

# **NUCLEAR ORGANIZATION AND REGULATION OF TRANSCRIPTION IN LYMPHOMAS AND NORMAL B- LYMPHOCYTES**

Yegor VASSETZKY, CNRS UMR 8126, Institut de Cancérologie Gustave Roussy



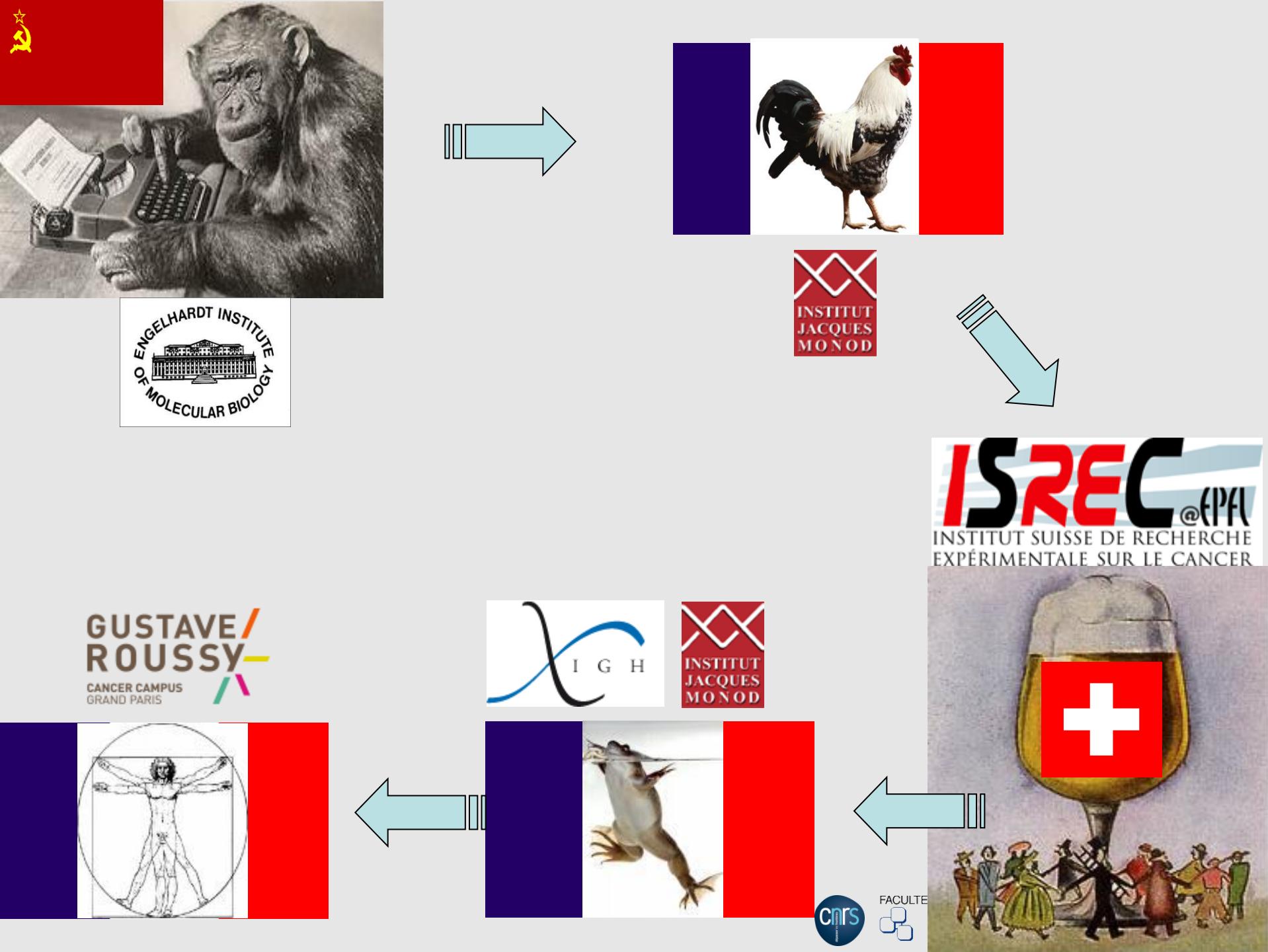
FACULTE DE MEDECINE  
PARIS-SUD



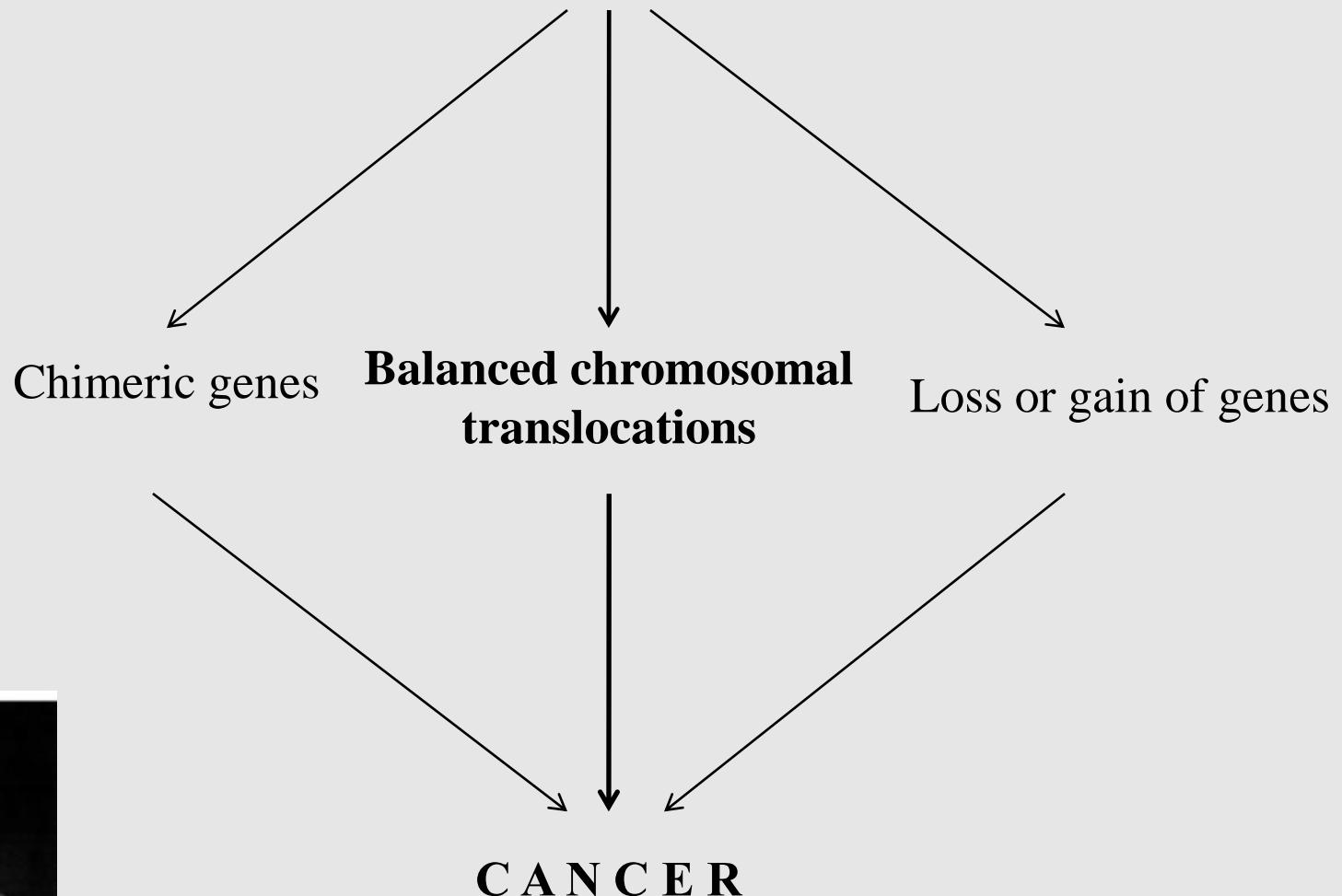


**Вадим Моисеевич Кавсан**

**1939 -- 2014**



# CHROMOSOMAL TRANSLOCATIONS



Theodor Boveri (1862-1915)



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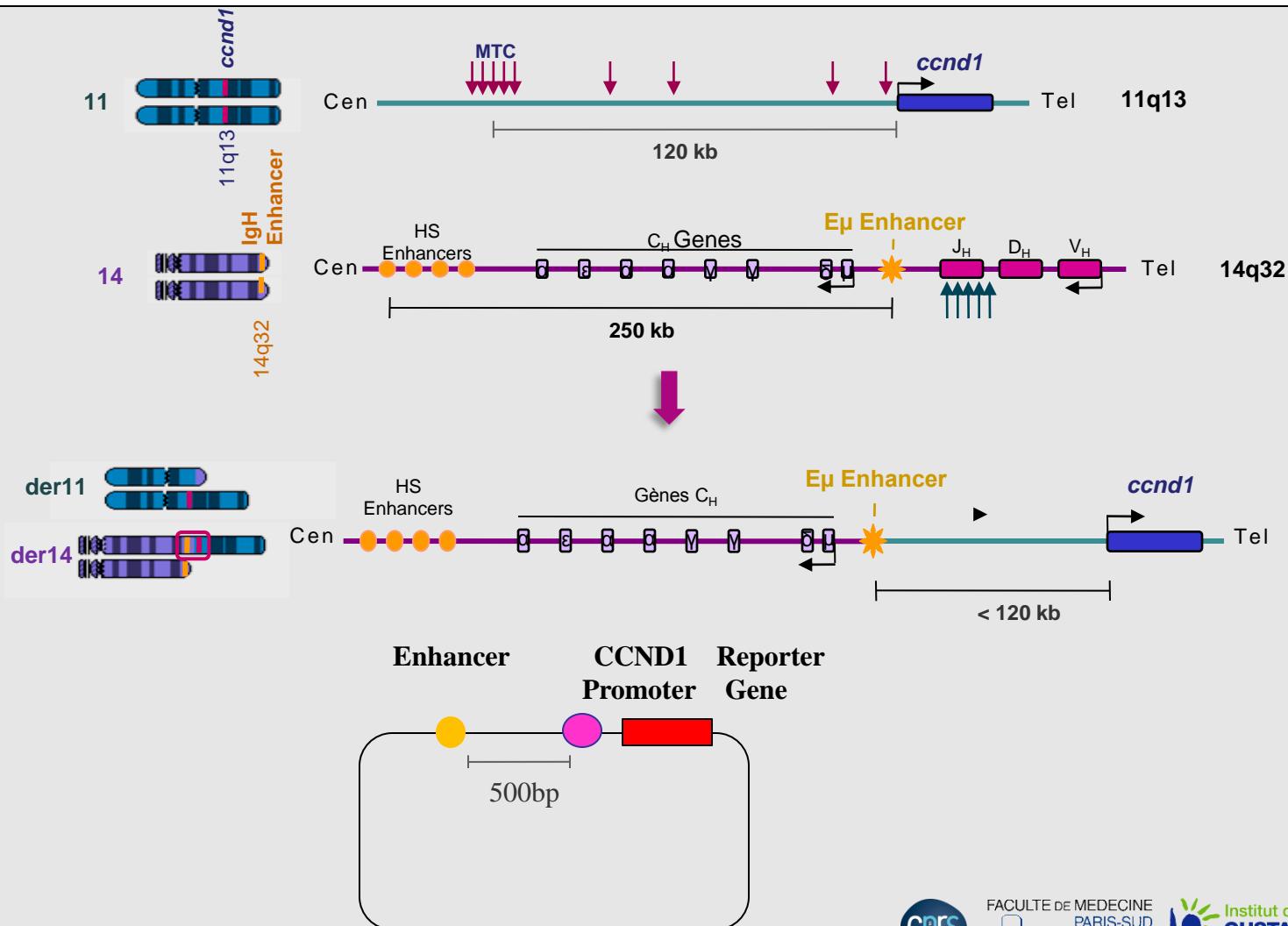


# MANTLE CELL LYMPHOMA (MCL)

- Non-Hodgkin lymphoma.
- Highest frequency between 60 and 65 years (4% before 40 years).
- Low occurrence (1 case in 200 000).
- No curative treatment; median survival rate is 3 to 4 years.

# MANTLE CELL LYMPHOMA (MCL)

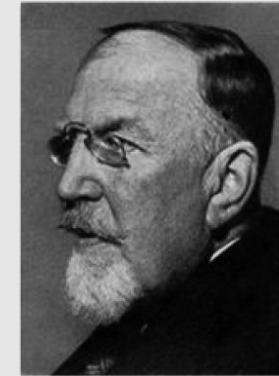
- t(11 ; 14)(q13 ; q32) translocation is found in 100% of MCLs by FISH (Fluorescent In Situ Hybridization).
- t(11 ; 14)(q13 ; q32) translocation leads to overexpression of cyclin D1 (CCND1) in B lymphocytes



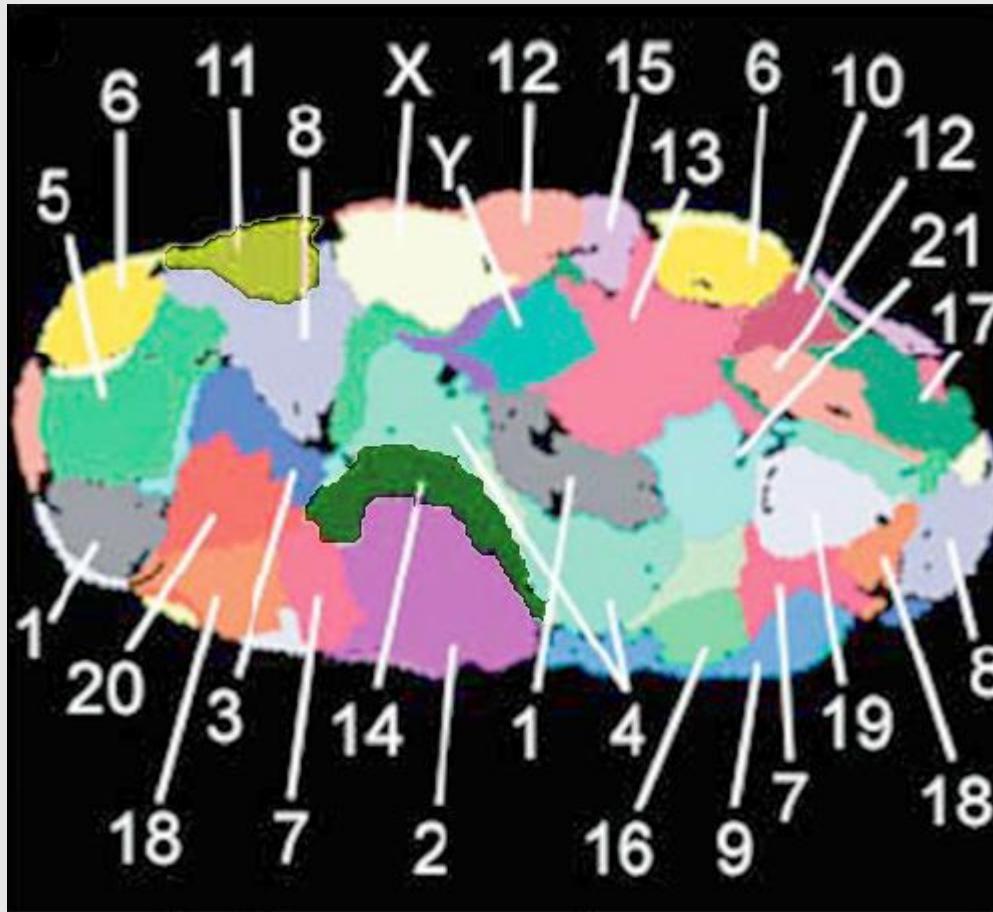
# NUCLEAR ARCHITECTURE



Theodor Boveri (1862-1915)



