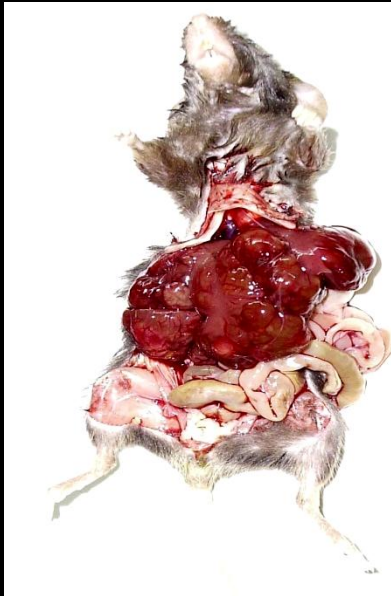


# *Butea monosperma in Cancer Treatment*



**Dr.G.Mathan**  
**Assistant Professor**  
**Department of Biomedical Science**  
**Bharathidasan University**  
**Tiruchirappalli -620 024**  
**Tamil Nadu, India**

# X 15- myc Transgenic Mouse Model for HCC



X15- myc Transgenic Mouse

➤ Recombinant bicistronic construct (Singh *et al*, 2003) (Kumar *et al*, 2001) US patent no: 6274788 B1.

➤ X-15 myc transgenic mice appeared to be an ideal model to study the disease process and also screening drugs.

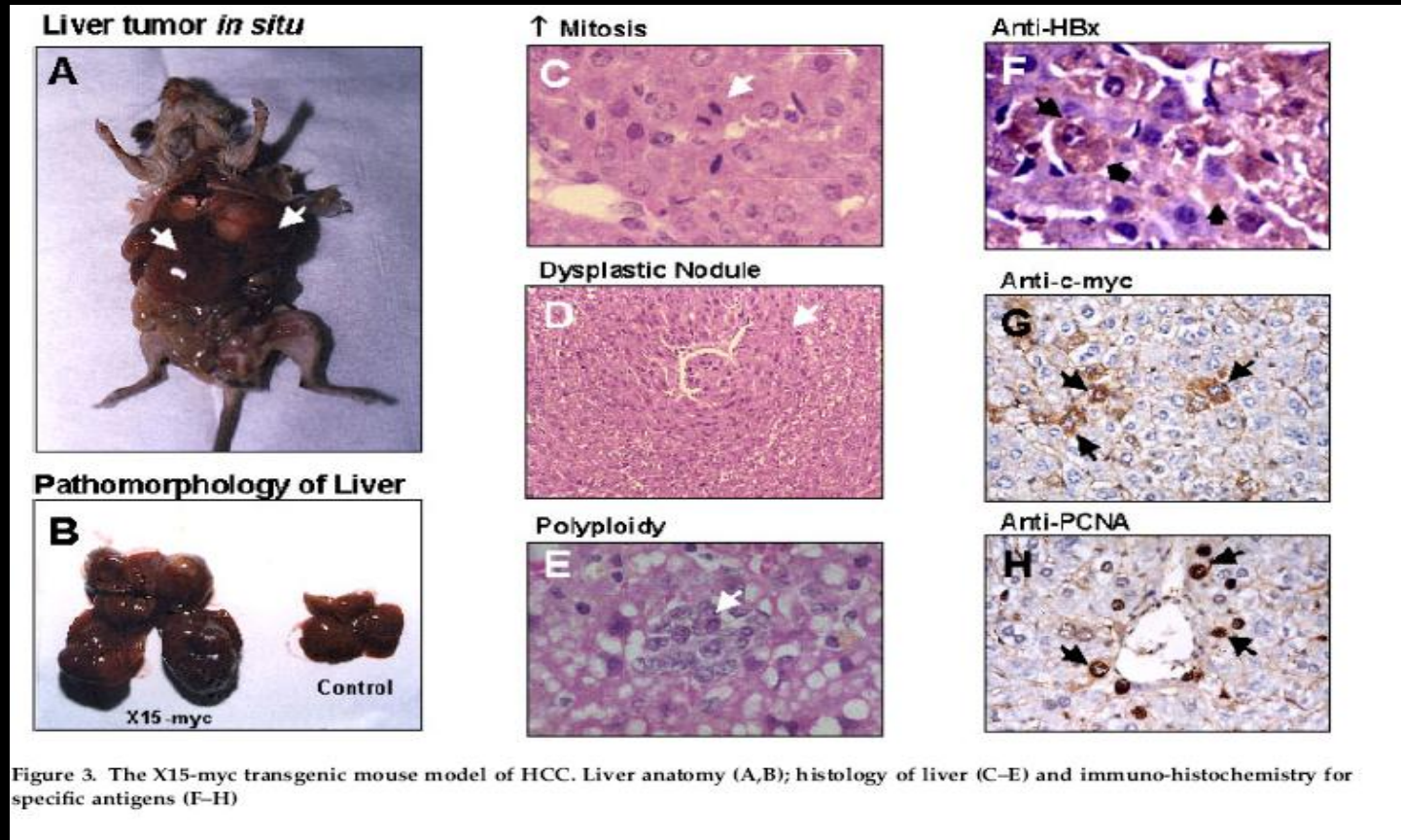


Normal Liver

X15- myc Tg Liver

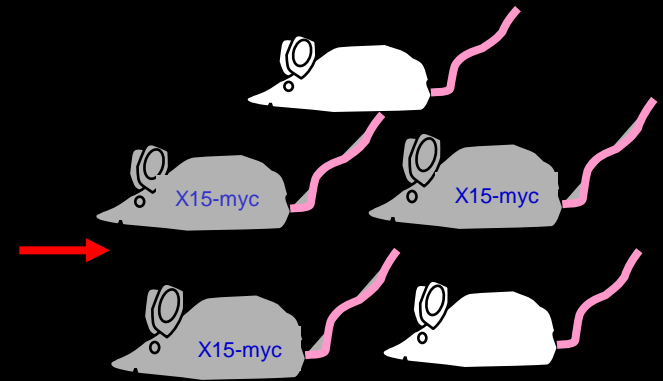
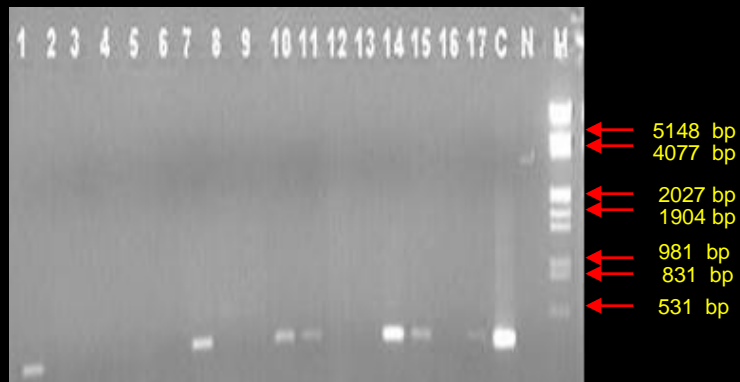
Transgene	expressed Promoters	used Mouse strain	Pathology	Reference
HBx and c-myc	Xp and Cp	C57BL/6 x SJL	HCC in the first half of animals' life span (3 to 5 months)	1. Singh M,et.al., 1998 2. Kumar V,et.al., 2001(us patent)

