

Cell cultures and molecular investigation on *Polygonum tinctorium* and *Indigofera tinctoria* plants to understand indican biosynthesis

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ABSTRACT

Indican biosynthesis in *Indigofera tinctoria* and *Polygonum tinctorium* plants is of topical interest due to its academic and industrial relevance for indigo dye production. Dye yield depends on indican content in the plant biomass. Cell culture and molecular biological investigations were carried out to assess indican biosynthesis in these plants. Tissue culture protocols were optimized for explant identification, decontamination, *in vitro* culture medium & suitable growth regulators and culture conditions for *P. tinctorium* at Okayama University of Science, and *I. tinctoria* at CSIR-NEERI. Four different growth hormones i.e. BA, Kin, NAA, 2,4-D, at 0.01 – 2 mg L⁻¹ culture medium, in random combination, and two different explants i.e. leaf and internode of both plants were experimented. In both plants, callus proliferation was better from leaf tissue with growth index (GI) up to 10 on MS agar gelled medium fortified with BA+NAA in comparison to BA+2,4-D. Suspension cell cultures of *I. tinctorium* were induced in MS liquid medium with only 2,4-D through 3 stages with GI up to 30. *In vitro* raised cell biomass of *I. tinctorium* presented higher indican synthesis ($p > 0.5$) in comparison to that of *P. tinctorium*. Both of these plants synthesize indican, but the differential response under *in vitro* is interesting. Total transcriptomes of both plants were worked out and annotated. Comparative analysis of transcriptome profile indicated > 80% genes are similar for the indican biosynthetic pathways. Complete alignment of both transcriptomes and validation for biosynthesis pathways specific genes is needed in both the plants to ascertain their differential