Carl Rabl (1853-1917)



Bolzer et al., Plos Biology (2005), 3 (5) e207

- The chromosomes are organized in the nucleus:
  - In a tissue specific manner
  - The organization is transmitted through the cell divisions
  - This organization is evolutionarily conserved
- The gene-rich regions occupy more central position in the nucleus

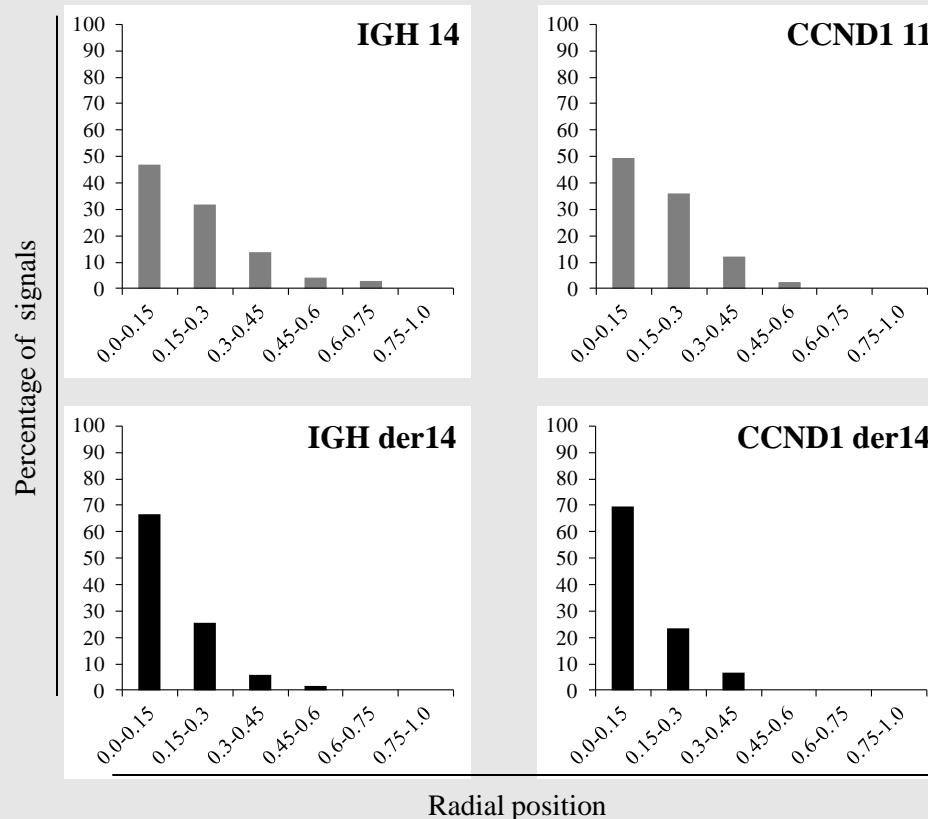
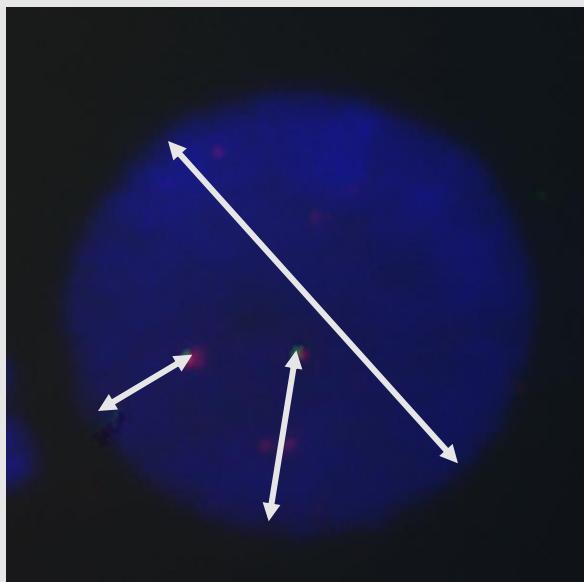


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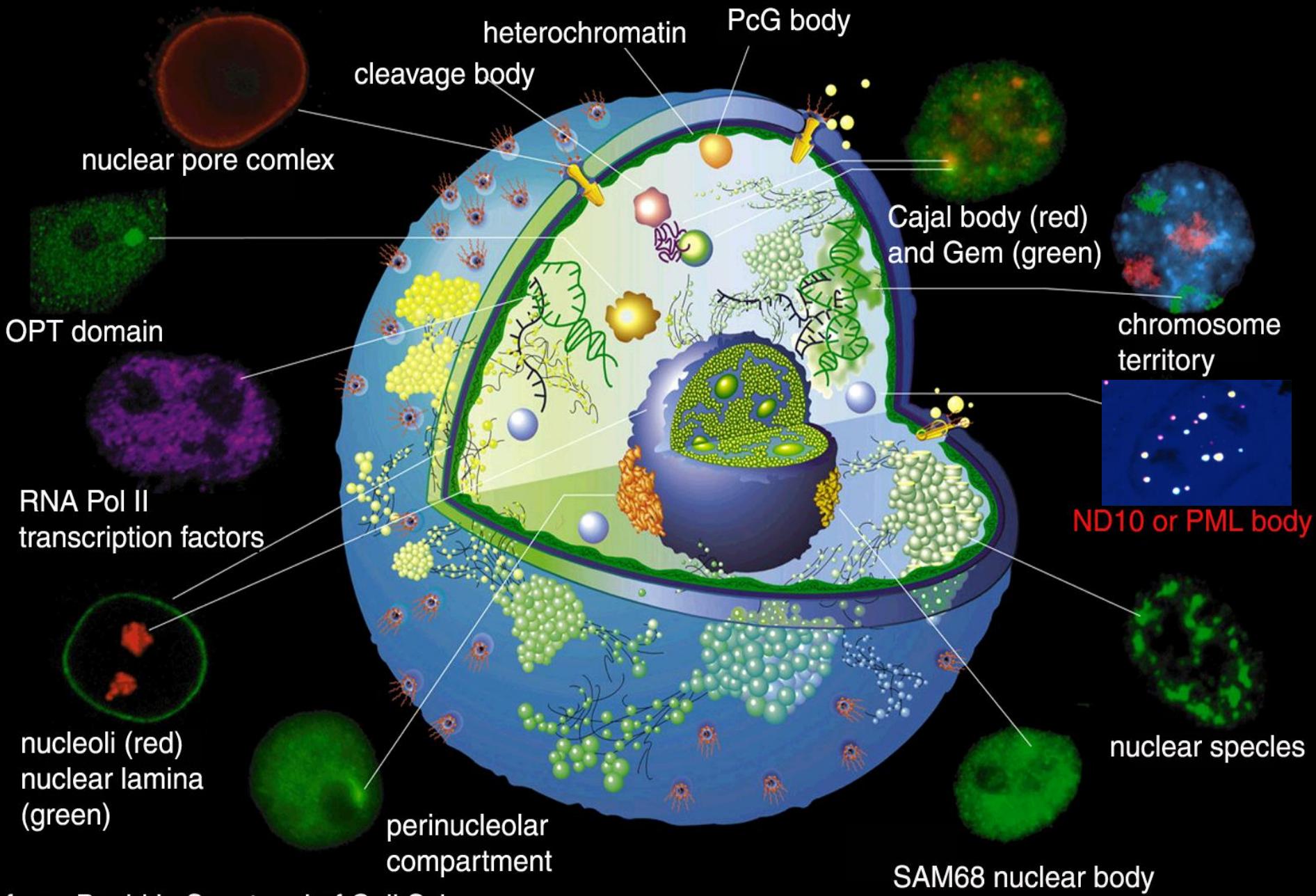


# THE 11q13 LOCUS IS RELOCALIZED WITHIN THE NUCLEUS IN MANTLE CELL LYMPHOMA



\*  
\*  
\*

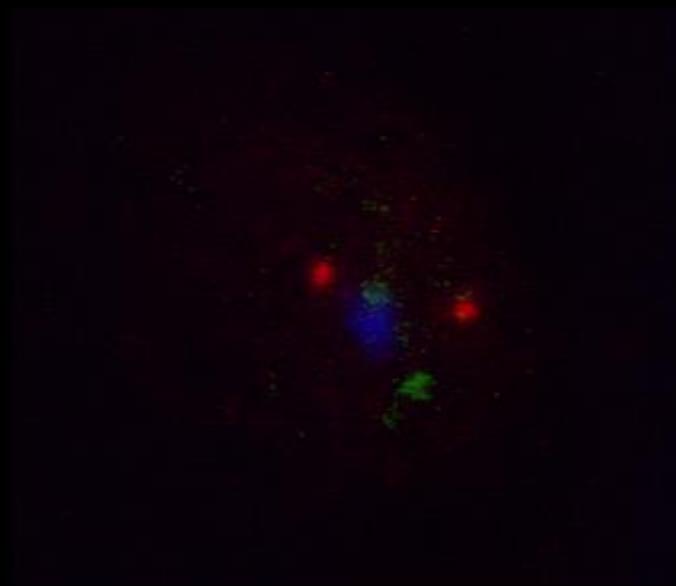
# NUCLEAR SUBSTRUCTURES



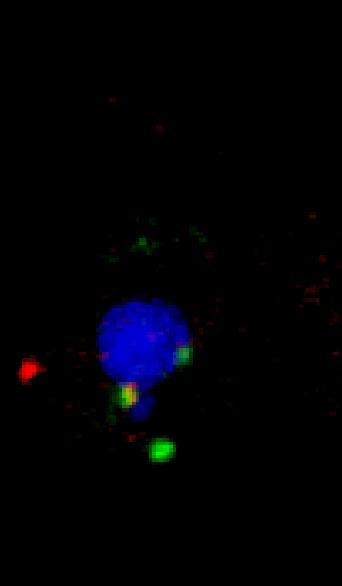
from: David L. Spector, J of Cell Sciense

# THE 11q13 LOCUS IS RELOCALIZED TOWARDS THE OUTER SURFACE OF THE NUCLEOLUS IN MCL

- 3D FISH ON INTERPHASE NUCLEI



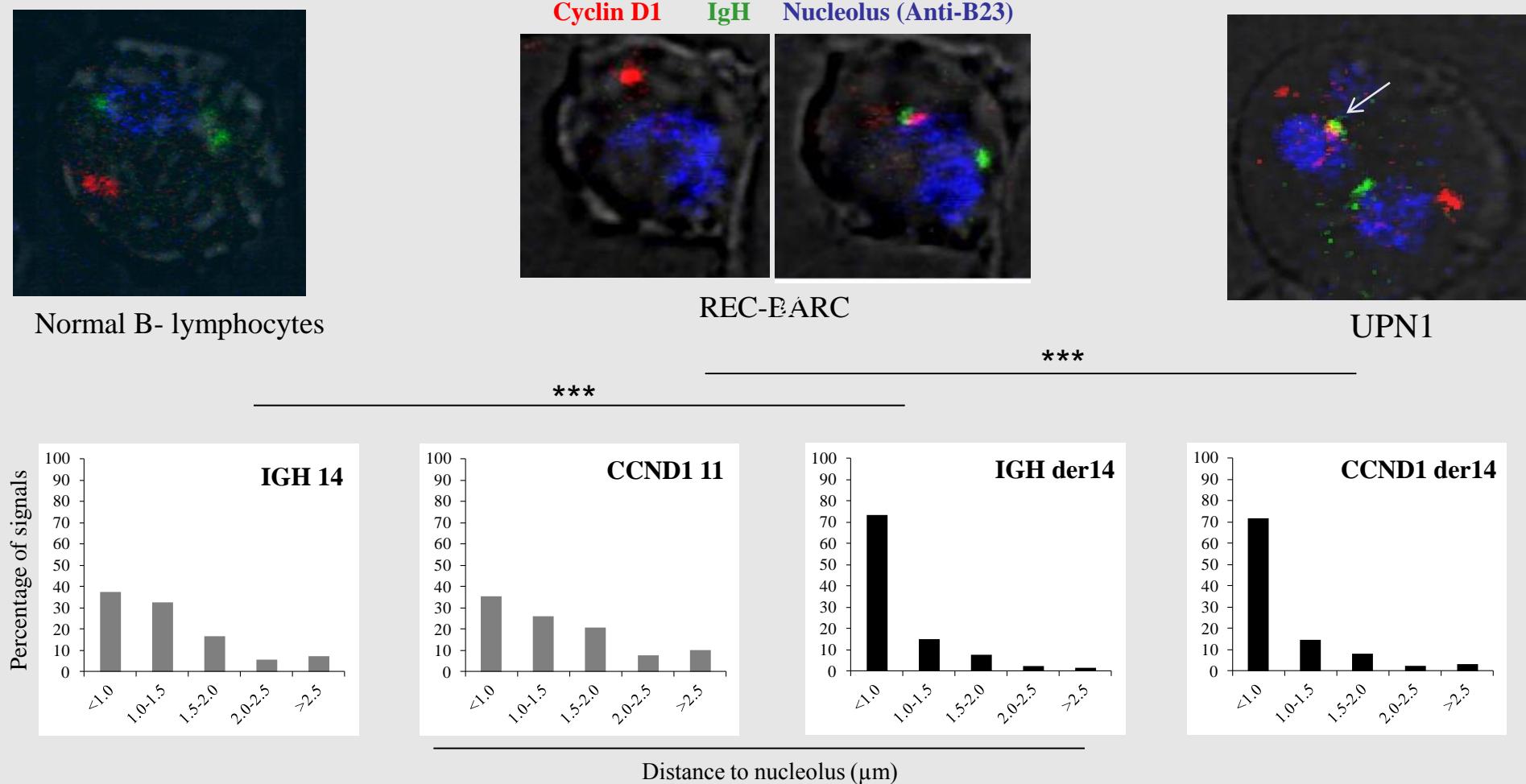
NORMAL  
LYMPHOCYTES



GRANTA  
(MCL)

Cyclin D1  
IgH  
NUCLEOLUS (Anti-B23)

# THE 11q13 LOCUS IS RELOCALIZED TOWARDS THE OUTER SURFACE OF THE NUCLEOLI IN MCL



## Statistical analysis:

- The relocalization event is highly reliable ( $p < 10^{-13}$ )
- The relocalization is linked to translocation; non-translocated CCND1 is an excellent control
- The relocalization is observed both in MCL cell lines and patients

# NUCLEOLUS AND REGULATION OF TRANSCRIPTION – LR1

Proc. Natl. Acad. Sci. USA  
Vol. 94, pp. 3605–3610, April 1997  
Biochemistry

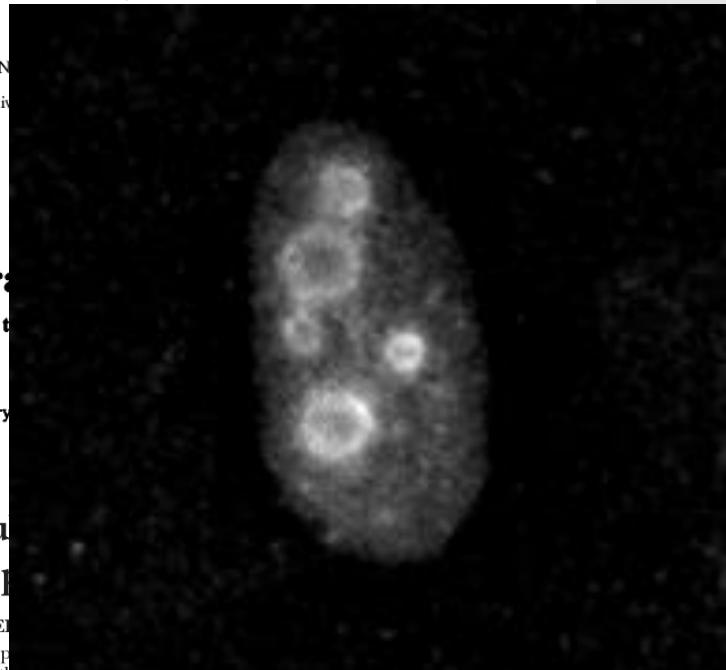
## Nucleolin is one component of the B cell-specific transcription factor and switch region binding protein, LR1

(Ig/rDNA/recombination)

L. A. HANAKAHI\*, LAURIE A. DEMPSEY\*, MING-JIE LI\*, AND NANCY MAIZELS†

Departments of \*Molecular Biophysics and Biochemistry and †Genetics, Yale University, New Haven, Connecticut 06511

Proc. Natl. Acad. Sci. USA  
Vol. 91, pp. 4915–4919, May 1994  
Biochemistry



## LR1 regulates c-myc transcription

(MYC gene/immunoglobulin/chromosome translocation)

APRIL BRYST‡ AND NANCY MAIZELS‡

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Vol. 13 November 1999

The FASEB Journal

THE JOURNAL OF BIOLOGICAL CHEMISTRY  
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Vol. 279, No. 12, Issue of March 19, pp. 10855–10863, 2004  
Printed in U.S.A.