# X15-myc transgenic mouse model

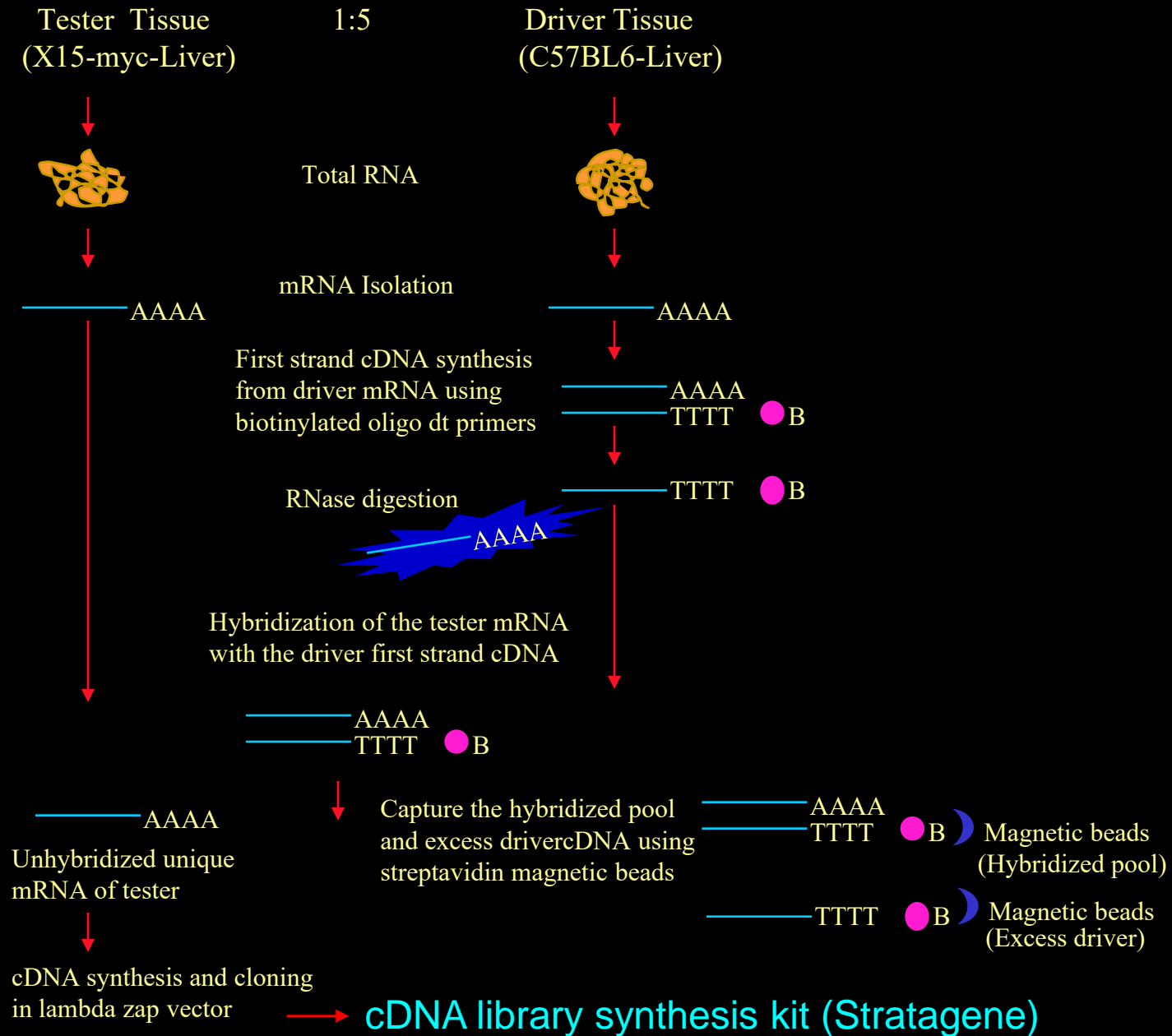


➤ Most of the pathomorphological and microscopic changes were similar to those observed with the HCC patients (Lakhtakia *et al*, 2003)

Analysis of differential gene expression  
in the tumors of X15-myc oncomouse.



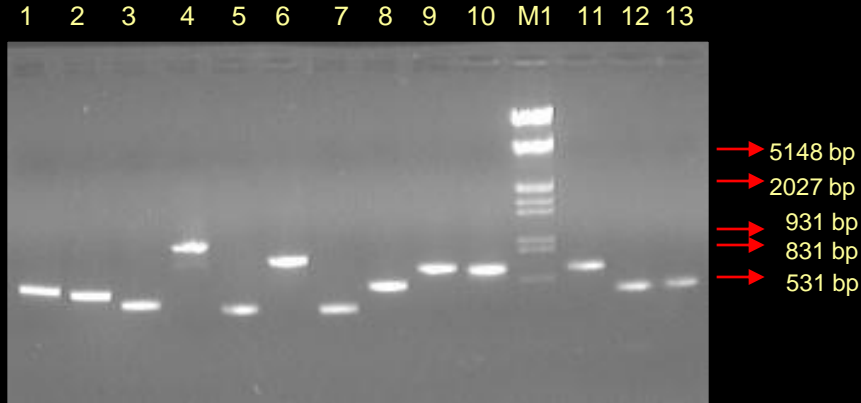
PCR analysis for X15-myc Transgenic positive animals  
 “ C “- Positive control; “ N” – Negative control; “ M “-  $\lambda$  DNA Marker



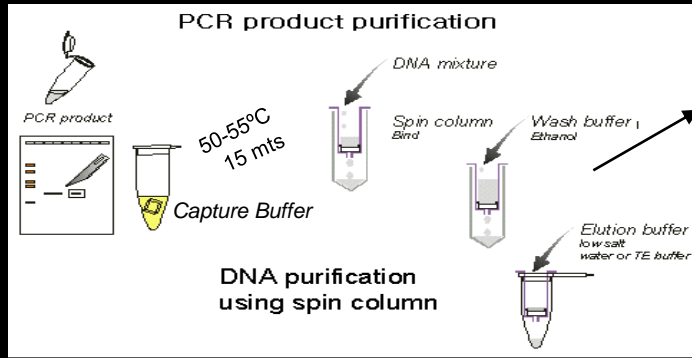
## Primers

M13 F: 5'-CGTTGTAAACGACGGCCAGTG -3'  
M13 R: 5'-CACAGGAAACAGCTATGACCATG-3'

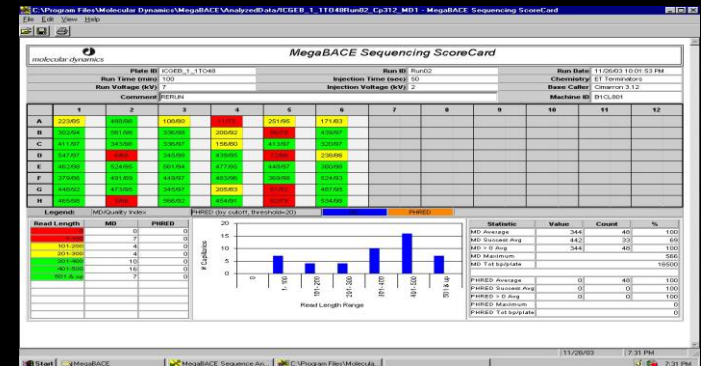
A.



B.



C.



A. Agarose gel analysis of PCR amplified cDNA inserts from phagemids. M1, DNA Marker,  
B. DNA band purification steps by GFX PCR, C. Chromatogram of MegaBACE  
sequencing score card.

Vector Screened & Changed in to FASTA format sequences



BLAST Analysis

<http://www.ncbi.nlm.nih.gov/blast/blast.cgi>



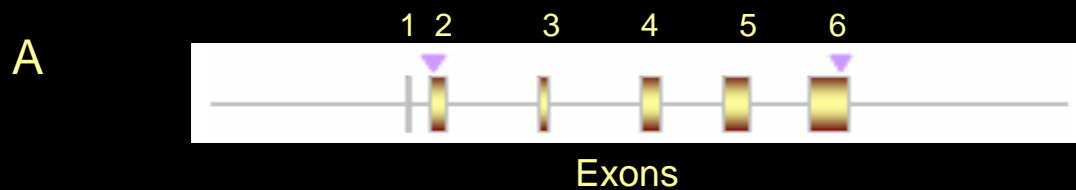
## List of differentially expressed genes.

Sequence analysis of transcripts revealed 19 discreet categories

Protein biosynthesis genes	100
Electron transport genes	81
Metabolism related genes	75
Transport molecule genes	36
Ubiquitin proteasome pathway genes	25
Protein metabolism genes	23
Signal transduction genes	22
RNA processing genes	18
Calcium/sugar/carbohydrate/copper binding genes	14
Cytoskeletal prganisation/ cell adhesion genes	17
Endocytosis /protein transport genes	13
Oxidative stress genes	13
Transcription genes	13
Cell cycle and growth differentiation genes	12
Immune response genes	11
Apoptosis and anti apoptosis genes	9
Blood cogulation genes	9
Molecular chaperone genes	8
Replication gene	1

List of top five differentially expressed genes obtained from X15-myc Transgenic mouse liver cDNA subtraction library.

<u>SYMBOL</u>	<u>GENE NAME</u>	<u>FREQUENCY</u>
Rps27a	Ribosomal Protein s27a	20
COX3	Cytochrome C Oxidase, Subunit III	16
ATP6	ATP Synthase F0 Subunit 6	14
COX2	Cytochrome C Oxidase, Subunit II	11
ND1	NADH dehydragenase, Subunit, type 1	11



**B**

Chromosome: 11, Location 11 A3.3  
 Filename : *Mus musculus* Rps27a-CDS  
 Sequence Size : 471  
 Translation Position : 1 - 471; Genetic Code : Universal

```

      10      20      30      40      50      60
ATGCAGATCTTTGTGAAGACCCCTTACGGGGAAAAACCATCACGCTCGAGGTTGAACCCCTCG
M  Q  I  F  V  K  T  L  T  G  K  T  I  T  L  E  V  E  P  S  20
      70      80      90     100     110     120
GACACTATAGAAAAATGTAAAGGCCAAGATCCAGGATAAGGAAGGAATTCCTCCTGATCAG
D  T  I  E  N  V  K  A  K  I  Q  D  K  E  G  I  P  P  D  Q  40
      130     140     150     160     170     180
CAGAGGCTGATCTTTGCTGGTAAGCAGCTGGAAGATGGCCGGACTTTGTCTGACTACAAC
Q  R  L  I  F  A  G  K  Q  L  E  D  G  R  T  L  S  D  Y  N  60
      190     200     210     220     230     240
ATTCAAAAGGAGTCCACCCCTTCATCTGGTGTGAGACTTCGGGGTGGTGCTAAGAAAAGG
I  Q  K  E  S  T  L  H  L  V  L  R  L  R  G  G  A  K  K  R  80
      250     260     270     280     290     300
AAGAAGAAGTCTTACACCACTCCCAAGAAGAACAAGCATAAGAGGAAGAAGGTTAAGTTG
K  K  K  S  Y  T  T  P  K  K  N  K  H  K  R  K  K  V  K  L  100
      310     320     330     340     350     360
GCTGTGCTGAAATACTATAAGGTGGATGAAAATGGCAAAATTAGCCGACTTCGTTCGAGAG
A  V  L  K  Y  Y  K  V  D  E  N  G  K  I  S  R  L  R  R  E  120
      370     380     390     400     410     420
TGTCCTTCTGATGAATGTGGTGTGAGATTTTCATGGGAAGCCACTTTGACAGGCATTAC
C  P  S  D  E  C  G  A  G  V  F  M  G  S  H  F  D  R  H  Y  140
      430     440     450     460     471
TGTGGCAAGTGTGTCTGACTTACTGCTTCAACAAACCAGAAGACAAGTAG
C  G  K  C  C  L  T  Y  C  F  N  K  P  E  D  K  *      156-AA
  
```

Structure of the Rps27a gene of *Mus musculus* and cDNAs. A, Organization and chromosomal location. In total, 6 exons are shown in boxes. Closed boxes designate the coding sequences. B, ORF of the Rps27a gene showing 156 amino acids in single letter code (An red arrowmark indicates the site of proteolytic cleavage that would be required to generate mature ubiquitin from the primary translation product).

