

Criterion-1: Curricular Aspects

Key Indicator – 1.3: Curriculum Enrichment Metric: 1.3.3

Programme: M.Sc. Genetics

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Annexure-I List of Students

Number of MSc students who have undertaken research projects in the department

Year	S.No	Names	Gender	Roll No	Numbers
2018	1	Arushi Shukla	F	61371	11
	2	Ayush Kumar	M	61372	
	3	Bharti	F	61373	
	4	Himani Singh	F	61374	
	5	Prerna Aggarwal	F	61375	
	6	Privakshi Gogoi	F	61376	
	7	Roopam Sharma	F	61377	
	8	Sabita Yadav	F	61378	
	9	Sonu Yadav	F	61379	
	10	Srashti Jyoti Agrawal	F	61380	
	11	Vikram Sen	M	61381	
2019	1	Bm Minhajuddin	М	61571	14
	2	Deepali Mishra	F	61572	
	3	Hungharla Hungyo	F	61573	· · · · ·
	4	Neelesh Prashant	М	61574	
	5	Priya	F	61575	2
	6	Rohit Kachhwaha	М	61576	
	7	Sayeed Qadir Danishiar	М	61577	
	8	Shikha Gautam	F	61578	
	/ 9	Shruti Khanna	F	61579	
	10	Simran Choudhary	F	61580	
	11	Vagisha Kulshreshtha	F	61581	
	12	Vaishali Kataria	F	61582	
	13	Zoha Sadaqat	F	61583	
	14	Yogita	F	61382	
2020	1	ANKITA MADDHESHIYA	F	61771	8
	2	AYUSH GOEL	М	61772	
	3	BHAWNA	F	61773	
	4	HIMANSHI GANGWAR	F	61774	
	5	RASHI ANAND	F	61776	
	6	RASHMI SHARMA	F	61777	
	7	SHREESH PRATAP SAMRAT	M	61778	
	8	SUKRIT MAHAJAN	M	61779	
2021	1	AMIT KUMAR BHATT	M	61971	12
	2	ANUPAM CHAWLA	M	61972	
	3	DEEPTI THAPLIYAL	F	61973	
	4	KHANGEMBAM			10
		NONGTHANGLEIMA DEVI	F	61974	
	5	MOHD AAQUIB ,	M	61975	
	6	NGATHINGWON KASAR	F	61976	
	7	OSHEEN TANEJA	F	61977	
	8	PRIYA KHADGAWAT	F	61978	
	9	SHABNAM	F	61979	
	10	SHIVANI GAHLOT	F	61980	

	11	SIDAK MINOCHA	M	61981	1
	12	YOGEETA GUPTA .	F	61983	
2022	1	ALISHA SRIVASTAVA	F	20/1256	14
	2	DEEPTI ROY	F	20/1258	
	3	KARTIKEYA	M	20/1261	
	4	MANASVI CHOPRA	F	20/1257	
	5	NEELAM	F	20/1254	
	6	NEERU YADAV	F	20/1252	
	7	PARUL KUMAR	M	20/1253	
	8	PRAGATI VIRMANI	F	20/1260	
	9	PREETI SINGH	F	20/1262	
	10	SHURUTI SOOD	F	20/1251	
	11	SUMITA RANI	F	20/1255	
	12	VANDANA	F	20/1259	
	13	VARSHA	F	20/1263	
	14	YATHARTHA KUMAR	M	19/1258	

Total Number : 59

Prof Jagreet Kaur Head, Department of Genetics (Sign and Stamp)

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Annexure-II

Sample Project Reports

Artificial miRNA Technology and Its Applications in Developing Stress Tolerant Crop Plants

Dissertation (Review) submitted to the Department of Genetics, University of Delhi, South Campus for the partial fulfilment of the degree of M. Sc. in Genetics



Alisha Srivastava

Department of Genetics University of Delhi - South Campus Benito Juarez Road, New Delhi 110021

(May 2022)

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CERTIFICATE

This is to certify that **Ms. Alisha Srivastava** has worked on the dissertation (Review) entitled **"Artificial miRNA technology and its applications in developing stress tolerant crop plants"** under my guidance. This dissertation being submitted to the Department of Genetics, University of Delhi, South Campus for the partial fulfilment of the degree of **M. Sc. in Genetics**. All the sources of information used in this review have been duly acknowledged. This dissertation has not been submitted to any other University for the award of any other degree.

