

10, Institutional Area, Vasant Kunj, New Delhi 110070

# MINUTES OF THE 50<sup>th</sup> MEETING OF ACADEMIC COUNCIL HELD ON 08<sup>th</sup> JANUARY 2022 AT 11:00 A.M.

The 50<sup>th</sup> meeting of Academic Council was held on 08<sup>th</sup> January 2022 at 11:00 hours on Microsoft Team Platform.

The following were present:

#### Members

Professor Prateek Sharma, Chairperson Mr Manoj Chugh Professor Shaleen Singhal Professor Arun Kansal Professor Anandita Singh Professor Ramakrishnan Sitaraman Professor Vinay Shankar Prasad Sinha Professor Nandan Nawn Dr Naqui Anwer Dr Sukanya Das Dr Anu Rani Sharma Dr Montu Bose Dr Chander Kumar Singh Dr Seema Sangita, Controller of Examination Mr Kamal Sharma, Secretary Special Invitees

Dr Shashi Bhushan Tripathi Dr Chaithanya Madhurantakam Mr Manish Shrivastava

Professor T C Kandpal, Professor Vivek Suneja, Professor Arun Kharat, Mr Rajesh Ayapilla, Mr Rahul Mittal and Professor Manipadma Datta could not attend the meeting.

Before starting the proceedings, the Registrar while welcoming the members, informed that Prof. Eklabya Sharma resigned from the services of the TERI SAS on 21 November 2021 and Prof. Prateek Sharma has been appointed as the Acting Vice Chancellor by the Chancellor with effect from 23 November 2021. Prof Prateek Sharma joined TERI SAS in August 2007 and has held the positions of Head, (DNR), Associate Dean, (Faculty of Applied Sciences) and Dean (Academic).

Prof Prateek Sharma welcomed the Board members before requesting the Registrar to take up the agenda items.

- Item No.1: To confirm the minutes of the Forty ninth Meeting of the Academic Council held on 17<sup>th</sup> July 2021. The Registrar informed that minutes of the Forty-Ninth Meeting of the Academic Council, held on 17<sup>th</sup> July 2021, were circulated to the members and few minor comments received from the internal members have been incorporated. Hence, the Council might confirm the minutes.
- **TS/AC/50.1.1** The Council resolved that the minutes of the 49<sup>th</sup> Academic Council Meeting held on 17<sup>th</sup> July 2021 be confirmed.
- Item No. 2: To consider and approve Promotion Rule of Students and Grade Conversion. The Controller of Examination explained to the members that TERISAS has received several requests from students and alumni for a grade conversion formula over the last few years. Hence, there is a need for a policy on grade conversion. In order to finalize a policy, three faculty members undertook the task of researching and reviewing policies followed in other institutions (detailed in Enclosure 1) in the context of the comments received on this matter in the past AC meetings. A range of different options were narrowed down to four, and their merits and demerits were assessed (detailed in Enclosure 2) and presented for consideration in the internal AC meeting. Based on the discussions in the internal AC meeting, a presentation on the proposal for grade conversion was made. This proposal is based on following principles:
  - CGPA and percentage systems of evaluation of students are fundamentally different in characteristics and there is no uniform and scientifically rigorous method of conversion. Hence, the final decision is based on precedence in other institutions and universities like ours.

- The policy of "Grade Conversion" may be delinked with "Continuation of Registration/eligibility of degree requirements" (reasons are elaborated in **Enclosure 2**)
- Set the notional conversion from CGPA to percentage with a formula of multiplication by a factor of 10.<sup>1</sup>
- CGPA of 6 and above may be notionally considered first division

An extensive discussion was carried out on the merits and demerits of this proposal

- 1. The presentation showed data on the minimum, maximum and average CGPA of each program for the last two years to demonstrate that only a selected few students would score a 90% or higher after conversion. It was pointed out that this may not be sufficient. The entire distribution may be studied for at least three years and may be compared with similar outcomes in other universities. In view of the time that it would take to undertake such a task, it was suggested that this approach be considered while drafting any future iterations of the policy.
- 2. There was a detailed discussion on specific details as well as merits and demerits of the proposal.
- 3. After reflecting on the circumstances, it was felt that the proposal would be an acceptable way forward in the best interest of the future of our students and alumni. Further, it was based on precedence set by other similar institutions. Majority of the AC agreed with this conclusion. Prof. Nandan Nawn expressed his disagreement regarding the second part of the conversion statement, that is, notional consideration of CGPA of 6 and above as first class.

#### Final conversion from CGPA to percentage policy text:

TERI-SAS adopts the following formula for <u>notional</u> conversion of CGPA to percentage terms for the purpose of internship/employment/scholarships/higher education, etc. for all its graduates since the inception of the university:

# *Percentage = 10\*[CGPA]*

A CGPA of 6 and above may be <u>notionally</u> considered to be first class.

<sup>&</sup>lt;sup>1</sup> This is option 1 in Enclosure 2. The merits and demerits are outlined there.

- **TS/AC/50.2.1** The Academic Council resolved to approve the final conversion from CGPA to percentage and first class of CGPA 6 and above as presented by the Controller of Examinations.
- Item No. 3: To approve the number of seats in various programmes. Prof Arun Kansal, Dean (Academic) informed that after internal discussion with the Head of Departments, it is proposed to reduce the number of seats to 20 from present 25 in respect of MSc (Geoinformatics) and MSc (Biotechnology) programmes due to the infrastructure constraint and lab space requirements. However, for MSc (Economics) it is proposed to increase the seats from present 40 to 60 due to the high number of applications received. For other programmes, there is no change in the number of seats. The Academic Council is requested to consider and approve the number of seats with respect to the above mentioned three programmes.
- **TS/AC/49.3.1** The Academic Council resolved to approve the revised number of seats in respect of the above mentioned three programmes.
- Item No. 4: To consider and approve courses to be offered in Second Semester of M.Sc. Biotechnology Programme. Dr Shashi Bhushan Tripathi presented and requested the Academic Council to approve the courses to be offered in second semester of M.Sc (Biotechnology) programme, as placed at Enclosure 3. He also further stated that each of the proposed course
  - is in accordance with the programme structure approved by the AC
  - has been reviewed by atleast two subject experts
  - has been discussed and recommended by the Board of Studies
- TS/AC/50.4.1 The Academic Council resolved to approve the courses to be offered in the second semester of MSc (Biotechnology) programme as placed at Enclosure 3.
- Item No. 5: To consider and approve revised course outline of "Climate Change and Law" offered to M.Sc. Climate Science and Policy. Dr Manish Shrivastava informed that this course was earlier offered by the Centre for Postgraduate Legal Studies with course code MPL134 which has been discontinued from AY 2021-22. Few suggestions and recommendations of BoS were incorporated in the course structure as placed at Enclosure 4 and requested the Academic Council for its approval.

- **TS/AC/50.5.1** The Academic Council resolved to approve the revised course outline of "Climate Change and Law" as placed at **Enclosure 4**.
- Item No. 6: To obtain concurrence on approval of resolution by circulation. The Registrar informed that in the 49th Academic Council meeting, it was decided to align the course credits to 15 hour per credit for all programmes as per the UGC rules. According to this decision, all the programmes has gone through their respective Board of Studies and carried out the required changes in each course(s) and the summary of the changes were circulated to the Academic Council members on 25 August 2021 for approval. After the Council's approval through circulation, it was implemented with effect from 31 August 2021.

The members noted the matter.

There being no other items for discussion, the meeting was adjourned with a vote of thanks to the Chair at 1440 hours.

Sd/ Kamal Sharma Registrar (Acting)

Enclosures:-

- Enclosure 1 Review of other institutions' grade conversion policies
- Enclosure 2 Summary of the background review of different options
- Enclosure 3 Courses to be offered in Second Semester of M.Sc. Biotechnology Programme
- Enclosure 4 Revised course outline of "Climate Change and Law"

Distribution:-

Electronic Copy:

- 1. Vice Chancellor, TERI School of Advanced Studies
- 2. All members of Academic Council
- 3. Website

Printed Copy: Registrar Office

#### **Enclosure 1**

#### **REVIEW OF OTHER INSTITUTIONS' GRADE CONVERSION POLICIES**

- 1. Different institutes have different grade conversion formulae. Policies in IIT Delhi, Delhi University, JNU, Mumbai University, IIT Bhubaneswar, IIT Dhanbad, IIT Hyderabad, IIITM Gwalior, AMU, and Amity University were reviewed.
  - a. Among these universities, IIT Delhi, Amity University, IIT Bhubaneswar, IIT Hyderabad, IIITM Gwalior multiply by a factor of 10. Most universities seem to use this straightforward conversion rule.
  - b. IIT Dhanbad uses this formula: Percentage = (CGPA-0.5) \*10.
  - c. Mumbai University: *Percentage* = 7.1\**CGPA* + 11.
  - d. Delhi University (undergraduate): *Percentage* = 9.5\*CGPA
  - e. JNU: Percentage = 5 + 10 \* CGPA
- 2. None of the institutions explain the logic behind the choice of conversion formula.
- 3. The review did not reveal any perfect way to map the CGPA to percentage as these two methods of assessment of a student are fundamentally different in characteristics. In fact, two institutions IIT Bhubaneswar and Amity University clearly state that there is no rigorous formula to convert CGPA to percentage terms. But, they accept method of multiplication of CGPA by 10 if a student requires a conversion of CGPA to percentage.

#### **References:**

- 1. IIT Delhi
  - a. They use a direct conversion of factor of 10
  - https://home.iitd.ac.in/uploads/IITD-Courses-of-Study-2021-22\_(08-09-2021).pdf
     pg 17 (4.2.2) indicates that different programs have different eligibility criteria. Regardless all use the same conversion rate.



अतुल व्यास Atul Vyas उप कुलसचिव Deputy Registrar



#### TO WHOM IT MAY CONCERN

The CGPA of all IIT Delhi graduates notionally be converted to percentage by multiplying the CGPA by a factor of 10. This is applicable for all graduates since 18.10.1982.

For the purpose of employment or requirement of any external body that IIT Delhi graduate wishes to join, a CGPA of 6.0 or above be taken as First Class.



Note: This certificate will not be issued to the individual candidates/ agencies for now onwards. It may be downloaded from the website.

# 2. Delhi University

a. http://exam.du.ac.in/pdf/11012018/11012018\_CGPA.pdf

UNIVERSITY दिल्ली विश्व	OF DELHI विद्यालय
Ref. No. Dean (Exams)/2017/ 912-4	20 <sup>th</sup> December, 2017
NOTIFICA	TION
It is notified for information to all concerned that as p	per the deliberations of the Committee
constituted to prepare the formula for conversion	of Cumulative Grade Point Average
(C.G.P.A., into percentage for the final year stud	lents of the Undergraduate Courses
under the Choice Based Credit System and in	pursuance of the approval of the
competen: authority, the University of Delhi sha calculating the final percentage of marks:	all adopt the formula as below for
Final Percentage of marks (%) = C.G.P.A. ba	sed on all six semesters x 9.5
The above formula shall be applicable w.e.f. from th	e Undergraduate examinations under
the Choice Based Credit System (CBCS) to be h	eld in May/June 2018 and onwards.
(for the first batch of students admitted under the Ch 2015)	cice Based Credit System in the year
Only the conversion formula will be printed on the G	rade Certificates/Transcripts.
	DEAN (EXAMINATIONS)
University of Delhi, Main Campus, Delhi- Tel:27001957/27667934, Website	10007 (India) duac in

- 3. JNU
  - $a. \ https://www.jnu.ac.in/sites/default/files/Conversion\%20 certificate.pdf$

fter deliberations, reso irector, Internal Quali ercentage of marks wh	olved to approve Formula-II, out ty Assurance Cell for conversio ich is given below;	of other formulas suggested n of Grade Point Average
ormula - II:-		
In this formula a a CGPA of 3 is e	CGPA of 5 is equivalent to 55% quivalent to 35% nversion table suggested by 1Q2	
CGPA	Procedure of Conversion	Result/Percentage
CGPA CGPA of 8-9	Procedure of Conversion 5+CGPA x 10	Result/Percentage
CGPA CGPA of 8-9 CGPA of 7-7.99	Procedure of Conversion 5+CGPA x 10 5+CGPA x 10	Result/Percentage         3           85%-95.99%         75%-84.99%
CGPA CGPA of 8-9 CGPA of 7-7.99 CGPA of 6-6.99	Procedure of Conversion 5+CGPA x 10 5+CGPA x 10 5+CGPA x 10	Result/Percentage         3           85%-95.99%         75%-84.99%           65%-74.99%         65%-74.99%
CGPA CGPA of 8-9 CGPA of 7-7.99 CGPA of 6-6.99 CGPA of 5-5.99	Procedure of Conversion           5+CGPA x 10           5+CGPA x 10           5+CGPA x 10           5+CGPA x 10	Result/Percentage         3           85%-95.99%         75%-84.99%           65%-74.99%         55%-64.99%
CGPA CGPA of 8-9 CGPA of 7-7.99 CGPA of 6-6.99 CGPA of 5-5.99 CGPA of 4-4.99	Procedure of Conversion           5+CGPA x 10	Result/Percentage         3           85%-95.99%         75%-84.99%           65%-74.99%         55%-64.99%           45%-54.99%         45%-54.99%
CGPA CGPA of 8-9 CGPA of 7-7.99 CGPA of 6-6.99 CGPA of 5-5.99 CGPA of 4-4.99 CGPA of 3-3.99	Procedure of Conversion           5+CGPA x 10	Result/Percentage         3           85%-95.99%         75%-84.99%           65%-74.99%         55%-64.99%           45%-54.99%         35%-44.99%
CGPA CGPA of 8-9 CGPA of 7-7.99 CGPA of 6-6.99 CGPA of 5-5.99 CGPA of 4-4.99 CGPA of 3-3.99 CGPA of 2-2.99	Procedure of Conversion           5+CGPA x 10	Result/Percentage           85%-95.99%           75%-84.99%           65%-74.99%           55%-64.99%           35%-44.99%           25%-34.99%
CGPA CGPA of 8-9 CGPA of 7-7.99 CGPA of 6-6.99 CGPA of 5-5.99 CGPA of 4-4.99 CGPA of 3-3.99 CGPA of 2-2.99 CGPA of 1-1.99	Procedure of Conversion           5+CGPA x 10           5+CGPA x 10	Result/Percentage           85%-95.99%           75%-84.99%           65%-74.99%           55%-64.99%           35%-44.99%           25%-34.99%           15%-24.99%