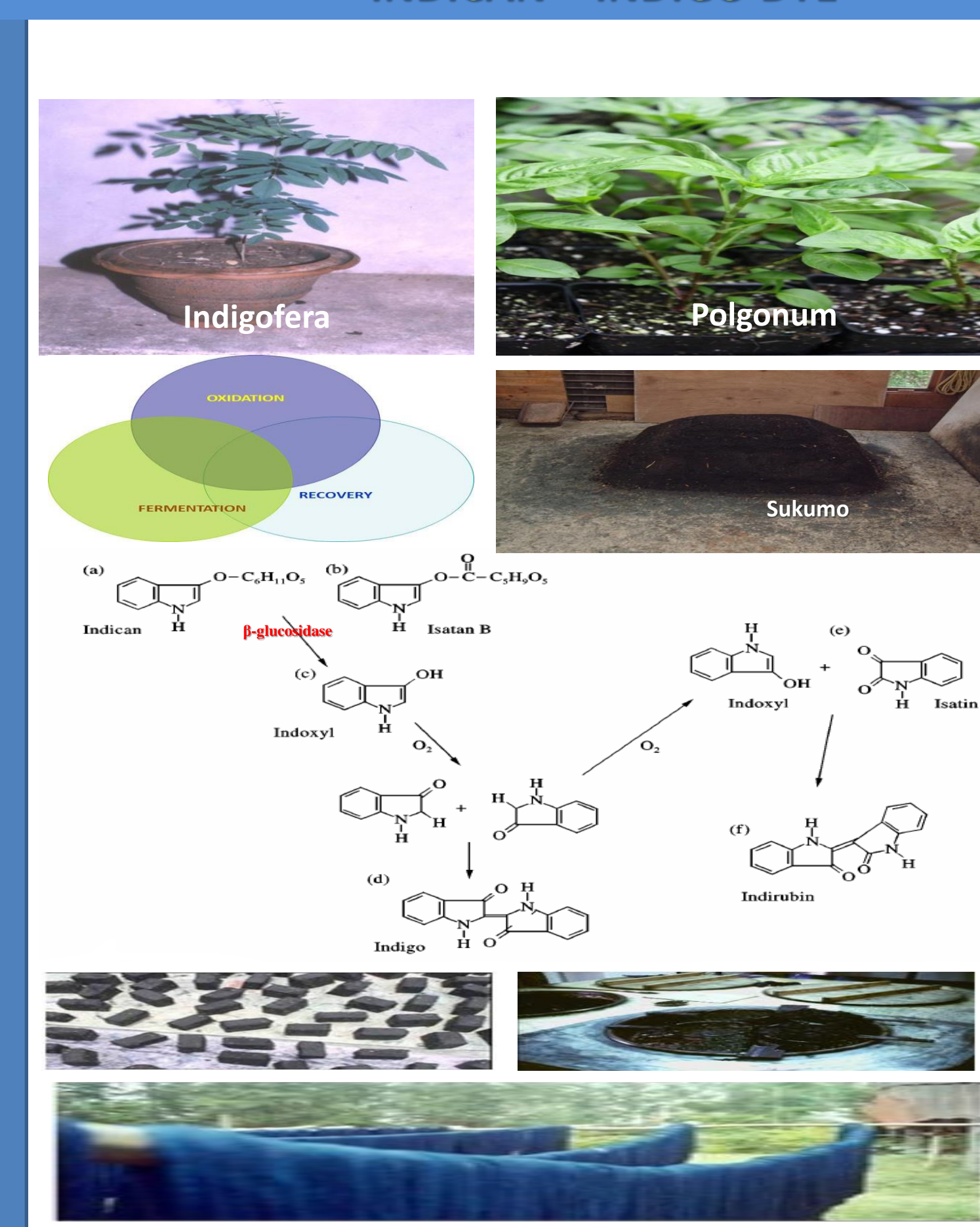
GENESIS



TISSUE CULTURE – IN VITRO BIOMASS PRODUCTION



INDICAN – INDIGO DYE

