UG Department of Microbiology
Add on Course: Bioinformatics and Computational Biology
Session 2019-20

# Course Coordinator Report

A free Add-On Course for UG students in the Department Microbiology, Shri Shivaji Education Society Amravati's Science College, Congress Nagar, Nagpur was held from 2<sup>nd</sup> August 2019 to 5<sup>th</sup> October 2019. The course title was "Bioinformatics and Computational Biology". It is the complete beginner to Expert Course was perfect for anyone who wants to learn Bioinformatics and Computational Biology.

The Bioinformatics and Computational Biology course is designed to provide a comprehensive introduction to bioinformatics and computational biology, focusing on sequence alignment, molecular modeling, and data analysis using various software tools. It is designed to equip students with essential skills in data analysis, computational techniques, and molecular modeling.

The course duration was 10 weeks (30 hours). Two theory classes were engaged on Friday & Saturday and one Practical was engaged in every week. The structure of marking system was 50 marks on theory paper and 40 marks on practical examination including 10 marks for internal. The question paper of theory examination was in MCQ type of 25 questions with four multiple choices. Practical examination was also taken on this course for 40 marks. Internal marks assessment was on the basis of regularity, attendance, assignment submission etc. All the 81 students were present in both theory and practical examination. The result was prepared and certificates were also distributed to the students.

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# UG Department of Microbiology Add on Course: Bioinformatics and Computational Biology Session 2019-20

To, The Principal SSES Amt's Science College, Congress Nagar, Nagpur-12

Subject: For permission to conduct the add on courses in the Department of Microbiology and Biotechnology – 2019-2020

Respected Sir,

This is to request you that, the teachers of our Microbiology and Biotechnology department have prepared the syllabus and modules of the 30 hours certificate courses for the session 2019-2020.

The details of the course module, syllabus and time table is submitted here with.

Hence please permit to run the add on courses and oblige me.

Thanking you

Date: - 21/06/2019

Yours sincerely

HEAD
Department of Microbiology
Science College, Congress Nagar,
NAGPUR.

Mrs MJMadhugn

Permitted Notices

# **UG Department of Microbiology**

# NOTICE

Date: 15/07/2019

All the students are informed that **U.G. Department of Microbiology** runs **Add on Course: Bioinformatics and Computational Biology** for the session 2019-20. Interested students of B.Sc. are requested to register their names to the course Coordinator Ms. Nupur Deshmukh on or before 30/07/2019.

# **UG Department of Microbiology**

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# U.G. DEPARTMENT OF MICROBIOLOGY, SCIENCE COLLEGE, CONGRESS NAGAR, NAGPUR

Accredited with CGPA of 3.51 at 'A+' Grade by NAAC, Bangalore
A College with Potential for Excellence
An Institutional Member of APQN
Recognized Center for Higher Learning & Research
A Mentor College under Paramarsh Scheme of UGC, New Delhi
A Mentor College under Paris Sparsh Scheme of Maharashtra State

# Add on Course for the Session 2022-23 Bioinformatics and Computational Biology

# **Course Introduction**

This course provides a comprehensive introduction to bioinformatics and computational biology, focusing on sequence alignment, molecular modeling, and data analysis using various software tools. It is designed to equip students with essential skills in data analysis, computational techniques, and molecular modeling.

# Course Objectives

- To understand the fundamental concepts of bioinformatics and computational biology.
- To develop proficiency in sequence alignment techniques.
- To learn the principles and applications of molecular modeling.
- To gain hands-on experience with bioinformatics software tools for data analysis.

Registration Date: 30/07/2019

Prof. Atul Bobdey Coordinator Dept. of Microbiology Prof. Mahendra Dhore Principal Science College, Nagpur

## Session 2019-2020

Course Co-ordinator: Ms. Nupur Deshmukh

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Instructional Strategies: Theory class, Practical, Video clips, Models etc.