- 4. Amity
  - a. <u>https://www.amity.edu/placement/Popup.asp?Eid=441</u> (Amity uses a factor of 10 for placement purposes)
  - b. <u>https://img0cf.b8cdn.com/images/course/70/1869170\_1552134911.pdf</u> (towards the end of the transcript, they state that conversion from CGPA to percentage has no rigor or rationale)
- 5. Mumbai University
  - a. <u>https://mu.ac.in/wp-content/uploads/2014/03/conversion-Circular\_2017.pdf</u>
  - b. I am not quite sure of the rationale of this formula.
- 6. IIT Bhubaneshwar
  - a. <u>https://www.iitbbs.ac.in/notice/noticeb\_1516190404.pdf</u>
  - b. They state that there is no formula for conversion but students can use multiplication by 10 if they need to.
- 7. IIT Dhanbad
  - a. <u>https://www.iitism.ac.in/assets/uploads/news\_events/admin/CGPA-to-</u> Percentage.pdf
- 8. IIT Hyderabad
  - a. <u>https://www.iith.ac.in/academics/assets/files/forms/CGPA-to-percentage-conversion.pdf</u>

- b. Section 3.5.4 indicates that D is passing grade which carries 4 points.
- 9. IIITM Gwalior
  - a. <u>https://www.iiitm.ac.in/images/2019/June\_2019/Academics\_2019/Conversion-</u> <u>Certificate-CGPA-1.pdf</u>
  - b. <u>https://www.iiitm.ac.in/images/2019/June\_2019/Academics\_2019/Prospecus2019</u> <u>final.pdf</u> (D is the passing grade with 4 points, and the minimum CGPA required is 5, see pg 25 to 27

# Enclosure 2

# SUMMARY OF THE BACKGROUND REVIEW OF DIFFERENT OPTIONS

Discussions in the previous AC meetings presented the dilemma of whether to benchmark the "traditional pass percentage" of 40% with a CGPA of 4 which is the passing grade of each course or with a CGPA of 6 which is the minimum requirement of award of degree. This dilemma is difficult to resolve. The "additional illustration" in the footnote gives one example of challenges faced.<sup>2</sup> Four different options were studied with the objective of trying to find a resolution, keeping in mind the feedback received in the previous AC meetings.

The following is a summary of 4 strategies reviewed and discussed in the internal meeting:

# **OPTION 1: Delink Conversion Formula and Promotion Criteria. Regardless of promotion criteria, use a rule of multiplication by 10 for conversion.** MERITS

- This approach bypasses the dilemma described above.
- Many universities and engineering institutions follow this approach. In the absence of availability of methodology for a scientifically rigorous conversion between CGPA and percentage, the next best approach may be to follow the norms in the sector.
- The choice of multiplication by 10 (as opposed to say 9.5 or 9) is justified by our moderate pattern of final CGPAs where a selected few students make it to a CGPA>9. Thus, only a selected few would score 90% or more after conversion.
- Ease of implementation, as there would be one uniform rule across cohorts.

Suppose CGPA 6 is benchmarked as 40% (traditional pass percentage) after 2018. Then using the same approach CGPA 5.5 ought to be benchmarked as 40% prior to 2018. Now, if we consider two students, both with CGPA of 6, one graduated in 2019 and another in 2017, then these two students would have different percentage marks. The former would have a percentage score of exactly 40% while the later would have a percentage score higher than 40%

Therefore, use of benchmarks based on "minimum CGPA for award of degree" in any method for percentage conversion or for estimating class divisions would not be consistent across cohorts.

<sup>&</sup>lt;sup>2</sup> Additional illustration: An example to explain the challenges of benchmarking with the minimum CGPA requirement for award of degree. The minimum CGPA for award of degree was 5.5 prior to 2018 and 6 thereafter.

- In the circumstance that no perfect solution that could address all the feedback of previous AC meetings could be found, this solution is the second best approach
- Divisions can be pegged at 1<sup>st</sup> class or 1<sup>st</sup> division for 60% and above, as is the traditional norm in Indian Universities.

#### DEMERITS

- This is not based on a strict scientific rationale but is based only on what several institutions in India appear to follow.
- The question of 10 and why not 9.5 or 9 also does not have a strict scientific rationale
- Choice of divisions also does not have a strict scientific rationale.

# **OPTION 2:** Allow award of degree to students CGPA of 4 and multiplication by 10 is the conversion formula with 4 = 40% as the passing benchmark.

This option was presented in last AC meeting, and was reviewed in light of the comments received MERITS

# • Dilemma of benchmarking the "traditional pass percentage" to 4 or 6 is resolved by creating an equivalence of grade point for passing a course and CGPA of award of degree. In this option, both would be set at 4.

• The formula of multiplication by 10 for percentage marks and also setting up the divisions as per the norms of traditional Indian university system of first class/division for 60% of higher is in alignment with this option

# DEMERITS

- As pointed out in the last AC meeting, reduction of the minimum CGPA required for award of degree from 6 to 4 may be perceived as reduction in the academic standards of the university.
- This will not provide a conversion for current and past cohort of students for whom the minimum CGPA required for award of degree cannot be changed retrospectively.

OPTION 3: Y = 14.5\*X - 47'', where Y is the equivalent percentage and X is the CGPA/SGPA and CGPA of 6.0 is made equivalent to 40% marks and CGPA of 10.0 is made equivalent to 98% marks<sup>3</sup>

CGPA	Equivalent %	Division	CGPA range	<b>Conversion Formula</b>
6.00	40.00			
6.50	47.25	Pass without	6.0 ≤ CGPA <6.69 40 < Percentage <50	
6.69	50.01			
6.70	50.15			
7.00	54.50	2nd division	$6.7 \leq CGPA < 7.4$ $51 \leq Percentage < 60$	Y = 14.5 * X - 47
7.40	60.30			
7.50	61.75			
8.20	71.90	1st division	7.5 ≤ CGPA < 8.4 61 ≤ Percentage < 74	$\mathbf{X} = \mathbf{CGPA}/\mathbf{SGPA}$
8.40	74.80			Y = Equivalent %
8.50	76.25			-
9.00	83.50	1st division with	10.0 ≤ CGPA ≥ 8.5	
9.50	90.75	Distinction	$75 \le Percentage \ge 98$	
10.00	98.00			

#### MERITS

- Elegant Formula, a good attempt at trying to achieve scientific rigour
- Benchmarks 40% to CGPA 6
- Has a clear indication of the divisions

#### DEMERITS

- Different conversion formula with similar methods would have to be created for past students who graduate with a CGPA of 5.5.
- There would be intertemporal inconsistencies. It would be difficult to justify why a particular CGPA has different percentage conversions prior to 2018 and post 2018.

<sup>&</sup>lt;sup>3</sup> Credit for this formula goes to Dr. Naqui Anwer

- In case the CGPA rule is changed in future, the conversion rule also will have to change. There would be a need to reach out to experts for creating such formula, leading to administrative costs.
- Pass grade of each individual course is D which carries 4 grade points is not addressed in this formula.

**OPTION 4:** Eliminate D grade. So, passing grades would be A+ to C. C has a score of 5. Also, CGPA of 5 would be set minimum requirement for award of degree. Following this, 50%, based on a formula of multiplication by 10 could be set as pass percentage.

#### MERITS

• Creates parity between the pass grade at a course level and the minimum CGPA required for graduation. Then, the dilemma of benchmarking the pass percentage to individual course vs final aggregate CGPA is resolved.

#### DEMERITS

- Eliminating D grade would not be in alignment with the norms in other universities.
- This will not provide a conversion formula for current and past cohorts of students for whom the minimum CGPA required for award of degree cannot be changed retrospectively.
- The alignment of divisions may not match the traditional Indian university norms.

#### **CONCLUSION OF INTERNAL MEETING:**

The meeting also acknowledged that options 2, 3 and 4 would have a range of inconsistencies across different cohorts that follow different rules of minimum CGPA which would be difficult to explain. **Hence, option 1 may be the way forward.** The demerits of option 1 were recognized and reflected upon in detail. It was also acknowledged that finding a strict scientific rationale that could also resolve internal consistencies as well as be administered with reasonable ease was, perhaps, not feasible. Hence adopting an option that many other universities and institutions followed may be an agreeable way forward.

## Enclosure 3

#### A. Approval of Semester 2 courses of M.Sc. Biotechnology Programme

Following eight courses of Semester 2 of M.Sc. Biotechnology Programme were presented to the Academic Council.

#### 1. Conservation genetics and genomics

**Observation:** No changes were suggested in this course and BoS approved the detailed structure

#### 2. Microbial pathogenesis

**Observation:** No changes were suggested in this course and BoS approved the detailed structure

3. Molecular Cell Biology: From Genes to Communities

Observation: There were no changes suggested and BoS approved the detailed structure

4. Genome organization and molecular markers

**Observation:** The course coordinator informed that one of the reviewers suggested changing the title of this course to "Genome Structure and Diversity: Concepts and Methodologies" to make it more meaningful and appropriate. The BoS accepted the observation of the reviewer. Further, it was requested that the total credit of the course be increased from 2 to 3 considering a large number of new concepts as suggested by the reviewers. The BoS accepted this proposal based on comments and opinions received by the course coordinator. It was decided that the overall credit of the Programme is also increased from 75 to 76 to accommodate this change in course credit.

5. Biotechnology Laboratory- Part 2

**Observation:** Laboratory was designed for common as well stream specific practicals, the BoS asked to segregate the experiments into Part A (which will be common for students of both the streams) and Part B1 as only for the Microbial Biotechnology stream while Part B2 will be only for Plant Biotechnology Stream.

#### 6. Introduction to Nanobiotechnology

**Observation:** The BoS members suggested, as this is an introductory course on Nanobiotechnology for Biotechnology students, a greater emphasis needs to be on the application of Nanobiotechnology in areas such as agriculture, health and medicine. The topics such as synthesis and characterization of Nanomaterials may be dealt with at a basic level.

7. Molecular Plant Physiology and Metabolism

**Observation:** It was suggested that the topic of respiration might be removed, as it is already part of the course in Biochemistry. A greater emphasis needs to be given to the topic of the Physiology of plant development and flowering.

8. Molecular microbiology and immunology

**Observation:** No changes were suggested in this course and BoS approved the detailed structure.

# B. Approval of the revised programme structure of the M.Sc. Biotechnology Programme

The name of the course, Genome organization and molecular markers, has been changed to Genome Structure and Diversity: Concepts and Methodologies. Further, the total credit of the course has been increased from 2 to 3. Consequently, the total credit of the Programme needs to be increased from 75 to 76. Accordingly, the revised programme structure outline of the M.Sc. Biotechnology Programme is provided below.

Year	Courses	Credits	Duration
First Year			
1st Semester	7 core courses of 2-7 credits each, and 2 core audit courses	21	15 weeks
2nd Semester	7 core courses of 2-7 credits and 1 course of 2 credits in the area of specialisation**	23	15 weeks
Second Year			
3rd Semester	4 core courses of 2-7 credits and 1 course of 2 credits in the area of specialisation**	16	15 weeks
4th Semester	Major project	16	15 weeks

#### **Programme outline for M.Sc. Biotechnology\***

\*In addition to above, a minimum 4 credits equivalent of elective courses (audit only) listed below need to be completed during the Programme which may be taken in any semester when

offered by the concerned Department and provided it doesn't conflict with any other course taken by the student. There is no upper limit for the number and credit equivalent for Elective courses. \*\*Specialisation specific practical component equivalent to 2 credits will carried out under Biotechnology Laboratory- Part 2 (2<sup>nd</sup> Semester) and Biotechnology Laboratory- Part 3 (3<sup>rd</sup> Semester) each.