Prof. M. V. Rajam Guide

Date: May 4, 2022

Date: May 4, 2022

Signature of the HoD

Abbreviations

amiRNA	-	Artificial micro RNA
DCL-1	-	Dicer like 1
EXPO1	-	Exportin-1
HEN-1	-	Hua enhancer
HIGS	-	Host-induced gene silencing
HYL-1	-	Hyponastic leaves
miRNA	-	Micro RNA
PTGS	-	Post-transcriptional gene silencing
RdRP	-	RNA-dependent RNA polymerase
RISC	-	RNA induced silencing complex
RNAi	-	RNA interference
ROS	-	Reactive oxygen species
SDN	-	Small RNA degrading nucleases
siRNA	-	Small interfering RNA
sRNAs	-	Small RNAs

Trafficking of Small Regulatory RNAs Between the Host Plants and Their Pathogens

DISSERTATION (REVIEW) SUBMITTED TO THE DEPARTMENT OF GENETICS,

UNIVERSITY OF DELHI IN PARTIAL FULFILMENT FOR THE DEGREE OF

Master of Science in Genetics



Shuruti Sood

Department of Genetics University of Delhi, South Campus New Delhi 110021

(May 2022)

Certificate

This is to certify that **Ms. Shuruti Sood** has worked on the dissertation (Review) entitled **"Trafficking of Small Regulatory RNAs Between the Host Plants and Their Pathogens"** in partial fulfilment for the **Degree of Master of Science in Genetics** under my guidance. All the sources of information used in this review have been duly acknowledged. This dissertation has not been submitted anywhere else for the award of any other degree.

Prof. M. V. RAJAM (Guide) Head of the Department

Abbreviations

AGO	-	Argonaute proteins
CMV	-	Cucumber mosaic virus
DCL	-	DICER like protein
dsRNA	-	Double-stranded RNA
EVs	-	Extracellular vesicles
EXPO	-	Exocyst Positive Organelle
GUS	-	β-Glucuronidase
HEN1	-	Hua Enhancer 1
HIGS	-	Host-induced gene silencing
hpRNA	-	Hairpin RNA
HYL1	-	Hyponastic leaves 1
miRNA	-	MicroRNA
natsiRNA	-	Natural antisense short interfering RNA
nt	-	Nucleotide(s)
pre-miRNA	-	Precursor miRNA
pri-miRNA	-	primary miRNA
PTGS	-	Post-transcriptional gene silencing
RISC	-	RNA-induced silencing complex
RNAi	-	RNA interference
SE	-	Serrate proteins
SIGS	-	Spray-induced gene silencing
siRNA	-	Small interfering RNA
sRNAs	-	small RNAs
TET	-	Tetraspanin
TGS	-	Transcriptional Gene Silencing
TRV	-	Tobacco rattle virus
VSRS	-	Viral suppressors of RNA silencing

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Abstract

Various non-coding RNAs have been known to play a regulatory role in cellular pathways. The structure and biogenesis of small RNAs (sRNAs) define their mode of action. Although plant-pathogen interactions are mediated by different types of sRNAs, the roles of the sRNAs are still to be explored. Plants express sRNAs for a number of processes such as defense, immune response and growth. On the contrary, pathogens or pests also take support of sRNA to suppress plant immune response. The movement of these sRNAs between plants and pathogens is a critical step in the process of cross-kingdom communication. Both pathogens and plants have developed ways of delivering the sRNAs across kingdoms, in a safe and efficient manner. This review majorly focuses on sRNA trafficking and cross-talk between distinct organisms and the ways of delivery of these sRNAs.

