 –
	CSB	antimicrobial, antioxidant and UV protective Mar.1	
		properties	
20	SBRL003	Identification of uzifly maggot tissue protein	Dec.14 -
	CSB	that induces toxicity in silkworm Bombyx mori	Nov.15
	Pilot study		
21	SBRL002	Male Accessory Gland proteome analysis and	Oct.14 -
	CSB	characterization of Oviposition Stimulating	Sep.16
	Pilot study	substances (OSS) from Tasar Silkworm,	-
	2	Antheraea mylitta	
22	SBRL001	Cloning and expression of three drought	Oct.14 -
	CSB	resistance genes in mulberry ( <i>Morus</i> spp.)	Sep.15
23	ARP 3158	Expression profiling of genes associated with	Oct.14 -
	CSB	resistance to <i>Beauvaria bassiana</i> in <i>Bombyx</i> Sep.17	
		mori strains (SBRL & CSGRC)	~~P.1.
24	ARP 3513	Molecular characterization of Indian isolate (s)	Jun.14 -
47	DST-JSPS	of Densovirus (DNV) and viral resistance gene	May'16
	001-0020		may 10
		in the host silkworm, Bombyx mori (SBRL	

		Kodathi, National Institute of Agrobiological	
		Sciences, Japan)	
25	SBRL004 CSB	Discerning (Exploring) the mechanism of resistance and its pathways using microarray technique in silkworm <i>Bombyx mori</i> associated with nucleopolyhedrovirus ( <i>Bm</i> NPV)	Apr.14 - Mar.15
26	AIB 3503 CSB	Identification of autumn-specific silkwormNov.13 -breeds/hybrids suitable for sub-tropical zonesOct.16of North and North West India (CSB funded -with CSR&TI Pampore, NSSO Bangalore,CSTRI Bangalore)CSTRI Bangalore	
27	ARP 3495 CSB	Development of immuno-molecular techniques for early diagnosis of major infectious pathogens of silkworm, Bombyx mori L.	
28	ARP 3494 DBT	Horr D.Host-parasiteinteraction:TranscriptomeJan.13 -responses to parasitism in the silkwormJuly16Bombyx mori.[Jan.13-Dec.16]	
29	ARP 3489 CSB	Identification and molecular characterization of major pathogens associated with flacherie disease in Antheraea mylitta (in collaboration with CTR&TI Ranchi)Oct.12 Sep.15	
30	ARP 3477 CSB	Therapeutic control of microsporidiosis in mulberry silkworm through characterization of methionine amino peptidase enzyme genes (MetAP2) in Nosema bombycis (in collaboration with CSR&TI Mysore)Jul.12 Jun.1	
31	AIT 3468 DBT	Development of RNA interference (RNAi) basedSep.nuclearpolyhedrosisvirusresistancetransgenicsilkmoth.[(Jointly with CDFD,Hyderabad,APSSRDI,Hindupur)(UnderCenterofexcellenceonGenomicsofSilkmothtoPhase IIIIIIIIII	
32	AIG 3473 CSB	Molecular characterization of the flacherie causing virus in Bombyx mori with specific reference to RdRp (RNA Dependent RNA polymerase) gene and the regulatory elements in the viral genome.Jan.12 - Dec.14	
33	PIG-3465 CSB	In the viral genome.2011-Isolation and characterization of Microsatellites in mulberry (Morus spp.)2014genome.2014	
34	ARP 3453 CSB	Identification, isolation and molecular characterization of major pathogens associated with flacherie disease in <i>Bombyx</i> <i>mori</i> .	Apr.11 – Mar.13