Semester 1					
Course No.	Course title	Туре	Number of Credits	No. of L-T-P	
BBP 105	Biotechnology Laboratory - Part 1	Core	7	7-0-196	
NRE 101	Communication Skills and Technical Writing	Audit	2*	16-12-0	
BBP 155	Principles of Genetic Engineering and Recombinant DNA Technology	Core	3	30-15-0	
NRE 113	Applied Mathematics	Audit and bridge course	0*	31-11-0	
BBP 158	Conceptual Foundations of Molecular Biology	Core	2	30-0-0	
BBP 154	Principles of Biochemistry and Biophysics	Core	2	30-0-0	
BBP 111	<b>Bioanalytical Techniques</b>	Core	3	39-6-0	
BBP 123	Plant and Animal Biotechnology	Core	2	30-0-0	
BBP 174	Bioinformatics and Computational Biology	Core	2	22-8-0	

Semester 2					
Course No.	Course title	Туре	Number of Credits	No. of L-T-P	
TBA	Conservation Genetics and Genomics	Core	2	30-0-0	
TBA	Biotechnology Laboratory -	Core*	7	0-0-210	

Semester	: 2			
Course No.	Course title	Туре	Number of Credits	No. of L-T-P
	Part 2			
TBA	Introduction to Nanobiotechnology	Core	2	22-8-0
BBP 130	Molecular Microbiology and Immunology	Core	2	30-0-0
BBP 112	Statistics for The Life Sciences	Core	3	28-14-0
BBP 114	Molecular Cell Biology - From Genes to Communities	Core	2	30-0-0
TBA	Genome Structure and Diversity: Concepts and Methodologies	Core	3	23-22-0
TBA	Molecular Plant Physiology and Metabolism	Specialization (Plant Biotechnology)	2	30-0-0
TBA	Microbial Pathogenesis	Specialization (Microbial Biotechnology)	2	15-15-0

\*Specialisation specific practical component equivalent to 2 credits will carried out under Biotechnology Laboratory- Part 2

Semester 3				
Course	Course title	Туре	Number	No. of L-T-P
No.			of Credits	
TBA	Biotechnology Laboratory -	Core*	7	
	Part 3			
BBP	Bioethics, IPR and	Core	3	
141	Regulations in Biotechnology			
TBA	Gene Expression Analysis	Core	2	
	and Transcriptomics			
TBA	Proteomics and Protein	Core	2	

Semester 3				
Course	Course title	Туре	Number	No. of L-T-P
No.			of Credits	
	Engineering			
TBA	Functional Genomics in	Specialization	2	
	Plants	(Plant		
		Biotechnology)		
TBA	Bioprocess Engineering and	Specialization	2	
	Environmental Biotechnology	(Microbial		
		Biotechnology)		

\*Specialisation specific practical component equivalent to 2 credits will carried out under Biotechnology Laboratory- Part 3

Elective courses* (Audit only)					
Course	Course title	Туре	Number of Credits	No. of L-T-P	
NRF	Environmental Chemistry	Flective	3	35-7-0	
131	and Microbiology	Liceuve	5	33-7-0	
NRE	Introduction to Sustainable	Elective	1	14-0-0	
165	Development				
TBA	Nanomaterials: Introduction	Elective	2		
	and Applications				
NRE	<b>Biodiversity Assessment</b>	Elective	3	17-15-20	
123	and Conservation				
NRE 168	Food Security and	Elective	3	23-16-6	
	Agriculture				
NRE 112	Multivariate Data Analysis	Elective	3	28-14-0	
NRE	Wildlife Conservation and	Elective	3	35-7-0	
151	Management				

\*Elective courses may be taken in any semester when offered by the concerned Department and provided it doesn't conflict with any other course taken by the student. There is no upper limit for the number and credit equivalent for Elective courses.

# Courses to be offered in Second Semester of M.Sc. Biotechnology Programme

Course title: Conservation Genetics and Genomics							
Course o	code: BBP	No. of credits: 2	L-T-P: 30-0-0	Learning hours	: 30		
Pre-requ	uisite course cod	le and title (if any):	Science graduate				
Departm	nent: Department	t of Natural and App	lied Sciences				
Course o	coordinator:		Course instructe	or:			
Contact	details:						
Course t	ype: Core		Course offered i	n: Semester 2			
Course of The broa and tech conserva genetic of sequencin sequencin on micro included. Course of 1. To in 2. To in 3. To fa 4. Appli	lescription: d objective of the miques of class tion of biodiversis composition of ming in generating ing and its application bial genetic dive bial genetic dive bial genetic dive bial genetic dive bial genetic dive	is course is to provid sical genetics, pop ity. The students will natural populations. g data for characte ations in genetic dive ersity such as 16S RI ents to concepts of cla ents to concepts of po dents to next generation generation sequence of	de the students a f ulation genetics Il be acquainted w Considering the erization of popu rsity assessment h NA sequencing an assical and modern opulation and cons ton sequencing pla lata for characteris	oundation on the and genomics ith various factors importance of n lations, a modul as been included. d metagenomics n genetics ervation genetics atforms sation of genetic re	conce as ap s that ext g le on Furth have a	pts, to oplied affect enerat genc er, top also b	ools in the ion ome oics een
Course o	contents						
Modul	Торіс				L	Т	Р
с 1	Principles of F	volution and Popul	ation genetics		6	0	0
-	Principles of E	f evolution and Natur	al selection		0	0	
	<ul> <li>Population a</li> </ul>	attributes and structur	e				
	Gene and ge changes in g random gene	enotype frequency: H gene frequency throug etic drift; Population	ardy-Weinberg Ec gh natural selection bottlenecks	quilibrium; n, migration and			

	Adaptive radiation; Speciation; Allopatric and Sympatric;     Convergent evolution:			
	<ul> <li>In-breeding depression &amp; mating systems</li> </ul>			
2	Introduction to conservation genetics	6	0	0
	• Introduction to conservation genetics,			
	• Concepts of gene pool (primary, secondary, tertiary)			
	• Natural variation: Phenotypic and genetic diversity including allelic richness; Analysis of genetic diversity			
	• In situ and ex situ conservation, core collections			
3	Principles of Genetics and mapping	12	0	0
	Genetics and inheritance: Laws and exceptions			
	Recombination and linkage mapping			
	• Quantitative genetics and mapping, polygenic inheritance, heritability; Linkage disequilibrium			
4	Genomics platforms for population and conservation genetics	6	0	0
	• Introduction to next generation sequencing platforms, Pyrosequencing, Illumina, Single molecule real time (SMRT) sequencing			
	• 16S RNA based analysis of microbial diversity and taxonomy, metagenomics			
	Nuclear and Organellar DNA for conservation and diversity			
	Total	30	0	0
Evaluat	ion criteria:			
1. Test	1- (Module 1) 30%			
2. Test	2- (Module 2) 30%			
3. Test	3- (Modules 3 and 4) 40%			
Learnin	g outcomes:			
1. 5	students will be able to use the principles of evolution and population ge	enetics (	Test 1	.)
2. E	Basic understanding of principles of germplasm conservation (Test 1-2)			
3. U	Jnderstanding of principles of genetics (Test 2-3)			

4. Basic understanding of next generation sequencing platforms and their application in

#### genetic diversity analysis (Test 3)

#### **Pedagogical Approach:**

- 1. Online/classroom lectures and discussions
- 2. Case studies and examples from original research articles

#### Skill Set:

- 1. Next generation sequencing platforms
- 2. Germplasm characterisation using principles of population genetics
- 3. 16S RNA sequencing and metagenomics analysis

#### **Employability:**

- 1. Forestry and wildlife research institutions
- 2. Academic organisations
- 3. Companies providing genotyping and sequencing services

# Materials:

#### **Suggested Readings**

- 1. A Primer of Ecological Genetics. Conner, J. K. and D. L. Hartl. Sinauer Associates. 2009
- 2. Conservation and the Genetics of Populations. 2nd edition. Allendorf, Luikart and Aitken. 2013.
- Adaptive radiations: From field to genomic studies. Scott A. Hodges, Nathan J. Derieg. Proceedings of the National Academy of Sciences Jun 2009, 106 (Supplement 1) 9947-9954; DOI: 10.1073/pnas.0901594106
- 4. Methods in Molecular Biology, vol. 376: Linkage Disequilibrium and Association Mapping: Analysis and Applications Edited by: A. R. Collins © Humana Press Inc., Totowa, NJ

# Additional information (if any):

#### **Student responsibilities:**

- 1. Class attendance.
- 2. Study of reading materials as specified by course instructor
- 3. Self-study

#### **Course reviewers:**

- 1. Prof. Sandip Das, Department of Botany, University of Delhi, New Delhi
- 2. Dr. R. Yasodha, Scientist G, Institute of Forest Genetics and Tree Breeding, Coimbatore

3. Dr. Ram Kumar Sharma, Scientist G, Institute of Himalayan Bioresource and Technology, Palampur, Himachal Pradesh

Course ti	tle: Microbial pathogenesi	S						
Course co	ode:	No. of credits	<b>s:</b> 2	L-T-P: 15-	Learni	ng h	ours	: 30
				15-0				
Pre-requi	site course code and title	e (if any): None	;					
Departme	ent:							
Course co	oordinator(s):		Course	instructor(s): In	nternal fa	cult	у	
		1	member	(s) and external	experts.			
Contact d	letails:							
Course ty	pe: Core		Course	offered in: Sem	ester 2			
Course de	escription: Microbial dise	eases impose sig	gnificant	t social and ecor	nomic bu	rden	s on	
human so	ciety. However, the insigh	nts gained from	both me	dicine and basic	: biology	thu	s far l	nave
led to a be	etter understanding of disea	ase mechanisms	s. This r	new knowledge	has great	ly h	elped	in
the prever	tion, management and cur	e of several dis	eases. 7	This course aims	to impa	rt an		
understan	ding of some of the curren	t paradigms in 1	microbia	l pathogenesis.				
The study	material for this course w	ill include textb	ooks, ca	use studies and a	rticles fr	om f	field	
journals.								
This is a h	ighly participatory course	with a significa	ant comp	ponent of self-st	udy of as	sign	ed	
material f	rom the literature and stude	ent presentation	ns of cas	e studies. Probl	em-base	d lea	rning	5
will be a c	critical component of the e	valuation proce	ss. Evol	utionary and eco	ological p	persp	pectiv	res
will be en	phasized to provide a trul	y integrative fra	amework	to understand h	10st-path	ogei	1	
interaction	ns and their consequences.							
Course of	ojectives:							
1. To pre	esent key aspects of the bio	ology of differen	nt pathog	gens and their in	teraction	is wi	th the	e
host.			_					
2. To ena	able synthesis of information	on in order to st	tudy cor	nmunicable dise	ases with	nin a	n	
evolut	ionary-ecological framewo	ork.						
Course co	ontents					-		
S.No	Торіс					L	Т	P
Module	Introduction							
1		1 1 1						
1	Dethe some cull V and	a evolution		of ' ami-''		<u>2</u> 1		
2	Pathogens and Koch's po	ostulates. Conti	ribution	or -omic scien	ices	1		
2	to our understanding of p	patnogens.	1.1	1.0		2		
5	Nodes of disease transm	ussion, epidemi	cs and the	he spread of anti	-	3		
N	microbial resistance							
Module	Niechanisms and Molec	cules						

2						
1	Heat nother can interpretions at the melecular level hert resistance and	4				
1	Host-pathogen interactions at the molecular level, host resistance and	4				
2	susceptibility genes and the determinants of outcomes.	2				
2	The microbiome in health and disease	3				
3	Diagnostics, vaccines and therapeutic agents	2				
Module	Case studies in microbial pathogenesis					
3						
1	SARS-CoV-2 and the WHO vaccine programme		3			
2	Anthrax – an acute zoonotic disease		3			
3	Tuberculosis – a chronic and worldwide problem		3			
4	Helicobacter pylori – commensal, pathogen and carcinogen		3			
5	ESKAPE pathogens: <i>Pseudomonas aeruginosa</i> – a wide-ranging		3			
	opportunist					
	Total	15	15	0		
Evaluatio	n criteria:					
Test $1-30$	0% weightage					
Test $2-30$	0% weightage					
Test $3-40$	0% weightage					
Learning	outcomes:					
1. Knowl	edge of basic concepts in the field (Tests 1-3).					
2. Ability	to critically analyze and synthesize primary data to develop coherent m e hypotheses (Tests $1-3$ )	odels	or fra	me		
3 Detaile	ed understanding of nathogens and their strategies for colonization imm	une ev	asion			
and die	enercal (Tests 2-3)	une ev	usion			
Pedagogi	$\mathbf{r}_{\mathbf{r}} = \mathbf{I} \mathbf{A} \mathbf{n} \mathbf{r}_{\mathbf{r}} \mathbf{r}} \mathbf{r}_{\mathbf{r}} \mathbf{r}_{\mathbf{r}} \mathbf{r}_$					
Online/off	line lectures and self-study assignments. Detailed discussion and studer	nt nres	entati	on		
of articles	from peer-reviewed journals in class for module 3	n pres	ciitati	011		
Skill Set	from peer reviewed journais in class for module 5.					
1 Critical	analysis of concepts, hypotheses and experimental design					
2 Formul	ation of experimental strategies for molecular genetic studies of model l	nost-ne	athoge	'n		
systems	ation of experimental strategies for molecular generic statics of model i	iost pt	11105			
3 Compa	rative analysis of preventive and therapeutic strategies					
Employal	nility.					
1 Acader	nic and industrial research on microbial pathogens					
2 Intellec	tual property firms					
2. Interfect	2. Interfectual property mins.					

- Life science teaching at school and undergraduate levels.
   Pathology laboratories.