The Convoluted Role of FOXO in Polyglutamine disorders: A review

Dissertation submitted in partial fulfilments of the requirements for the Degree of MASTER OF SCIENCES in

GENETICS

By DEEPTI ROY

Under the supervision of DR SURAJIT SARKAR



Department of Genetics University of Delhi New Delhi-110021

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Abstract

Polyglutamine diseases are a family of neurodegenerative conditions which arise from CAG triplet repeat expansion in a specific gene. This mutated gene produces a pathogenic protein containing a critically expanded glutamine tract. These diseases impact every 10 out of 1,00,000 people, so developing novel, effective therapeutics is vital. Previously, SIRT1 has been a prime target for drugs, but new research has emphasized the potential role of FOXO in neuroprotection, especially in neurological disorders. The Forkhead Box O (FoxO) transcription factor family is a crucial player in an evolutionarily conserved pathway downstream of insulin and insulin-like growth factor receptors. They are also well-known for their ability to coordinate upstream stress signals (like increased oxidative stress in the cell and reduced glucose levels) to activate fate-determination processes like survival or apoptosis. Owing to the potential FOXO presents, this article explores the role of FOXO in polyglutamine disorders its potential as a therapeutic target.

Keywords: Polyglutamine disorders, FOXO, neurodegenerative diseases, *drosophila* melanogaster, insulin pathway

Functional Ramifications of *Drosophila* Toll Pathway beyond Immunity and Development

Dissertation (Review) submitted to the

Department of Genetics University of Delhi, South Campus

For the partial fulfilment of the degree of M. Sc. in Genetics



Neeru Yadav

Department of Genetics University of Delhi - South Campus Benito Juarez Road, New Delhi 110021

(2022)

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 - 7.1. G protein-coupled receptor kinase 2 (Gprk2)
 - 7.2. Glycogen synthase kinase (GSK3β)
 - 7.3. c-Jun N-terminal kinase (JNK)
 - 7.4. let-7
- 8. Conclusion and future perspectives
- 9. Acknowledgements
- 10. References

Abbreviations

4E-BP	-	eIF4E binding protein
AD	-	Alzheimer's disease
ALS	-	Amyotrophic Lateral Sclerosis
AMP	-	Anti-microbial peptides
APOE4	-	Apolipoprotein E
A-T	-	Ataxia-Telangiectasia
ATM	-	Ataxia telangiectasia mutated gene
Αβ42	-	Amyloid β-protein 42
Bcl-2	-	B-cell lymphoma 2
CHMP2B	-	Charged multivesicular body protein 2b
CNS	-	Central nervous system
DD	-	Death domains
Dif	-	Dorsal- related immunity factor
dsRNA	-	Double stranded ribonucleic acid
FoxO	-	Forkhead Box O
FTD	-	Frontotemporal dementia
GNBP1	-	Gram-negative binding protein 1
GNBP3	-	Gram-negative binding protein 3
Gprk2	-	G protein-coupled receptor kinase
GSK3	-	Glycogen synthase kinase 3
Hid	-	Head involution defective
IFN-γ	-	Interferon γ
IL-18	-	Interleukin 18
IL-1β	-	Interleukin 1 beta
IL-6	-	Interleukin 6
IMD	-	Immune deficiency
		i

JAK-STAT	-	Janus kinase/signal transducers and activators of transcription
JNK	-	c-Jun N-terminal kinase
IncRNA-CR33942	-	long non-coding ribonucleic acid CR33942
mHtt	-	Mutant huntingtin protein
miRNA	-	Micro-ribonucleic acid
ModSP	-	Modular serine protease
mRNA	-	Messenger ribonucleic acid
NF-kB	-	Nuclear factor-kB
NFTs	-	Neurofibrillary tangles
NLP-29	-	Neuropeptide-like protein-29
PAMPs	-	Pathogen-associated molecular pattern molecules
PD	-	Parkinson's disease
PGRP-SA	-	Peptidoglycan recognition protein SA
PGRP-SD	-	Peptidoglycan recognition protein SD
PRRs	-	Pattern recognition receptors
PSH	-	Persephone
RIPK1	-	Receptor-interacting protein kinase 1
RNAi	-	RNA (ribonucleic acid) interference
ROS	-	Reactive oxygen species
Ser129	-	Serine 129
SPE	-	Spätzle processing enzyme
TGF-	-	Transforming growth factor β
TLR	-	Toll like receptor
TNF-α	-	Tumor necrosis factor-a
TOR	- '	Target of rapamycin
UTR	-	Untranslated Region
XBP1	-	X-Box Binding Protein 1