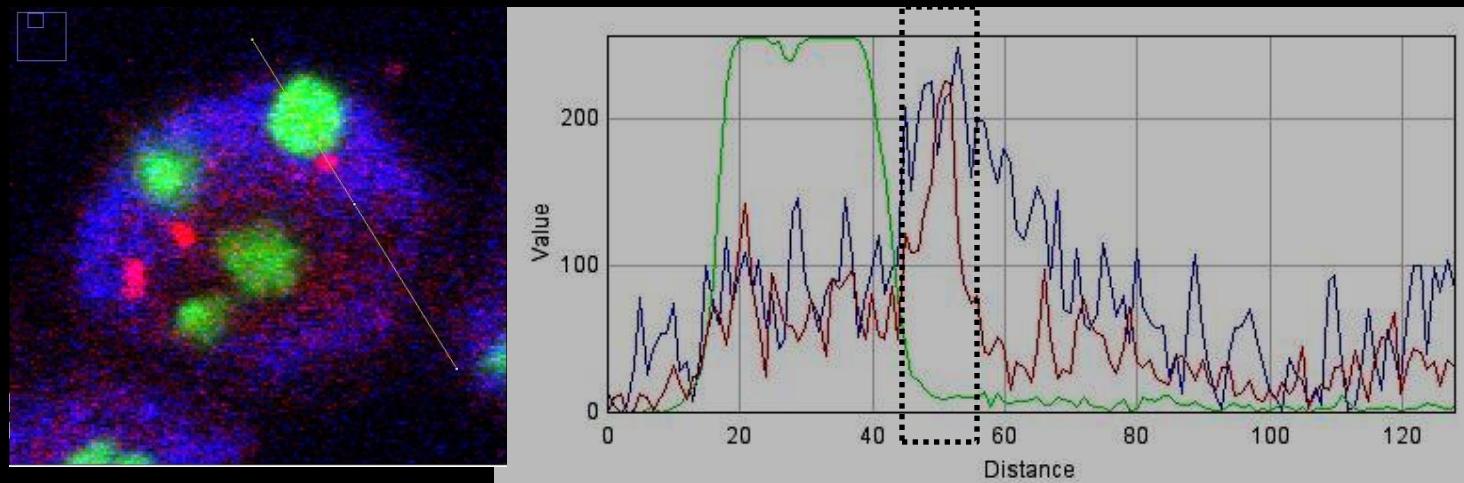
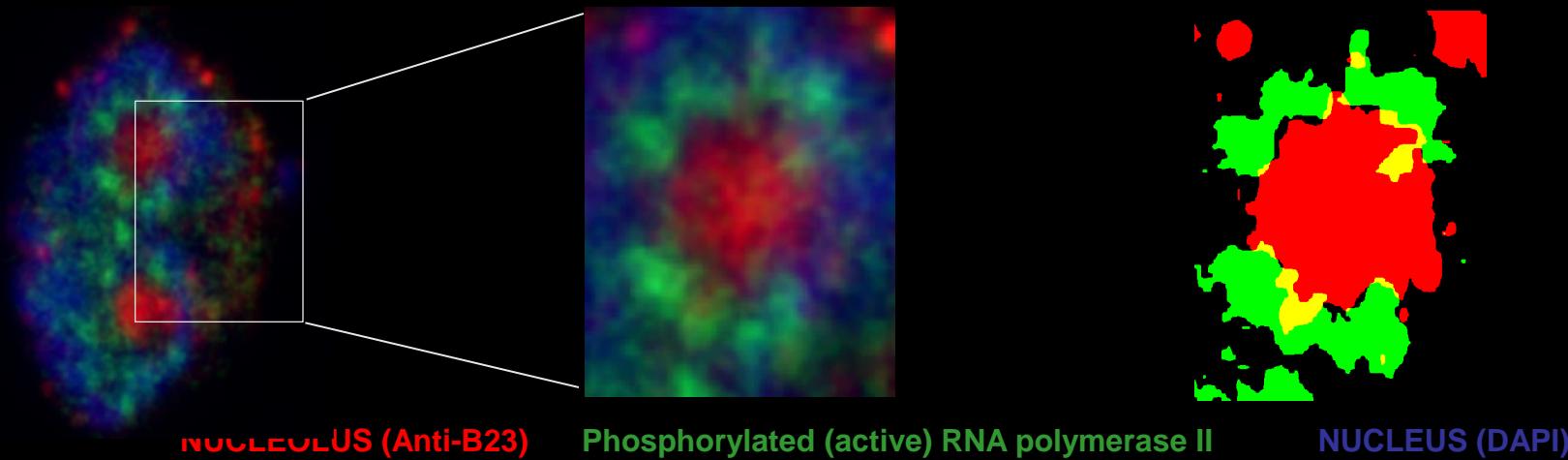
## Identification of Nucleolin as an AU-rich Element Binding Protein Involved in *bcl-2* mRNA Stabilization\*

Received for publication, August 18, 2003, and in revised form, December 4, 2003  
Published, JBC Papers in Press, December 16, 2003, DOI 10.1074/jbc.M309111200

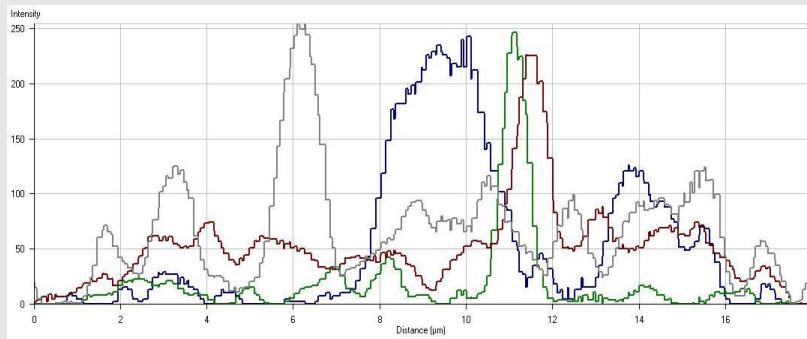
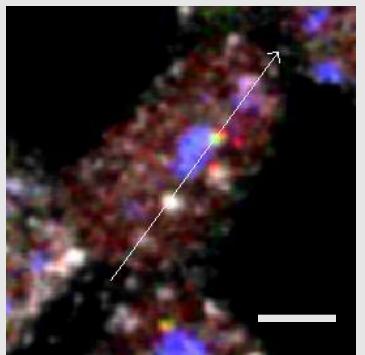
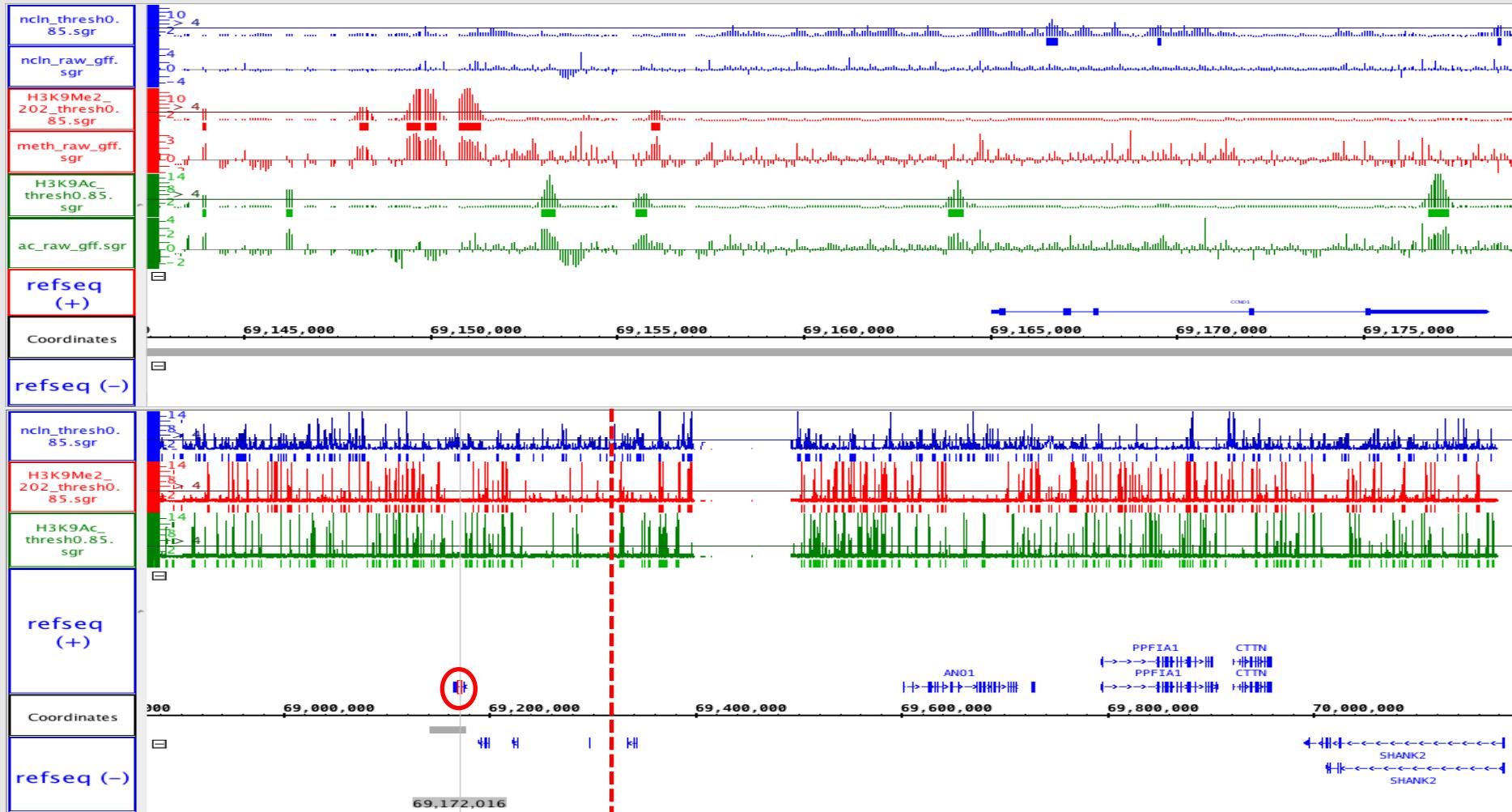
Tapas K. Sengupta‡§, Sumita Bandyopadhyay‡§, Daniel J. Fernandes‡, and Eleanor K. Spicer‡¶||

From the ‡Department of Biochemistry and Molecular Biology and ¶Department of Pharmaceutical Sciences, Medical University of South Carolina, Charleston, South Carolina 29425