5. Management and/or supervision of laboratory-based research in academic/industrial/medical settings.

# Materials:

# **Required texts**

- 1. B.A. Wilson et al. Bacterial Pathogenesis: A Molecular Approach. ASM Press, ed. 4, 2019.
- 2. J.C. Herron, S. Freeman. Evolutionary Analysis. Pearson Education, India. ed. 5, 2013.
- 3. B. Tungland. Human Microbiota in Health and Disease: From Pathogenesis to Therapy. Elsevier Science, 2018.
- 4. Viral Pathogenesis, From Basics to Systems Biology. Academic Press, ed. 3, 2016.
- 5. N. Bergman. Bacillus anthracis and Anthrax. Wiley-Blackwell, 2010.
- 6. P. Sutton & H. Mitchell (eds.). *Helicobacter pylori* in the 21st Century (Advances in Molecular and Cellular Biology Series). CABI, 2010.
- 7. S.E. Hasnain *et al. Mycobacterium tuberculosis*: Molecular Infection Biology, Pathogenesis, Diagnostics and New Interventions. Springer Singapore, 2019.
- 8. B.H.A. Rehm (ed.). *Pseudomonas*: Model Organism, Pathogen, Cell Factory. Wiley-VCH, 2008.
- 9. R. Rappuoli & F. Bagnoli (eds). Vaccine Design: Innovative Approaches and Novel Strategies. Horizon Scientific Press, 2011.
- 10. S. Pan & J. Tang (eds.). Clinical Molecular Diagnostics. Springer, 2021.
- 11. L. Pirofski & A. Casadevall. *mBio* 11(4), e01175-20 (2020). doi: 10.1128/mBio.01175-20
- 12. E.Janik et al. Medicina, 56(11), 591 (2020). doi: 10.3390/medicina56110591
- 13. S. Suerbaum & P. Michetti. *New England Journal of Medicine* 347(15), 1175-86 (2002). doi: 10.1056/NEJMra020542
- 14. L.I. Rankine-Wilson *et al. Microbiology (Reading)*167(4):001041 (2021). doi: 10.1099/mic.0.001041
- 15. I. Jurado-Martín *et al. Int J Mol Sci.* 22(6), 3128 (2021). doi: 10.3390/ijms22063128.
- 16. Y. Taoufik et al. Front Immunol. 2021 12, 692598 (2021). doi:
- 10.3389/fimmu.2021.692598.
- 17. M.J. Culyba & D. van Tyne. *PLoS Pathog* 17(9), e1009872 (2021). doi: 10.1371/journal.ppat.1009872

Case studies

**Suggested readings** 

Journals

Other readings Additional information (if any):

#### **Student responsibilities:**

- 1. Class attendance (online/offline).
- 2. Study/self-study/presentation of course materials as specified by the instructor.
- 3. Ensuring functionality of essential IT hardware & software at their preferred location(s).

#### **Course reviewers:**

- Prof. Vijaya Satchidanandam, Department of Microbiology and Cell Biology, Indian Institute of Science, Bengaluru (superannuated) and Adjunct Professor, St. John's Medical College, Sarjapur Road, Bengaluru – 560034
- Dr. S. Ramachandran, Chief Scientist, Professor of the AcSIR in the Faculty of Biological Sciences, Room 130, CSIR-Institute of Genomics and Integrative Biology, Mathura Road, Near Sukhdev Vihar Bus Depot New Delhi 110 025

Course title: Molecular Cell Biology: From Genes to Communities									
Course code:No. of credits: 2L-T-P: 30-0-Learn			Learni	ing h	ours	: 30			
Pre-requisite course code and title (if any): None									
Departme	ent:								
Course coordinator(s):     Course instructor(s):									
Contact d	etails:		~						
Course ty	pe: Core		Course	offered in: Sem	ester 2				
Course de	escription: This course wi	ll highlight the	physiol	ogical versatility	that un	derlie	es the	;	
ability of o	organisms to adapt to vary	ing needs of the	eir respe	ctive developme	ental stag	ges,			
environme	ental stimuli and ecologica	l niches. Advar	nced and	l contemporary t	themes in	n mo	lecula	ar	
and cell bi	ology will be highlighted	as indicated. The	he cours	e is divided into	three m	odule	es to	11	
facilitate t	he analysis of living system	ms at progressi	vely mo	re complex level	ls. This	cour	se wi	11	
neip stude	nts gain new knowledge if	h, and develop	their ow	n perspectives o	n the rap	bialy			
Comment	i neid of modern biology.								
	opecuves:	f callular proces	accor of p	rograggivaly oor	nnlay la	uala			
1. To pre	ble synthesis of isolated in	formation in o	sses at p	noluza biologia	al phono	veis.	in a		
2. TO Cha	tually relevant manner			ularyze biologica	ai piteno	mena	a III a		
3 To del	ineate the overarching role	e of evolutionar	ry consid	lerations at mult	inle leve	als of	•		
comple	exity		ry consic	crations at man		15 01			
Course co	ontents								
S.No						L	Т	Р	
Module	The genetic material							_	
1	8								
1	The evolution of complete	xity.				3	0	0	
2	The dynamic nature of th	ne genome				3	0	0	
	Recombination, gene cor	nversion, extrac	chromos	omal elements,					
	horizontal gene transfer,	transposition, t	ransduc	tion, phase varia	tion,				
	DNA rearrangements and	d the vertebrate	adaptiv	e immune system	m				
3	Epigenetics					3	0	0	
	Epigenetic mechanisms of	of gene regulati	ion, non-	-coding RNAs in	1				
	gene regulation and cellu	ılar defence.							
Module	Cellular processes – fro	om molecules t	o cells						
1	Model organisms – over	view of <i>F_coli</i>	S cervi	siae (veast) C		2	0	0	
1	wodel organisms – over	view of E. coli,	S. Cervi	siae (yeast), C.		2	U	U	

	elegans, D. melanogaster, and A. thaliana			
2	Spatio-temporal gene regulation	5	0	0
	Molecular processes underlying the eukaryotic cell cycle, cell			
	signalling and responses, regulatory networks and cross-talk between			
	cellular pathways, protein secretion and localization.			
3	A systems view of regulatory processes in biology.	3	0	0
	Types of regulatory mechanisms, bistability, intrinsic and extrinsic			
	noise, synthetic biology.			
Module	Organisms to ecosystems			
3				
1	Microbial interactions	5	0	0
	Gene transfer, barriers to gene transfer, quorum sensing, host-			
	microbe interactions, symbiosis and pathogenesis.			
2	Microbial communities and the microbiota – an evolutionary-	6	0	0
	ecological synthesis. The self versus non-self recognition			
	conundrum.			
	Total	30	0	0
<b>Evaluatio</b>	n criteria:			
Test 1 – 3	0% weightage			
Test $2-3$	0% weightage			
Test $3-4$	0% weightage			

#### Learning outcomes:

- 1. Detailed knowledge of specific aspects of model living systems in consonance with topics in the outline (Tests 1-3).
- Ability to critically analyze and synthesize primary data to develop coherent models (Tests 1-3).
- 3. Understanding implicit evolutionary arguments underlying the analysis of organisms from the genetic to community levels (Tests 1-3).

#### **Pedagogical Approach:**

Online/offline lectures emphasizing the detailed discussion of research/review articles from scientific journals in class.

#### **Skill Set:**

1. Design of molecular biology/genetic engineering experiments.

2. Critical analysis of molecular biology/genetic engineering experimental design and results.

3. Formulation of experimental strategies for molecular genetic studies of simple model organisms.

4. Analysis and design of complex regulatory networks.

## **Employability:**

- 1. Academic and industrial research involving molecular biology approaches.
- 2. Intellectual property firms.
- 3. Life science teaching at school and undergraduate levels.
- 4. Management and/or supervision of laboratory research in academic/industrial settings.

#### Materials:

# **Required texts**

- 1. J. D. Watson., *et al.* Molecular Biology of the Gene. Pearson, Cold Spring Harbor, ed. 7, 2014.
- 2. S. Brenner. Phil. Trans. R. Soc. B, 365, 207-212 (2010).
- 3. M. W. Gray et al. Science, 330, 920-921 (2010).
- 4. A. Rokas. *Nature*, 443,401-402 (2006).
- 5. T.D.P. Brunet, W.F. Doolittle. Biol Philos 33, 2 (2018). doi: 10.1007/s10539-018-9614-6.
- 6. I. R. Henderson, S. E. Jacobsen. *Nature* 447, 418-424 (2007).
- 7. V.L. Chandler. Cell, 128 (4), 641-645 (2007).
- 8. B. Alberts, et al. Molecular Biology of the Cell. Garland Science, New York, ed. 5, 2008.
- 9. D. G. Gibson et al. Science, 329, 52-56 (2010).
- 10. R.J. Hall, et al. Front Microbiol 11:1569. doi: 10.3389/fmicb.2020.01569 (2020).
- 11. J. -H. Hehemann et al. *Nature*, 464, 908-912 (2010).
- 12. N. C. Reading, V. Sperandio. FEMS Microbiol Lett, 254, 1-11 (2006).
- 13. O.P. Duddy, B.L. Bassler. *PLOS Pathogens* 17(1): e1009074. doi:
- 10.1371/journal.ppat.1009074 (2021).
- 14. E. K. Costello et al. *Science*, 336, 1255-1262 (2012)

# Case studies

#### **Suggested readings**

1. J. E. Krebs *et al.* Lewin's GENES XII. Jones and Bartlett Publishers, Inc., Burlington, ed. 12, 2017

#### Journals

Other readings

#### Additional information (if any):

#### **Student responsibilities:**

1. Class attendance (online/offline).

- 2. Study/self-study of course materials as specified by the instructor.
- 3. Ensuring functionality of essential IT hardware & software at their preferred location(s).

#### **Course reviewers:**

- Prof. Vijaya Satchidanandam, Department of Microbiology and Cell Biology, Indian Institute of Science, Bengaluru (superannuated) and Adjunct Professor, St. John's Medical College, Sarjapur Road, Bengaluru – 560034
- Dr. S. Ramachandran, Chief Scientist, Professor of the AcSIR in the Faculty of Biological Sciences, Room 130, CSIR-Institute of Genomics and Integrative Biology, Mathura Road, Near Sukhdev Vihar Bus Depot New Delhi 110 025

Course title: Genome Structure and Diversity: Concepts and Methodologies									
Course code: BT XXX	No. of credit	t <b>s:</b> 3	L-T-P: 23-22-	Learning	hours:	45			
			0						
Pre-requisite course code and title (if any): None									
Faculty:		Departn	nent: Department	of Natural	and App	lied			
Sciences									
Course coordinator(s):		Course i	nstructor(s):						
Contact details:									
Course type: Core		Course of	offered in: Semes	ter 2					
Course description:									
The extraordinary diversity among living	ng organisms is	reflective	of structural and t	functional d	iversity	of			
genomes. The tree of life is strident evi	dence of evolut	ionary pro	cesses underlying	biological	variatio	n.			
Genome elucidation studies are crucial	for gaining insi	ights into t	he molecular basi	s of morpho	ological				
diversity and trait variation. This advar	iced course prov	vides a cor	nceptual framewor	k on genon	ne				
architecture and experimental methods	for analysis of	its compoi	nents and sequenc	es. In the fi	rst modu	ıle,			
students will gain insights on features of	of diverse genor	nes, hierar	chies of genome of	organisation	, variab	ility			
in genome complexity and content and	dynamic nature	e of genom	es at varying leve	ls of resolu	tion. In t	the			
second module, a critical appraisal of the	raditional marke	er techniqu	ies and modern, so	ophisticated	genoty	ping			
platforms vis-à-vis relative efficiencies	in polymorphis	sm detectio	on will be discusse	ed. Introduc	tion to r	next			
generation, genomics based, genotypin	g platforms wil	l sensitize	the students to fro	ontier areas	of resear	rch			
directed towards sustainable agriculture	e, generation of	climate re	silient crops and h	nealthcare p	roducts.				
Third module is designed to inform the	students, by wa	ay of inter	esting case studies	s, applicatio	n of ma	rkers			
in sectors of plant, animal and microbia	al biotechnology	y. Through	this course, stude	ents will gai	n a holis	stic			
perspective on "genotype-phenotype as	ssociation" by in	ntegration	of core principles	related to d	iverse				
disciplines as molecular genetics, geno	mics and evolut	tion.							
Course objectives:									
1. Building perspectives on structure	and variability	in genome	s and its constitue	nts					
2. Illustrating the relationship betwee	n genotypic and	l phenotyp	ic variation						
3. Introducing versatile methodologie	es, concepts and	applicatio	ons of molecular m	narker techr	iques				
Course contents									
Module1 Genome Structure and G	Organization (1	Prokaryot	tes and Eukaryot	es) L	, T	P			