35	AIT 3448	Studies on differential gene expression	Oct.10 –		
55	CSB	profiling of Bombyx mori	Mar.15		
	COD	Nucleopolyhedrovirus (BmNPV) resistant and	Mar.10		
		susceptible <i>Bombyx mori</i> silkworm strains.			
36	AIT 3446	Cloning, expression and characterization of Jun.10			
	DBT	yolk protein receptors from Indian silkworms. May'13			
37	ARP 3429	Biology of microsporidians infecting silk moth	Jun.09 –		
	DBT	[Bombyx mori and Antheraea mylitta]	May'12		
38	AIT 3428	Molecular mechanism of stress in silkworms	2009-		
	DBT	Bombyx mori and Samia cynthia ricini	2012		
39	AIT 3427	Studies on diapauses related gene expression	Jun.09 –		
	DBT	in diapauses induced eggs of multivoltine	Jun.12		
		silkworm races of <i>Bombyx mori</i> .			
40	AIG 3426	Identification and mapping of DNA markers	Jun.09 –		
	DBT	linked to NPV resistance in silkworm <i>Bombyx</i>	Jun.12		
		mori L. [Jointly with APSSRDI, Hindupur]			
41	Pilot Project CSB	PCR-based detection of silkworm diseases	2006-2009		
42	CSB	Functional Characterization of a brain-	2006-2010		
		specific lipophorin receptor variant from the			
		silkworm, Bombyx mori			
43	DBT	Phylogeography of Antheraea mylitta (tropical	Oct.05 –		
		tasar silkworm) and Antheraea assamensis	Oct.08		
		(muga silkworm) (In collaboration with CDFD,			
		Hyderabad; CMERTI, Jorhat and CTRTI,			
		Ranchi).			
44	ARP-3351	Studies on the morphology, life cycle and	Jul.05 –		
	CSB	pathogenicity of some microsporidia infecting May'08			
		silkworm, Bombyx mori L. and their			
		identification (Collaborative project with			
		CSR&TI, Berhampore)			
45	CSB	Construction of genetic linkage maps and QTL	2005-2008		
		analysis of economically important traits in			
10	005	mulberry.	A 0 F		
46	CSB	Characterization of Eri silkworms (Samia	Apr.05 –		
		<i>ricini</i> ) with morphological characters and	Mar.08		
		molecular markers (In collaboration with CEMRTI, Jorhat).			
47	DBT	Identification of DNA markers for baculovirus	Dec.04 –		
<b>Τ</b>		resistance in silkworm, <i>Bombyx mori</i> L.	Mar.08		
		(Network project with collaboration from	mar.00		
		CDFD, Hyderabad; CSRTI, Mysore; APSSDI,			
		Hindupur and KSSDI, Bangalore).			
48	CSB	Molecular characterization of microsporidian	Jun.04 –		
		infecting commercial silkworms	May'09		
			J		

49	CSB	Genome Analysis in Muga silkworm host 2004-2007 plants (Som & Soalu): DNA profiling of certain elite genotypes using molecular markers, development of mapping population and linkage map.(In Collaboration with CMERTI, Jorhat)	
50	DBT	Genetic analysis on <i>Wolbachia</i> affecting the natural parasites of domesticated silkworm, <i>Bombyx mori</i> and possibility on the analysis of its use as a vector) (In collaboration with Bangalore University, Bangalore).	
51	CSB	Studies on genetic basis of hardiness in silkworm and MAS program.Apr.01- Mar.04	
52	CSB	Molecular genetics of differential growth and yield potential in silkworm Bombyx moriApr.01- Mar.04	
53	CSB	Dissection of mulberry genome: further Apr.01- studies on genetic variability and Mar.04 characterization of genes associated with productivity.	
54	CSB	Molecular breeding in silkworm using DNA Apr.98 - markers. Mar.01	
55	CSB	Molecular characterization of mulberry and silkworm germplasm – Phase IApr.96 Mar.9	
56	DBT	RFLPpolymorphismanalysisforNPVDec.95 –resistance in silkworm, Bombyxmori.Nov.98	
57	DBT	MolecularcharacterizationofsilkwormDec.94varietiesusing micro and minisatelliteDNADec.97	
58	CSB	Genome analysis of silkworm, <i>Bombyx mori</i> Jan.94 - Mar.98	

### 3. Training

SBRL, Kodathi conducts training programmes that include project dissertation works and molecular biology techniques for M.Sc. / B. Tech. / M.Tech. students on charge basis depending on the type of programme. Under specified projects research fellows are appointed for the project period and are permitted to do Ph.D. under the University of Mysore as per the University rules and regulations.