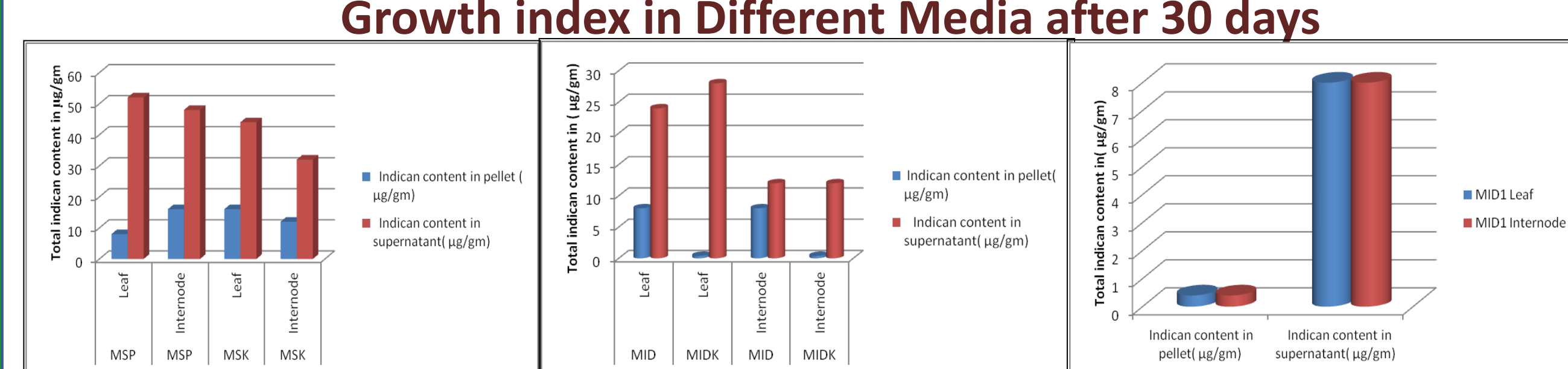
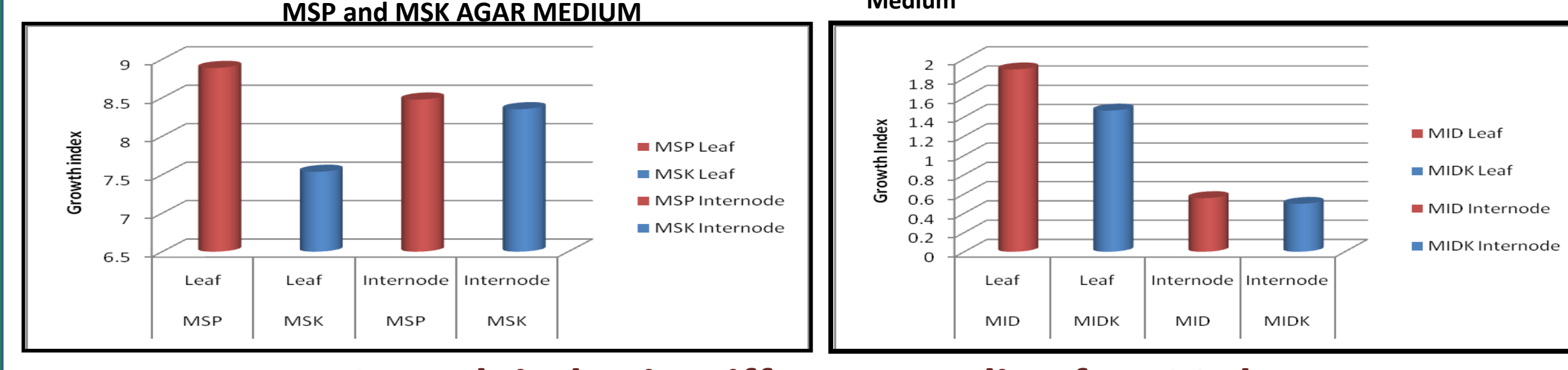
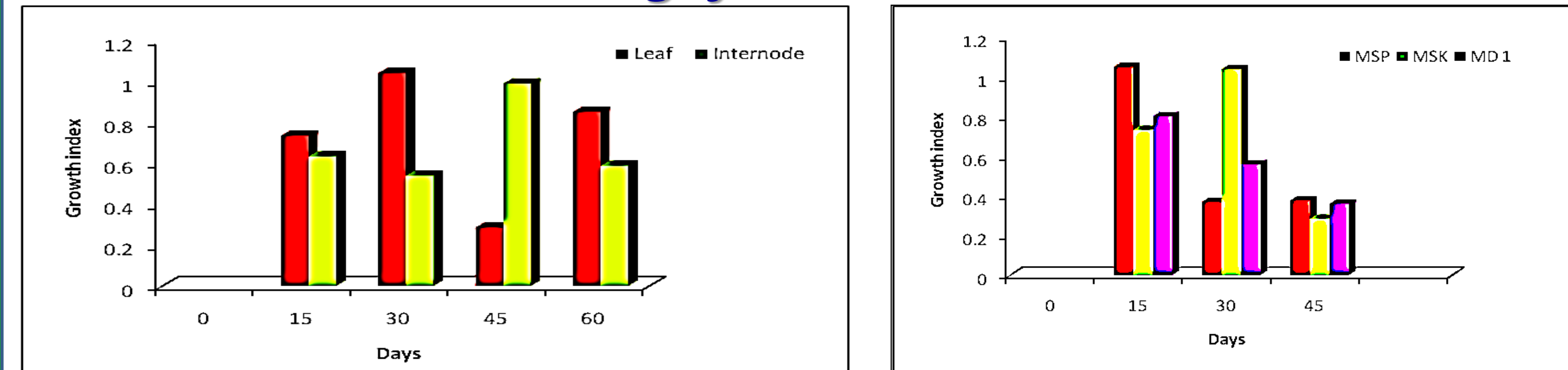