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ABSTRACT

Neurodegenerative diseases that are characterized by progressive loss of neurons and synaptic junctions are increasingly affecting the health of the society at large. For decades, the brain and immune response were studied exclusively as the central nervous system was thought to be an immune privileged region. Growing evidences have now linked hyperactivation of the immune response to various neurodegenerative diseases. However, the molecular pathway(s) that bridges these two events is yet to be explored. To study the same, one can exploit the functional homology between humans and *Drosophila* in both neurodegeneration and immune response. Recent studies have reported the involvement of Toll pathway in mediating neurodegeneration but the conduit of events are still enigmatic. Finding the conventional and non-conventional channels that connect the Toll pathway with neurodegeneration would help to develop better therapeutic interventions. In the present review, we have summarized the role of the Toll pathway in various *Drosophila* models of neurodegeneration; non-conventional interactions of the Toll pathway in different physiological conditions. Lastly, I have proposed a conciliatory pathway that links the Toll pathway to neurodegeneration.

Keywords: Neurodegeneration, Drosophila, Innate immunity, Toll pathway

DISSERTATION REPORT

On

Drug screening and standardization of effective drug concentration against poly(Q) toxicity in *Drosophila* poly(Q) models.

By

SHABNAM M.Sc. Genetics



Faculty of Interdisciplinary Sciences Department of Genetics University of Delhi, South Campus

Supervised By Dr. Surajit Sarkar Assistant Professor, Department of Genetics University of Delhi, South campus New Delhi-110021

CERTIFICATE

This is to certify that Ms. Shabnam, student of M.Sc. Genetics (Final), has completed the dissertation work titled "**Drug screening and standardization of effective drug concentration against poly(Q) toxicity in** *Drosophila* **disease models.**" under the guidance and supervision of Dr. Surajit Sarkar, Department of Genetics, University of Delhi South Campus, New Delhi-110021.

Dr. Surajit Sarkar Department of Genetics University of Delhi, South campus New Delhi-110021

Date:

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Genotype-Phenotype association of seed Glucosinolate trait in *Brassica juncea*

DISSERTATION REPORT

Submitted to University of Delhi for the partial fulfilment of

> MASTER OF SCIENCE IN GENETICS

> > 2018-2020



SUBMITTED BY

HIMANSHI GANGWAR

Under the guidance of

Prof. AKSHAY K. PRADHAN

DEPARTMENT OF GENETICS UNIVERSITY OF DELHI SOUTH CAMPUS NEW DELHI-110021

Declaration

I, **Himanshi Gangwar**, hereby declare that my dissertation entitled "**Genotype-Phenotype** association of seed Glucosinolate trait in *Brassica juncea*" is the bonafide record of the original research work carried out by me under the supervision and guidance of **Prof. Akshay K. Pradhan**. It has not been submitted earlier elsewhere for the award of any degree, diploma, or fellowship.

I certify that all sources of information and data are fully acknowledged in the dissertation Report.

Date: May 29, 2020

Himanshi Gangwar

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Review of literature

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Materials and methods

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[TO INVESTIGATE THE ACTIVITY OF TP53 GENE UNDER STRESS]

DISSERTATION PROJECT REPORT SUBMITTED TO THE UNIVERSITY OF DELHI IN PARTIAL FULFILLMENT FOR THE DEGREE OF

Master of Science In Genetics

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June 2021

Certificate

This is to certify that **Ms Khangembam Nongthangleima Devi** has worked on the research topic entitled "TO INVESTIGATE THE ACTIVITY OF TP53 GENE UNDER STRESS" in partial fulfillment for the degree of **Master of Science in Genetics** under my supervision. The dissertation work embodies the original work of the candidate and has not been submitted in full or part to any other university for the award of any other diploma or degree.