Evaluation Strategies: Oral discussions and Final MCQ examination

### Course Outcomes

Upon completion of this course, students will be able to:

- Apply bioinformatics tools for sequence analysis and alignment.
- · Perform molecular modeling and understand its applications in biological research.
- Analyze biological data using computational methods.
- Utilize bioinformatics software for various biological data analysis tasks.
  - Duration of course: Ten weeks (30 Hours)

Session 2019-2020

Course Co-ordinator: Ms. Nupur Deshmukh

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Ms. Nupur Deshmukh Course-Coordinator

Add on Course

# Session 2019-2020

# Module: The Structure of Syllabus and system of evaluation

	Theory Papers and Practical		Total Marks		
Course			Internal	Practical	
Certificate Course in Bioinformatics and computational biology	Theory paper- bioinformatics and computational biology  * Theory examination will be of MCQ pattern having 25 questions each with equal marks.	50	10	40	
	* Practical examination will be based on performance evaluation in the laboratory and hands-on-training	100			

Ms. Nupur Deshmukh Add on Course Coordinator

Dr. Amitabh Halder

IQAC Coordinator
Internal Quality Assurance Cell (IQAC)

S. S. E. S. A. Science College Science College, Nagpur. Congress Nagar, Nagpur.

Prof. Mahendra Dhore

Principal

Principal

S. S. E. S. Amravati's

## Course Units and Practical Sessions

### Unit 1: Introduction to Bioinformatics

- · Topics:
- · Definition and scope of bioinformatics
- Biological databases (GenBank, EMBL, PDB)
- · Basic bioinformatics tools and software
- · Practical 1:
- · Navigating biological databases
- · Retrieving sequence data from NCBI

# Unit 2: Sequence Alignment

- · Topics:
- Types of sequence alignment (global, local)
- Algorithms for sequence alignment (Needleman-Wunsch, Smith-Waterman)
- Multiple sequence alignment (ClustalW, MUSCLE)
- · Practical 2:
- · Performing pairwise and multiple sequence alignments
- · Interpreting alignment results

## Unit 3: Molecular Modeling

- · Topics:
- · Basics of molecular modeling
- · Homology modeling
- Molecular docking and dynamics
- Practical 3:
- Building a homology model using software (e.g., SWISS-MODEL)
- Molecular docking using AutoDock

### Unit 4: Data Analysis and Visualization

- · Topics:
- · Bioinformatics data analysis techniques
- Statistical tools for data analysis
- · Visualization of bioinformatics data
- · Practical 4:
- · Analyzing sequence data using R/Bioconductor
- Visualizing data using software (e.g., PyMOL, R)

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# UG Department of Microbiology Add on Course: Bioinformatics and Computational Biology Week-wise teaching plan: (Session 2019-20)

Week	Hrs.	Syllabus	
Week 1	2	Definition and scope of bioinformatics	
	2	Biological databases (GenBank, EMBL, PDB)	
Week 2	2	Basic bioinformatics tools and software	
	2	Types of sequence alignment (global, local)	
Week 3	2	Algorithms for sequence alignment (Needleman-Wunsch, Smith-Waterman)	
	2	Multiple sequence alignment (ClustalW, MUSCLE)	
Week 4	2	Basics of molecular modelingHomology modeling	
	2	Homology modeling	
Week 5	2	Molecular docking	
	2	dynamics	
Week 6	2	Bioinformatics data analysis techniques	
Week 7	2	Statistical tools for data analysis	
Week 8	2	Visualization of bioinformatics data	
Week 9	1	Navigating biological databases Retrieving sequence data from NCBI	
	1	Performing pairwise and multiple sequence alignments Interpreting alignment results	
Week 10	1	Building a homology model using software (e.g., SWISS-MODEL) Molecular docking using AutoDock	
	1	Analyzing sequence data using R/Bioconductor  Visualizing data using software (e.g., PyMOL, R)	