Identities = 155/156 (99.4%), Positives = 155/156 (99%)  
 Gap frequency = 0/156 (0%) , PI= 9.68

mRps27a 1	MQIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQ	40
hRps27a 1	MQIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQ	40
	UBQ	
mRps27a 41	QRLIFAGKQLEDGRTLSDYNIQESTLHLVLRLRGGAKKR	80
hRps27a 41	QRLIFAGKQLEDGRTLSDYNIQESTLHLVLRLRGGAKKR	80
	UBQ	
mRps27a 81	KKKSYTTPKKNKHKRKKVKLAVLKYYKVDENGKISRLRRE	20
hRps27a 81	KKKSYTTPKKNKHKRKKVKLAVLKYYKVDENGKISRLRRE	20
	Ribosomal_s27	
mRps27a 121	CPSDECGAGVFMASHFDRHYCGKCCLTYCFNKPEDK	156
hRps27a 121	CPSDECGAGVFMASHFDRHYCGKCCLTYCFNKPEDK	156
	Ribosomal_s27	

Alignment of the primary sequences of human and mouse Rps27a. Identical residues are shown in same color, while the non-identical regions are marked in different colors. Blue and rose boxes denote the regions of ubiquitin and Ribosomal\_s27 respectively.



- The Rps27a Highly conserved very basic protein due to presence of 80 basic AA in the Carboxyl terminal (CEP) of Ubiquitin.

- UBQ-76 AA conserved eukaryotic protein- Diverse cellular functions

- UBQ-Poly ubiquitin chain (UBb, UBc) or Fused to unrelated protein (UBA)

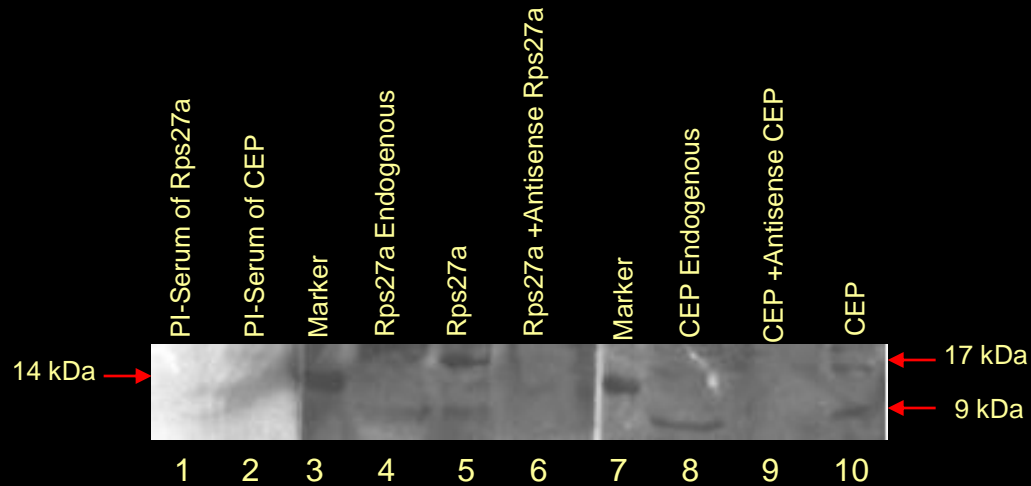
- UBA —————→ UBA80 (Rps27a)or UBA52 (RpL40). Conserved in man, yeast and plants

[Salvensan et al, Nucleic Acid Research (1987), Ozkaynak et al, EMBO J (1987), Lund et al, J. Bio. Chem (1985)]

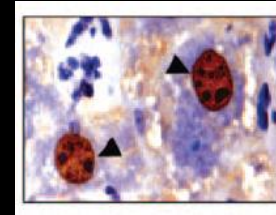
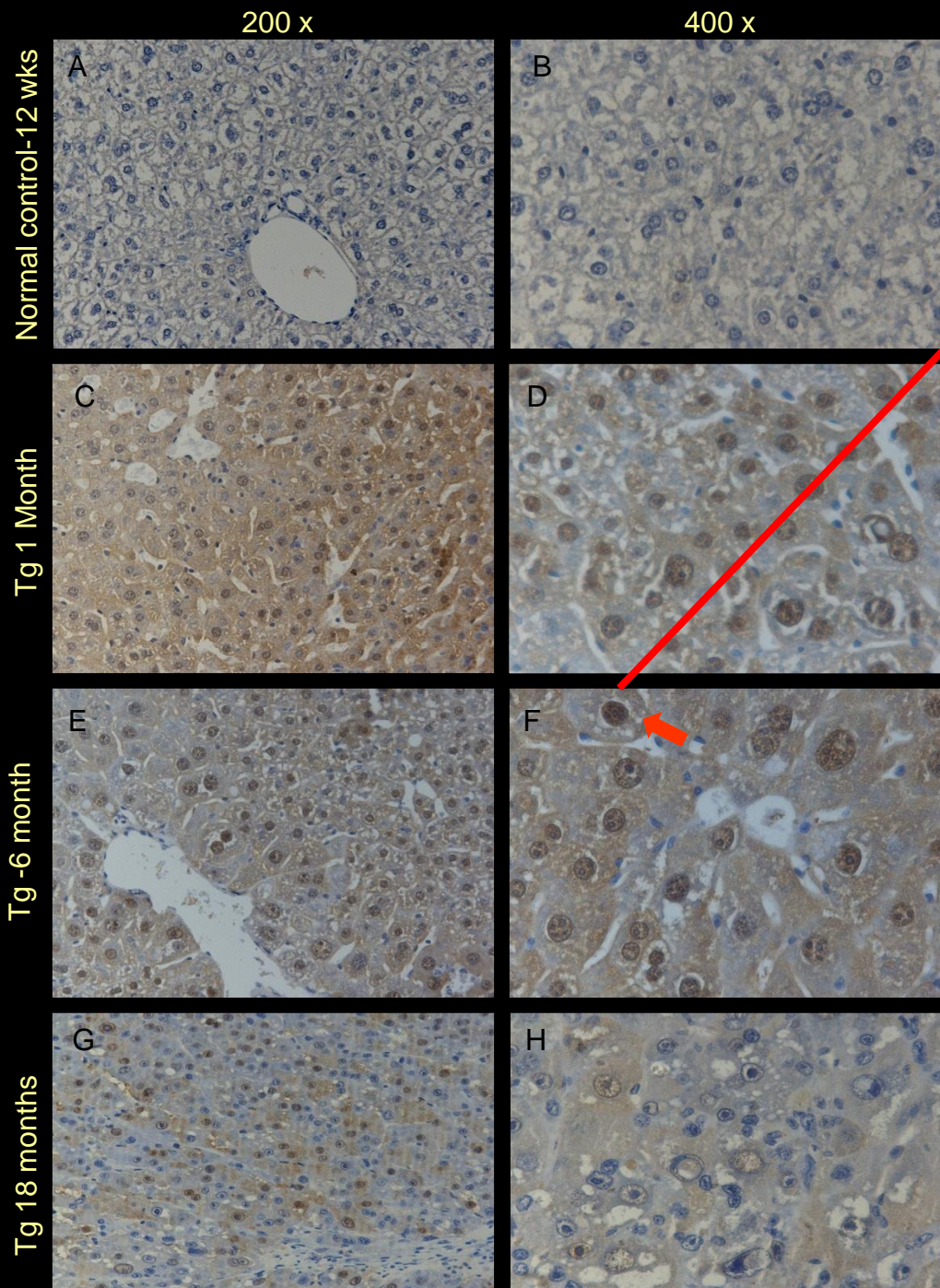
- CEP analysis shows strong homology at the AA level with Zinc finger protein.

- CEP can associate with ribosome binding specifically ribosome or messenger RNA.

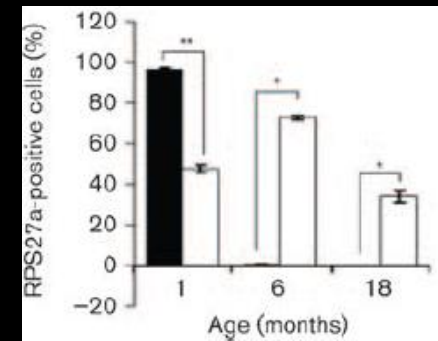
- Function of UBQ-CEP fusion genes has not been fully elucidated



Detection of Rps27a and CEP expression in Huh7 cell lines. Total cells were lysed after 48hr post transfection with Rps27a (lanes 5), anti-Rps27a (lanes 6), CEP (lanes 10) and anti-CEP (lanes 9).