# TRANSCRIPTION FACTORIES ARE LOCATED IN THE PERINUCLEOLAR REGION



# EPIGENETIC MARKS IN THE TRANSLOCATION REGION

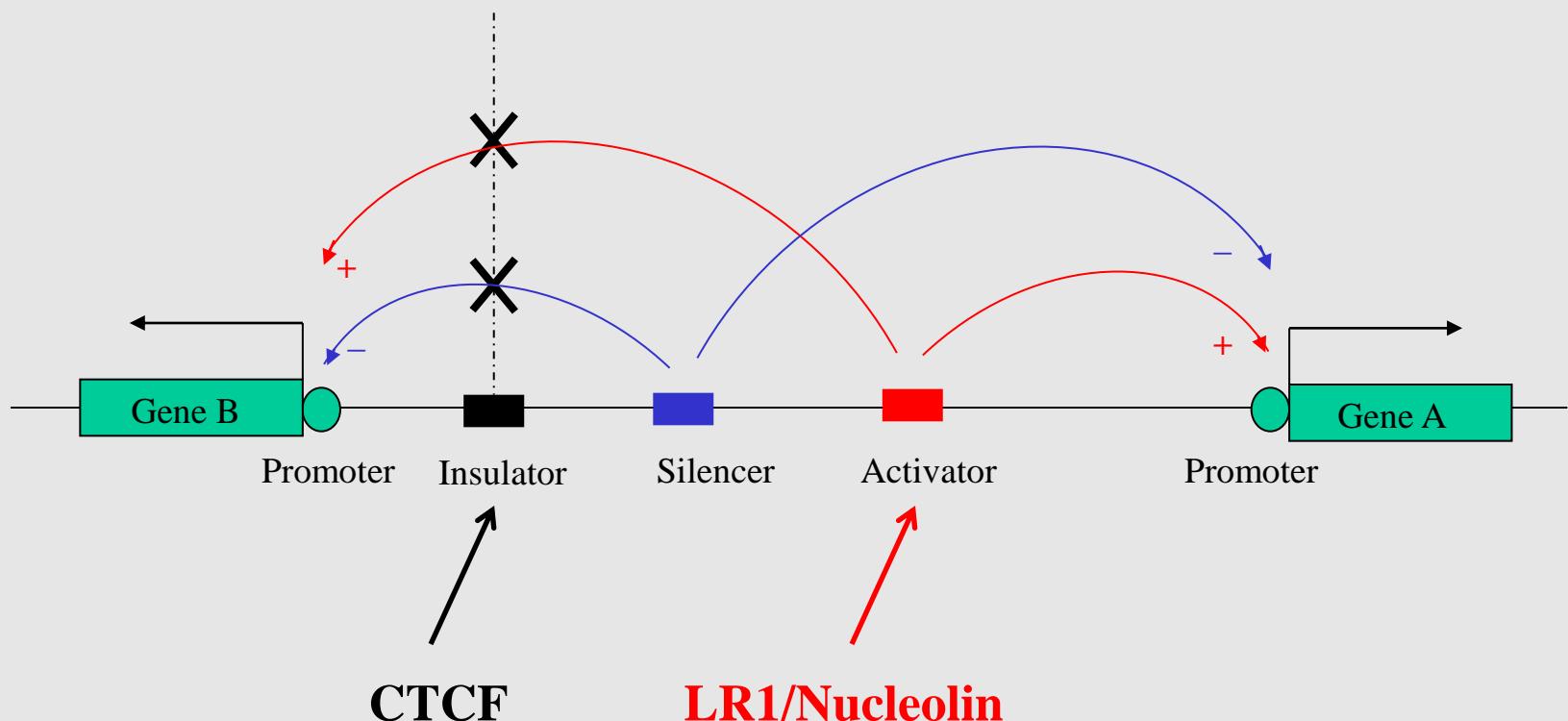


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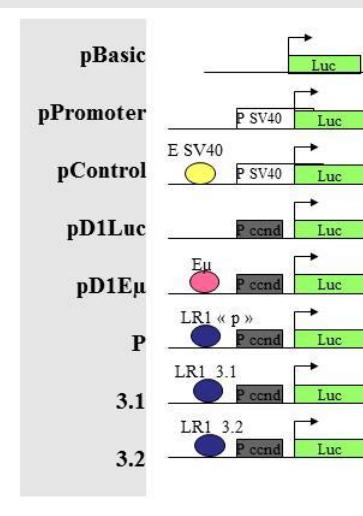
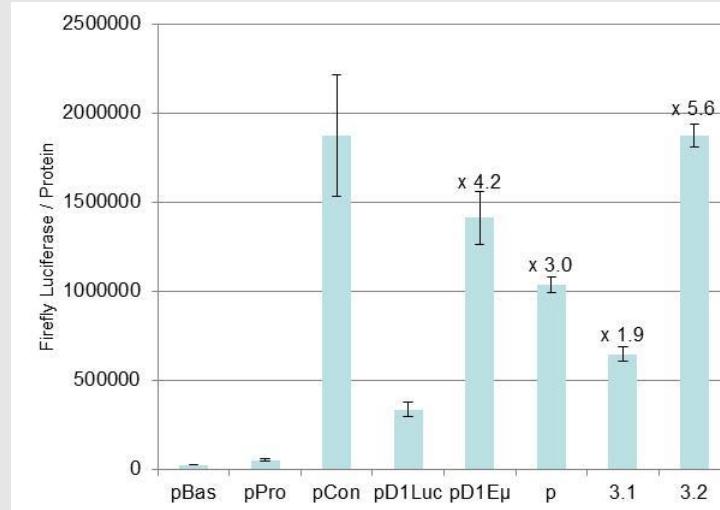
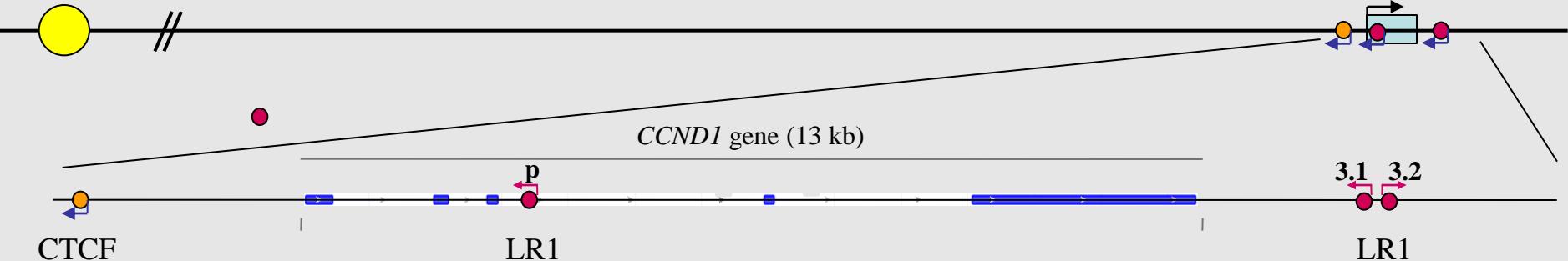
Interactions  
Moléculaires  
et Cancer

# REGULATION OF TRANSCRIPTION:



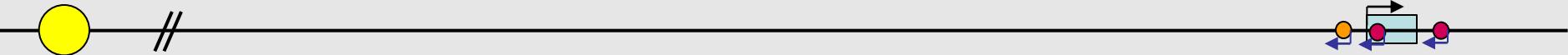
# A NUCLEOLIN-BINDING SITE ACTIVATES TRANSCRIPTION FROM THE CCND1 PROMOTER

IGHm Enhancer



# NUCLEOLIN, CTCF AND TRANSCRIPTIONAL REGULATION OF THE *CCND1* LOCUS

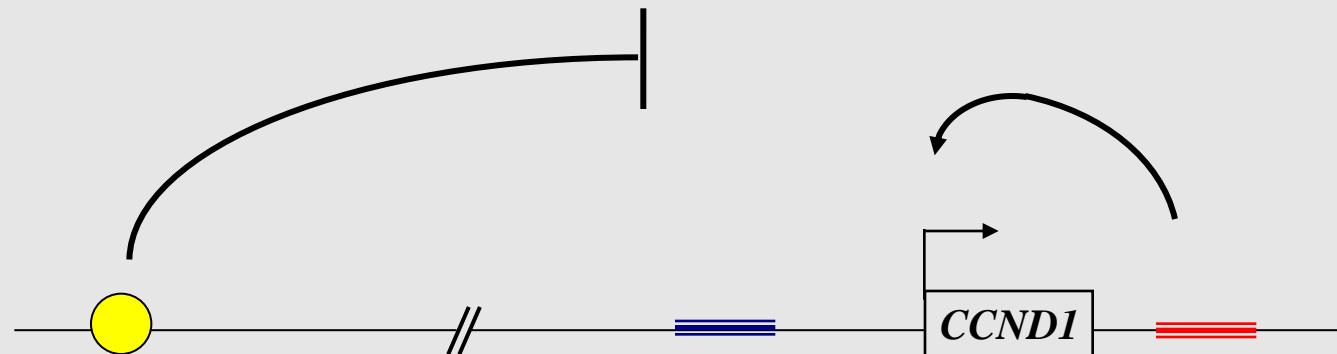
IGHm Enhancer



IGHm Enhancer

CTCF

Nuc



IGHm Enhancer

Allinne et al., Blood, 2014



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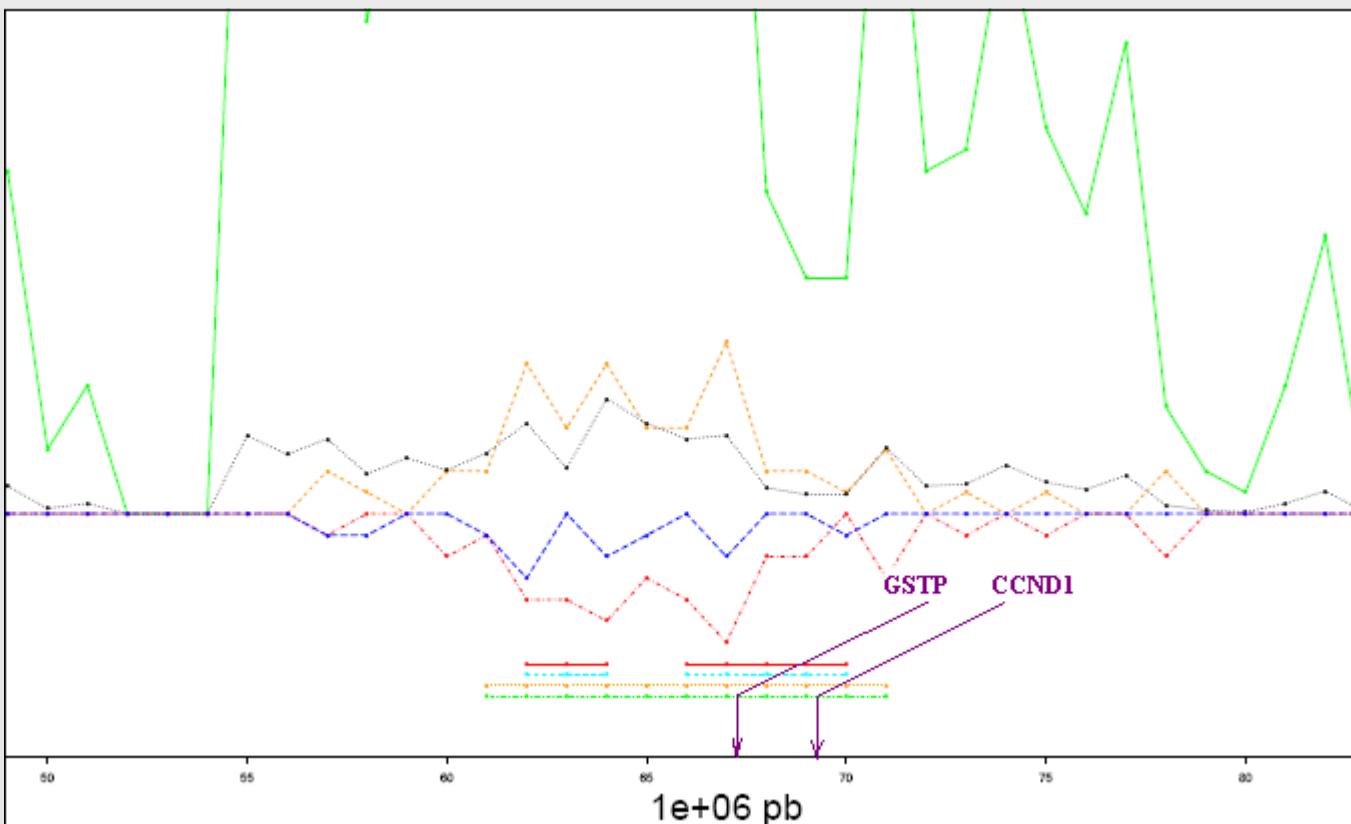
# ARE OTHER 11q13 OR 14q32 GENES ACTIVATED AFTER THE TRANSLOCATION?

## Curves

- Ensembl genome
- Tested list
- Hazard estimation
- Upregulated genes
- Downregulated genes

## Statistical results

- Units where Tested list > Hazard estimation
- Statistically validated ► Binomial test on the unit
- Optimized regions
- Significant ► Wilcoxon test on the small region



# 11q13 GENES ACTIVATED AFTER THE TRANSLOCATION

Start point	Gene	Expression rate	Gene name
60739115	<b>CD6</b>	0,850392574	CD6 molecule
60869930	<b>CD5</b>	0,959535821	CD5 molecule
61100654	DAK	0,530690076	dihydroxyacetone kinase 2 homolog (S. cerevisiae)
62201016	AHNAK	2,183624534	AHNAK nucleoprotein
62360675	<b>MTA2</b>	0,511737498	metastasis associated 1 family, member 2
62457747	BSCL2	0,771842816	Bernardinelli-Seip congenital lipodystrophy 2 (seipin)
62623518	SLC3A2	0,806756116	solute carrier family 3
63341934	PLA2G16	1,236042113	phospholipase A2, group XVI
63448922	RTN3	0,941432897	reticulon 3
64037300	<b>BAD</b>	0,607078976	BCL2-associated agonist of cell death
64494383	RASGRP2	0,618417066	RAS guanyl releasing protein 2 (calcium and DAG-regulated)
64513861	PYGM	0,851019014	phosphorylase, glycogen, muscle
64692180	PPP2R5B	0,452563551	protein phosphatase 2, regulatory subunit B', beta isoform
64794910	SNX15	0,556354468	sorting nexin 15
65337943	SSSCA1	0,493153003	Sjogren syndrome/scleroderma autoantigen 1
65479489	KAT5	0,478519186	K(lysine) acetyltransferase 5
65686728	DRAP1	0,777999239	DR1-associated protein 1 (negative cofactor 2 alpha)
66059373	TMEM151A	0,59365937	transmembrane protein 151A
66278119	BBS1	0,546155142	Bardet-Biedl syndrome 1
66330935	<b>CTSf</b>	0,49155074	cathepsin F
66610883	RCE1	0,526977415	RCE1 homolog, prenyl protein peptidase (S. cerevisiae)
67195935	RPS6KB2	0,40202797	ribosomal protein S6 kinase, 70kDa, polypeptide 2
67205518	CORO1B	1,284980925	coronin, actin binding protein, 1B
67259239	PITPNM1	0,401006288	phosphatidylinositol transfer protein, membrane-associated 1
67351066	<b>GSTP1</b>	1,546682584	glutathione S-transferase pi 1
67820326	CHKA	0,679197547	choline kinase alpha
68080108	LRP5	0,549308072	low density lipoprotein receptor-related protein 5
68522088	CPT1A	1,355308335	carnitine palmitoyltransferase 1A (liver)
68671319	IGHMBP2	0,607078976	immunoglobulin mu binding protein 2
<b>69455873</b>	<b>CCND1</b>	<b>4,651669229</b>	<b>cyclin D1</b>
69480331	<b>ORAOV1</b>	0,503355776	oral cancer overexpressed 1
70244612	CTTN	0,495420482	cortactin
71145457	DHCR7	0,515050676	7-dehydrocholesterol reductase
71639768	RNF121	0,386244877	ring finger protein 121
71900602	FOLR1	1,229322176	folate receptor 1 (adult)
75526212	<b>UVRAG</b>	0,664071631	UV radiation resistance associated gene

der11

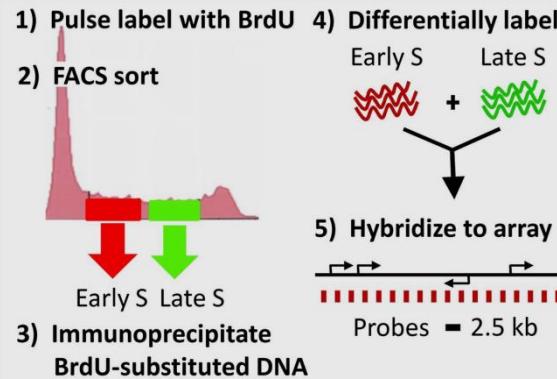
der14



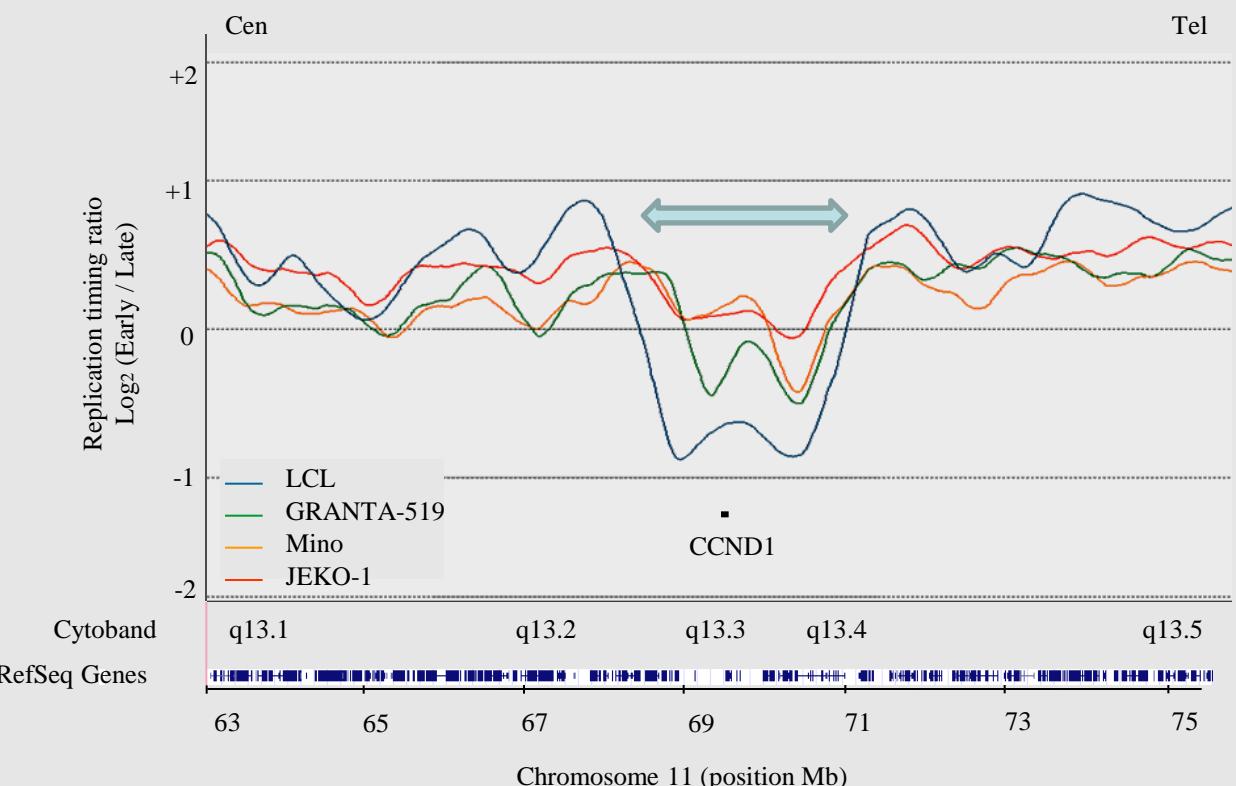
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# REPLICATION TIMING CHANGES BOTH ON der11 AND der14 AFTER THE TRANSLOCATION



From Ryba et al., 2012



Collaboration: MN Prioleau



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# OTHER CANCERS?