1	Genome diversity (Viral, Bacterial, Archaeal, Eukaryotic, Auxiliary DNA	6	6	
	structures viz. Plasmids and Organellar genome); Hierarchies of Genome			
	<b>Organization</b> (Genomic sequences, chromatin, nucleosomes, packaging,			
	3D genomes and chromosome territories); Dynamic genomes and			
	variability in Genome Content (Genome sizes and complexity, C-Value			
	paradox. Unique and repeat DNA sequences: Tandem and Interspersed			
	repeats. Mobile Elements. Micro- and Mini-satellites. hyper-variable			
	VNTRs. Whole Genome Multiplications and Fractionation. DNA			
	rearrangements. SNPs and Structural variation (Microscopic and sub-			
	microscopic, Copy Number Variation, Presence Absence Variation,			
	Inversions. Mobile Element Insertion and Deletions. Homologous			
	Exchange Variation): Variability in gene categories and structure			
	(Protein coding genes, non-protein coding genes, Intron-less and			
	interrupted genes (Structure of exons, introns, variability in number, size,			
	GC-content): Intron types: poly-cistronic genes, overlapping genes (+/+ and			
	+/- strand): cis-regulatory regions (Promoter, bi-directional promoters,			
	Enhancers, Insulators, Terminators), case studies to illustrate structural			
	variations as basis for phenotypic diversity, notion of genome maps			
Module 2	Genome analysis by Genetic markers			
2	Molecular Markers and DNA fingerprinting techniques	12	12	
4	Molecular Markers and Divis inger printing teeningues	14	14	
2	Definition of trait, classification genetic markers, molecular basis of	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs,	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism,	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs,	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the genome); <b>Modern Genotyping platforms</b> Array based genotyping	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the genome); <b>Modern Genotyping platforms</b> Array based genotyping (Affymetrix axiom, Affymetrix genechip, Illumina Infinium Bead Chip;	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the genome); <b>Modern Genotyping platforms</b> Array based genotyping (Affymetrix axiom, Affymetrix genechip, Illumina Infinium Bead Chip; NGS based genotyping methods (GBS, DArT-seq, RAD-seq, ddRAD,	12	12	
2	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the genome); <b>Modern Genotyping platforms</b> Array based genotyping (Affymetrix axiom, Affymetrix genechip, Illumina Infinium Bead Chip; NGS based genotyping methods (GBS, DArT-seq, RAD-seq, ddRAD, REST-seq); de-novo sequencing and/or WGS (PacBio. HiC. 10X	12	12	
	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the genome); <b>Modern Genotyping platforms</b> Array based genotyping (Affymetrix axiom, Affymetrix genechip, Illumina Infinium Bead Chip; NGS based genotyping methods (GBS, DArT-seq, RAD-seq, ddRAD, REST-seq); de-novo sequencing and/or WGS (PacBio. HiC. 10X Chromium, Oxford nanopore, HiSeq4000/NovaSeq6000, IonTorrent)	12		
2 Module 3	Definition of trait, classification genetic markers, molecular basis of dominant and co-dominant markers; Restriction Fragment Length Polymorphism, MAAP (Multiple Arbitrary Amplicon Profiling) and other PCR based markers (DNA Amplification Fingerprinting, Arbitrarily Primed PCR, Randomly Amplified Polymorphic DNA, SSRs, STMS, SCARs, Inter-SSRs, Amplified Fragment Length Polymorphism, Selectively Amplified Microsatellite Polymorphic Loci, Inter retrotransposon amplified polymorphism, retrotransposon-microsatellite amplified polymorphism, Intron spanning markers, SNP based marker assays (CAPs, dCAPs, dHPLC, molecular beacons, 5'nuclease assay/TaqMan assays, FEN based Invader reactions), Eco-TILLING (Targeting induced local lesions in the genome); <b>Modern Genotyping platforms</b> Array based genotyping (Affymetrix axiom, Affymetrix genechip, Illumina Infinium Bead Chip; NGS based genotyping methods (GBS, DArT-seq, RAD-seq, ddRAD, REST-seq); de-novo sequencing and/or WGS (PacBio. HiC. 10X Chromium, Oxford nanopore, HiSeq4000/NovaSeq6000, IonTorrent) <b>Applications and key concepts related to marker technology: Case</b>	12	12	

<b>3 Diversity analysis in plants</b> Geographical diversity, center of origin domestication, gene pools, pan-genomes and super-pangenomes; methods (numerical taxonomy and phenetics), conservation of plant genetic resources; <b>Diversity analysis in microbes</b> (Microbiomes, structures and functions, 16S to metagenomics); <b>Molecular Breeding</b> (MAS, Genomics Assisted Breeding); plant variety protection; DNA barcoding; hybrid purity	5	4	
fingerprinting; LD/ Haplotype mapping, GWAS in context to natural populations (animals and plants), <b>human diseases</b> (mapping human diseases, risk prediction, discovery of drug targets and improving	; L Ç		
healthcare), genomic selection in plants and animals			
Total	23	22	
Evaluation criteria:         1. Test 1:       30%         2. Test 2:       30%         3. Test 3:       40%			
<ol> <li>Learning outcomes:</li> <li>An understanding on structure and variability in genomes and its constituents (Test</li> <li>Ability to rationalize deployment of genotyping techniques for relevant applications</li> <li>Understanding genetic and molecular basis of phenotypic variation (Test 1-3)</li> </ol> Pedagogical Approach:	1-3) (Test 1	-3)	
Lectures and tutorials in online or offline mode with a major emphasis on the detailed di original research articles	scussior	n of	
<ul> <li>Skill Set:</li> <li>Generating and interpreting DNA fingerprints and profiles for forensics</li> <li>Developing natural and synthetic microbiomes as biofertilizers, biopesticides, health</li> <li>Testing Hybrid purity</li> <li>Diagnosing varieties, cultivars, accessions and land races</li> <li>Ascertaining clonal fidelity for tissue culture raised regenerants</li> <li>Applying MAS (Marker Assisted Selection) strategies in breeding programmes</li> <li>Introducing transgenes for development of new plant varieties</li> <li>DNA bar-coding technology</li> <li>Evaluating gene-flow in transgenic field trials</li> <li>Formulating appropriate conservation strategies</li> <li>Innovating genome interrogation methods for unarticulated research problems</li> </ul>	care pro	oducts	
1. Forensic Science laboratories, molecular diagnostic testing laboratories			22

- 2. Genotyping and sequencing companies
- 3. Agri-biotechnology and seed companies
- 4. Tissue culture and horticulture companies
- 5. Law firms and knowledge processing organizations (IP management consultancy)
- 6. Regulatory bodies and funding agencies

#### Materials:

#### Suggested readings (Representative)

- 1. Krieg, N.R., Ludwig, W., Whitman, W.B., Hedlund, B.P., Paster, B.J., Staley, J.T., Ward, N. and Brown, D. (eds., 2010). Bergey's Manual of Systematic Bacteriology, 2nd ed., vol. 4, Springer-Verlag, New York, NY
- 2. Dale, J.W., Schantz, M.V. and Plant, N. (2011). From Genes to Genomes: Concepts and Applications of DNA Technology. Third edition. John Wiley & Sons, UK.
- 3. Brown, T. A. (2017). Genomes 4. CRC Press, Taylor & Francis Group, USA.
- 4. Krebs J.E, Goldstein E.S., Kilpatrick S. T. (2018) Lewin's GENES XII. Jones and Bartlett Learning. USA
- 5. Meksem K., Kahl G. (2005) The Handbook of Plant Genome Mapping: Genetic and Physical Mapping, Wiley-VCH Verlag GmbH & Co. KGaA, Weinheim
- 6. Varshney R., Pandey M., Chitikineni A (2018) Plant Genetics and Molecular Biology. Advances in Biochemical Engineering / Biotechnology series number 164. Springer Nature, Switzerland
- 7. Varshney R., Roorkiwal M., Sorells M (2017) Genomic Selection for Crop improvement: New Molecular Breeding strategies for crop improvement. Springer Nature, Switzerland
- 8. Research and review articles on relevant topics
- Scherer, S., & Visscher, P. (2016). Genome-Wide Association Studies: From Polymorphism to Personalized Medicine (K. Appasani, Ed.). Cambridge: Cambridge University Press. doi:10.1017/CBO9781107337459

#### Student responsibilities:

- 1. Class attendance
- 2. Study of course materials as specified by the instructor
- 3. Self-study

#### **Course reviewers:**

1. Prof. Surekha Katiyar-Agarwal

Department of Plant Molecular Biology

University of Delhi, South Campus, New Delhi- 110021, India

2. Dr. Neeti Sanan Misra Group Leader: Plant RNAi, International Centre for Genetic Engineering and Biotechnology Aruna Asaf Ali Marg New Delhi-110 067, India

	atory – Part 2						
Course code: BBP No. of	f credits: 7 L-T-P: 0-0-210 Learning hours	:210					
Pre-requisite course code and tit	le (if any): None						
Department: Department of Natural and Applied Sciences							
Course coordinator: Course	instructor:						
Contact details:							
Course type: Core/Specialization	<b>Course offered in:</b> Semester 2						
Course description:							
The objective of this laboratory co	urse is to introduce students to experiments related	to					
biotechnology. The course is desig	ned to teach students the utility of set of experimer	ıtal					
methods in biotechnology in a pro-	blem-oriented manner. The list of experiments give	n in e	ach				
module is representative and inclu-	des experiments. Part A will be common for both th	ne					
streams. Part B1 is only for Micro	bial Biotechnology stream whereas Part B2 is only	for P	lant				
Biotechnology stream. The instruc	tor may choose experiments for student's laborator	y trair	ning				
as per requirements.							
Course objectives:							
1. To introduce the students to sta	andard techniques of molecular biology.						
2. To impart intensive hands-on-t	training using molecular tools in a research project	mode.					
3. To train the students in designi	ing experiments with appropriate controls.						
Course contents							
Module Topic		Т					
Suggested and sticel			P				
Suggested practical			Р				
PART A: Common to both	streams		<b>P</b> 154				
PART A: Common to both	streams		<b>P</b> 154				
PART A: Common to both           I-         Genotyping methods a	streams and analysis of data-		<b>P</b> 154				
Suggested practical           PART A: Common to both           I- Genotyping methods a           1. Genotyping of natural p	streams and analysis of data- populations with ISSR markers		<b>P</b> 154				
Suggested practical           PART A: Common to both           I- Genotyping methods a           1. Genotyping of natural p           2. Genotyping of natural p	streams and analysis of data- populations with ISSR markers populations with SSR markers		<b>P</b> 154				
Suggested practical           PART A: Common to both           I- Genotyping methods a           1. Genotyping of natural p           2. Genotyping of natural p           3. Analysis of molecular d	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes		<b>P</b> 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis		<b>P</b> 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core		<b>P</b> 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         collections       6.	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core		<b>P</b> 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         6.       Genotyping of mapping         7.       Genotyping of mapping	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core g populations with SSR markers populations with ISSR markers		P 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         6.       Genotyping of mapping         7.       Genotyping of mapping         8.       Construction of linkage	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core g populations with SSR markers g populations with ISSR markers maps with marker data		P 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         6.       Genotyping of mapping         7.       Genotyping of mapping         8.       Construction of linkage         9.       Identification of OTLs a	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core g populations with SSR markers g populations with ISSR markers maps with marker data using mapping populations		P 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         6.       Genotyping of mapping         7.       Genotyping of mapping         8.       Construction of linkage         9.       Identification of QTLs u         10.       GWAS using SNP data	streams         and analysis of data-         populations with ISSR markers         populations with SSR markers         lata using MS Excel- marker attributes         lata using GeneAlex- Cluster analysis         lata using Corehunter or PowerCore- Core         g populations with SSR markers         g populations with SSR markers         maps with marker data         using mapping populations		P 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods a         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         6.       Genotyping of mapping         7.       Genotyping of mapping         8.       Construction of linkage         9.       Identification of QTLs of         10.       GWAS using SNP data         11.       Marker-trait association	streams and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core g populations with SSR markers g populations with ISSR markers maps with marker data using mapping populations as in natural populations		P 154				
Suggested practical         PART A: Common to both         I-       Genotyping methods at 1.         1.       Genotyping of natural p         2.       Genotyping of natural p         3.       Analysis of molecular d         4.       Analysis of molecular d         5.       Analysis of molecular d         6.       Genotyping of mapping         7.       Genotyping of mapping         8.       Construction of linkage         9.       Identification of QTLs of 10.         10.       GWAS using SNP data         11.       Marker-trait association         12.       ISSR fingerprinting for	streams         and analysis of data- populations with ISSR markers populations with SSR markers lata using MS Excel- marker attributes lata using GeneAlex- Cluster analysis lata using Corehunter or PowerCore- Core         g populations with SSR markers g populations with ISSR markers maps with marker data using mapping populations         ns in natural populations clonal uniformity testing		P 154				

1		 	
II- 1. 2. 3. 4. 5. 6. 7. 8. III	Molecular biology techniques- Isolation of total cellular RNA from diverse plant tissue samples, qualitative and quantitative assessment Synthesis of first strand cDNA using M-MuLV reverse transcriptase RT-PCR for analysing spatio-temporal expression pattern of candidate genes Designing artificial miRNAs Analysis of relative expression levels using qRT PCR qRT PCR for protein coding/ miRNA genes Quantitation of relative expression levels (delta delta CT method) by Livak and Schmittgen and RQ method by Knight et al. (2009) Overlapping PCR for joining promoter elements to CDS for construction of artificial gene		
1.	<ul> <li>Macromolecular analysis by Dynamic Light Scattering (DLS)</li> <li>a. To detect aggregate formations of a protein using DLS</li> <li>b. To detect the size of a protein molecule and to analyse the protein- ligand complex through DLS analysis.</li> <li>2. ELISA Assays: <ul> <li>a. To determine the Ag conc. by sandwich ELISA</li> <li>b. To determine the Ab capture by Ab capture ELISA method.</li> <li>c. To determine the Ag conc. by Ag capture ELISA method.</li> <li>d. To perform Dot-ELISA to detect an antigen.</li> </ul> </li> </ul>		
DADT	R1. Microbial Biotochnology		56
 I ANI I.	Immuno-Techniques and Assays-		50
1.	<ul> <li>Immunodiffusion and Immuno-precipitation assays:</li> <li>a. To study immunodiffusion techniques by single radial Immunodiffusion.</li> <li>b. To perform Ouchterlony double diffusion.</li> </ul>		
2.	To determine antibody concentration by using quantitative precipitin		
3.	<ul> <li>assay.</li> <li>Antibody Titrations:</li> <li>a. To detect titre value of antibodies, present in serum due to the infection of Salmonella genus causing enteric or Typhoid Fever by guertitative tube agglutination test.</li> </ul>		
	<ul><li>b. To detect the titre value of antibodies, present in test serum by using quantitative tube agglutination test</li></ul>		