Prof.Dr. Tapasya Srivastava (Supervisor)

Head of the Department

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PROJECT WORK

ON

CLONING OF THE DNA BINDING DOMAIN 0F P53 IN His tag PROTEIN EXPRESSION VECTOR, pET28a(+)



DISSERTATION

submitted to

Department of Genetics, University of Delhi South Campus

BY

NGATHINGWON KASAR

M.Sc GENETICS(2021)

Supervisor: Dr.Pradeep K. Burma Professor, Department of Genetics University of Delhi- South Campus New Delhi-110021

Certificate

This is to certify that Ms. Ngathingwon Kasar has carried out her dissertation project on the topic "Cloning of the DNA Binding Domain of p53 protein in His tag protein expression vector pET28a(+)" for the partial fulfillment of Degree of Masters of Science in Genetics from University of Delhi under my guidance and supervision. The content of this thesis is original and has not been submitted in full or in part in this University or elsewhere for award of any degree or diploma.

Dr.Pradeep K.Burma Supervisor

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- 2. OBJECTIVE AND EXPERIMENTAL DESIGN
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Expression profiling of Ayurveda Prakriti-specific Susceptibility Genes in Rheumatoid Arthritis - A Pilot Study

DISSERTATION PROJECT REPORT SUBMITTED TO THE UNIVERSITY OF DELHI IN PARTIAL FULFILLMENT FOR THE DEGREE OF

Master of Science In

Genetics



Amit Kumar Bhatt

Department of Genetics University of Delhi, South Campus New Delhi – 110021 INDIA

10 June 2021

Certificate

This is to certify that **Mr. Amit Bhatt** has worked on the research topic entitled "**Expression Profiling of Ayurveda Prakriti-specific Susceptibility Genes in Rheumatoid Arthritis -A Pilot Study**" in partial fulfillment for the degree of **Master of Science in Genetics** under my supervision. The dissertation work embodies the original work of the candidate and has not been submitted in full or part to any other university for the award of any other diploma or degree.

Prof. B.K. Thelma (Supervisor)

Head of the Department

LIST OF COMMONLY USED ABBREVIATIONS

- 1. bp Base pairs
- 2. Ct Cycle Threshold
- 3. GWAS Genome Wide Association Study
- 4. MCT Micro Centrifuge Tube
- 5. NTC No Transcription Control
- 6. NFW Nuclease Free Water
- 7. PCR Polymerase Chain Reaction
- 8. RT-PCR Real-Time Polymerase Chain Reaction
- 9. RA Rheumatoid arthritis
- 10. RPM Rotations/revolutions per minute
- 11. RT Room Temperature
- 12. SD Standard Deviation

NOTATIONS

- 1. ^OC Degree Celsius
- 2. sec-Second
- 3. min Minute
- 4. kDa Kilo Dalton
- 5. mL Milli Litre
- 6. μ L Micro Litre
- 7. mg Milligram
- 8. ng Nanogram

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Overexpression and purification of Mtg3p truncates, containing either deletion of mitochondrial targeting sequence (MTS) or both MTS and C-terminus, from *Escherichia coli*.

Dissertation project report

Submitted to Department of Genetics University of Delhi South Campus University of Delhi South campus New Delhi-110021, India New Delhi-110021

Submitted by: **Rashmi Sharma** M.Sc. (F) Genetics Semester- 4th Roll no- 61777 Supervised by: Dr. Kaustav Datta Assistant Professor Department of Genetics University of Delhi, South Campus

Certificate

This is to certify that Ms. **Rashmi Sharma** has carried out her dissertation project entitled "**Overexpression and purification of Mtg3p truncates, containing either deletion of mitochondrial targeting sequence (MTS) or both MTS and C-terminus, from** *Escherichia coli.*" towards the partial fulfillment of Degree of Masters of Science in Genetics from University of Delhi under my guidance and supervision. The content of this thesis is original and has not been submitted in full or in part in this University or elsewhere for award of any degree or diploma.