- A multitude of translocations involving the acrocentric chromosomes (13, 14, 15, 21, 22)
- The most frequent variant of Burkitt lymphoma has a t(8;14)(q24;q32) translocation.

*Is the nucleolus involved in this case?*

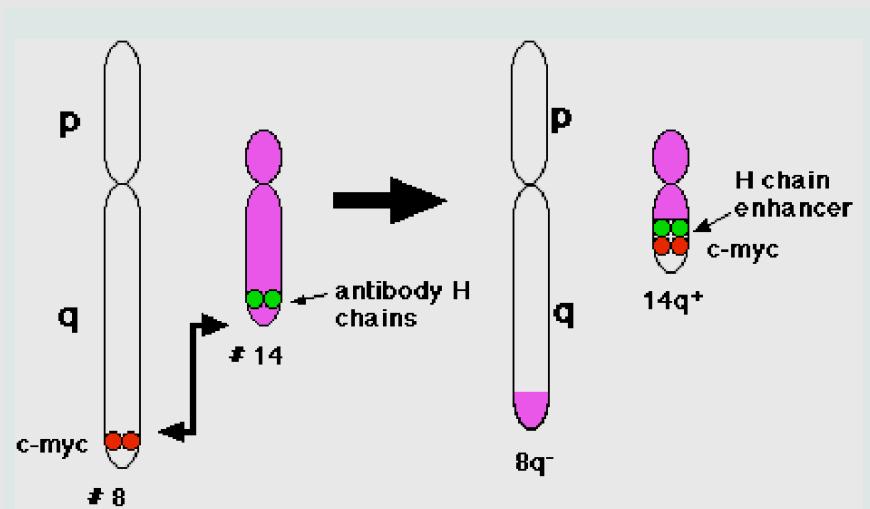


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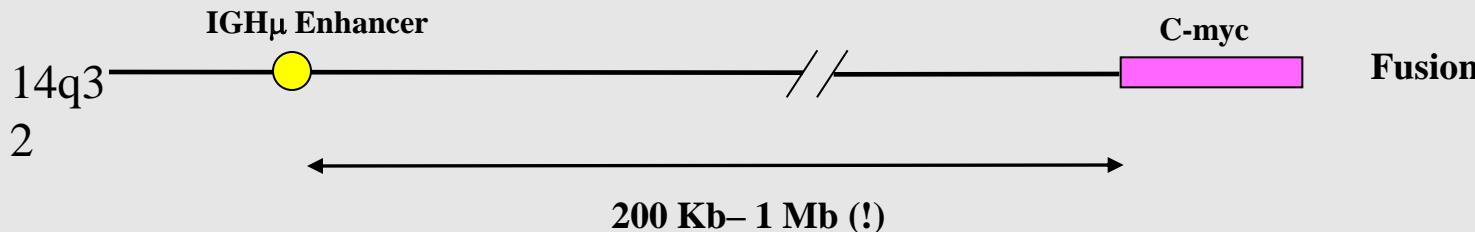
# BURKITT LYMPHOMA

- Burkitt lymphoma is a monoclonal proliferation of B lymphocytes characterized by small non-cleaved cells that are uniform in appearance and that produce a diffuse pattern of tissue involvement.
- Most Burkitt lymphomas carry a translocation of the c-myc oncogene from chromosome 8 to either the immunoglobulin (Ig) heavy-chain region on chromosome 14 [t(8;14)] or chromosome 22 (lambda light chain) [t(8;22)].
- Burkitt lymphoma is a very rare form of cancer in France, with about 100 new cases occurring each year.
- High occurrence in EBV-infected children in Africa and HIV patients in Europe

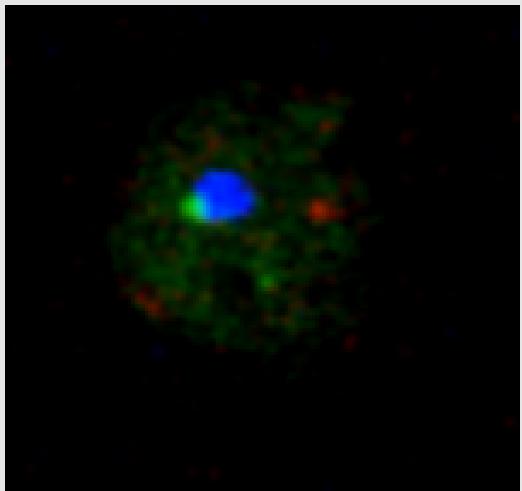


# C-MYC ACTIVATION IN BURKITT LYMPHOMA: IS IGH $\mu$ ENHANCER REALLY INVOLVED?

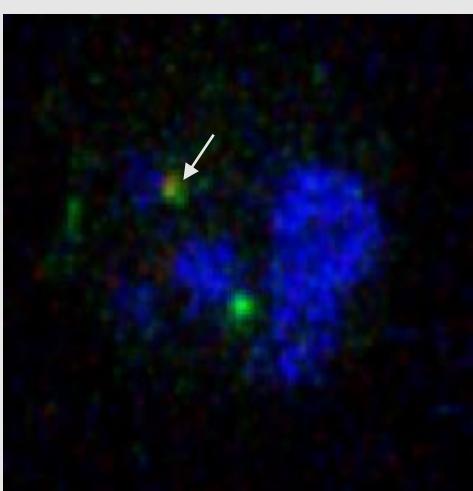
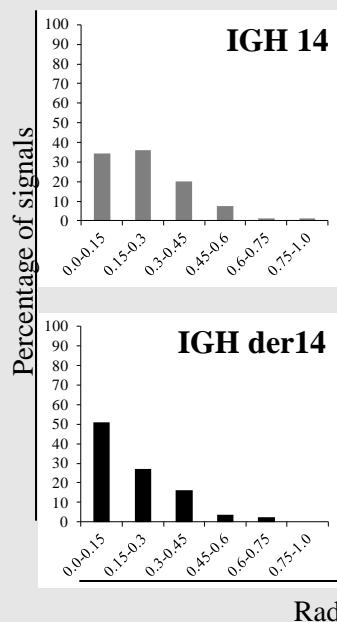
- t(8 ; 14) translocation induces overexpression of *c-myc* in B-cells



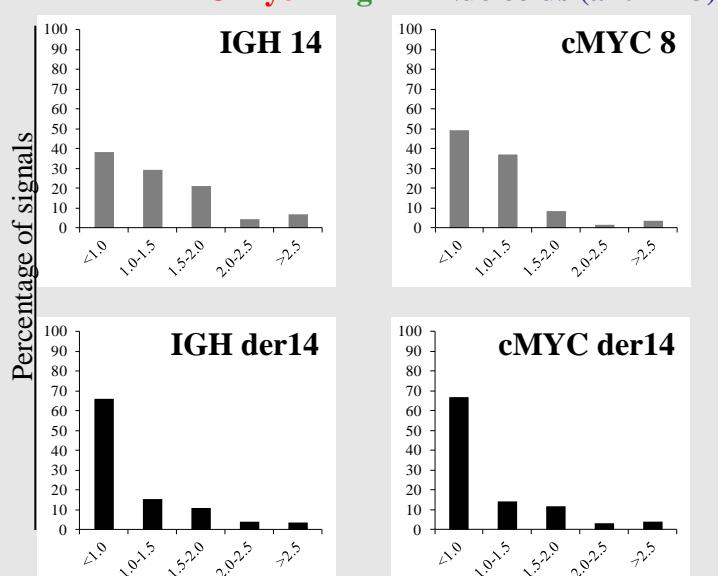
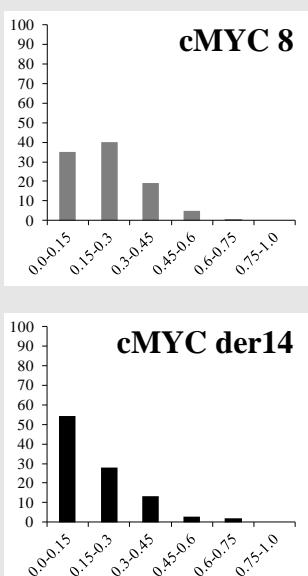
# TRANSLOCATED *c-myc* LOCUS IS LOCALIZED IN THE PERINUCLEOLAR REGION IN BURKITT LYMPHOMA



Normal B-lymphocytes



P3HR1 Burkitt cell line



# NUCLEOLIN IS A FACTOR OF TRANSCRIPTION THAT REGULATES *C-MYC*

*Proc. Natl. Acad. Sci. USA*  
Vol. 94, pp. 3605–3610, April 1997  
Biochemistry

## Nucleolin is one component of the B cell-specific transcription factor and switch region binding protein, LR1

(Ig/rDNA/recombination)

L. A. HANAKAHY\*, LAURIE A. DEMPSEY\*, MING-JIE LI\*, AND NANCY MAIZELS\*†‡

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*Proc. Natl. Acad. Sci. USA*  
Vol. 91, pp. 4915–4919, May 1994  
Biochemistry

## LR1 regulates *c-myc* transcription in B-cell lymphomas

(*MYC* gene/immunoglobulin/chromosome translocation/Burkitt lymphoma)

APRIL BRYST† AND NANCY MAIZELS‡

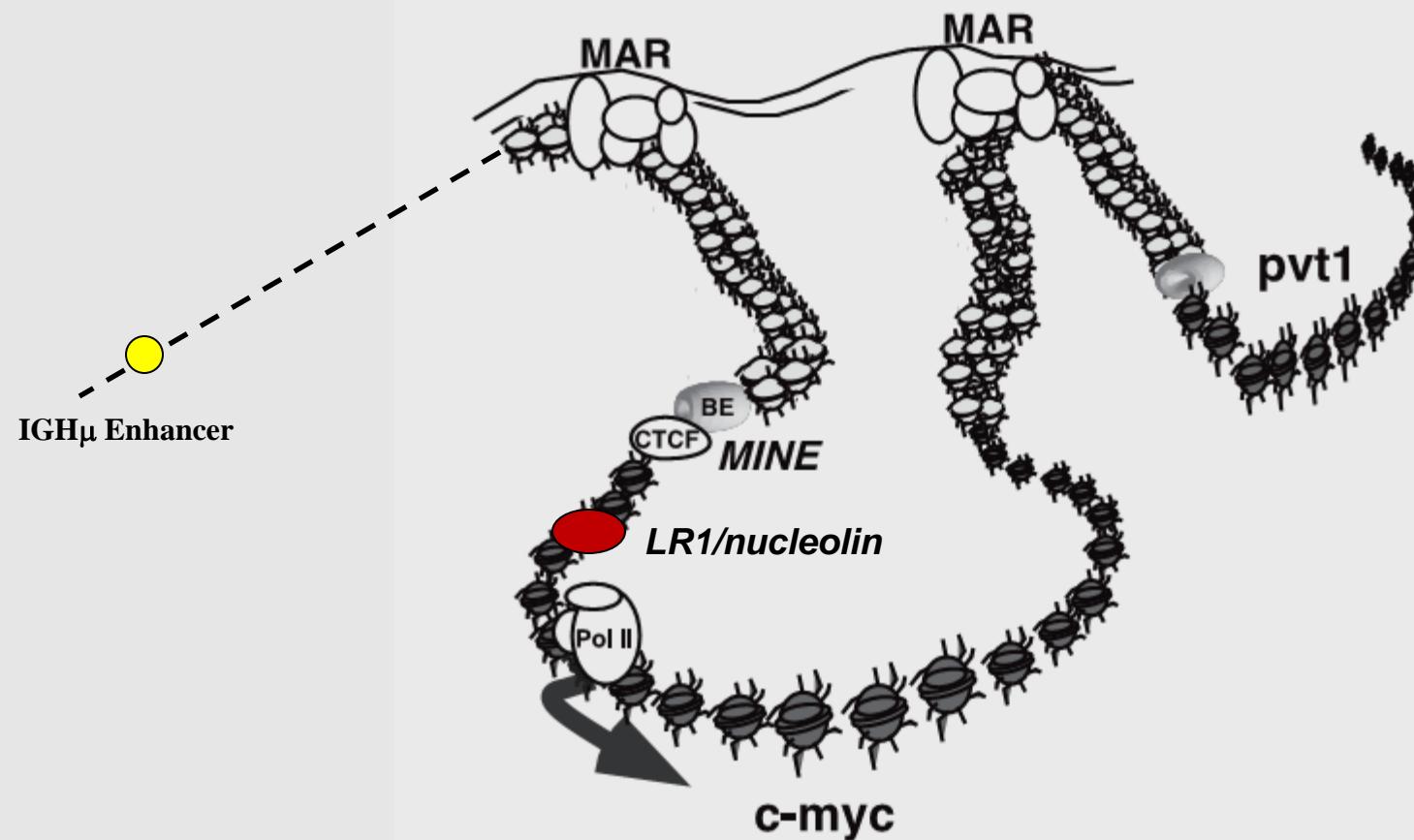
Department of Molecular Biophysics and Biochemistry, Yale Medical School, 333 Cedar Street, New Haven, CT 06510