Positive staining is noted more in nuclear and perinuclear (red arrow)

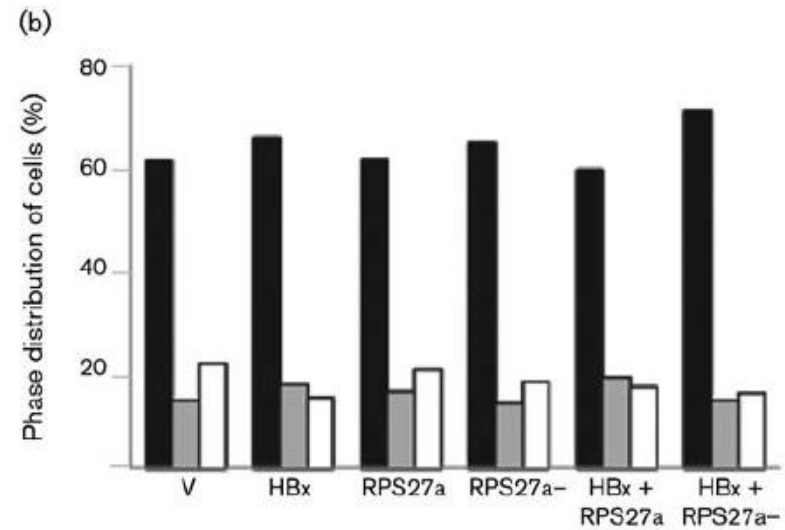
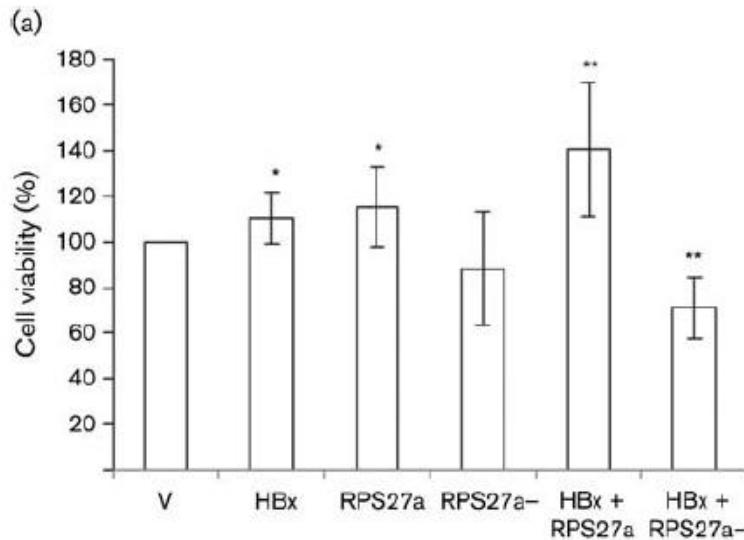


transgenic mice (white bars). Cells ( $n=500$ ) were counted randomly from 10 different areas. \* $P<0.01$ ; \*\* $P<0.05$ .

Rps27a expression in X15-myc Tg mice liver tissues.



# Co-expression of Rps27a and HBx leads to improved survival and proliferation of cells



G0/G1	61.0	65.6	61.3	64.4	59.4	71.0
S	13.4	16.7	15.2	12.8	18.1	13.3
G2/M	20.6	14.0	19.7	17.2	16.2	15.0

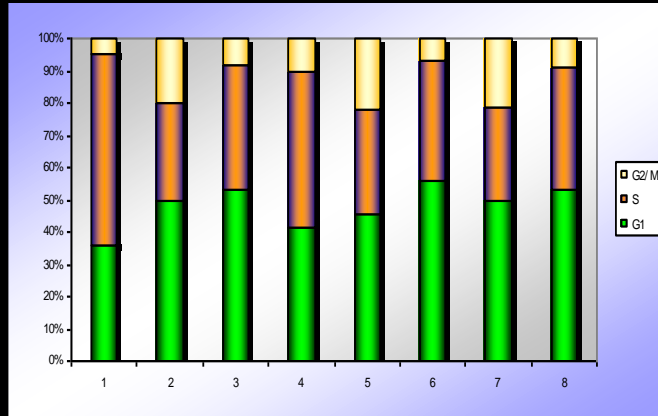
\* $P < 0.01$  \*\* $P < 0.001$ , compared with controls.

**Fig. 3.** Cooperation between RPS27a and HBx for cell survival, proliferation and size. Huh7 cells were transfected with expression vectors for HBx, RPS27a or RPS27a-, either alone or in combinations as indicated, and their viability, size and cell-phase distribution were monitored. (a) Cell viability by MTT assay. The bar graph shows means  $\pm$  SD of ten independent observations. (b) Flow cytometry of asynchronous cells to show percentage distribution of cells in different phases of the cell cycle (black bars, G0/G1; grey bars, S; white bars, G2/M). Results are means of three independent observations. (c, d) The