	II-	Techniques in microbiology-			
	1.	Isolation and identification of a probiotic strain from a fermented drink			
	2.	16S rRNA amplification and sequencing of a mixed culture.			
	3.	Isolation and assay of phages from the environment.			
	4.	Examination of bacterial motility using soft agar medium			
	5.	Sporulation of bacteria			
	6.	Evaluating environmental bacterial isolates for antibiotic production			
	PART	B2: Plant Biotechnology			
	1.	Selfing and emasculation, setting up of controlled crosses			56
	2.	Making rooted cuttings in Sweet Basil (effect of different rooting mixtures)			
	3.	Effect of salt stress/ABA on stomatal conductance/proline			
		concentration			
	4.	Seed viability testing and grow out test			
	5.	Pollen viability testing			
	6.	Histochemical staining for transgene expression			
	7.	Plant genetic transformation			
	8.	Generation of Arabidopsis transgenics by floral dip method			
	9.	Micrografting			
	10.	Root system architecture analysis			
Eva	luation	criteria:			
1. A	ttendan	ce: 5%			
2. P	reparatio	on of lab record(s) throughout the semester: 25%			
3. E	nd seme	ster evaluation: 70% (Following components would be included)			
	a) Spot	ting: 15 %			
1	b) Viva	-voce: 15 %			
	c) Exp	eriment(s) assigned on the day of the exam: 40%			
Lea	rning o	utcomes:			
1.	Ability t	o conduct experiments with adequate safety precautions.			
2	Canacity	to compare and evaluate various approaches in solving a given ext	herime	ntal	
<u></u>	problem				
3.	Ability t	o design and interpret molecular biology experiments.			
4.	Proficie	icy in defining a research problem, drawing logical inferences from	result	ts and	
	docume	nting outcomes in systematic manner.			
Ped	agogica	<b>Approach:</b> Laboratory experiments, demonstration, writing and e	xperir	nents	
resu	lt analys	Sis.	T		
Skil	l Set:				
~					

1. Able to work in biotechnology lab and perform experiments

# **2.** Able to analyses experimental data and critical thinking.

# **Employability:**

1. Academic and industrial research

2. Industries based on biotechnology, pharmacy, and agriculture.

# Materials-

- 1. Study material and laboratory protocol will be provided by course instructor.
- 2. "Biochemistry Laboratory: Modern Theory and Techniques" Rodney Boyer, second Edition, Pearson Education, 2012.
- 3. "Analytical Techniques in Biochemistry and Molecular Biology" Rajan Katoch, Springer, 2011.
- 4. "Molecular cloning: A laboratory manual" Sambrook, Joseph. & Russell, David W. & Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y : Cold Spring Harbor Laboratory, 2001.
- 5. "DNA and protein sequence analysis. A Practical approach" Bishop M.J., Rawlings C.J. (Eds.)1997.

# Website

1. https://nptel.ac.in/

# Journals

1. Peer reviewed relevant scientific journals.

# **Advanced Reading Material**

Will be provided by instructor, if require.

# Additional information (if any)

List of experiments given in each module are representative, instructor may choose any of them for student's laboratory training as per requirements.

# **Student responsibilities**

- 1. Class attendance.
- 2. Study of course materials as specified by the instructor.

3. Regular submission of given class assignments.

# Reviewers

- 1. Prof. Bijoy Neog, Department of Life Sciences, Dibrugarh University, Dibrugarh, Assam
- 2. Dr. Rupesh Chaturvedi, Ramalingaswami Fellow, National Agri-food Biotechnology Institute, Mohali, Punjab

Course title: Introduction to Nanobiotechnology									
Course co	de:	No. of credits: 2	L-T-P: 22-	Learni	ng ho	urs:			
			08-0	30					
Pre-requi	Pre-requisite course code and title (if any):								
Faculty:		Department: Department	ment of Natural	and Appl	ied				
Sciences									
Course co	ordinator:	Course instructor							
Contact d	etails:								
Course ty	pe: Core	<b>Course offered in:</b> S	emester 2						
Course de	escription:								
Nanotechr	ology is an interdisciplinary f	ield and attracts studer	nts from various	disciplin	es. Tł	nis			
course pro	vides basic overview of nanor	materials and their app	lications. This co	ourse beg	ins w	ith a			
review of	various types of nanomaterials	s and an introduction to	o general termino	ologies.					
Subsequer	tly the course covers synthesi	s methodologies, phys	ical and chemica	al charact	erizat	ion			
of nanoma	terials. Finally, case studies il	lustrating application of	of nanomaterials	in divers	e fiel	ds			
will be dis	cussed.								
Course of	ojectives:								
1. To und	lerstand the nature and proper	ties of nanomaterials.							
2. To pro	vide scientific understanding	of application of nanor	naterials and nar	notechnol	logy i	n			
agricul	ture, health and environmenta	al conservation							
Course co	ntents				-	-			
S.No	Торіс			L	Т	P			
1.	Introduction to nanomater	ials;		4	2	0			
	• Various types of nane	omaterials, Three-dime	ensional, two-						
	dimensional, one-din	nensional and zero-dim	nensional						
	nanomaterials.								
	• Carbon nanotubes, G	raphene, Carbon dots,	metal						
	nanoparticles, metal	oxide-based nanomater	rials,						
	semiconductor nanon	naterials, quantum dots	s, hybrid						
	nanoparticles,								
	• Bio-nanomaterials, p	olymer nanoparticles,	lipid nanoparticle	es					
	etc.								
	• Synthesis methodolo	gies, Top down and bo	ottom up						
	approaches for nanor	naterial synthesis.							

2.	Properties of nanomaterials	4	4	0
	Structural properties, chemical properties, surface			
	functionalization, physical properties.			
	Characterization of nanomaterials by various analytical			
	methods, optical characterization and spectroscopy such as			
	FT-IR, UV-Vis, DLS, Zetapotential, structural			
	characterization by X-Ray Diffraction, XPS and advanced			
	microscopy (TEM, SEM, AFM) etc.			
3.	Nanobiotechnology in healthcare;	8		
	• Role of nanobiotechnology in the area of infectious & non-			
	infectious diseases			
	Nanopharmaceuticals			
	Diagnosis, sensors and biosensors			
	• Delivery vehicles, biomedical applications of nanomaterials.			
	Multimodal nanoparticles, targeted drug delivery, theranostics			
4.	Nanobiotechnology for Agriculture: Nanotechnology based tools to	6	2	0
	enhance agricultural productivity			
	• Nanobased Agri and Food Products, food preservation and			
	toxicity			
	Nanopesticides and Nanofertilizers			
	Nano-biostimulants and soil enhancers			
	Nano-enabled technologies and abiotic stress management			
	Nanobiotechnology for Crop improvement			
	Precision Delivery Systems			
	• Diagnostics and sensing			
	Nanotechnology for environment: contamination detection			
	and remediation			
		22	8	
Evaluation	on criteria:			
1. Test 1	and 2: 20% weightage to each			
2. Test 3	6 (end semester): 50% weightage			
3. Assig	nment: 10% weightage			

#### Learning outcomes:

- 1. Familiarity with working principles, tools and techniques in the field of nanomaterials.
- 2. Understanding of the strengths, limitations and potential uses of nanomaterials.

# Materials:

# **Suggested readings:**(1–7)

1. A. L. Rogach, *Semiconductor nanocrystal quantum dots synthesis, assembly, spectroscopy and applications* (Springer, Wien; London, 2008).

2. E. Gazit, *Plenty of room for biology at the bottom: an introduction to bionanotechnology* (Imperial College Press ; Distributed by World Scientific Pub. in the USA, London : Hackensack, NJ, 2007).

3. G. E. J. Poinern, *A laboratory course in nanoscience and nanotechnology* (CRC Press, Taylor & Francis Group, Boca Raton, 2015).

4. C. A. Mirkin, C. M. Niemeyer, Eds., *More concepts and applications* (Wiley-VCH, Weinheim, 2007), *Nanobiotechnology*.

5. A. K. Mishra, Ed., *Application of nanotechnology in water research* (Wiley, Scrivener Publishing, Hoboken, New Jersey, 2014).

6. K. R. Nill, *Glossary of biotechnology and nanobiotechnology terms* (Taylor & Francis, Boca Raton, 4th ed., 2006).

7. J. Kim, Ed., *Advances in nanotechnology and the environment* (Pan Stanford, Singapore, 2012).

8. P. N. Prasad. Nanophotonics (Wiley, New York, 2003).

# Websites

# Journals Other readings Additional information (if any): None

# Student responsibilities:

- 1. Study of course materials as specified by the instructor
- 2. Timely submission of given class assignment

Course reviewed by:

 Dr. Amit K Dinda, MD, Ph.D Professor Department of Pathology All India Institute of Medical Sciences, New Delhi President, Indian Society of Renal & Transplant Pathology (ISRTP) Secretary, Indian Society of Nanomedicine (ISNM) Fellow, Electron Microscopy Society of India (EMSI) dindaaiims@gmail.com

2. Dr Indrajit Roy, Ph.D Associate Professor Department of Chemistry, University of Delhi, Delhi-110007. indrajitroy11@gmail.com

Course ti	tle: Molecular F	Plant Physiology and	Metabolism				
Course code: BBP No. of credits: 2 L-T-P: 30-0-0 Learning hours			<b>s:</b> 30				
Pre-requi	isite course cod	e and title (if any):	Science graduate				
Departm	ent: Departmen	t of Natural and App	olied Sciences				
Course co	oordinator:		Course instruct	or:			
Contact d	letails:						
Course ty	v <b>pe:</b> Specializati	on	Course offered	in: Semester 2			
This course of special comprehe photomor	se is designed for ization in the M nsive knowledg phogenesis, hor <b>bjectives:</b> To provide a for	or the students who h .Sc. Biotechnology I e of molecular plant mones, water relatio	have opted for Plan Programme. The c physiology. The r ns, photosynthesis	nt Biotechnology ourse aims to pro nain topics includ and stress physic vsiological processo	as the wide a de ology.	strea	m
2. 3. 4. Course co	Knowledge of p Familiarity with	alant stress physiology secondary plant meta	and tolerance mech	nanisms			
Module	Topic				L	Т	Р
	Photomorphoge Circadian rhyth	enesis: Role of light ms, Phytochrome, C	in growth and dev Cryptochrome and	elopment, Phototropins	4	0	0
	Phytohormones biological func Gibberellins, E Salicylic acid a	: Biosynthesis, mod tions, perception and thylene, Abscisic Ad nd Jasmonic acid	e and mechanism l signaling (Auxin cid, Brassinosteroi	of action, s, Cytokinins, ds), Polyamine,	6	0	0
	Physiology of p apical, basal & and shoot apica induction and d	blant development ar radial patterning; D l meristem; M levelopment	nd flowering: Emb evelopmental cont lolecular mechani	oryogenesis, rol of root sm of floral	7	0	0
	Plant nutrients: relationships, h	Uptake and utilizati ydroponics	on, Solute transpo	ort, Plant water	2	0	0

	Physiology of biotic and abiotic stress, Molecular plant-pathogen interactions	3	0	0	
	Photosynthesis ( $C_3$ , $C_4$ and CAM), photorespiration	3	0	0	
	Metabolism of secondary metabolites in plants, Phenolics, Terpenoids and Alkaloids, biochemical and physiological significance	3	0	0	
	Biological N <sub>2</sub> fixation, Plant growth promoting Rhizobacteria, Amino acid metabolism, Urea cycle	2	0	0	
	Total	30	0	0	
Evaluation criteria:					

- 2. Test 2- (Module 3-5) 30%
- 3. Test 3- (Modules 5-9) 40%

#### Learning outcomes:

- 1. An understanding of photomorphogenesis and plant hormones (Test 1)
- 2. An understanding of floral induction and water relations and stress tolerance mechanisms (Test 1-2)
  - 3. An understanding of electron transport, secondary metabolites, and nitrogen metabolism (Test 2-3)
  - An ability of making hypotheses related to plant metabolism and development (Test 1-3)

#### **Pedagogical Approach:**

- 1. Online/classroom lectures and discussions
- 2. Case studies and examples from original research articles

# Skill Set:

- 1. Developing and screening mutants with novel traits
- 2. Ability to develop strategies for genetic improvement of crops having climate resilience

#### **Employability:**

- 1. Academic and research organisations
- 2. Tissue culture facilities and horticulture companies
- 3. Agri-biotechnology and seed companies
- 4. Pharmaceutical and drug research companies

# Materials:

#### **Suggested Readings**

- 1. Plant Physiology, Sixth Edition" by Lincoln Taiz and Eduardo Zeiger
- 2. Biochemistry & Molecular Biology of Plants by Bob Buchanan, Gruissen W and Jones R L

# Additional information (if any):

#### **Student responsibilities:**

- 1. Class attendance.
- 2. Study of reading materials as specified by course instructor
- 3. Self-study

#### **Course reviewers:**

- 1. Dr. B. P. Shaw, Scientist G, Institute of Life Sciences, Bhubaneswar, Odisha
- 2. Dr. Santan Barthwal, Scientist F, Forest Research Institute, Dehradun, Uttarakhand

Course title: Molecular Microbiology and Immunology								
Course code: BBP	No. of credits: 2	L-T-P: 30-0-0	Learning hours: 30					
Pre-requisite course code and title (if any): BBP161								
Principles of Biochemistry	Principles of Biochemistry and Biophysics (semester 1)							
Department: Department of Natural and Applied Sciences								
Course coordinator: Course instructor								
Contact details:								
Course type: CoreCourse offered in: Semester 2								

#### **Course description:**

The course is designed to provide students with basic concepts, principles and applications of molecular microbiology and immunology. The course aims to introduce microbial systems to the students and molecular basis of microbial pathogenicity and resistance. Various mechanisms employed by microbes against host immune responses will be covered at a molecular level. The course will provide information on microbial growth patterns and pathogens. Further, basic concepts in immunological responses including adaptive T cell and B cell responses will be described including T cell receptor/MHC interactions and antigen-antibody interactions. The section on molecular assays and techniques will comprehensively provide concepts and principles of major microbiological and immunological methods employed regularly in research laboratories. Finally, through this course, actual concepts and insights within cancer biology and potential anti-cancer therapies are provided.