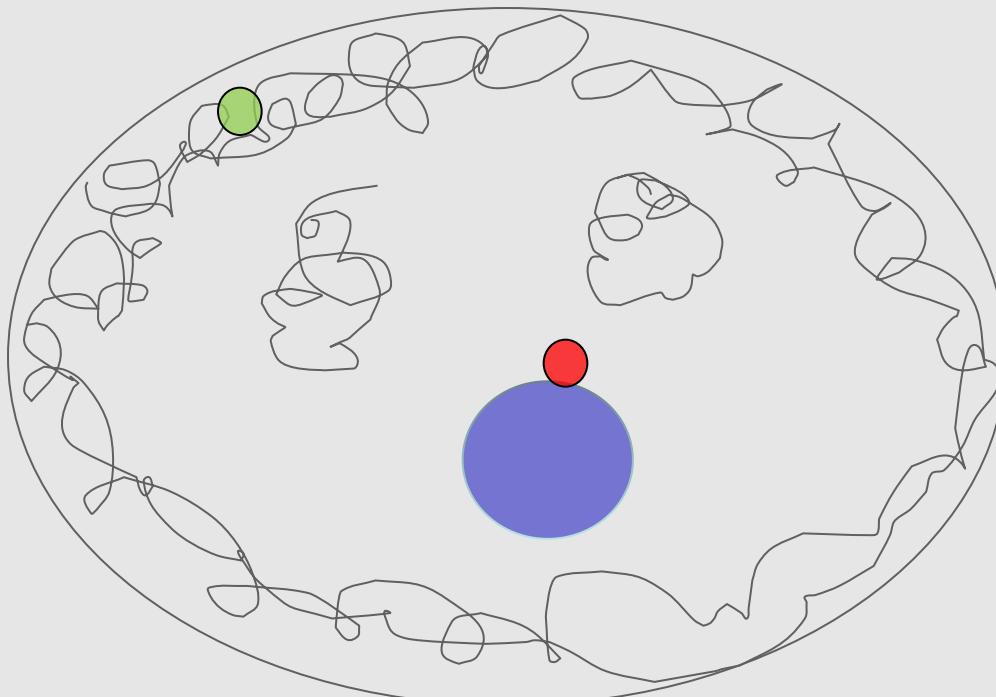
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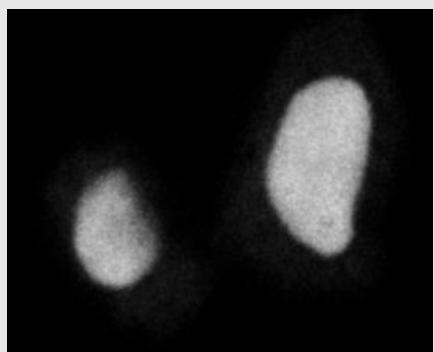
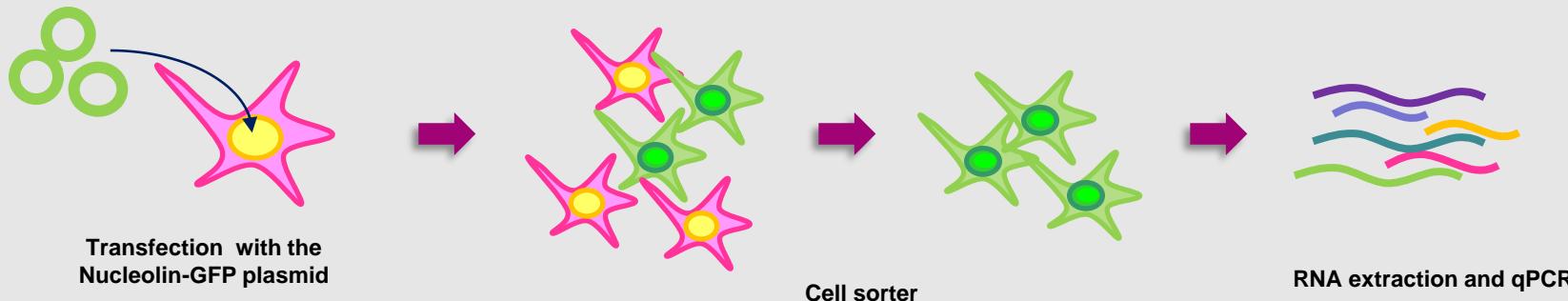
# AN INSULATOR AND A NUCLEAR MATRIX ATTACHMENT SITE SEPARATE *c-myc* FROM THE IGH $\mu$ ENHANCER



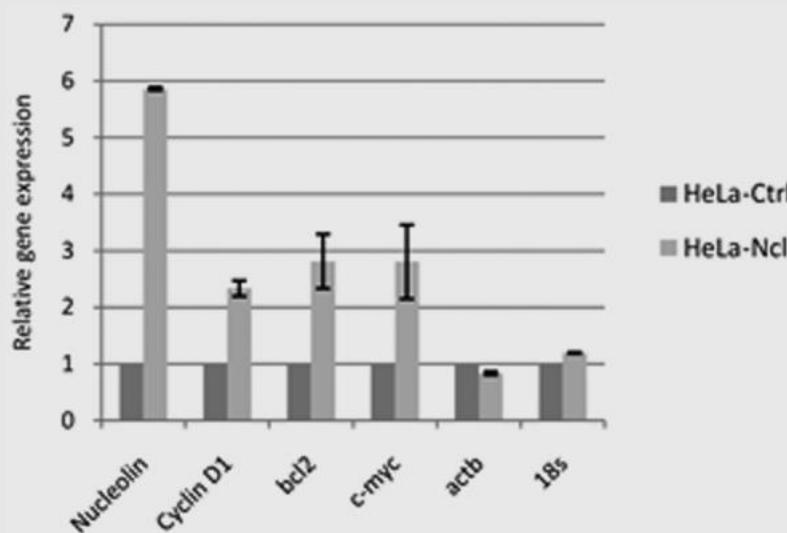
# INTRANUCLEAR RELOCALIZATION AND GENE ACTIVATION IN CANCER



# IF THE MOUNTAIN WILL NOT COME TO MAHOMET, MAHOMET MUST GO TO THE MOUNTAIN



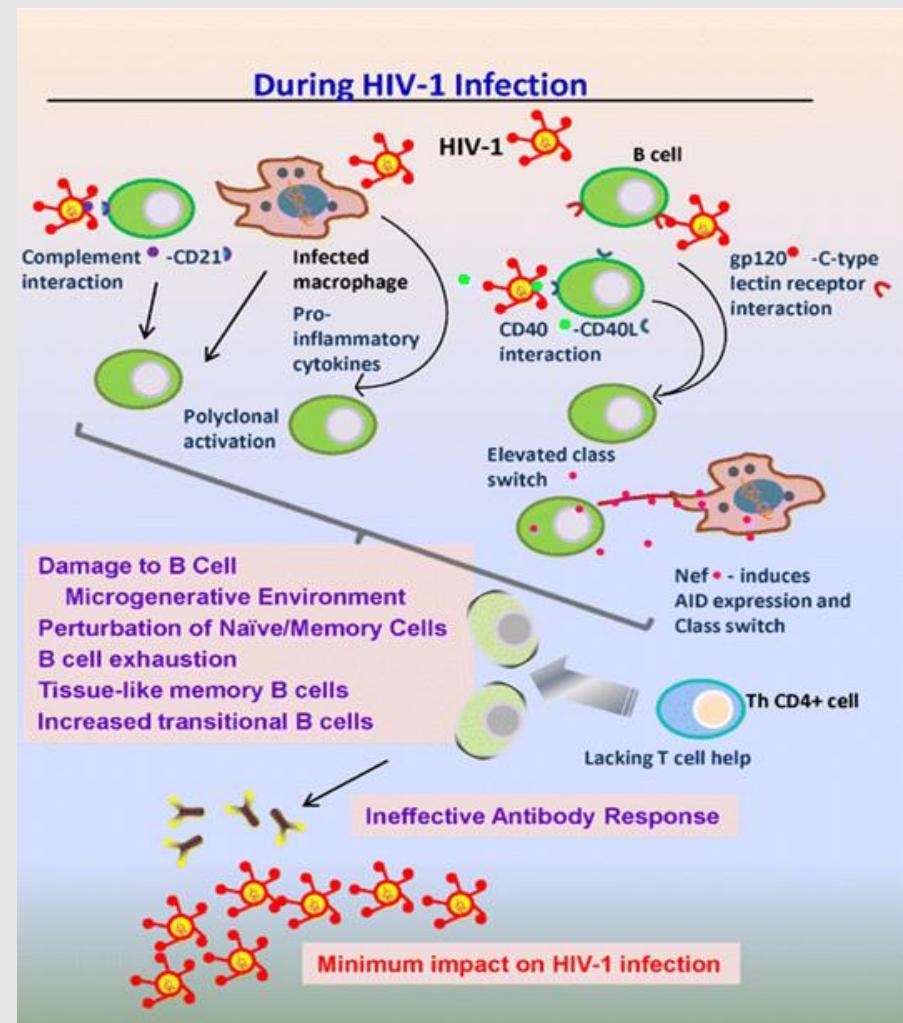
Nucleolin-GFP



→ Ectopic overexpression of nucleolin leads to its delocalization in the nucleus and overexpression of endogenous *CCND1*, *c-myc* and *bcl-2*

# HIGH OCCURRENCE OF BURKITT LYMPHOMA IN HIV PATIENTS: WHY?

- ✓ gp-120 can interact with CD21 expressed on B cells (Moir.S and al 2000)
- ✓ HIV-1 causes B-cell hyperactivation (Schnittan and al,1984)
- ✓ Elevated class switch in B lymphocytes
- ✓ Induces B cell to proliferate (Nair MPN and al 1988)
- ✓ Causes B cell abnormal response
- ✓ The production of autoantibodies
- ✓ Aberrant B-cell surface markers: Change in B cell receptors.



Shen ,2011



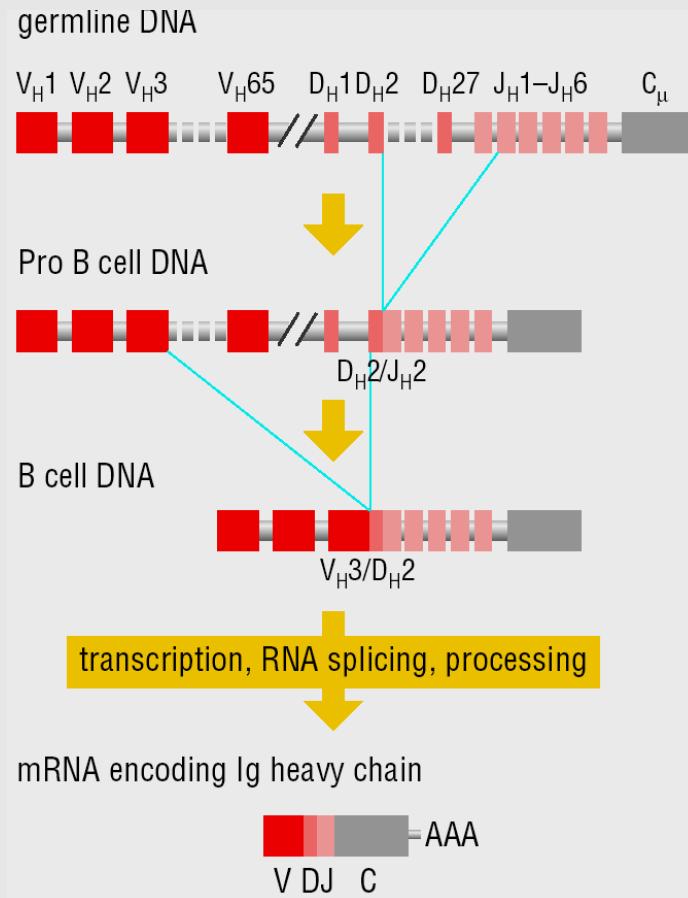
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Interactions  
Moléculaires  
et Cancer

# SOMATIC CELL

# B LYMPHOCYTE



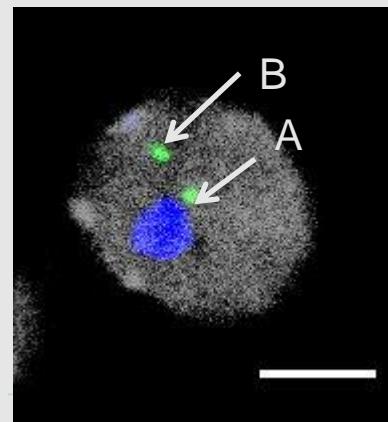
Recombination

Repair

Recombination

Repair

# WHAT IS THE ROLE OF THE NUCLEOLUS IN NORMAL B-LYMPHOCYTES?



Gray-nucleus, green-IgH, blue-B23

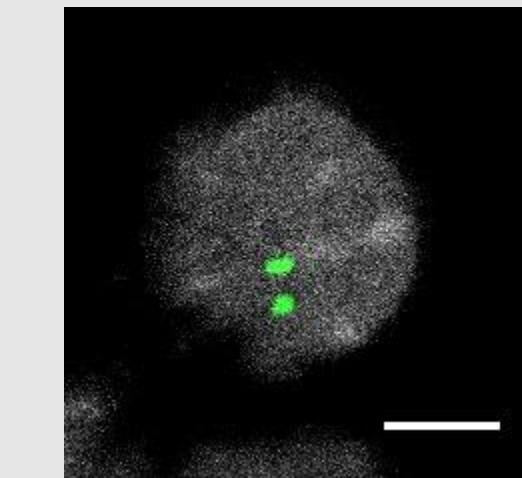
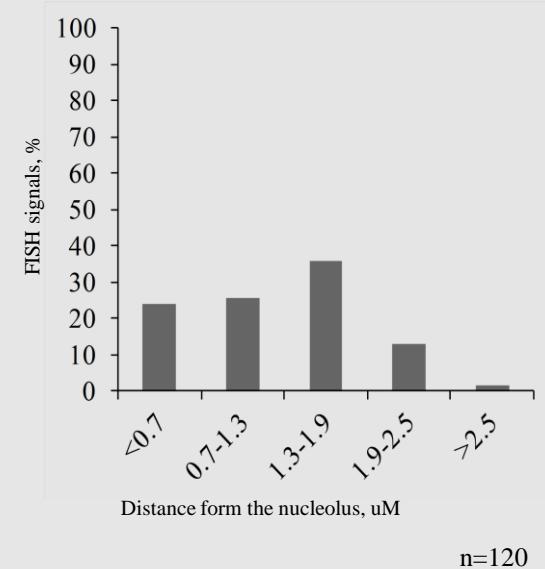
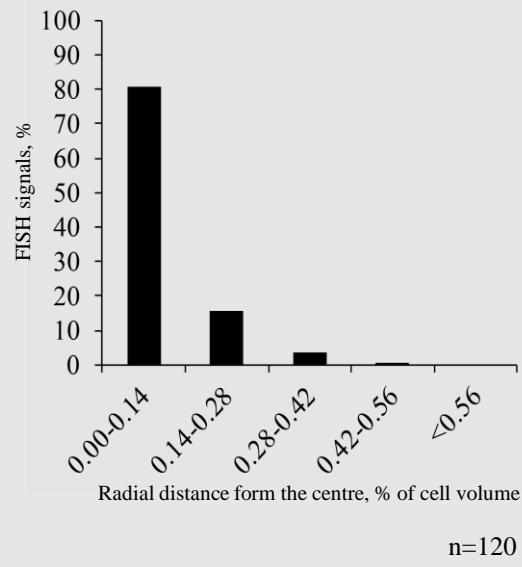
One IgH allele is always present in the perinucleolar region in normal B-lymphocytes