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Week	Hrs.	Syllabus		
Week 1	Definition and scope of bioinformatics			
	2	Biological databases (GenBank, EMBL, PDB)		
Week 2	2	Basic bioinformatics tools and software		
	2	Types of sequence alignment (global, local)		
Week 3 2 Algorithms for sequence alignment (Needleman-Wuns Waterman)		Algorithms for sequence alignment (Needleman-Wunsch, Smith-Waterman)		
	2	Multiple sequence alignment (ClustalW, MUSCLE)		
Week 4	2	Basics of molecular modelingHomology modeling		
	2	Homology modeling		
Week 5	2	Molecular docking		
	2	dynamics		
Week 6	2	Bioinformatics data analysis techniques		
Week 7	2	Statistical tools for data analysis		
Week 8	2	Visualization of bioinformatics data		
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Week 10	1	Building a homology model using software (e.g., SWISS-MODEL) Molecular docking using AutoDock		
	1	Analyzing sequence data using R/Bioconductor  Visualizing data using software (e.g., PyMOL, R)		

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(Session 2019-20)

Time Table

w.e.f. 02/08/2019

Day	Theory
Friday	Nupur Deshmukh (R. no B9) Theory 4.00 PM - 5.00 PM
Saturday	Nupur Deshmukh (R. no B9) practical, 4.00 PM - 5.00 PM
	Nupur Deshmukh (R. no B9) Theory, 4.00 PM - 5.00 PM



# **UG Department of Microbiology**

# **EXAMINATION NOTICE**

Date:07/10/2019

All the students enrolled for Add on Course: Bioinformatics and Computational Biology for the session 2019-20 are informed that Theory and Practical Exam of the course is scheduled on 11/10/2019. All the appearing students are informed to remain present in Microbiology Laboratory at 10:30 – 11:30AM AM for Theory Exam and at 12:30PM – 5:30PM for Practical Exam.

SEAL THOUSE

# List of the Students: Add on Course- Bioinformatics and Computational Biology (Session 2019-2020)

Sr. No.	Name of Student	Signature
1)	Adase Aniket	Amblet
2)	Admane Samiksha	Som.
3)	Agashe Rashmi	Doy!
4)	Anantwar Pranjal	fragely
5)	Arghode Isha	SetraAzghad
6)	Armarkar Khushi	Chushi
7)	Bagde Sarvesh	SBagele
8)	Bobde Sakshi	Sameh
9)	Borkar Vrunda	Que,
10)	Burchunde Mahek	MBurchende
11)	Chaudhari Nidhi	W. Chaudhas
12)	Chauhan Anajali	Angelo
13)	Chavhan Aarya	Alhaucen
14)	Chavhan Sakshi	June
15)	Chavhan Sneha	Sthavary
16)	Chopkar Shruti	Church:
17)	Choure Muskan	M.chause
18)	Dave Mayank	M. chause
19)	Dehury Padmabati	( 0
20)	Deshpande Shivani	Deshpand

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7)	Bagde Sarvesh	SBOgele
8)	Bobde Sakshi	Sameh
9)	Borkar Vrunda	Que,
10)	Burchunde Mahek	MBurchende
11)	Chaudhari Nidhi	W. Chaudhas
12)	Chauhan Anajali	Angelo
13)	Chavhan Aarya	Alhaucen
14)	Chavhan Sakshi	June
15)	Chavhan Sneha	Sthawary
16)	Chopkar Shruti	Shusti.
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25)	Dube Vaishnavi	Jaistre
26)	Fulzele Sakshi	fulzele
27)	Gaikwad Snehal	Suint
28)	Gajbe Mansi	Der '
29)	Gaure Tarushi	Jarushi
30)	Gorlawar Sakshi	gague.
31)	Gour Aishwarya	Aleone
32)	Halmare Sharwari	Pols
33)	Hatwar Rajashree	Jashue.
34)	Hirapure Teneshwari	- ceneshuras
35)	Jadhav Payal	PauplJadlav
36)	Jadhav Ritika	Die.
37)	Jaronde Vaibhav	Vell.
38)	Jeevaji Nazish	Mazish
39)	Jogi Sanket	Loubs -
40)	Kali Vedanti	redant
41)	Kapse Prachi	Pkapse
42)	Karpate Harshali	flashalt.
43)	Khode Aditi	Adith.
44)	Kothale Khushi	Kothale