Polygonum tinctorium

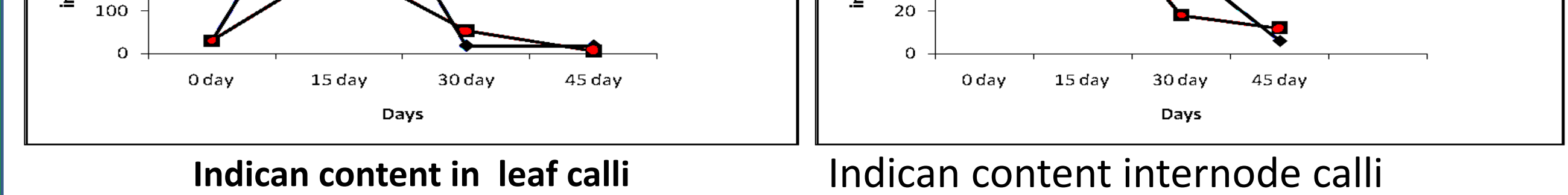
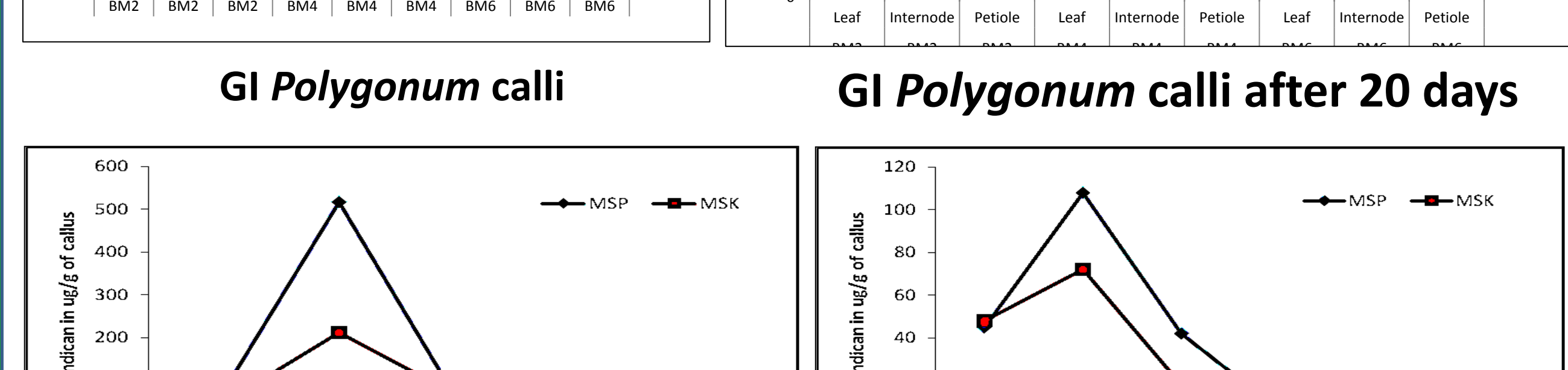
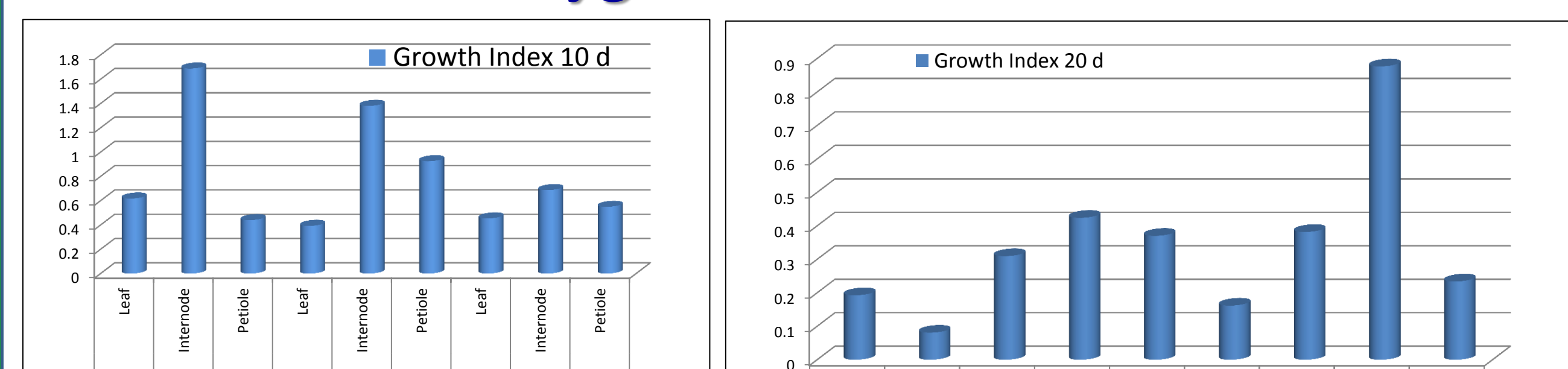
METHODS

MSP- Murashige and Skoog medium with 0.5mg/L BA + 2mg/L NAA
MSK- Murashige and Skoog medium with 0.5mg/L Kn + 2mg/L NAA
MID- Murashige and Skoog medium with 0.25mg/L 2,4-D
MIDK- Murashige and Skoog medium with 0.25mg/L 2,4-D + 0.1mg/L Kn
MIDI- Murashige and Skoog medium with 0.5mg/L 2,4-D

Indigofera tinctoria



Polygonum tinctorium



Indigofera and Polygonum transcriptome deposited in NCBI GenBank database

Sample: SAMN03015981 - Transcriptomic analysis of Indigo dye plant, *Indigofera tinctoria* L., using Illumina HiSeq2000 sequencing platform (SRR156381)

Organism: *Indigofera tinctoria*

Attributes: cultivar: tinctoria; bioserial: provider: CSIR-NEERI, Nehru Marg, Nagpur - 440020, India; tissue: Leaf

Library: IT_Lib (SRR156381)

Strategy: RNA-Seq

Source: TRANSCRIPTOMIC

Selection: cDNA

Layout: SINGLE

Platform: Illumina (SRR156381)