> Dr. Kaustuv Datta Supervisor

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5. Material and methods

Abbreviations

Abbreviations	Full forms
a.a.	Amino acid
Amp	Ampicllin
CaCl ₂	Calcium Chloride
cpGTPase	Circularly permuted GTPase
dNTP	Deoxyribonuclotide
DMSO	Dimethylsulfoxide

EtBr	Ethilidium bromide
Kan	Kanamycin
LB	Lysis Broth
MgCl ₂	Magnesium Chloride
mtLSU	Mitochondrial ribosomal Large Subunit
mtSSU	Mitochondrial ribosomal Small Subunit
O/N	Overnight
RT	Room temperature
SDS	Sodium dodecyl sulfate
Tet	Tetracycline
w/o	Without
wrt	With respect to
Zn^{2+}	Zinc

1. INTRODUCTION

1.1 Ribosomes

It is known that mitochondrial ribosomes or mitoribosomes have bacterial origin, but they have substantial divergence from their ancestors and are prominently distinct from their cytosolic counterparts (Amunts et al, 2014).

Mitoribosomes are made up of nuclear encoded proteins and mitochondrial encoded rRNAs. The ordered assembly of the functional structure in the matrix involves an intricate play between synthesis of mitochondrial encoded rRNA and expression and import of cytoplasmic factors. This process, as of yet, remains elusive. A myriad of factors such as rRNA modifying enzymes, RNA helicases, GTPases and chaperones are seen to be involved in this process, but the functional details at many points. (Kressler et al.,2010 ; De Silva et al.,2015).

GTPases represent the major class of players essential for the ribosomal assembly in bacterial and the mitochondria as well. The ribosome assembly GTPases function either by acting as sensors of the GTP/GDP ratio (thus, coupling energy status of the cell to ribosomal assembly) or drive the assembly by channeling the energy released from GTP hydrolysis. They may act via binding to the RNA and/or the ribosomal protein. (Britton, 2009; Karbstein, 2007).

Developing a Yeast One Hybrid (Y1H) assay for studying interaction between transcription factors from plant and their putative *cis*-elements

Dissertation project report submitted to

Department of Genetics University of Delhi South Campus New Delhi-110021

Submitted by: Shreesh Pratap Samrat M.Sc. (F) Genetics Semester- 4th Roll no- 61778 Supervised by: **Pradeep Kumar Burma** Professor Department of Genetics University of Delhi South campus New Delhi-110021, India

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Cloning of Membrane Glycoprotein CD133 gene in *E. coli* DH5α from non-small cell lung carcinoma cell line A549

Submitted By Sukrit Mahajan

TO THE UNIVERSITY OF DELHI FOR PARTIAL FULFILMENT OF THE DEGREE

MASTER OF SCIENCE IN GENETICS



Under the Supervision of

Dr. Tapasya Srivastava

Assistant Professor

Department of Genetics

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DECLARATION BY THE STUDENT

I hereby declare that the thesis 'Cloning of Membrane Glycoprotein CD133 gene in *E. coli* DH5 α from non-small cell lung carcinoma cell line A549' is an original work carried out by me, under the guidance of Dr. Tapasya Srivastava, Assistant Professor, Department of Genetics, University of Delhi and has not been submitted to University of Delhi or any other university, for the award of any degree.

Date: 31.5.2020

Place: New Delhi

Sukrit Mahajan

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LIST OF ABBREVIATIONS

- 1. ABC ATP Binding Cassette
- 2. CD Cluster of Differentiation
- 3. CSC Cancer Stem Cell
- 4. DEPC Diethyl pyro carbonate
- 5. DMEM Dulbecco's Modified Eagle Medium
- 6. DMSO Dimethyl sulfoxide
- 7. dNTP deoxyribonucleoside triphosphate
- 8. EtBr Ethidium bromide
- 9. FBS Fetal Bovine Serum
- 10. FLIP Flice Like Inhibitory Protein
- 11. GBSC Glioblastoma Stem Cell
- 12. HDAC Histone deacetylase
- 13. HEK Human Embryonic Kidney
- 14. LB Agar Luria Bertani Agar
- 15. PBS Phosphate Buffer Saline
- 16. PCR Polymerase Chain Reaction
- 17. PI3K Phosphoinositide 3-kinase
- 18. TGF Transforming growth factor
- 19. TNF Tumour necrotic factor
- 20. NSCLC Non-Small Cell Lung Cancer
- 21. OCT4 Octamer Binding Transcription Factor 3/4 (OCT4)
- 22. SOX2 SRY-box containing gene 2

To study the role of accessory factors in mitochondrial function in *Saccharomyces cerevisiae*