45)	Kshirsagar Sharvari	Vhincagas
46)	Kulkarni Kinjal	AP
47)	Kulkule Sakshi	Sulgyle.
48)	Kumar Bhavish	Blurney
49)	Kumbhare Pratik	Dhund .
50)	Lokhande Anjali	tokhande.
51)	Mathpal Himanshi	Jumand'
52)	Meshram Ruruja	tmedian
53)	Nannawre Vaishnavi	Mishnake.
54)	Nilatkar Sejal	Sejal.
55)	Ninawe Harsh	Hise -
56)	Nishane Samiksha	SNidere
57)	Palandurkar Pratiksha	Profestia.
58)	Pandey Supriya	Palent
59)	Pangul Amisha	Angha.
60)	Paralkar Anuradha	- Anuadha-
61)	Pathak Samruddhi	Sawsudhi
62)	Patil Sakshi	Rivilia
63)	Patra Kalpana	Laypens.
64)	Poddar Bhavana	2
65)	Ramteke Shruti	Spadden Zinteku -
66)	Rangari Shruti	Sund!
67)	Rangu Dipti	Drangy
68)	Rathod Atharva	Atta

21)	Dhakate Utkarsha	Officers.
22)	Dhoble Vaishnavi	Drobbe
23)	Dhoke Priyal	Dheble
24)	Dhote Janhvi	Thom,
25)	Dube Vaishnavi	Jaislie
26)	Fulzele Sakshi	fulzele
27)	Gaikwad Snehal	Snivel
28)	Gajbe Mansi	Der '
29)	Gaure Tarushi	Jarushi
30)	Gorlawar Sakshi	gagie.
31)	Gour Aishwarya	Alions
32)	Halmare Sharwari	Pals
33)	Hatwar Rajashree	Jashue.
34)	Hirapure Teneshwari	- ceneshuras
35)	Jadhav Payal	Paup Jadhar
36)	Jadhav Ritika	Die.
37)	Jaronde Vaibhav	Well.
38)	Jeevaji Nazish	Nazish
39)	Jogi Sanket	(Paulod)
40)	Kali Vedanti	Vedant
41)	Kapse Prachi	Pkapse
42)	Karpate Harshali	Hastralit.
43)	Khode Aditi	Adith.
44)	Kothale Khushi	Kothale

69)	Renge Shruti	Skenge-
70)	Roy Saptaparna	Suptarna
71)	Sahare Harshal	Henshal
72)	Sarda Sakshi	Sanda.
73)	Sawalkar Sejal	Safet.
74)	Sawane Sharayu	Saware
75)	Sharma Shubhangi	Jubhari
76)	Sharma Swati	Swah
77)	Trivedi Shikha	Strivedy
78)	Wagh Asmita	Denysu.
79)	Waghmare Danshika	waghman
80)	Wasnik Rutik	WPutik.
81)	Yelekar Radha	Radhayeleran



Mes. Nupus Deshnukh

# UG Department of Microbiology Add on Course: Bioinformatics and Computational Biology Week-wise teaching plan: (Session 2019-20)