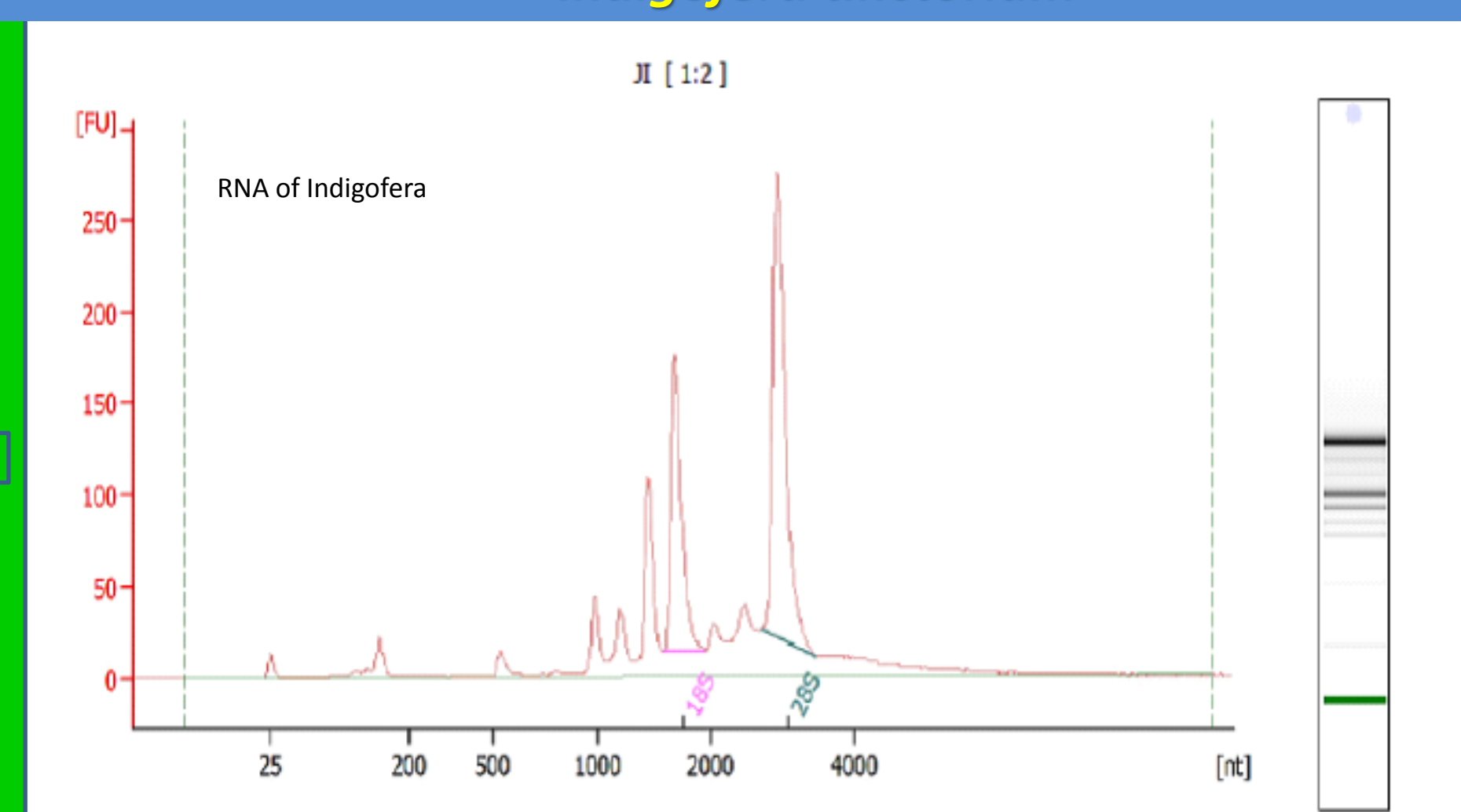
Instrument model: Illumina HiSeq 2000

Spot descriptor: 1 forward

Experiment attributes: Authors: Bijaya K. Sarangi, Sanjog T. Thul and Yoshiko Mina; Title: Indigo dye plant, *Indigofera tinctoria* L., using Illumina HiSeq2000 sequencing platform (SRR156381); Total: 1 run, 26.7M spots, 4.8G bases, 3Gb

The cDNA transcriptome sequence of indigo dye plant, *Indigofera* & *Polygonum* has been inventoried in GenBank (Acc. No. SAMN03015981 & SRX692542).