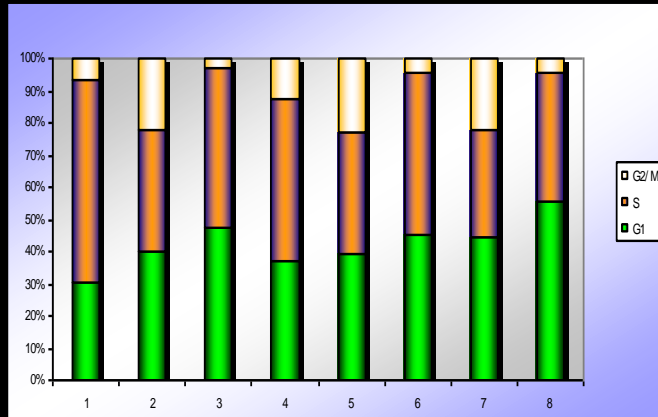
12hrs

A



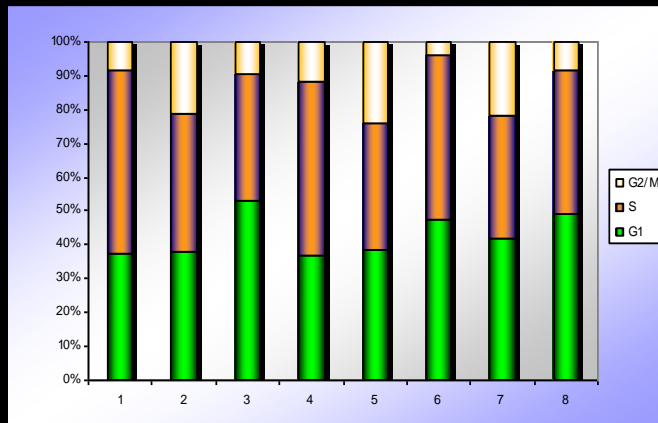
24hrs

B



42 hrs

C

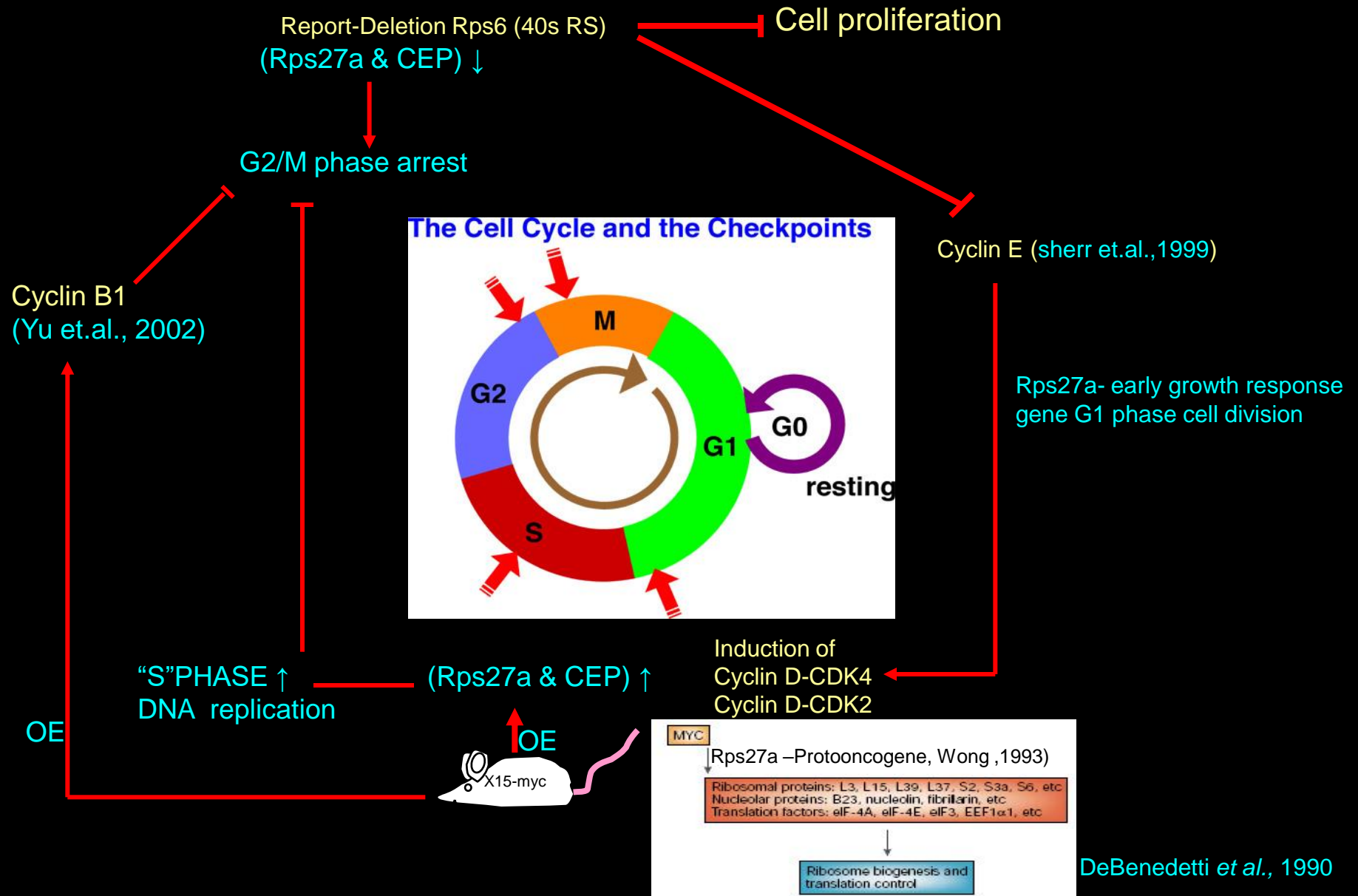


## Constructs

1. Rps27a
2. Antisense Rps27a
3. Rps27a + Antisense Rps27a
4. CEP
5. Antisense CEP
6. CEP + Antisense CEP
7. Rps27a + Antisense Rps27a + Antisense CEP
8. pSG5

## Effect of Rps27a and CEP on cell cycle.

Huh7 cells were transfected with Rps27a, CEP, anti-Rps27a and anti-CEP constructs at 60-70 % confluency and cells were serum synchronized and collected at various time points of 12hrs (A) , 24hrs (B) and 42 hrs (C). The percentages of cells at different phases of the cell cycle were analysed by FACS and values are shown on the left (Y-axis) bar diagram.



In yeast, UB13 deletion-Slow growth of phenotype (1.6 hr-6.8hr) (Finley et.al., 1989)  
 Many small & Large Rib.Sub unit Protein ↑ variety of p.Tumor- Ex RPL36A. OE-Over Expression

## Summary

- Rps27a, a ubiquitin precursor protein fused to the 80 amino acid carboxyl extension protein (CEP)
- The flow cytometry analysis revealed that Rps27a overexpressing cells accelerated cells to enter G1 phase to S phase with enhanced cell proliferation
- Immunohistochemical staining of Rps27a expression was moderately reduced from 3 months to 18 months old Transgenic liver tissues.

Overall, The sensitivity of over expressed Rps27a in HCC might be a general biomarker for tumor proliferation.

# Evaluation of the antitumorogenic/anti-angiogenic activities of the floral isolates of *B. monosperma*



*Butea monosperma* (Flame of the Forest)

Family	:	Fabaceae
Orgin	:	India
Type/Uses	:	flowering tree
Size	:	50feet
Growth Rate	:	Slow growing at first
Lighter Requirments	:	full sun
Water Requirments	:	average, drier in the winter
Min.Temp.	:	mid 30° s
Flower	:	late winter, spring

Test Material	A003	F008	F009	Adriamycin	Mitomycin	Tamoxifen	5-Fluorouracil
<b>Concentration used:</b>	100 mg/L	100 mg/L	100 mg/L	1x10 <sup>-5</sup> M	1x10 <sup>-5</sup> M	1x10 <sup>-5</sup> M	2x10 <sup>-5</sup> M
<b><i>Cell lines</i></b>							
Breast – MCF-7	6	-	-	72	-	-	-
Breast – T47D	4	22	0	34	-	-	-
Breast – ZR75-1	0	-	-	46	-	-	-
Cervix – SiHa	0	9	0	-	-	-	19
CNS - SK N MC	23	57	2	-	-	27	-
CNS - SK N SH	43	-	-	82	-	-	-
CNS - SNB78	2	-	-	20	-	-	-
Colon - Colo205	87	19	0	-	-	-	40
Liver - Hep2	51	58	35	-	88	74	-
Lung - A549	19	27	11	58	-	17	-
Lung - NCI H23	0	-	-	-	59	-	-
Oral - KB	16	-	-	-	-	-	9
Ovary – NIH OVCAR3	0	-	-	-	31	-	-
Ovary – OVCAR5	5	37	6	-	-	-	-
Prostate – DU145	0	-	-	-	69	-	-

Percent growth inhibition of human cancer cell lines with the aqueous extract (A003) of flowers of *Butea monosperma* and its fractions (F008 and F009) [Mathan *et.al.*,] [Communicated]

# Hepatoprotective Herbal Extracts

- Traditional Herbal Medicine & CAM becoming popular among cancer patients (Molassiotis *et al.*, 2005; Yates *et al.*, 2005)
- Commonly used herbal preparations for treating Liver Diseases-Silymarin (Mayer *et al.*, 2005), TJ-9 (Oak *et al.*, 1995), Liv-52 (Huseini *et al.*, 2005), Phyllanthus (Liu *et al.*, 2001), Glycyrrhizin (Kumada *et al.*, 2002), curcumin (Aggarwal *et al.*, 2003; Campbell *et al.*, 2005), *Calotropis procera* (Choedon & Mathan, 2006).
- RRL (Regional Research Laboratory, CSIR), Jammu

➤ The traditional Indian medical system of Ayurveda, the flowers of *B. monosperma* have been used in the treatment of hepatic disorders and viral hepatitis (Schuppan *et al.*, 1999; Dhiman *et al.*, 2005)

➤ Chemopreventive potential of methanolic extract of *B. monosperma* flower on chemically induced oxidative damage in rats were reported (Sehrawat *et al.*, 2006)

➤ The mechanism of its action has not been elucidated.

Flowchart of *Butea monosperma*  
floral extracts and its fractions.



*Butea monosperma*



Flowers ( Dried)

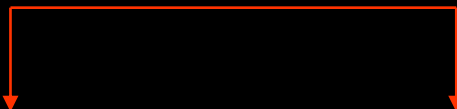


Extraction

A003 Aqueous (Active *in vivo*)



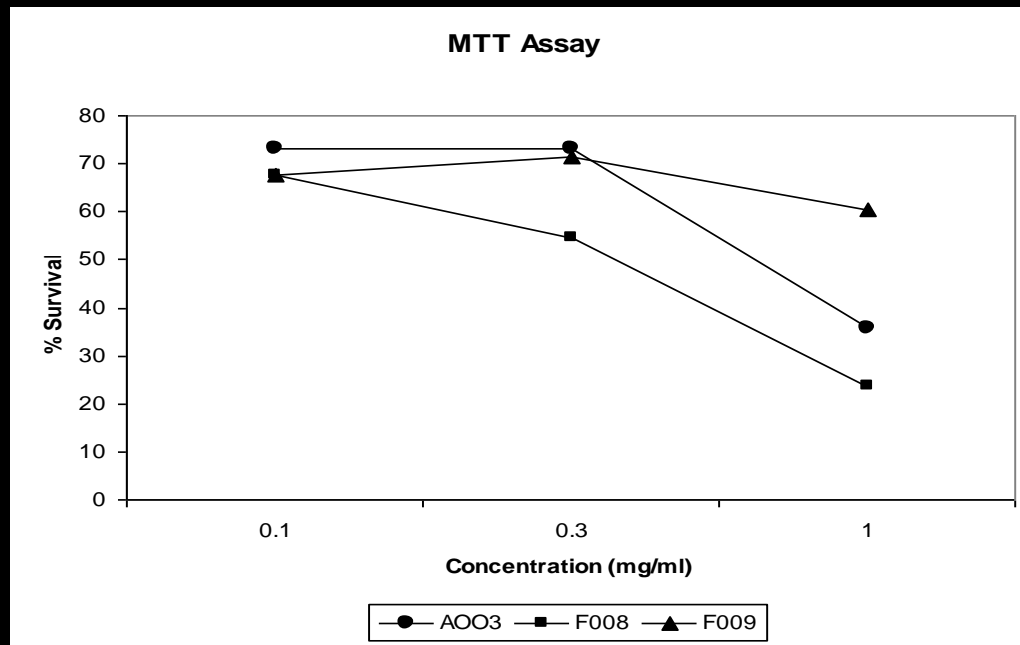
Fractination



F008 n-Butanol  
(Active *in vitro*)

F009 Aqueous  
Active *in vitro* &  
*in vivo*)