→ *A role in regulation of transcription (repression/activation)?*

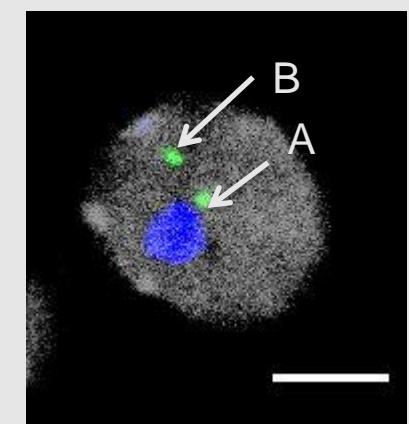
→ *A role in allelic exclusion?*

→ *A role in somatic hypermutation and class switch recombination?*

# BOTH IgH ALLELES ARE LOCATED IN THE CENTER OF THE NUCLEUS; ONE ALLELE IS ALWAYS LOCATED CLOSE TO THE NUCLEOLUS



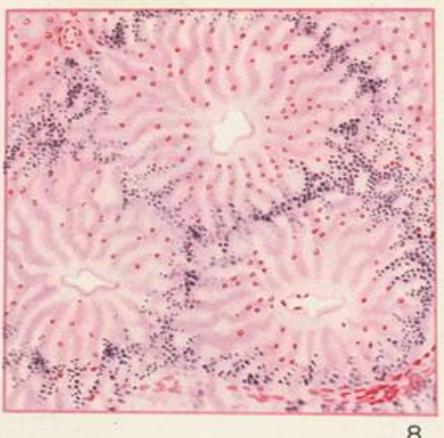
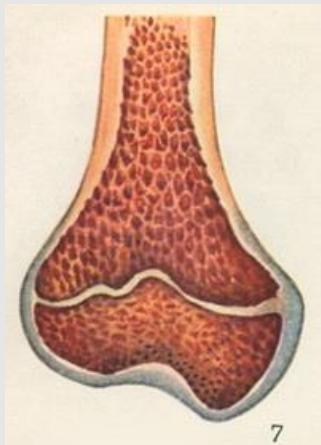
Gray-nucleus, green-IgH, blue-B23



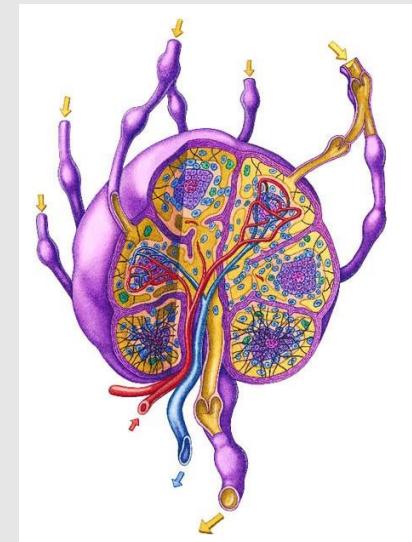
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# B-CELL DIFFERENTIATION

Bone marrow



Lymph node



B-cell maturation

V-D-J Recombination

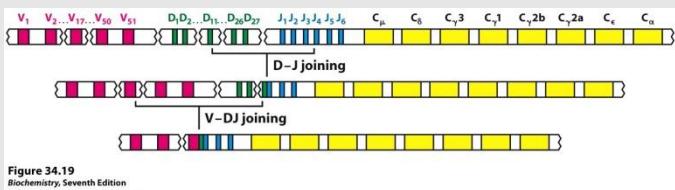
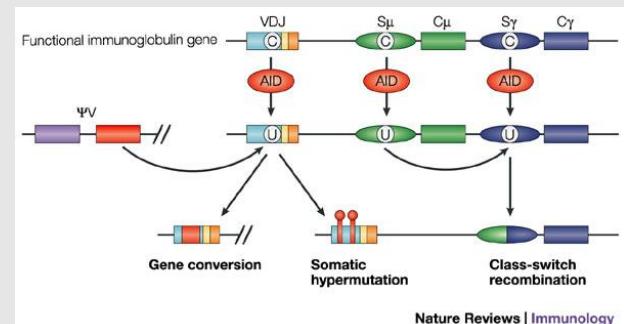


Figure 34.19  
Biochemistry, Seventh Edition  
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Class switch recombination and  
Somatic hypermutation



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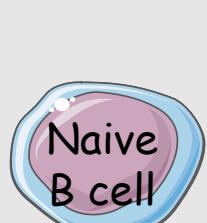
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Interactions  
Moléculaires  
et Cancer

# AN IN VITRO SYSTEM OF HUMAN B-LYMPHOCYTE DIFFERENTIATION

Stim1

IL-2, CD40L, CpG and Anti Fab'2



Day 0

Human naive B-cells from blood.

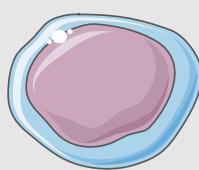


Day 1

Activated B-cells prior proliferation

Stim2

IL2, IL4 and IL10



Day 4

Proliferating B-cells.



Day 5

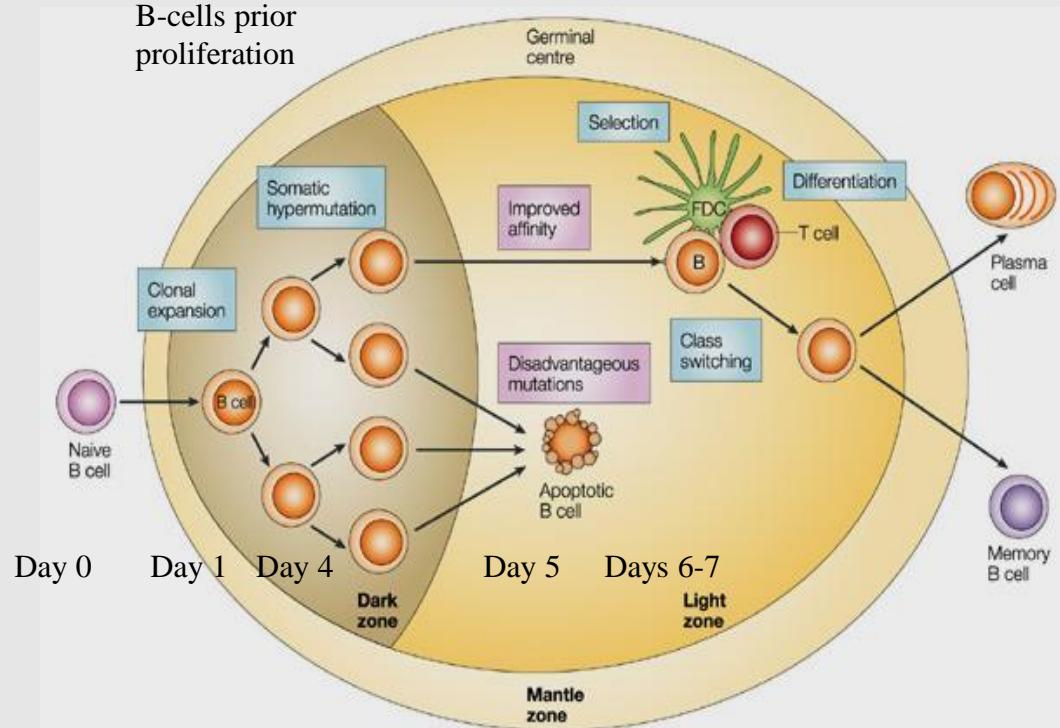
Somatic Hypermutation



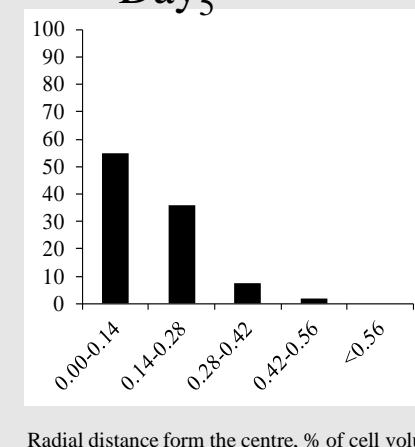
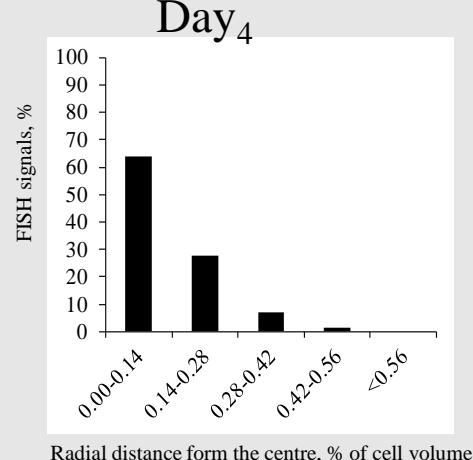
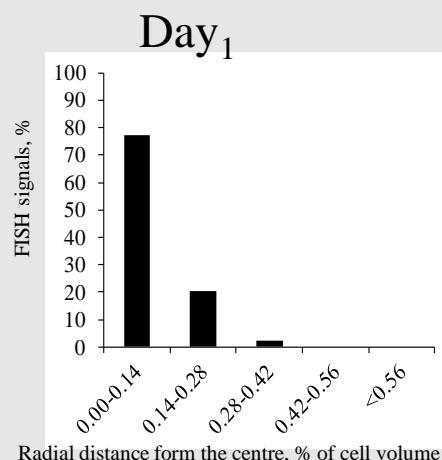
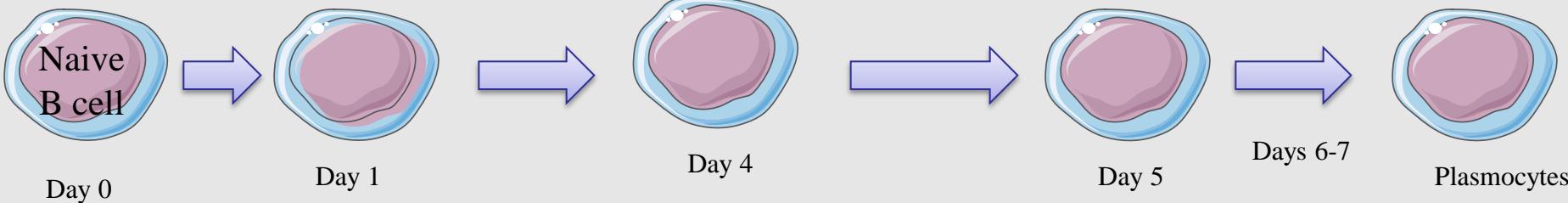
Days 6-7

Plasmocytes

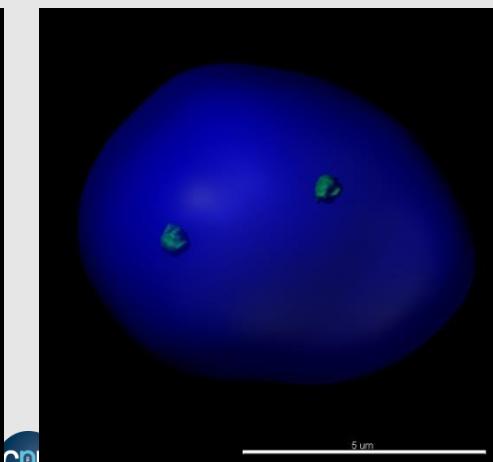
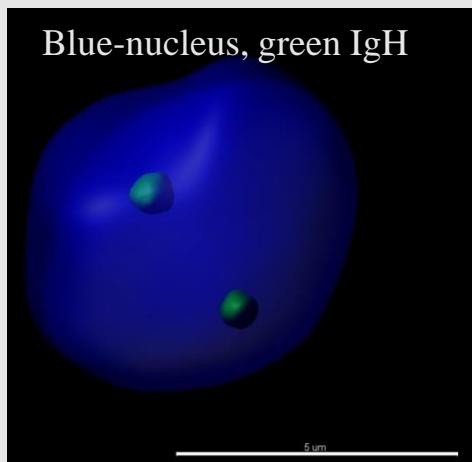
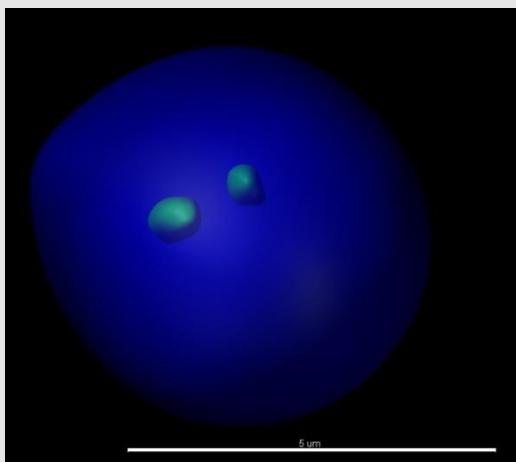
90% of cells undergo apoptosis.  
Remaining live cells undergo CSR.



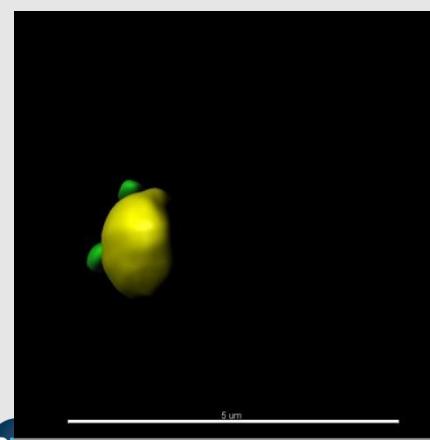
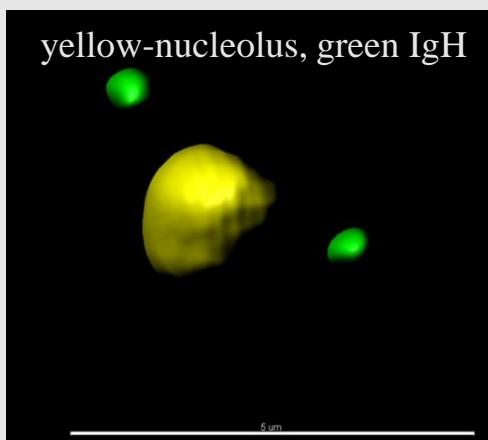
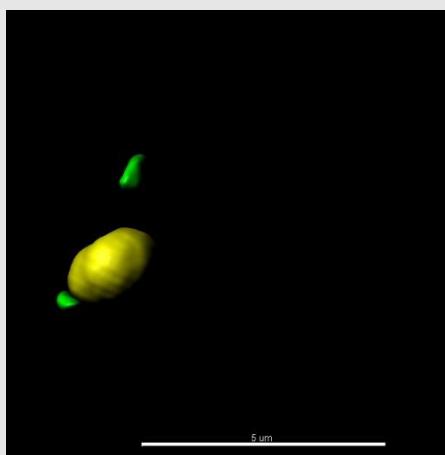
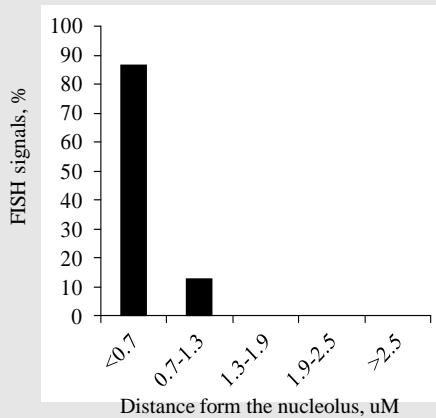
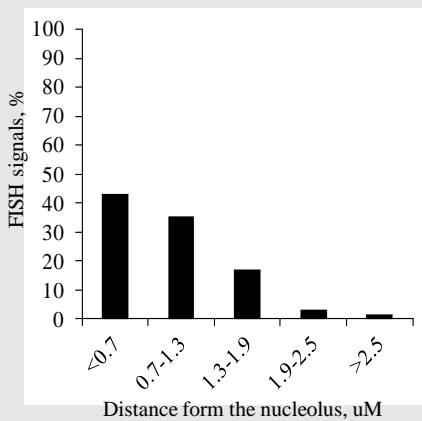
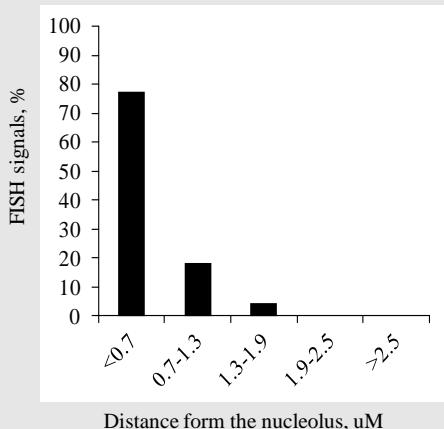
# IgH RADIAL POSITIONING DURING B-LYMPHOCYTE DIFFERENTIATION