Dissertation project report

Submitted to Department of GeneticsUniversity of Delhi South Campus University of Delhi South campus New Delhi-110021, India New Delhi-110021

Submitted by: **Priya Khadgawat** M.Sc. (F) Genetics Semester- 4th Roll no- 19252753008 Supervised by: Dr. Kaustav Datta Assistant Professor Department of Genetics

Certificate

This is to certify that Priya Khadgawat has carried out her dissertation project entitled "To govern the role of accessory factors in mitochondrial function in *Saccharomyces cerevisiae*" towards the partial fulfillment of Degree of Masters of Science in Genetics from University of Delhi under my guidance and supervision. The content of this thesis is original and has not been submitted in full or in part in this University or elsewhere for award of any degree or diploma.

> Dr. Kaustuv Datta Supervisor

University of Delhi, South Campus

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Mitochondrial translation and regulation
MRX8
G-Domain
Mitochondrial Ribosomes
MTG3
IRC3

II. Objectives:

- 1) To observe the growth phenotype of wild type and knock-out MRX8.
- 2) To perform a mating assay with Δ MRX8 strain to obtain it's mating type.

3) To determine the primary consequence of GKSAAA mutant on mrx8p by analyzing protein stability.

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- 7) To create stable clones in a suitable expression system for MTG3 truncates

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- 7) To create stable clones in a suitable expression system for MTG3 truncate

V. Material and methods

Abbreviations:

Abbreviation	Full
a.a	Amino acid
Amp	AmpicIlin
CaCl ₂	Calcium Chloride
cpGTPase	Circularly permuted GTPase
DMSO	Dimethylsulfoxide
EtBr	Ethilidium bromide
Kan	Kanamycin
LB	Luria-Bertani
MgCl ₂	Magnesium Chloride
d.d.H ₂ O	Double Distilled Water
YPG	Yeast, Peptone, and Glycerol
YPD	Yeast, Peptone, and Dextrose
rpm	rotations per minute
O/N	Overnight
RT	Room temperature
SDS	Sodium dodecyl sulfate
PAGE	Polyacrylamide Gel Electrophoresis
Tet	Tetracycline
H_2O_2	Hydrogen Peroxide
Ab	Antibody
Zn ²⁺	Zinc

Development of transgenic tomato for early blight disease by targeting *Alternaria solani* specific genes *CHS* and *Hog1* (HOG pathway MAP Kinase) using artificialmiRNA (amiRNA) bicistronic construct

DISSERTATION PROJECT REPORT SUBMITTED TO THE UNIVERSITY OF DELHI IN PARTIAL FULFILLMENT FOR THE DEGREE OF

Master of Science In Genetics



Osheen Taneja

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2021

Certificate

This is to certify that **Ms. Osheen Taneja** has worked on the research topic entitled "Development of transgenic tomato for early blight disease by targeting *Alternaria solani* specific genes *CHS* and *Hog1* (HOG pathway MAP Kinase) using artificial-miRNA (amiRNA) bicistronic construct" in partial fulfillment for the degree of Master of Science in Genetics under my supervision. The dissertation work embodies the original work of the candidate and has not been submitted in full or part to any other university for the award of any other diploma or degree.

Prof. M. V. RAJAM (Supervisor) Head of the Department

LIST OF COMMONLY USED ABBREVIATIONS

- 1. LBA Luria Broth Agar
- 2. LB Luria Broth
- 3. GTE Glucose Tris EDTA
- 4. rpm Rotations per minute
- 5. mins Minutes
- 6. MCT Micro Centrifuge Tube
- 7. RT- Room Temperature
- 8. RE Restriction Enzyme
- 9. FD- Fast Digest
- 10. amiRNA artificial micro RNA
- 11. dNTP- Deoxyribonucleotide triphosphate
- 12. bp Base pairs
- 13. PDA- Potato Dextrose Agar
- 14. PCA Potato Carrot Agar

NOTATIONS

- 1. ^{O}C Degree Celsius
- $2. \ sec-Second$
- 3. min Minute
- 4. kDa Kilo Dalton
- 5. mL Milli Litre
- 6. μ L Micro Litre
- 7. mg Milligram
- 8. ng Nanogram

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