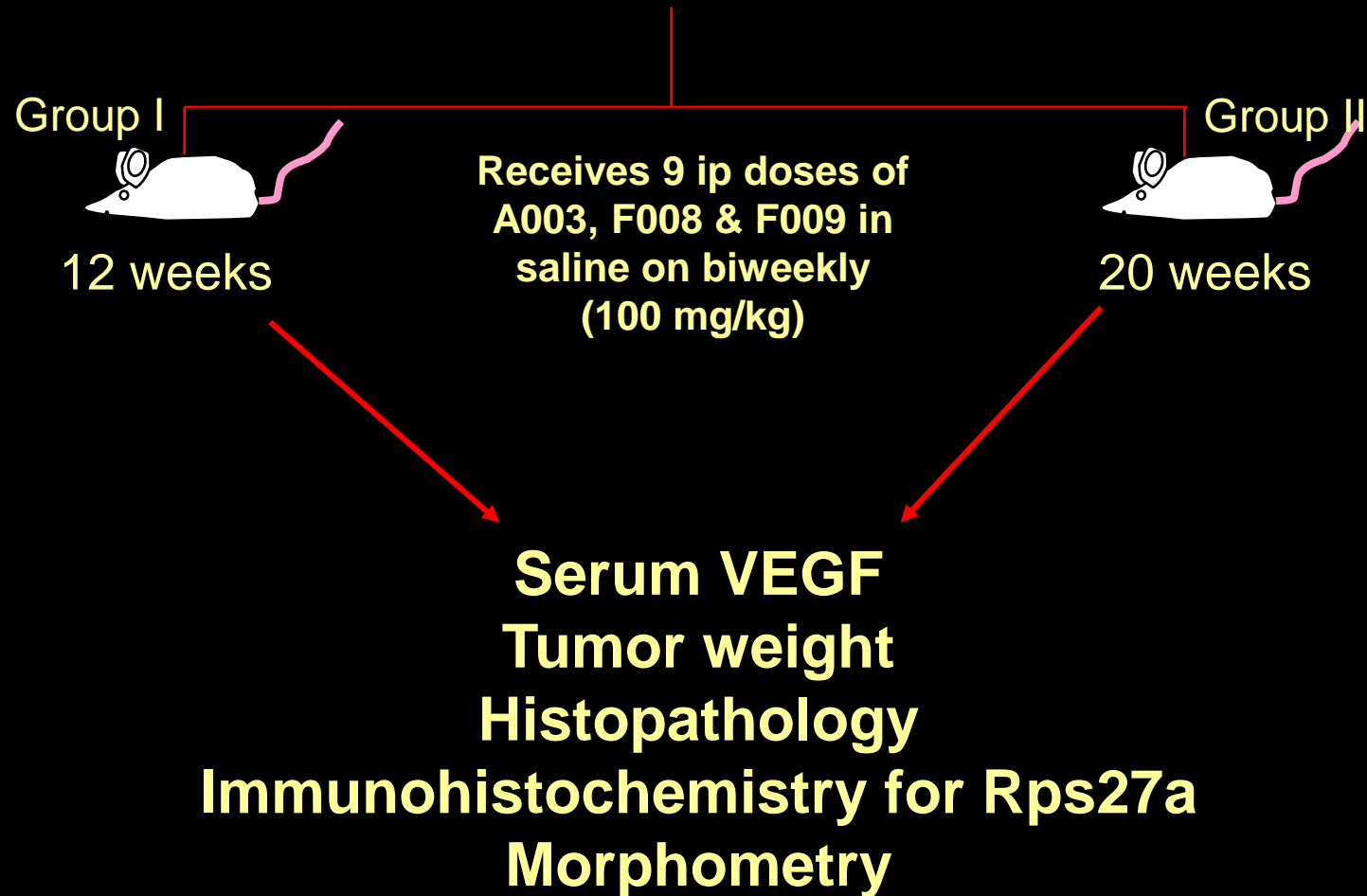




Cytotoxic effect of *Butea monosperma* extract on cancer cell lines. Huh-7 cells were incubated with different concentrations of *Butea monosperma* and analyzed for cell viability at 24h or 48h ( $n=6$ ), mean $\pm$ SE. a.  $P<0.001$ ; b.  $P<0.01$

- IC<sub>50</sub> in Huh7 cells – F009- less toxic (65% cell survival at 1g/L)
- A003 & F008 (0.5 – 1g/L) & Doxorubicin (0.1g/L)

# X 15-myc Transgenic mouse model (4 weeks old matched age group, n=6)



**Anti- tumorigenic effect of *B. monosperma* in transgenic mice**

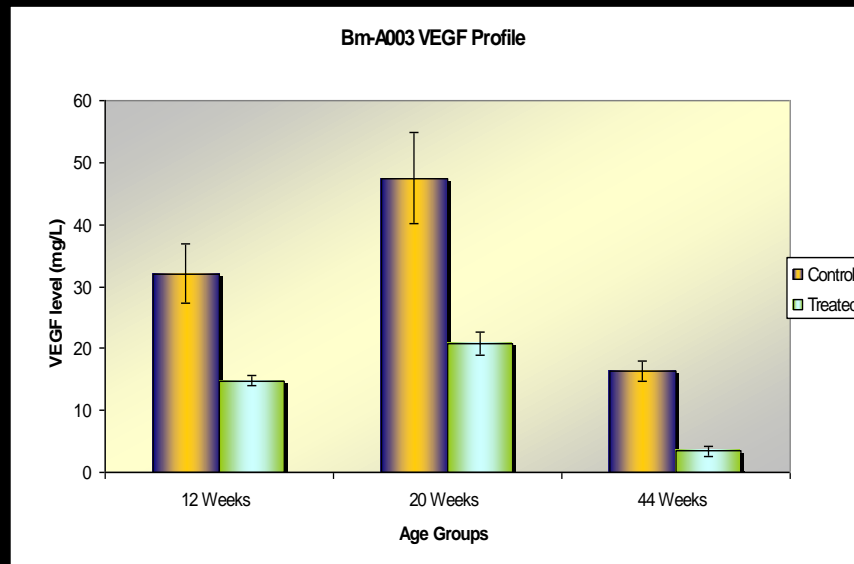
**A.**

		Group	
Period (wks)		Treated( n=6)	Control (n=6)
A003	12	14.8±0.8 <sup>b</sup>	32.1±4.8
	20	20.8±1.9 <sup>a</sup>	47.5±7.3
	44	3.45±0.8 <sup>a</sup>	16.3±1.6
F009	12	13.5±0.6 <sup>NS</sup>	14.4±0.5
	20	16.2±0.6	17.1±0.8 <sup>NS</sup>
	Non transgenic control	7.00 ± 0.55	

**Tg-2 to 4 fold↑**

Values are shown as mean± SE. <sup>a</sup> p<0.001; <sup>b</sup> p<0.01; <sup>NS</sup> p = 0.15

**B.**



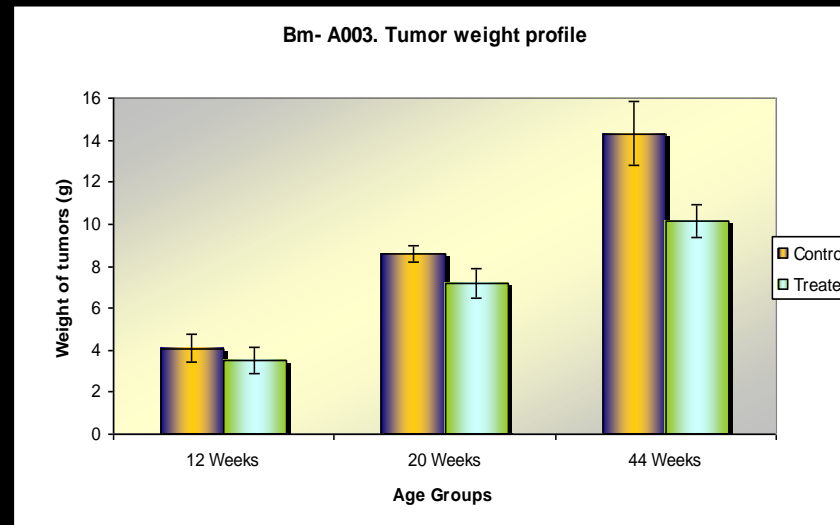
**A.** Serum VEGF levels (mg/L) in the X-15 *myc* mice after treatment with the aqueous extract (A003) or aqueous fraction (F009) of *Butea monosperma* (BM). **B.** BM-A003 treated, 20 and 44 wks old, Transgenic animals are showing the significant reduction (p<0.001) in their VEGF level.