# Theory Exam Multiple Choice Questions (MCQs) Pattern

- 1. What is the primary purpose of bioinformatics?
- · a) To study physical processes
- . b) To analyze biological data
- · c) To understand chemical reactions
- d) To diagnose diseases
- · Answer: b) To analyze biological data
- 2. Which database is commonly used for nucleotide sequences?
- a) PDB
- b) GenBank
- · c) PubMed
- d) Swiss-Prot
- · Answer: b) GenBank
- 3. What does the Needleman-Wunsch algorithm perform?
- · a) Local sequence alignment
- b) Global sequence alignment
- · c) Protein structure prediction
- · d) Molecular docking
- · Answer: b) Global sequence alignment
- 4. Which software is commonly used for multiple sequence alignment?
- a) BLAST
- b) ClustalW
- c) AutoDock
- d) PyMOL
- Answer: b) ClustalW
- 5. What is homology modeling?
- a) Predicting gene expression
- b) Aligning DNA sequences
- · c) Building a 3D model of a protein
- · d) Analyzing metabolic pathways
- . Answer: c) Building a 3D model of a protein
- 6. Which tool is used for molecular docking?
- a) MUSCLE
- b) AutoDock
- c) BLAST
- . d) R
- Answer: b) AutoDock
- 7. What is the role of BLAST in bioinformatics?
- · a) Data visualization
- · b) Sequence alignment
- · c) Molecular modeling

- · d) Statistical analysis
- · Answer: b) Sequence alignment
- 8. Which of the following is a bioinformatics visualization tool?
- a) ClustalW
- b) PyMOL
- c) BLAST
- · d) Needleman-Wunsch
- Answer: b) PyMOL
- 9. What type of data does the PDB database contain?
- · a) DNA sequences
- b) Protein structures
- · c) Metabolic pathways
- · d) Gene expression profiles
- · Answer: b) Protein structures

## 10. What is the Smith-Waterman algorithm used for?

- · a) Global sequence alignment
- b) Local sequence alignment
- · c) Phylogenetic analysis
- · d) Protein structure prediction
- · Answer: b) Local sequence alignment

# 11. Which software can be used for analyzing sequence data in R?

- · a) AutoDock
- b) BLAST
- · c) Bioconductor
- d) PyMOL
- · Answer: c) Bioconductor

### 12. Which technique is used for predicting protein-ligand interactions?

- · a) Sequence alignment
- b) Homology modeling
- c) Molecular docking
- d) Data visualization
- Answer: c) Molecular docking

### 13. What is a primary goal of molecular modeling?

- a) To edit genes
- b) To predict molecular structures
- c) To visualize metabolic pathways
- · d) To sequence DNA
- · Answer: b) To predict molecular structures

### 14. Which bioinformatics tool is used for comparing an input sequence to a database?

- a) BLAST
- b) AutoDock
- c) PyMOL
- d) SWISS-MODEL
- Answer: a) BLAST

### 15. What does the term 'bioinformatics' encompass?

- a) Only sequence alignment
- · b) Computational analysis of biological data
- · c) Physical experiments on cells
- · d) Chemical synthesis of biomolecules

- · Answer: b) Computational analysis of biological data
- 16. Which software is used for protein structure visualization?
- a) BLAST
- b) ClustalW
- c) PyMOL
- d) MUSCLE
- Answer: c) PyMOL
- 17. What is the main application of the SWISS-MODEL tool?
- a) Sequence alignment
- · b) Protein structure prediction
- · c) Data analysis
- · d) Molecular docking
- · Answer: b) Protein structure prediction

# 18. Which bioinformatics technique involves aligning three or more sequences?

- · a) Pairwise alignment
- · b) Multiple sequence alignment
- c) Molecular docking
- · d) Homology modeling
- · Answer: b) Multiple sequence alignment

# 19. What is the purpose of the EMBL database?

- a) Storing protein structures
- b) Storing nucleotide sequences
- c) Analyzing metabolic pathways
- · d) Visualizing gene expression
- Answer: b) Storing nucleotide sequences

# 20. Which software can be used for creating 3D models of biomolecules?

- · a) ClustalW
- b) SWISS-MODEL
- · c) Bioconductor
- d) MUSCLE
- Answer: b) SWISS-MODEL

## 21. What does the term 'genome' refer to?

- a) A single gene
- · b) The complete set of genes or genetic material
- c) A single protein
- · d) A metabolic pathway
- Answer: b) The complete set of genes or genetic material