Indigofera tinctoria



Overall Results for sample 9:

RNA Area: 1,331.7
 RNA Concentration: 505 ng/ul
 rRNA Ratio (28S / 18S): 1.8

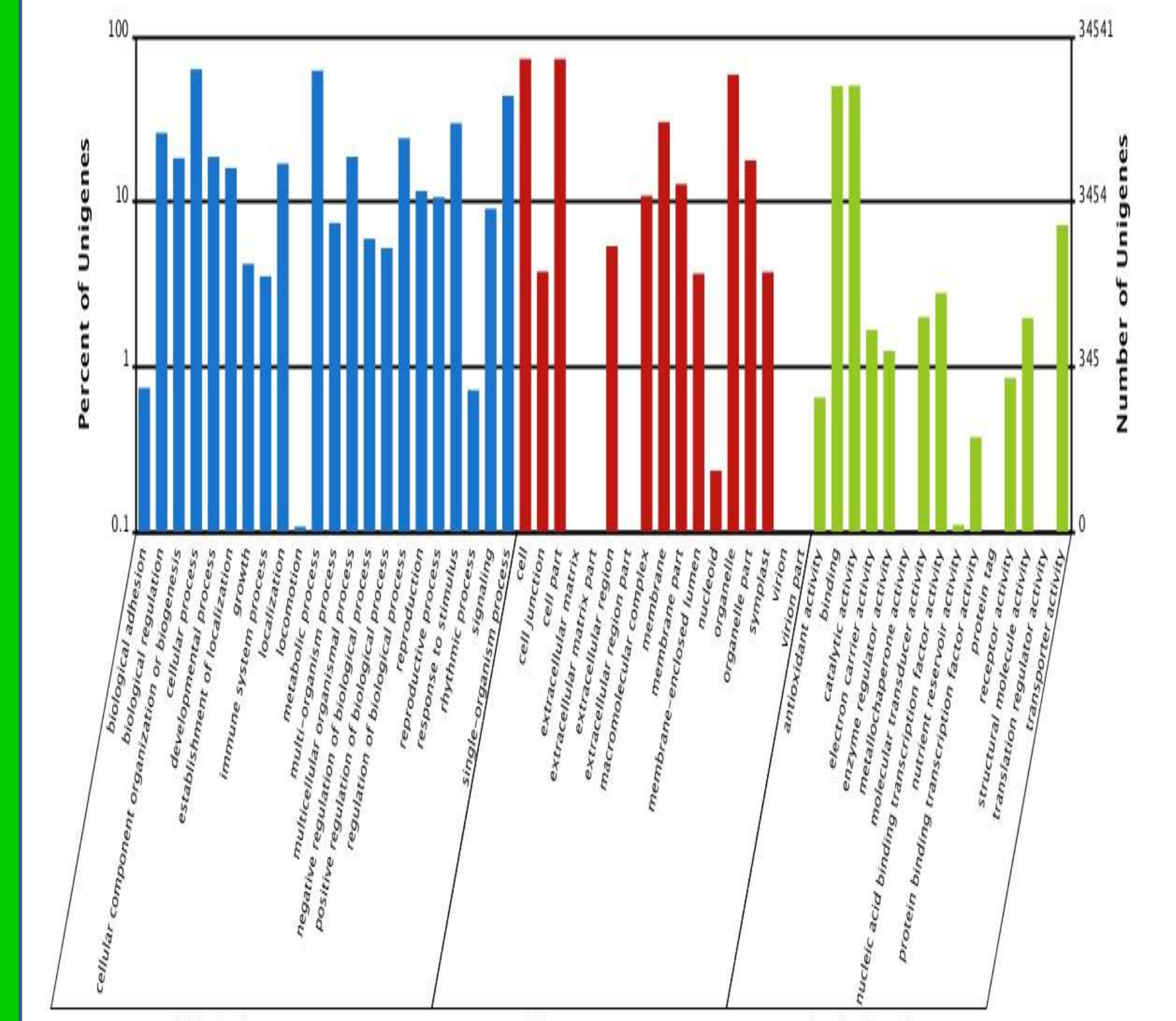
RNA Integrity Number (RIN): 8.4 (B.02.07, Anomaly Thresholds manually accepted)