**A.**

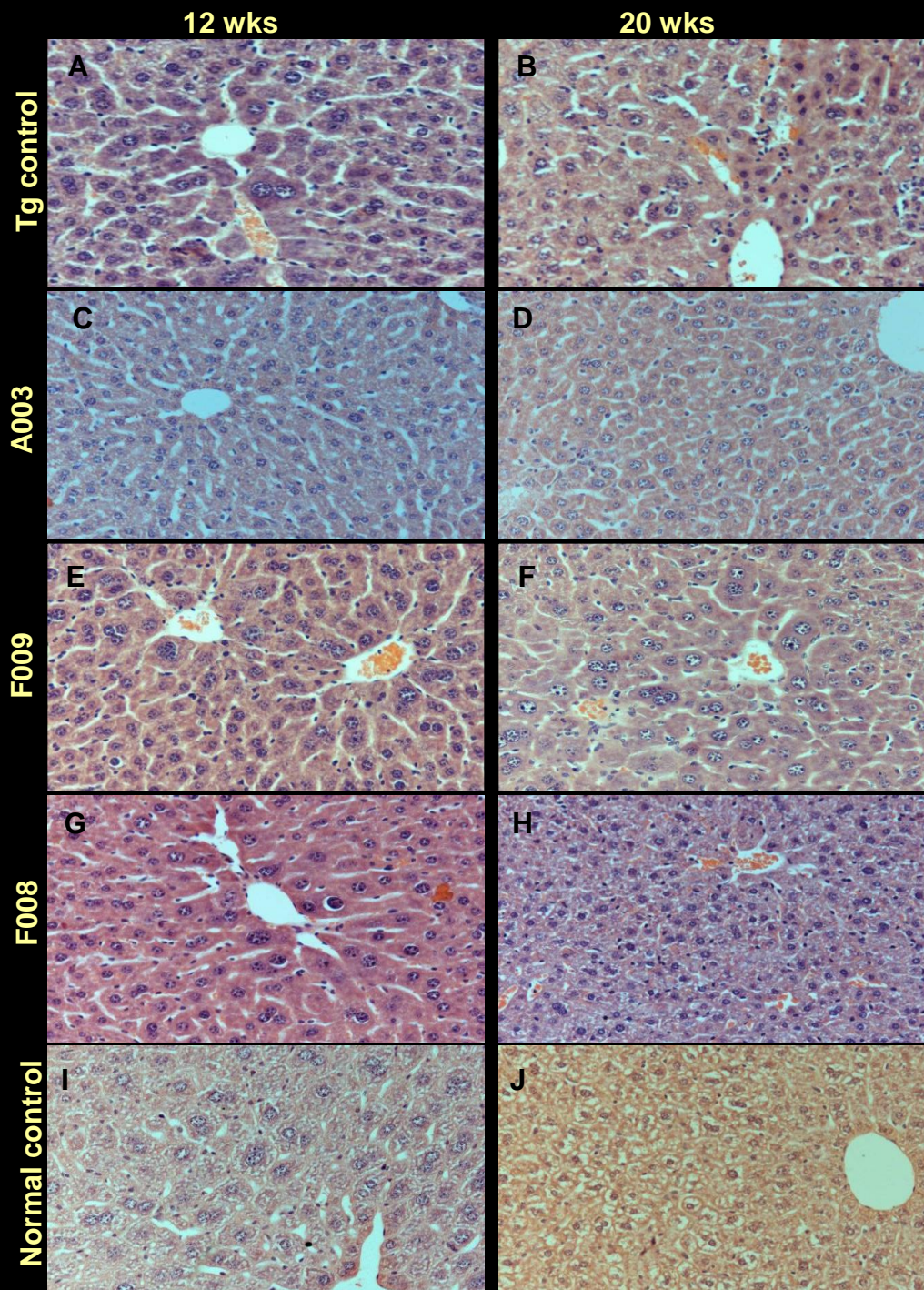
		Group	
Period (wks)		Treated( n=6)	Control (n=6)
A003	12	3.50±0.63	4.08±0.66
	20	7.17±0.72 <sup>b</sup>	8.57±0.39
	44	10.15±0.81 <sup>a</sup>	14.32±1.53

Values are shown as mean± SE. <sup>a</sup>p<0.001; <sup>b</sup>p<0.01

**B.**



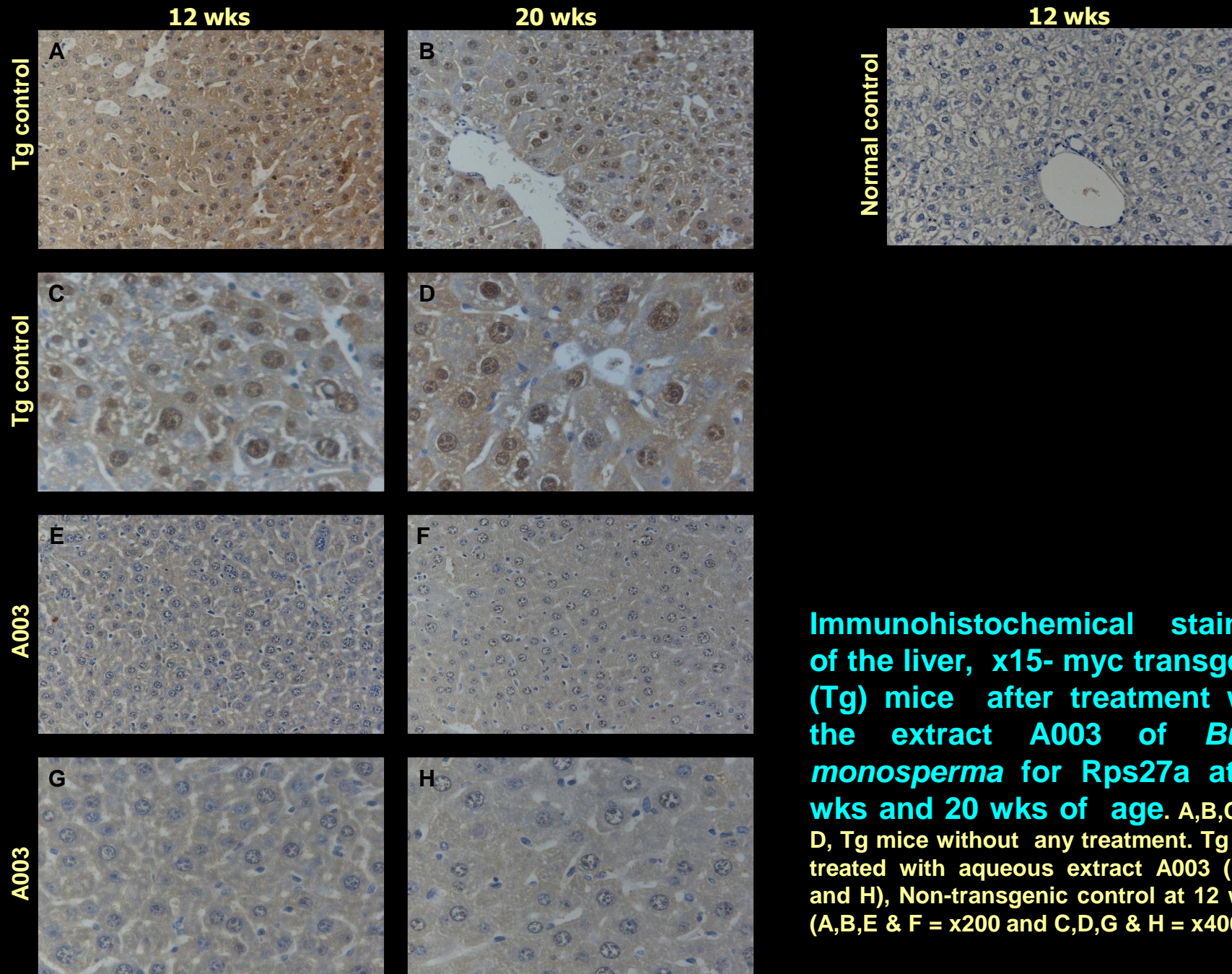
**A.** Liver tumor weight (g) of X-15 *myc* mice after treatment with the aqueous extract (A003) of *Butea monosperma* (BM). **B.** BM-A003 treated, 44 wks old, Transgenic animals are showing the significant reduction ( $p<0.001$ ) in their tumor weight.



**Histological analysis of the liver of X15-myc transgenic (Tg) mice at 12 wks and 20 wks of age after treatment with the extract and fractions of *Butea monosperma*.**

**(A&B)**, Tg mice without any treatment. Tg mice treated with aqueous extract A003 **(C&D)**, aqueous fraction F009 **(E&F)** and butanolic fraction F008 **(G&H)**; **(I&J)**, Non-transgenic control at 12 wks and 20 wks. **(All x200)**





**Immunohistochemical staining of the liver, x15- myc transgenic (Tg) mice after treatment with the extract A003 of *Butea monosperma* for Rps27a at 12 wks and 20 wks of age. A,B,C and D, Tg mice without any treatment. Tg mice treated with aqueous extract A003 (E,F,G and H), Non-transgenic control at 12 wk (I) (A,B,E & F = x200 and C,D,G & H = x400)**

## 12- wks

Parameters	Treatment Groups				
	Normal Control	Tg Control	A003	F008	F009
Area	26.9±11.8 <sup>a</sup>	45.54±22.0	22.7±10.6 <sup>a</sup>	42.46±26.4 <sup>c</sup>	42.36±23.6 <sup>c</sup>
Diameter(mean)	5.53±1.20 <sup>a</sup>	7.26±1.81	5.00±1.30 <sup>a</sup>	6.84±2.10 <sup>a</sup>	6.90±1.93 <sup>a</sup>
Perimeter	18.05±3.72 <sup>a</sup>	24.18±6.89	16.40±4.10 <sup>a</sup>	22.26±6.75 <sup>a</sup>	22.50±6.50 <sup>a</sup>
Size (length)	6.20±1.45 <sup>a</sup>	8.60±2.45	5.60±1.40 <sup>a</sup>	7.60±2.34 <sup>a</sup>	7.65±2.20 <sup>a</sup>
Size (width)	5.24±1.00 <sup>a</sup>	6.50±1.64	4.80±1.25 <sup>a</sup>	6.53±1.98	6.50±1.85
Feret (mean)	5.80±1.18 <sup>a</sup>	7.62±2.00	5.22±1.30 <sup>a</sup>	7.09±2.13 <sup>a</sup>	7.14±2.00 <sup>a</sup>
Density(mean)	124.7±10.8 <sup>a</sup>	119.43±11	132.22±9.8 <sup>a</sup>	108.7±12.3 <sup>a</sup>	126.43±15 <sup>a</sup>
Density (std.dev)	13.83±2.63 <sup>a</sup>	16.03±4.43	17.00±4.20	18.40±6.62 <sup>a</sup>	20.45±7.16 <sup>a</sup>
Heterogeneity	0.064±0.05 <sup>a</sup>	0.109±0.1	0.149±0.10 <sup>a</sup>	0.154±0.12 <sup>a</sup>	0.209±0.14 <sup>a</sup>

Values are shown as mean± SE (n=500). 'a' = p<0.001; 'b' = p<0.01; 'c' = p<0.05.