### 22. Which algorithm is used for local sequence alignment?

- · a) Needleman-Wunsch
- b) Smith-Waterman
- c) ClustalW
- d) MUSCLE
- · Answer: b) Smith-Waterman

# 23. Which bioinformatics tool is used for sequence similarity searching?

- a) AutoDock
- b) PyMOL
- c) BLAST
- d) R

Answer: c) BLAST

# 24. Which type of software is MUSCLE?

- a) Sequence alignment tool
- . b) Molecular modeling tool
- · c) Data visualization tool
- d) Statistical analysis tool
- Answer: a) Sequence alignment tool

# 25. What is the main focus of bioinformatics?

- · a) Chemical synthesis
- b) Biological data analysis
- c) Physical processes
- d) Clinical diagnostics
- · Answer: b) Biological data analysis

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# UG Department of Microbiology Add-on Course: Bioinformatics and Computational Biology (Session 2019-20)

# Practical Exam Question Paper:

Subject : Bioinformatics and Computational Biology

Center : S.S.E.S.A's Science College, Nagpur

Time : 5 hrs per day

Dates : 11/10/2019

Max. Marks: 40

Q.1. Building a homology model using software (e.g. SWIS	S-MODEL) 10
Q.2. Analyzing sequence data using R/Bioconductor	10
Q.3. Viva-Voce	10
Q.4. Practical Record	10

**Total Marks 40** 

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# **UG Department of Microbiology** Add on Course- Bioinformatics and Computational Biology (Session 2019-2020) **OMR Answer Sheet**



# Shri Shivaji Education Society, Amravati's SCIENCE COLLEGE



Congress Nagar, Nagpur-12 (M.S.), India

Accredited with CGPA of 3.51 st 'Ar' greds by NAAC, Bengalore
A "College with Potential for Excelbence" identified by UGC New Gelhi. Institutional Member of APQR
Recognized Contre for Higher Learning and Research
Mentor College under 'PARAMARSH Scheme', UGC, New Delhi

# U.G. DEPARTMENT OF MICROBIOLOGY

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UG Department of Microbiology

Mark List: Add on Course- Bioinformatics and Computational Biology

(Session 2019-2020)

Sr. No.	Name of Student	Marks obtained out of 50 (Theory)	Marks obtained out of 40 (Practical)	Marks obtained out of 10 (Internal)	Total Marks 100	Grade
1)	Adase Aniket	50	37	10	97	0
2)	Admane Samiksha	48	36	10	94	0
3)	Agashe Rashmi	42	35	10	87	A+
4)	Anantwar Pranjal	48	36	10	94	0
5)	Arghode Isha	44	35	10	89	A+
6)	Armarkar Khushi	50	34	10	94	0
7)	Bagde Sarvesh	48	35	10	93	0
8)	Bobde Sakshi	42	34	10	86	A+
9)	Borkar Vrunda	48	36	10	94	0
10)	Burchunde Mahek	50	38	10	98	0
11)	Chaudhari Nidhi	46	39	10	95	0
12)	Chauhan Anajali	42	34	10	86	A+
13)	Chavhan Aarya	44	35	10	89	A+
14)	Chavhan Sakshi	46	39	10	95	0

15)	Chavhan Sneha	48	36	10	94	0
16)	Chopkar Shruti	48	36	10	94	0
17)	Choure Muskan	44	35	10	89	A+
18)	Dave Mayank	46	39	10	95	0
19)	Dehury Padmabati	42	34	10	86	A+
20)	Deshpande Shivani	48	36	10	94	0
21)	Dhakate Utkarsha	48	36	10	94	0
22)	Dhoble Vaishnavi	50	38	10 ,	98	0
23)	Dhoke Priyal	46	39	10	95	0
24)	Dhote Janhvi	48	36	10	94	0
25)	Dube Vaishnavi	42	34	10	86	A+
26)	Fulzele Sakshi	46	39	10	95	0
27)	Gaikwad Snehal	50	38	10	98	0
28)	Gajbe Mansi	48	36	10	94	0
29)	Gaure Tarushi	50	38	10	98	0
30)	Gorlawar Sakshi	48	36	10	94	0
31)	Gour Aishwarya	44	35	10	89	A+
32)	Halmare Sharwari	42	34	10	86	A+