#### **Course objectives:**

1. To introduce students to pathogens and microbial systems that have commercial applications.

2. Providing students with fundamentals of microbial growth and kinetics.

3. Familiarizing students with concepts of microbial drug resistance and the underlying molecular basis.

4. Acquainting students with basic concepts of immunology, with a focus on the molecular bases underlying TCR/pMHC and antibody/antigen interactions

5. Familiarizing students with various molecular techniques employed in microbiology and immunology.

6. Acquainting students with molecular mechanisms underlying cancer development and anti-cancer therapies.

Course contents			
Module Topic	L	Т	P
Module 1: Microbes and Microbial applications			
Pathogens (classification/structure and function), Infection life cycles of viruses			
and bacteria, Pathogen host interactions, Viral vectors in gene and cancer	2	0	0
therapy, Molecular compounds of microbial origin for agriculture, industry,	4	0	0
and pharmaceuticals			
Module 2: Microbial Kinetics			
Microbial growth and kinetics, batch, and continuous process, Microbial strain	3	0	0
improvement for pharmacologically active agents	5	0	0
Module 3: Microbial Pathogenicity and Resistance			
Molecular basis of pathogenicity and resistance of bacteria against host			
immune responses, Drug resistance: Mycobacterium tuberculosis (MDR-TB			
and XDR-TB) and Streptococcus pneumoniae, Mechanisms employed by	4	0	0
bacterial toxins (cholera, diphtheria, and tetanus), Microbial transformation of			
antibiotics			
Module 4: Antibodies and Antigens		Γ	1
Immunoglobulins- structure and function, Antigenic Determinants (isotype, allotype,			
idiotype), Antigens (types of antigens, characteristics of an antigen), Adjuvants,			
Cellular and Humoral immunity, Antigen presentation, TCR, pMHC, Monoclonal	4	0	0
antibodies (mAbs), Hybridoma technology, characterization of mAbs through epitope	-	0	0
mapping, Immune evasion mechanisms of virulent pathogens, raising antibodies in an			
animal system, Antibody and Vaccine engineering, Complement system.			
Module 5: Molecular Assays and Techniques			
5.1: Antibody Titration Techniques: Immuno assay systems, Immuno precipitin	2	0	0
reactions, ELISA, RIA, RID,	3	0	0
5.2: Immunotechniques: Yeast one hybrid, Yeast two hybrid, TAP- TAG Technology,	2		
Synthetic lethal screens, Pull down assays, expression library screening, AFM	2		
5.3: Fluorescent antibody techniques: Bimolecular fluorescence complementation			
(BiFC), Fluorescence resonance energy transfer (FRET) and Fluorescence correlation	2		
spectroscopy, Label transfer, Quantitative immunoprecipitation combined with knock-	3		
down (QUICK),			
5.4: Protein-Protein Interaction studies: PPI maps, Protein Chips for diagnostics, SPR,	•		
MST, ITC and nanoDSF, Static Light Scattering (SLS)	3		
5.5: Immunocytochemistry (cryo-sectioning, resin embedding, freeze-shattering and	2		
freeze fracture), Negative Staining, Immunogold labelling, Electron Microscopy	4		

Module 6: Cancer Biology			
Tumorigenesis, Invasion and Metastasis, Immunosuppressive mechanisms,	4		
Anti-cancer agents and Therapies	4		
Total	30	0	0
Evaluation criteria:			
1. Test 1 30%			
2. Test 2 30%			
3. Test 3 (end semester) 40%			
Learning outcomes:		-	
1. Acquaintance of basic microbial structure and microbial diversity. Grasp of various m	icrobia	al syster	ns and
applications of microbial compounds of commercial interest (Tests 1, 2 & 3)			
2. An insight into the growth patterns of microbes (Test 1 & 2).			
3. An understanding of mechanisms behind microbial pathogenicity and resistance. Stud	ents w	ill be ab	le to
outline key aspects of immune reactions and host responses against pathogens. (Test 2).			
4. Grasp of basic concepts of immunology: a. Able to define molecular machinations of	cellula	ar and h	umoral
immune responses, roles played by diverse immune cells. b. Understanding of molecular	basis	of	
immunological tolerance and autoimmunity. (Test 2).			
5. Knowledge of principles underlying the assays and techniques employed in immunolo	ogy and	l microl	biology
(Test 3).			
6. A detailed understanding of mechanistics of cancer biology (Test 3).			
Pedagogical Approach:			
1. Online/Offline teaching.			
2. Providing case studies to support the concepts.			
<b>3.</b> Peer-reviewed research articles to discuss various modules in the course.			
4. Peer-review reading			
Skill Set:			
1. Analytical skills based on case studies provided.			
2. Knowledge of immunological and microbiological applications in various	s secto	rs.	

3. Knowledge of techniques employed.

# **Employability:**

The course will provide skillsets and knowledge that may play key role to get employed in Universities, R & D industry, Medical centres/Colleges, Research Institutes and Diagnostic centres apart from specialized units like pharma, breweries, dairy and agri sectors.

#### Materials: Suggested Readings

- 1. Schroeder HW Jr, Cavacini L. Structure and function of immunoglobulins. *J Allergy Clin Immunol*. 2010;125(2 Suppl 2):S41-S52. doi:10.1016/j.jaci.2009.09.046
- 2. Peleg AY, Hogan DA, Mylonakis E. Medically important bacterial-fungal interactions. Nat Rev Microbiol. 2010 May;8(5):340-9. doi: 10.1038/nrmicro2313. Epub 2010 Mar 29. PMID: 20348933.
- 3. Vermelho AB, Supuran CT, Guisan JM. Microbial enzyme: applications in industry and in bioremediation. *Enzyme Res.* 2012;2012:980681. doi:10.1155/2012/980681
- Pham JV, Yilma MA, Feliz A, Majid MT, Maffetone N, Walker JR, Kim E, Cho HJ, Reynolds JM, Song MC, Park SR, Yoon YJ. A Review of the Microbial Production of Bioactive Natural Products and Biologics. Front Microbiol. 2019 Jun 20;10:1404. doi: 10.3389/fmicb.2019.01404. PMID: 31281299; PMCID: PMC6596283.
- Singh S, Kumar NK, Dwiwedi P, Charan J, Kaur R, Sidhu P, Chugh VK. Monoclonal Antibodies: A Review. Curr Clin Pharmacol. 2018;13(2):85-99. doi: 10.2174/1574884712666170809124728. PMID: 28799485.
- Lu RM, Hwang YC, Liu IJ, Lee CC, Tsai HZ, Li HJ, Wu HC. Development of therapeutic antibodies for the treatment of diseases. J Biomed Sci. 2020 Jan 2;27(1):1. doi: 10.1186/s12929-019-0592-z. PMID: 31894001; PMCID: PMC6939334.
- Madhurantakam C, Rajakumara E, Mazumdar PA, Saha B, Mitra D, Wiker HG, Sankaranarayanan R, Das AK. Crystal structure of low-molecular-weight protein tyrosine phosphatase from Mycobacterium tuberculosis at 1.9-A resolution. J Bacteriol. 2005 Mar;187(6):2175-81. doi: 10.1128/JB.187.6.2175-2181.2005. PMID: 15743966; PMCID: PMC1064030.
- Madhurantakam C, Chavali VR, Das AK. Analyzing the catalytic mechanism of MPtpA: a low molecular weight protein tyrosine phosphatase from Mycobacterium tuberculosis through sitedirected mutagenesis. Proteins. 2008 May 1;71(2):706-14. doi: 10.1002/prot.21816. PMID: 17975835.
- Madhurantakam C, Duru AD, Sandalova T, Webb JR, Achour A. Inflammation-associated nitrotyrosination affects TCR recognition through reduced stability and alteration of the molecular surface of the MHC complex. PLoS One. 2012;7(3):e32805. doi: 10.1371/journal.pone.0032805. Epub 2012 Mar 14. PMID: 22431983; PMCID: PMC3303804.
- Neiers F, Madhurantakam C, Fälker S, Manzano C, Dessen A, Normark S, Henriques-Normark B, Achour A. Two crystal structures of pneumococcal pilus sortase C provide novel insights into catalysis and substrate specificity. J Mol Biol. 2009 Oct 30;393(3):704-16. doi: 10.1016/j.jmb.2009.08.058. Epub 2009 Aug 31. PMID: 19729023.
- Duru AD, Sun R, Allerbring EB, Chadderton J, Kadri N, Han X, Peqini K, Uchtenhagen H, Madhurantakam C, Pellegrino S, Sandalova T, Nygren PÅ, Turner SJ, Achour A. Tuning antiviral CD8 T-cell response via proline-altered peptide ligand vaccination. PLoS Pathog. 2020 May 4;16(5):e1008244. doi: 10.1371/journal.ppat.1008244. PMID: 32365082; PMCID: PMC7224568.

- 12. Borek F. The fluorescent antibody method in medical and biological research. *Bull World Health Organ*. 1961;24(2):249-256.
- Slastnikova TA, Ulasov AV, Rosenkranz AA, Sobolev AS. Targeted Intracellular Delivery of Antibodies: The State of the Art. Front Pharmacol. 2018 Oct 24;9:1208. doi: 10.3389/fphar.2018.01208. PMID: 30405420; PMCID: PMC6207587.
- 14. Bertram JS. The molecular biology of cancer. Mol Aspects Med. 2000 Dec;21(6):167-223. doi: 10.1016/s0098-2997(00)00007-8. PMID: 11173079.
- Liu L, Wannemuehler MJ, Narasimhan B. Biomaterial nanocarrier-driven mechanisms to modulate anti-tumor immunity. Curr Opin Biomed Eng. 2021 Dec;20:100322. doi: 10.1016/j.cobme.2021.100322. Epub 2021 Jul 30. PMID: 34423179; PMCID: PMC8372976.
- 16. Günther G. Multidrug-resistant and extensively drug-resistant tuberculosis: a review of current concepts and future challenges. *Clin Med (Lond)*. 2014;14(3):279-285. doi:10.7861/clinmedicine.14-3-279

**Note:** Further updated reference and review articles will be provided during the lectures **Additional information (if any): Not Applicable** 

#### Student responsibilities:

- 1. Study of course material as specified by the instructor.
- 2. Proactive involvement in studying, reviewing and analysing the accessible scientific literature in online/offline modes.