**Mean values and standard deviation for the parameters of area, diameter (mean), perimeter, size (length & width), feret (mean) Density (mean), density (std.dev) and heterogeneity (in millimeter) of the liver cell nuclei of 12 wks old, X15- myc transgenic mice treated with *Butea monosperma* (A003) and its fractions (F008 and F009) with their relevant Tg control.**

## 20- wks

Parameters	Treatment Groups				
	Normal Control	Tg- Control	A003	F008	F009
Area	26.85±11.8 <sup>a</sup>	47.96±24.6	23.95±15.0 <sup>a</sup>	26.85±14.5 <sup>a</sup>	41.64±24.6 <sup>a</sup>
Diameter(mean)	5.526±1.20 <sup>a</sup>	7.370±1.90	5.071±1.60 <sup>a</sup>	5.439±1.52 <sup>a</sup>	6.760±2.10 <sup>a</sup>
Perimeter	18.04±3.72 <sup>a</sup>	24.20±6.24	16.56±5.17 <sup>a</sup>	17.84±4.94 <sup>a</sup>	22.14±6.91 <sup>a</sup>
Size (length)	6.194±1.45 <sup>a</sup>	8.433±2.32	5.669±1.75 <sup>a</sup>	6.230±1.75 <sup>a</sup>	7.567±2.40 <sup>a</sup>
Size (width)	5.237±1.00 <sup>a</sup>	6.879±1.70	4.844±1.60 <sup>a</sup>	5.088±1.46 <sup>a</sup>	6.395±2.00 <sup>a</sup>
Feret (mean)	5.763±1.18 <sup>a</sup>	7.684±1.95	5.286±1.64 <sup>a</sup>	5.691±1.56 <sup>a</sup>	7.039±2.15 <sup>a</sup>
Density(mean)	124.7±10.8 <sup>a</sup>	113.52±13	122.4±10.2 <sup>a</sup>	105.6±17.2 <sup>a</sup>	121.7±11.1 <sup>a</sup>
Density (std.dev)	13.83±2.63	14.25±4.35	17.86±5.03 <sup>a</sup>	21.83±8.54 <sup>a</sup>	16.44±4.26 <sup>a</sup>
Heterogeneity	0.064±0.05	0.073±0.07	0.160±0.12 <sup>a</sup>	0.198±0.16 <sup>a</sup>	0.126±0.10 <sup>a</sup>

Values are shown as mean± SE (n=500). 'a' = p<0.001; 'b' = p<0.01; 'c' = p<0.05.

**Mean values and standard deviation for the parameters of area, diameter (mean), perimeter, size (length & width), feret (mean) Density (mean), density (std.dev) and heterogeneity (in millimeter) of the liver cell nuclei of 20wks old, X15- myc transgenic mice treated with *Butea monosperma* (A003) and its fractions (F008 and F009) with their relevant Tg control.**



## SUMMARY

➤ *in vitro* cytotoxicity and *in vivo* transgenic animal studies suggested that aqueous extract (A003) and fractions (F008 and F009) of *Butea monosperma* flowers is not only hepatoprotective but also carries anti-proliferative, anti-tumorigenic and anti-angiogenic properties.

➤ The chemopreventive action of fractions, F008 and F009 was less prominent as compared to the aqueous extract A003 suggesting either loss or inactivation of some of the key constituents in these fractions.

➤ Immunohistochemical analysis of Rps27a in the transgenic animals treated with aqueous extract A003 revealed a marked reversal of pathological manifestation including no staining for Rps27a in the liver.

Overall, the aqueous extract of *Butea monosperma* flower has the potential for developing new cancer therapeutics

(19) World Intellectual Property Organization  
International Bureau

PCT

(43) International Publication Date  
30 November 2006 (30.11.2006)(11) International Publication Number  
WO 2006/126067 A1(51) International Patent Classification:  
A61K 36/48 (2006.01) A61P 35/00 (2006.01)For Review: Engineering and Biotechnology, Arun, Asat  
A1 Mang, New Delhi 110 067 (IN).(21) International Application Number:  
PCT/IN2006/001355(34) Agents: BHATTACHARYYA, Goutam et al; K & S  
Parnas, 34-C, D-4 Tana, DF Central Avenue, Sakinaka  
Gurgaon, New Delhi 110 062 (IN).

(22) International Filing Date: 24 May 2006 (24.05.2006)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Date:  
15 May 2005 (15.05.2005) 26 May 2005 (26.05.2005) IN(71) Applicants (for all designated States except (59))  
COUNCIL OF SCIENTIFIC AND INDUSTRIAL  
RESEARCH (ININ); Raji Bhag, New Delhi 110 001  
(IN) INTERNATIONAL CENTRE FOR GENETIC  
ENGINEERING AND BIOTECHNOLOGY (ININE)  
Aruna Asat Ali Mang, New Delhi 110 067 (IN)

(72) Inventors: and

(73) Inventors/Applicants (for US only): SAXENA, Ajit  
Kumar (ININ); Regional Research Laboratory, Canal  
Road, Jammu 180 001 (IN) GUPTA, Bishan, Datt  
(ININ); Regional Research Laboratory, Canal Road,  
Jammu 180 001 (IN) KAPAH, Raj, Kishan (ININ)  
Regional Research Laboratory, Canal Road, Jammu 180  
001 (IN) MUTHIAL, Shanmugavel (ININ); Regional  
Research Laboratory, Canal Road, Jammu 180 001 (IN)  
MONDHE, Dilip, Manikrao (ININ); Regional Research  
Laboratory, Canal Road, Jammu 180 001 (IN) RAJNA,  
MERNA, BALESHWAR (ININ); Regional Research  
Laboratory, Canal Road, Jammu 180 001 (IN) QAZI,  
Ghulam, Nahi (ININ); Regional Research Laboratory,  
Canal Road, Jammu 180 001 (IN) KUMAR, Vijay  
(ININ); Laboratory and Central Biotechnology Engineering and  
Biotechnology, Arun Asat Ali Mang, New Delhi 110 067  
(IN) MATHAN, Ganesham (ININ) International Centre(81) Designated States (unless otherwise indicated, for every  
class of national protection available): AR, AG, AL, AM,  
AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN,  
CO, CR, CU, CZ, DE, DK, DM, DO, EC, EE, EG, ES, FI,  
FR, GB, GR, GH, GM, GU, HP, HT, IL, IN, IS, IT, JP, KR,  
KG, KM, KN, KP, KR, KZ, LC, LI, LR, LS, LT, LU, LV,  
LY, MA, MD, MG, MK, MN, MW, MX, MY, NA, NG, NI,  
NO, NZ, OM, PG, PI, PL, PT, RO, RU, SC, SD, SE, SG,  
SK, SL, SM, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US,  
UZ, VC, VN, YU, ZA, ZW, ZY

Declaration under Rule 4.17:

— as is the applicant's entitlement to claim the priority of the  
earlier application (Article 4.17(a))

Published:

— with international search report  
before the expiration of the time limit for amending the  
claims and to be republished in the event of receipt of  
amendmentsFor two-letter codes and other abbreviations, refer to the "Glossary  
of Symbols and Abbreviations" appearing in the beginning  
of each regular issue of the PCT Gazette.

(54) TITLE: A PHARMACEUTICAL COMPOSITION FOR THE TREATMENT OF HEPATOCELLULAR CARCINOMA

(57) Abstract: The present invention relates to anticancer activity against hepatocellular carcinoma (HCC) and fraction isolated from leaves of *Butea monosperma*. Part of the present invention relates to anticancer activity against hepatocellular carcinoma of a composition containing purified flavonoid glycosides such as rutin and isobutyrin in the range of 2 to 9% by weight, isolated from the leaves of *Butea monosperma* by extracting the leaves with polar solvent like ethanol, methanol, acetone, ethane, or water, removing the non-polar components by filtration of the extract with solvents such as ethyl acetate, methyl ethyl ketone, chloroform or ethyl acetate, removing the residue in water, extracting with non-polar and drying the aqueous part.

## PATENTS

1. Saxena, A. K., Gupta, B. D., Kapahi, B. K., Shanmugavel, M., Mondhe, D. M., Qazi, G. N., **Mathan, G.**, and Kumar, V. Anticancer activity of an extract from *Butea monosperma* and the fraction thereof. Patent application #20060280817; Class: 424757000 (USPTO), 2006.

## ACKNOWLEDGEMENT

**SCHOOL OF LIFE SCIENCES,  
BHARATHIDASAN UNIVERSITY**

**ICGEB, NEW DELHI**

**CSIR, NEW DELHI**



**PROF. CHELLAM BALASUNDARAM**  
**Prof. VIJAY KUMAR Ph.D, FNAS**

THANK YOU