# (ii) The powers and duties of officers and employees

The brief information on powers and duties of officers and employees

working at SBRL are given below. For more details kindly visit http://csb.gov.in/right-to-information/rti-2/central-office/

SN	Designation	Power and duties		
1	Director	Over all in-charge of Research and		
		Administration of the laboratory		
2	Scientist-D	Research and development works connected with		
		technical aspects and administration		
3	Scientist-C	Research and development works connected with		
		technical aspects and administration		
4	Scientist-B	Research and development works connected with		
		technical aspects and administration		
5	Asst. Director	Over all in-charge of administration, Accounts etc.		
6	Superintendent	FAS, PRS and processing all the purchase		
		proposals, settlement of TA/TTA/LTC/Medical		
		reimbursement bills, Maintenance of SR, dead		
		stock/Assets register etc.		
7	Technical	Technical assistance and support in all Research		
	Assistants	and Development works		
8	Asst.	Diary, despatch, circulars circulations and		
	Technician	operating Xerox machine		
9	Staff Car Driver	Driving car for official work for staff and visiting		
		experts/dignitaries, maintenance of vehicle and its		
		utilization record		
10	Multi Tasking	Maintenance of cleanliness in around laboratory		
	Staff	and office premises		

### **Research Scholars on contract basis**

11	Junior Research	Assisting in Research and Development works	
	Fellow/ Research	related to research projects	
	Scholar		

### **ORGANIZATION CHART**

(Attached as Annexure 2)

# (iii) The procedure followed in the decision making process, including channels of supervision and accountability;

As per the rules and regulations laid down by Central Silk Board and

Government of India. For more details kindly visit (http://csb.gov.in/right-to-information/rti-2/central-office/)

# (iv) The norms set by it for the discharge of its functions;

As per the rules and regulations laid down by Central Silk Board and Government of India. For more details kindly visit (http://csb.gov.in/right-to-information/rti-2/central-office/)

# (v) The rules and regulations, instructions, manuals and records, held by it or under its control or used by its employees for discharging its functions;

General Financial Rules, Fundamental Rules and Supplementary Rules Part-I, II & III and CCA/CCS Conduct Rules etc.

# (vi) A statement of the categories of documents that are held by it or under its control;

Accounts :	Cash book, Day book, Ledger, Financial statement of Accounts, Assets register	
Establishment :	Personal files, Service Records,	
	Correspondence files	
General Administration :	Purchase files, stock registers, dispatch And dairy registers, general correspondence files	

(vii) The particulars of any arrangement that exists for consultation with, or representation by, the members of the public in relation to the formulation of its policy or implementation thereof.

-NA-

(viii) A statement of the boards, councils, committees and other bodies consisting of two or more persons constituted as its part or for the purpose of its advice and as to whether meetings of those boards, councils, committees and other bodies are open to the public, or the minutes of such meetings are accessible for public.

### Boards, councils, committees

### 1. Research Advisory Committee

<b>Dr. N. K. Krishnakumar,</b> Retired Deputy Director General (Horticulture) Indian Council of Agricultural Research (ICAR) New Delhi	Chairperson
<b>Dr. Sanjay Ghosh</b> Institute of Bioinformatics and Applied Biotechnology Bangalore	Member
<b>Dr. Mohan</b> Principal Scientist Division of Entomology, NBAIR, Bangalore	Member
<b>Prof. Nataraj Karaba</b> Dept. of Crop Physiology, UAS, GKVK,	Member
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# Bangalore

2	
<b>Dr. K. Vijayan</b> Scientist D (Retired) Central Silk Board Bangalore	Member
<b>The Director</b> CSR&TI, Mysore	Member
<b>The Director</b> CSGRC, Hosur	Member
<b>The Director</b> SBRL, Kodathi Bangalore-560035	Member- Convenor
2. Research Council	
Director	Chairperson
All Scientists	Members
3. Stores Purchase Committee	
Dr. K.M. Ponnuvel, Scientist- D	Member
Dr. A. Ramesha, Scientist- C	Member
Dr. K. S. Tulsi Naik, Scientist- C	Member
Dr. Pawan Shukla, Scientist- C	Member
Dr. Himanshu Dubey, Scientist- B	Member
Dr. Manjula S., Superintendent	Member
4. Institute Bio-safety Committee	
<b>Dr. P. J. Raju</b> Director, APSSRDI, Andhra Pradesh	Chairperson
<b>Director</b> Seri-biotech Research Laboratory, Bangalore	Member Convener