Result Flagging Color:
 Result Flagging Label: RIN: 8.40

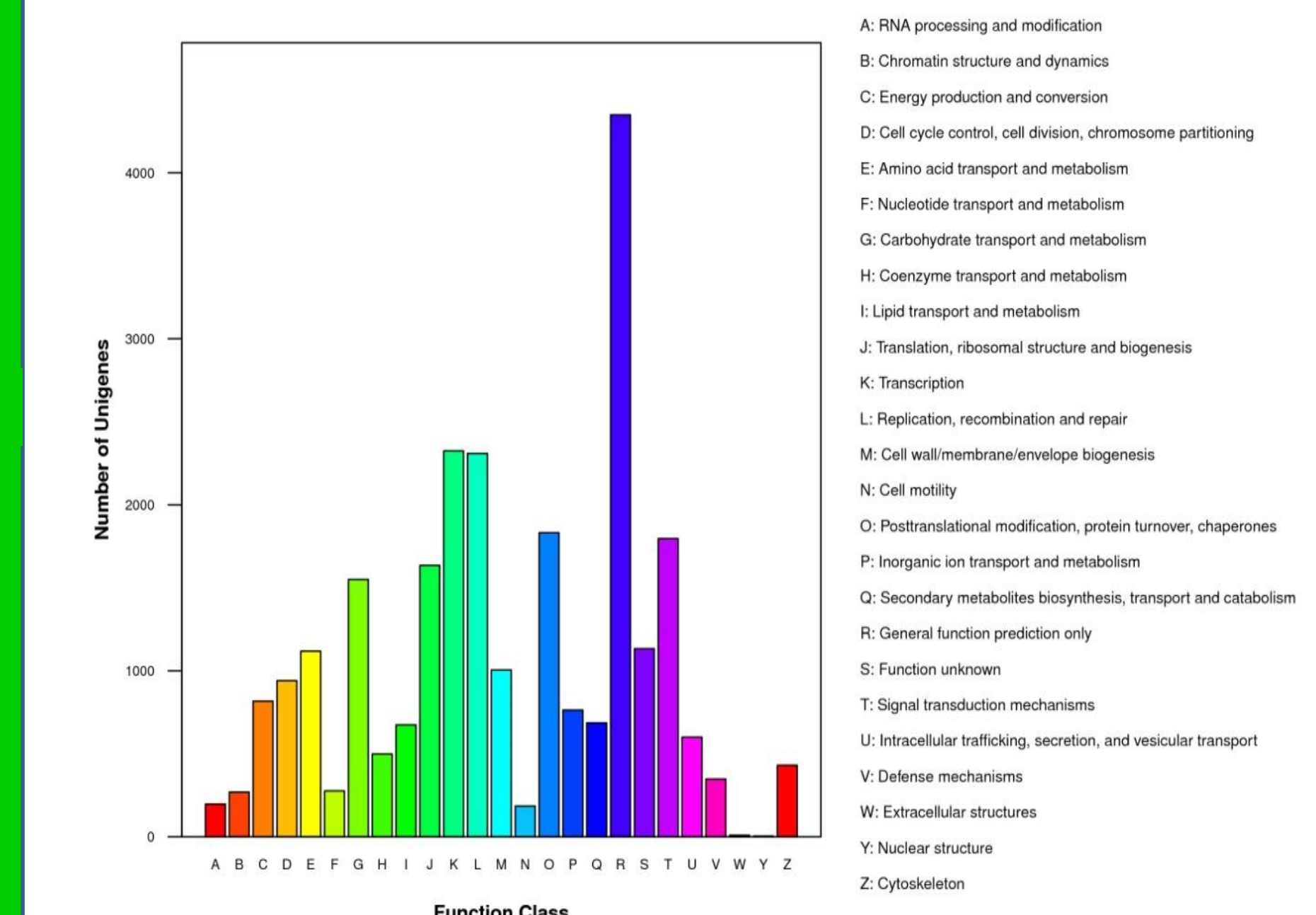
Fragment table for sample 9:

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,581	1,996	210.4	15.8
28S	2,682	3,463	374.8	28.1