#### **Course reviewers:**

**1. Prof. Adnane Achour,** Structural and Biophysical Immunology, Department of Medicine, Solna, Karolinska Institute, Stockholm, Sweden

**2. Dr. Rajakumara Eerappa,** Associate Professor, Department of Biotechnology, Indian Institute of Technology, Hyderabad, India

# Enclosure 4

# Revised course outline of "Climate Change and Law" offered to M.Sc. Climate Science and Policy

Course title: Climate Change and Law					
Course code	No. of credits: 2	L-T-P distribution: 24-	6-0 I	Learning ho	<b>urs</b> : 30
Pre-requisite cours	e code and title (if a	ny): None			
Department: Natur	al and Applied Science	ces			
Course coordinator	r (s):	Course instructor (s):			
Contact details:					
Course type	Elective				
Course offered in	Semester 2				
<b>Course Description</b>	l				
Climate change is o	one of the main chall	enges facing humanity tod	lay. It ha	is severe imj	olications
for the social, econo	omic, and political lif	e of people around the wo	orld. Its a	ascent as a g	lobal and
national policy age	nda has been driven	by the developments in	internati	ional law. I	ncreasing
recognition of the in	npacts of climate char	nge is also forcing other do	omains of	f law to take	note of it
and respond. Accord	dingly, the legal four	ndations of global and nat	tional go	vernance sys	stems are
going through sign	ificant changes. Thi	s course is aimed to intr	roduce the	he students	with the
processes that gover	in the legal responses	both at the national and in	nternatio	nal level to t	his grave
crisis. In addition, t	he course also looks	at the impact of this phen	omenon	on other bra	anches of
law like the law of the	he sea and human rig	hts.			
Course objectives	1 ( 1' C)	• , ,• 1 1 .•	1.1	1	1. (
1. To provide a	n understanding of th	e international and nationa	ll laws re	elating to	climate
change.	d the commission of the	ahanian annias and under (	1	ation of 1 a col	
2. To understan	id the compliance me	chamsin envisaged under t	ne mtern	lational legal	regime,
3 To opolyco th	erence to mula.	abanga an athar branchas a	flow		
Course content		mange on other branches o	<b>I</b> law	Т	р
Module 1. Introdu	ction				1
Sources of Intern	national law		-		
Sources of Intern     Science and law	nauonai iaw				
Science and law					
• Key concerns: e	quity, CBDK & RC	, polluter pays principle,			
binding characte	r climate vs develor	e, transparency, legally			
Modulo 2. Evolu	tion of Internetic	nol Logol Dogimo on	6	2	
Mouule 2: Evolu	uon of miternatio	nai Legai Regime on	U	$\angle$	

Climate Change			
• Intergovernmental Negotiating Committee-UN Framework			
Convention on Climate Change – Kyoto Protocol - Paris			
Agreement- Katowice Package-Glasgow Rulebook			
• Legal challenges of Top – Down and Bottom – Up approach,			
NDCs – Enforcement			
Montreal Protocol and the Kigali Amendment			
• International Organizations and Institutions: WMO, IPCC,			
• Current debates: Future of CBDR principle, legality of net-			
zero emission targets; now legally binding the Paris			
Agreement is, equity and legal implications of global goal of			
Module 3: Legal response to climate change in select countries	4		
The European Union	•		
United States of America			
United Kingdom			
Germany			
Module 4: Legal foundation of India's response to climate	6	4	
change	Ũ	-	
India's obligations under International law and NDCs			
NAPCC and its missions			
• Mitigation: Various acts, regulations, and authorities			
governing energy sector, urbanization, agriculture,			
buildings, transport, industry, Forests			
• Adaptation and Resilience: Disaster Management Act			
(2005), EIA, regulatory requirements/guidelines for			
resilience/disaster risk reduction in key sectors,			
Minimum Standards of Environmental Services in			
entergencies, and various sectoral acts relevant for			
• SAPCCs and their implementation: strengths and weakness of			
• SALCUS and then implementation, strengths and weakness of a federal governance systems Climate action by cities, powers			
and possibilities of climate action by cities			
<ul> <li>Climate action by non-state actors: the limits of CSR</li> </ul>			
<ul> <li>Does India need a climate change law per se?</li> </ul>			

Module 5: Impact on other areas of law and litigation (case studies)	4				
• Climate Change and the Law of the sea: Implications for sovereignty, Marine biodiversity.					
Climate change as a human rights issue					
• Trade issues – Technology Transfer – IPRs					
Litigation: Role of NGT					
Total	24	6			
Evaluation criteria					
• Class participation : 10 %					
• Term Paper : 25 % (module 1 and 2. Learning outcom	es 1 and	d 2)			
• Presentations : 25 % (Module 3 and 4: Learning outcom	e 3)				
• Test 3 : 40% (Module 1-5, Learning outcomes 2	t 3 : 40% (Module 1-5, Learning outcomes 1-3)				
Learning outcomes					
By the end of the course, it is expected that the students will:					
1. Be familiar with the international legal regime on climate change	•				
2. Be able to appreciate the concerns raised on the ground of equit	y and th	ne negotiati	ng position		
of developing countries.					
3. Be able to appreciate the functioning, context, and determina	ants of	effectivene	ss of legal		
regimes to address climate change	regimes to address climate change				
Materials					
Essential texts:					
The United Nations Framework Convention on Climate Change, 1992					

- The Kyoto Protocol, 1997
- The Paris Agreement, 2015

# **Reference Books:**

- Bodansky, D., Brunnee, J. and Rajamani, L. (2017), International Climate Change Law, Oxford: OUP.
- Carlarne, Cinnamon P., Gray, Kevin R., and Tarasofsky, Richard (eds) (2016), The Oxford Handbook of International Climate Change Law, Oxford: Oxford University Press.

# Module 1.

• Bodansky et al. (2017), Chapters 1-3

• French, D. and Rajamani, L. (2013), "Climate Change and International Environmental Law: Musings on a Journey to Somewhere", *Journal of Environmental Law*, 25 (3): 437-461.

# Module 2.

- Bodansky et al. (2017), Chapters 4-7
- Carlarne, Cinnamon (2014), "Delinking International Environmental Law and Climate Change", *Michigan Journal of Environmental and Administrative Law*, 4: 1. Available at: https://repository.law.umich.edu/mjeal/vol4/iss1/1
- Bodansky, D. (2016), "The Legal Character of the Paris Agreement", *Review of European, Comparative and International Environmental Law*, 25 (2): 142-150.

# Module 3.

- Siddi, M. (2020), *European Green Deal: assessing its current state and future implementation*, FIIA Working Paper #114, May 2020.
- Skjærseth, J.B. (2021), "Towards a European Green Deal: The evolution of EU climate and energy policy mixes", *Int Environ Agreements* **21**, 25–41. https://doi.org/10.1007/s10784-021-09529-4
- Averchenkova, A., Fankhauser, S. & Finnegan, J.J. (2021), "The impact of strategic climate legislation: evidence from expert interviews on the UK Climate Change Act", *Climate Policy*, 21:2, 251-263, DOI: 10.1080/14693062.2020.1819190
- Mildenberger, M. (2021), "The development of climate institutions in the United States", *Environmental Politics*, 30:sup1, 71-92, DOI: 10.1080/09644016.2021.1947445
- Lockwood, M. (2021), "A hard Act to follow? The evolution and performance of UK climate governance", *Environmental Politics*, 30:sup1, 26-48, DOI: 10.1080/09644016.2021.1910434
- Flachsland, C. & Levi, S. (2021), "Germany's Federal Climate Change Act", *Environmental Politics*, 30:sup1, 118-140, DOI: 10.1080/09644016.2021.1980288
- Mehryar, S. & Surminski, S. (2021), "National laws for enhancing flood resilience in the context of climate change: potential and shortcomings", *Climate Policy*, 21:2, 133-151, DOI: 10.1080/14693062.2020.1808439

# Websites:

- 1. www.congress.gov
- 2. https://ec.europa.eu
- 3. www.legislation.gov.uk

#### 4. www.bundesregierung.de

## Module 4

- Dubash, N.K., Khosla, R., Kelkar, U., and Lele, S. (2018), "India and Climate Change: Evolving Ideas and Increasing Policy Engagement", *Annual Review of Environment and Resources*, 43:1, 395-424.
- Upadhyaya, P., Shrivastava, M.K., Gorti, G., & Fakir, S. (2021), "Capacity building for proportionate climate policy: Lessons from India and South Africa", *International Political Science Review*, 42(1):130-145. doi:10.1177/0192512120963883
- Pillai, A.V. & Dubash, N.K. (2021), "The limits of opportunism: the uneven emergence of climate institutions in India", *Environmental Politics*, 30: sup1, 93-117, DOI: 10.1080/09644016.2021.1933800
- Pahuja, N., Pandey, N., Mandal, K., & Bandyopadhyay, C. (2014). *GHG Mitigation in India: An Overview of the Current Policy Landscape*, Working Paper. Washington, DC: World Resources Institute. Available online at http://www.wri. org/publication/ghgmitigation-ind-policy.
- Dutta, M. (2021), Adapting to Climate Change from a Gender and Human Rights Law Perspective: A Policy Review of India, Available at SSRN: https://ssrn.com/abstract=3993409
- Divan, S., Yadav, S. & Sawhney, R.S. (2021), *Legal Opinion: Directors' obligations to consider climate change-related risk in India*, available at https://ccli.ubc.ca/wp-content/uploads/2021/09/CCLI\_Legal\_Opinion\_India\_Directors\_Duties.pdf
- Gogoi, E. (2017), *India's state action plans on climate change: towards meaningful action*, Oxford Policy Management, New Delhi
- GoI (2000), National Agricultural Policy, Government of India, New Delhi
- GoI (2005), *National electricity policy*, Ministry of Power, Government of India, New Delhi. Available at https://powermin.nic.in/en/content/national-electricity-policy
- GoI (2007), *National urban housing and habitat policy*, Ministry of Housing & Urban Pverty Alleviation, Government of India, New Delhi. Available at https://nhb.org.in/Urban\_Housing/HousingPolicy2007.pdf
- GoI (2008), *National Water Mission under National Action Plan on Climate Change*, Ministry of Water Resources, Government of India. Available at http://mowr.gov.in/sites/default/files/Mission\_Doc\_Vol22880755143\_0.pdf
- GoI (2010), *National mission for a green India*, Ministry of Environment and Forests, Government of India. Available at http://www.indiaenvironmentportal.org.in/files/green-india-mission.pdf
- GoI (2010), National mission for sustainable agriculture, Department of Agriculture

and Cooperation, Ministry of Agriculture, Government of India. Available at http://agricoop.nic.in/sites/default/files/National%20Mission%20For%20Sustainable %20Agriculture-DRAFT-Sept-2010.pdf

- GoI (2010), National mission for sustaining the Himalayan Ecosystem under National Action Plan on Climate Change (NAPCC), Department of Science and Technology, Ministry of Scince and Technology, Government of India, New Delhi. Available at http://www.knowledgeportal-nmshe.in/Pdf/NMSHE\_MissonDocument.pdf
- GOI (2011), *Disaster management in India*, Ministry of Home Affairs. Government of India
- GoI (2012), *National water policy*, Ministry of Water Resources, Government of India, New Delhi. Available
  - at http://mowr.gov.in/sites/default/files/NWP2012Eng6495132651\_1.pdf
- GOI (2015), *India's intended nationally determined contribution*. Press Information Bureau, Government of India
- Gupta, A.K., Nair, S.S., & Singh, S. (2013), *Environmental legislation for disaster risk management*, National Institute of Disaster Management & Deutsche Gesellschaft für internationale Zusammenarbeit GmbH (GIZ)

# Websites

- http://moef.oov.in
- https://nidm.gov.in
- http://climatecasechart.com/climate-change-litigation/non-us-jurisdiction/national-greentribunal/

# Module 5

- Bodansky et al. (2017), Chapters 8-9
- Bodansky, D. (2021), "Climate Change: Reversing the Past and Advancing the Future", *AJIL Unbound*, *115*, 80-85. doi:10.1017/aju.2020.89
- Robinson, S. & Carlson, D. (2021), "A just alternative to litigation: applying restorative justice to climate-related loss and damage", *Third World Quarterly*, 42:6, 1384-1395, DOI: 10.1080/01436597.2021.1877128
- Peel, J., & Osofsky, H. (2018), "A Rights Turn in Climate Change Litigation?", *Transnational Environmental Law*, 7(1), 37-67. doi:10.1017/S2047102517000292
- Engler, C. (2020), "Transboundary Fisheries, Climate Change, and the Ecosystem Approach: Taking Stock of the International Law and Policy Seascape" *Ecology & Society*, 25:4 (43).
- Alter, K. J., (2017), *The Future of International Law*, iCourts Working Paper Series, No.

101, Northwestern Public Law Research Paper No. 17-18, Available at SSRN: https://ssrn.com/abstract=3015177

- McAdam, J. (2020), "Protecting People Displaced by the Impacts of Climate Change: The UN Human Rights Committee and the Principle of Non-refoulement", *American Journal of International Law*, *114*(4), 708-725. doi:10.1017/ajil.2020.31
- DeSombre, E. R. (2000), "The Experience of the Montreal Protocol: Particularly Remarkable, and Remarkably Particular", *UCLA Journal of Environmental Law & Policy*, 19(1): 49.
- Savaresi, A. (2016), "A Glimpse into the Future of the Climate Regime: Lessons from the REDD+ Architecture", *Review of European, Comparative and International Environmental Law*, 25 (2): 186–196.
- David, D. C., (2013), "Climate Change and the Oceans", in Harry N. Scheiber and Jin-Hyun Paik, (Eds), *Regions, Institutions, and the Law of the Sea: Studies in Ocean Governance*, Leiden: Brill Press.
- McInerney-Lankford, S. (2009). "Climate Change and Human Rights: An Introduction to Legal Issues", *Harvard Environmental Law Review*, 33: 431 437.

# Journals for further references

- Journal of International Environmental Agreements
- Climate Policy
- Climate Change and Law Review
- American Journal of International Law

**Employability:** This course exposes the students to the legal foundations of policy and action on climate change, as well as the political and economic drivers of legal framework. The students are well prepared for the jobs related to policy research and compliance.

**Student responsibilities** 

Attendance, pre-reads, critical engagement, feedback, discipline, etc.

# **Course reviewers:**

1. Dr. Anwar Sadat, Assistant Professor, Indian Society of International Law, New Delhi.

2. Dr. Jacob Joseph, Assistant Professor, National University of Advanced Legal Studies, Kochi.