**Prof. Upendra Nongthomba** Dept. of Molecular Reproduction 18

Member

and Developmental genetics Indian Institute of Science, Bangalore	
<b>Dr. R. Ashokan</b> Principal Scientist Dept. of Biotechnology, IIHR, Hesaraghatta. Bangalore	DBT Nominee
<b>Dr. H. K. Basavaraja</b> Retd. Scientist, Central Silk Board, Bangalore	Member
<b>Dr. Raghunath</b> Medical Officer (MBBS), Govt. Medical Hospital, Kodathi, Sarjapura Road, Bangalore	Member
<b>Dr. K. M. Ponnuvel</b> Scientist D, Seri biotech Research Laboratory, Bangalore	Member (Internal Expert)
<b>Dr. A. R. Pradeep</b> Scientist D, Seri biotech Research Laboratory, Bangalore	Member (Internal Expert)

# (ix) A directory of its officers and employees.

A directory of Officers and employees is enclosed (Attached as <u>Annexure 3</u>)

# (x) The monthly remuneration received by each of its officers and employees, including the system of compensation as provided in its regulations.

The monthly remuneration received by each Officer / employee is enclosed (Attached as <u>Annexure 4</u>)

# (xi) The budget allocated to each of its agency, indicating the particulars of all plans, proposed expenditures and reports on

disbursements made.

-NA-

(xii) The manner of execution of subsidy programmes, including the amounts allocated and the details of beneficiaries of such programmes.

-NA-

(xiii) Particulars of recipients of concessions, permits or authorisations granted by it.

-NA-

(xiv) Details in respect of the information, available to or held by it, reduced in an electronic form.

Available on the SBRL website.

(xv) The particulars of facilities available to citizens for obtaining information, including the working hours of a library or reading room, if maintained for public use

-NA-

# (xvi) The names, designations and other particulars of the Public Information Officers

**Details of Central Public Information Officer (CPIO)** 

Dr. K. M. Ponnuvel, Scientist D & In-charge Seri-Biotech Research Laboratory, CSB Campus, Sarjapur Road, Carmelram Post, Kodathi, Bangalore - 560 035, Karnataka. Email- sbrl@rediffmail.com, sbrlban.csb@nic.in Phone- +91 80 29519997

# **Details of Appellate Authority**

Shri. R. R. Okhandiar, IFS Member Secretary, Central Silk Board, Bangalore. Phone- +91 80 26680190; Email- ms.csb@nic.in

# (xvii) Such other information as may be prescribed; and thereafter update these publications every year.

Annual report

# List of research publications

- 1. Esvaran VG, Ponnuvel S, Jagadish A., Savithri, HS, Subramanya, HS, Ponnuvel KM. Cloning, Expression and Characterization of Spore Wall Protein 5 (SWP5) of Indian Isolate NIK-1S of Nosema bombycis. The Protein Journal, 2022, 41(6), 596-612.
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- 3. Raghavendar G, Gupta T, Ramesha A, Sivaprasad V, Ponnuvel KM. A real-time qPCR method for early detection of Bombyx mori Bidensovirus (BmBDV) infection in silkworm. Animal Gene. 2022 Jul 14:200132.
- 4. Ponnuvel KM, de Miranda JR, Terenius O, Li W, Ito K, Khajje D, Shamitha G, Jagadish A, Dubey H, Mishra RK. Genetic characterisation of an Iflavirus associated with a vomiting disease in the Indian Tropical tasar silkworm, *Antheraea mylitta*. Virus Res. 2022 Jan 30:198703.
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- 8. Jagadish A, Khajje D, Tony M, Nilsson A, de Miranda JR, Terenius O, Dubey H, Mishra RK, Ponnuvel KM (2021). Development and optimization of a TaqMan assay for Nosema bombycis, causative