33)	Hatwar Rajashree	50	38	10	98	0
34)	Hirapure Teneshwari	46	39	10	95	0
35)	Jadhav Payal	50	38	10	98	0
36)	Jadhav Ritika	44	35	10	89	A+
37)	Jaronde Vaibhav	42	34	10	86	A+
38)	Jeevaji Nazish	48	36	10	94	0
39)	Jogi Sanket	48	36	10	94	0
40)	Kali Vedanti	48	36	10	94	0
41)	Kapse Prachi	50	38	10	98	0
42)	Karpate Harshali	44	35	10	89	A+
43	Khode Aditi	46	39	10	95	0
44	Kothale Khushi	50	38	10	98	. 0
45	Kshirsagar Sharvari	42	34	10	86	A+
46	Kulkarni Kinjal	50	38	10	98	0
47	Kulkule Sakshi	48	36	10	94	0
48	Kumar Bhavish	50	38	10	98	0
4	Kumbhare Pratik	48	36	10	94	0
5	Lokhande Anjali	50	38	10	98	0

51)	Mathpal Himanshi	50	38	10	98	0
52)	Meshram Ruruja	48	36	10	94	0
53)	Nannawre Vaishnavi	42	34	10	86	A+
54)	Nilatkar Sejal	50	38	10	98	0
55)	Ninawe Harsh	48	36	10	94	0
56)	Nishane Samiksha	50	38	10	98	0
57)	Palandurkar Pratiksha	48	36	10	94	0
58)	Pandey Supriya	48	36	10	94	0
59)	Pangul Amisha	50	38	10	98	0
60)	Paralkar Anuradha	46	39	10	95	0
61)	Pathak Samruddhi	50	38	10	98	0
62)	Patil Sakshi	50	38	10	98	0
63)	Patra Kalpana	42	34	10	86	A+
64)	Poddar Bhavana	44	35	10	89	A+
65)	Ramteke Shruti	50	38	10	98	0
66)	Rangari Shruti	48	36	10	94	0
67)	Rangu Dipti	50	38	10	98	0
68)	Rathod Atharva	42	35	10	87	A+

9)	Renge Shruti	44	35	10	89	A+
70)	Roy Saptaparna	48	36	10	94	0
71)	Sahare Harshal	42	34	10	86	A+
72)	Sarda Sakshi	46	39	10	95	0
73)	Sawalkar Sejal	50	38	10	98	0
74)	Sawane Sharayu	48	36	10	94	0
75)	Sharma Shubhangi	42	34	10	86	A+
76)	Sharma Swati	50	38	10	98	0
77	Trivedi Shikha	44	35	10	89	A+
78	Wagh Asmita	48	36	10	94	0
79	Waghmare Danshika	48	36	10	94	0
80	Wasnik Rutik	42	34	10	86	A+
8	Yelekar Radha	46	39	10	95	0





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# SCIENCE COLLEGE, CONGRESS NAGAR, NAGPUR

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# CERTIFICATE

Mr./Ku. Adase Aniket

is awarded with certificate on successful completion of the

course entitled, Certificate course in "Bioinformatics & Computational Biology"

Session 2019-20 under Add-on course conducted for 30 hours from 02/08/2019 to 05/10/2019 by Department of Microbiology, SSESA's, Science College, congress Nagar, Nagpur 440012.

He/She has passed the Examination with '0' Grade.

Ms. Nupur Deshmukh

Coordinator, Department of Microbiology

Prof. M. P. Dhore

Principal, Science College, Nagpur