Total RNA



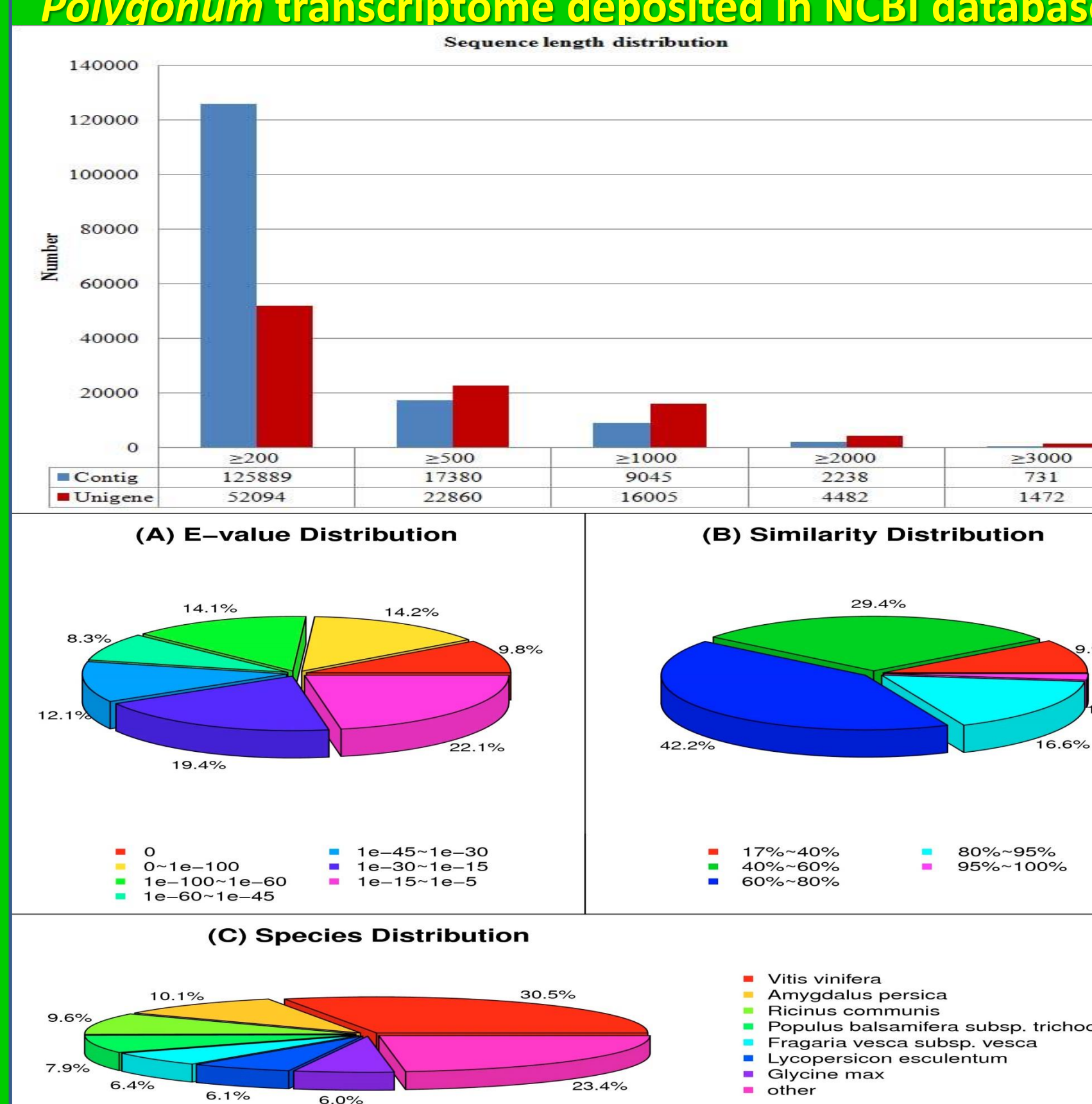
At least 23,721 unigenes mapped onto 128 pathways involved in secondary metabolite biosynthesis using the KEGG pathway database.



Unigenes mapped onto 128 pathways using the KEGG pathway database involved in the biosynthesis of secondary metabolites

A total of 60,395 indigo biosynthesis putative unigenes were obtained from transcriptomes of *Indigofera tinctoria* (L.) through cDNA sequencing and assembly study.

Polygonum transcriptome deposited in NCBI database



A total of 96,913 indigo biosynthesis putative unigenes were obtained from transcriptomes of *Polygonum tinctorium* through cDNA sequencing and assembly study.

CONCLUSION - FUTURE SCOPE

INDICAN BIO-SYNTHESIS STRATEGIES

- Clone β-glucosidase gene (*Polygonum* / microbial) to expression vector under strong promoter
- Identify genes and enhance tryptophan synthesis
- Metabolomics

Biotechnological

- Mass culture of tissue with high inherent indican content
 - Selection of clonal variants
 - Induced mutation to select mutant cell lines with enhanced indican content
 - Rapid cell growth and differentiation with wild genes through indirect gene transfer
 - Enhance quantitative traits through para sexual hybridization
 - Genetic transformation
- Contact:** minami@dbc.ous.ac.jp ; bk_sarangi@neeri.res.in

INDIGO DYE PRODUCTION STRATEGIES

- Biomass Fermentation:**
- Complete extraction of indican from biomass
 - Reduce fermentation duration
 - Reduce biomass : water ratio
- Ferment Oxidation:**
- Mechanical aeration arrangement
 - Optimize DO concentration to prevent rate limiting
 - Reduce oxidation time