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- Shambhavi H. Prabhuling, Pooja Makwana, A.R. Pradeep, K. Vijayan and R. K. Mishra (2021) Release of mediator enzyme β-hexosaminidase and modulated gene expression accompany hemocyte degranulation in response to parasitism in the silkworm *Bombyx mori*. Biochemical Genetics. 59(4)997-1017.
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- 23. Vijaya Gowri Esvaran, Anupama Jagadish, Olle Terenius, Siripuk Suraporn, Rakesh K Mishra and Kangayam Ponnuvel\* (2020) Targeting essential genes of *Nosema* for the diagnosis of pebrine disease in silkworms. Annals of Parasitology 66: 303-310; \* corresponding author
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- 33. Pawan Shukla1\*, Ramesha A. Reddy1, Kangayam M. Ponnuvel, Gulab Khan Rohela, Aftab Ahmad Shabnam, S. S. Chauhan, Mrinal Kanti Ghosh, Rakesh Kumar Mishra (2018) Comparative analysis of gene expression profiles among contrasting mulberry varieties under cold stress condition. J. Exp. Biol. Agric. Sci. Vo.6(6):p 973-982. 1Co-First authors
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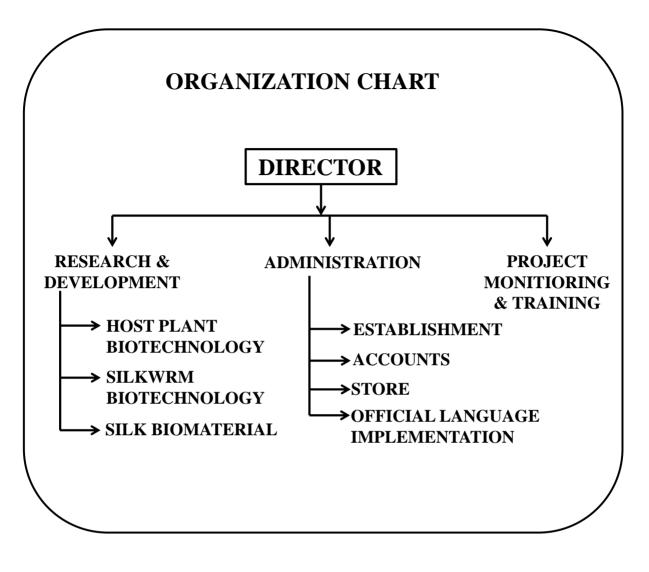
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# Directory of officers and employees

S.		DESIGNATIO	Contact details
N.	NAME	N	
1	Dr. K.M. Ponnuvel	Scientist-D & Director (Incharge)	Phone: +91 80 29519997 Email: sbrlban.csb@nic.in sbrl@rediffmail.com
2	Dr. K.S. Tulsi Naik	Scientist-C	tulsinaik.csb@gov.in
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12	Shri. Srinivas Raju K.V.	Field Assistant	srinivasrajuster@gmail.com
13	Shri K. Nanjundappa	Staff car driver	NA
14	Shri Kenchappa	MTS	NA
15	Shri N. Pillappa	SFW (TS)	NA
16	Shri E. Shivanna	SFW (TS)	NA
17	Smt. Kanthamma	SFW (TS)	NA
18	Shri. R. Amaresha	SFW (TS)	NA
19	Shri L.P. Sampangi Rao	Time Scale farm worker	NA

# Monthly remuneration received by each Officer / employees

S.N.	NAME	DESIGNATION	Gross Salary (Rs.)
1	Dr. K.M. Ponnuvel	Scientist-D & In-charge	201,006.00
2	Dr. K.S. Tulsi Naik	Scientist-C	132,036.00
3	Dr. A. Ramesha	Scientist-C	132,036.00
4	Dr. Pawan Shukla	Scientist-C	132,036.00
5	Dr. G. Subrahmanyam	Scientist-C	132,036.00
6	Dr. Rajal Debnath	Scientist-C	132,036.00
7	Dr. Himanshu Dubey	Scientist-B	111,081.00
8	Mrs. Manjula S.	Supdt. (Admin)	120,468.00
9	Shri. K.M. Humayun	STA	113,868.00
10	Shri R.N. Sreekantaiah	STA	107,598.00
11	Shri. G. Raghavendar	Field Assistant	47,868.00
12	Shri. Srinivas Raju K.V.	Field Assistant	47,868.00
13	Shri K. Nanjundappa	Staff car driver	90,768.00
14	Shri Kenchappa	MTS	62,718.00
15	Shri N. Pillappa	SFW (TS)	37,838.00
16	Shri E. Shivanna	SFW (TS)	37,838.00
17	Smt. Kanthamma	SFW (TS)	37,838.00
19	Shri. R. Amaresha	SFW (TS)	37,838.00
20	Shri L.P. Sampangi Rao	Time Scale farm worker	16,450.00

# [As on 30.11.2022]