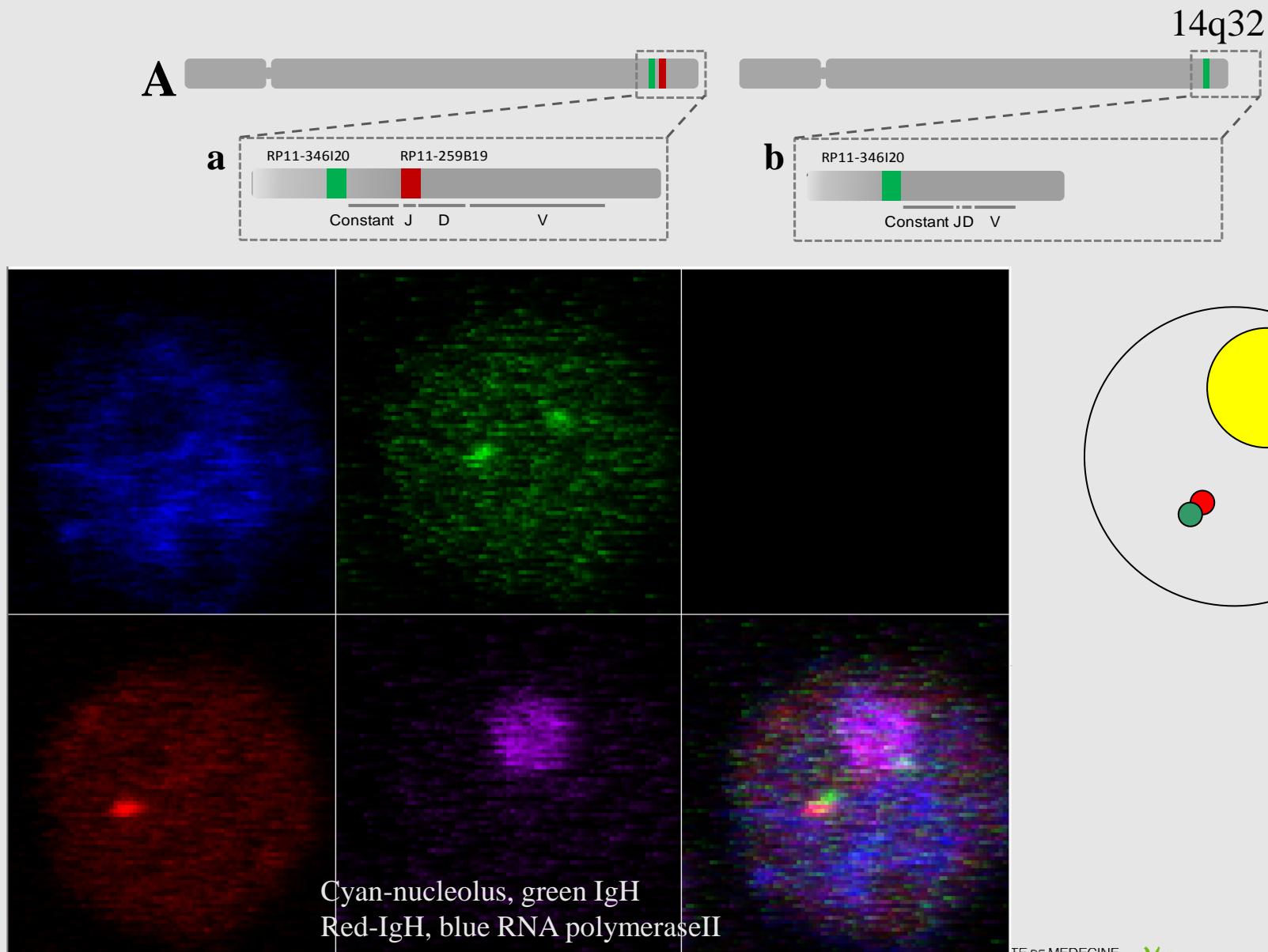
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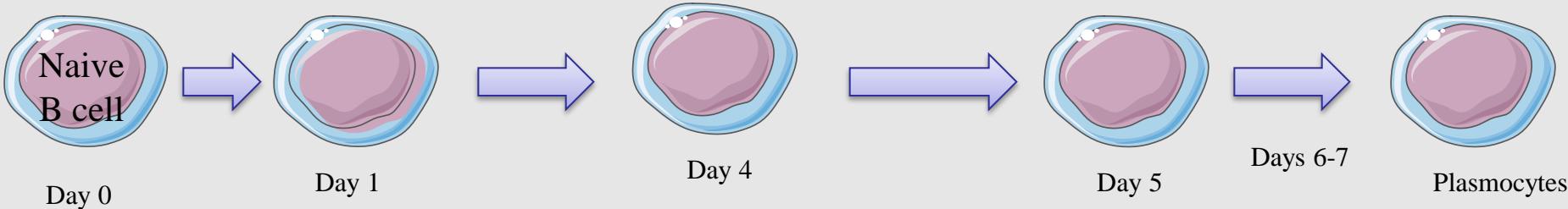
# IgH PROXIMITY TO THE NUCLEOLUS CHANGES B-LYMPHOCYTE DIFFERENTIATION



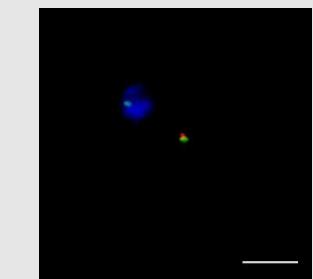
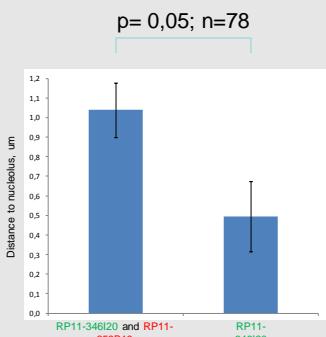
# DETECTING THE PRODUCTIVE ALLELE



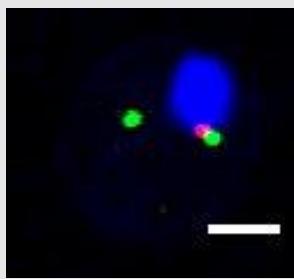
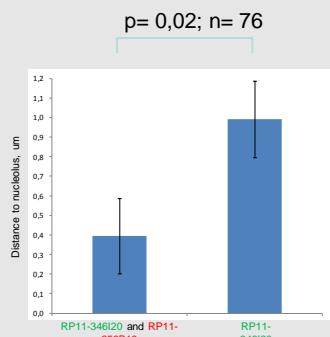
# PRODUCTIVE IgH ALLELE ASSOCIATES WITH THE NUCLEOLUS ON DAY5



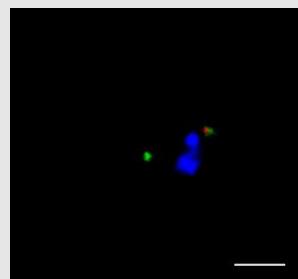
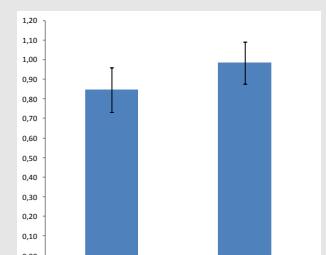
Day 0



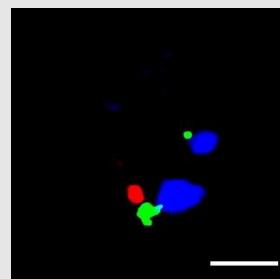
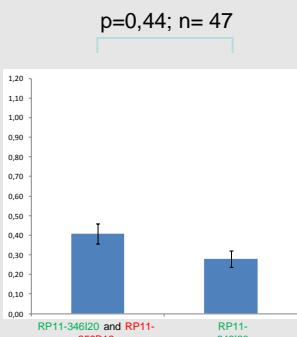
Day 1



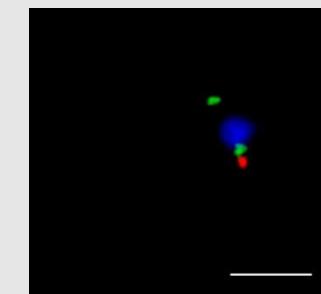
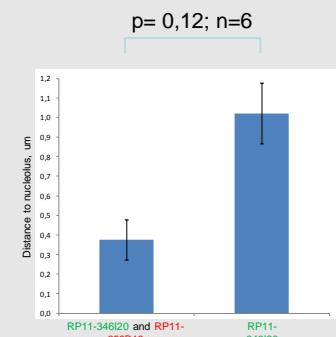
Day 4



Day 5



Plasmocytes



blue-nucleolus, green IgH, red - IgH

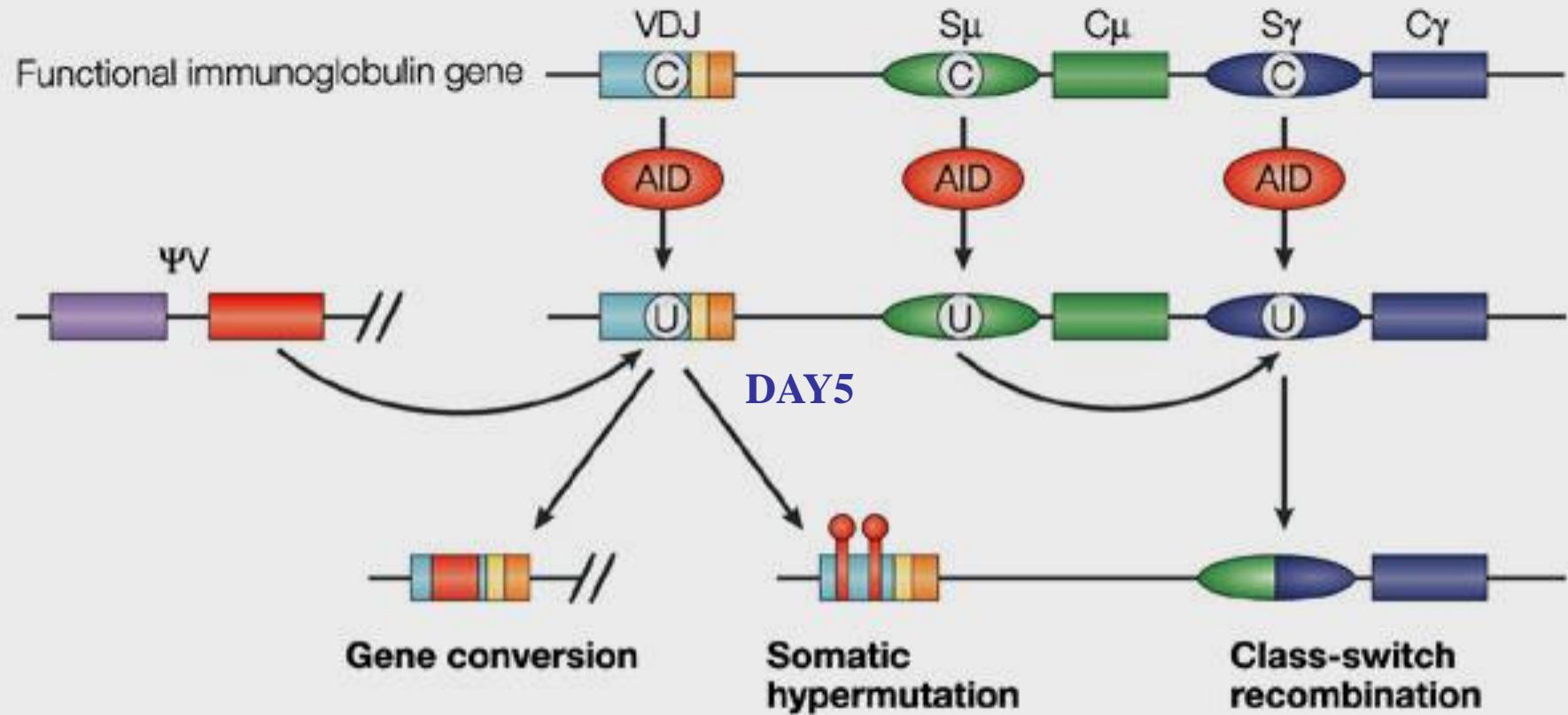


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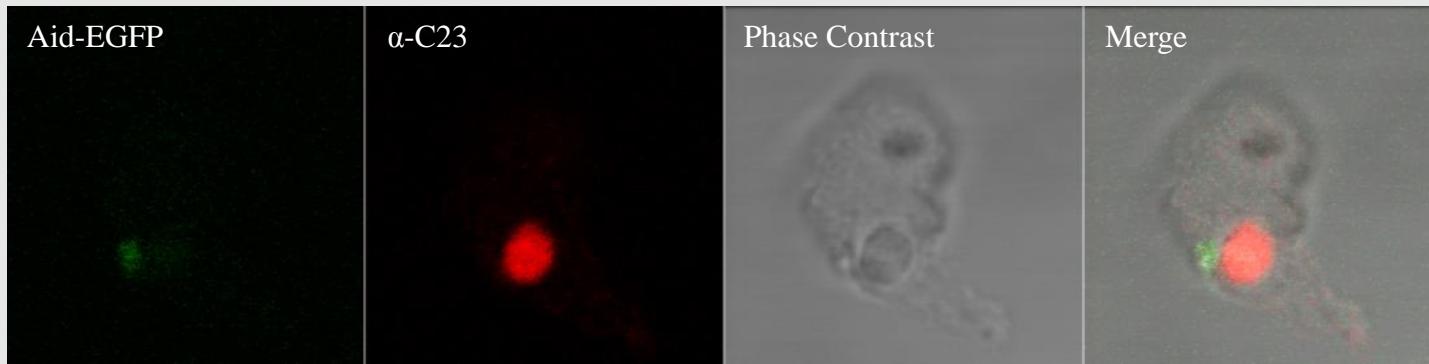
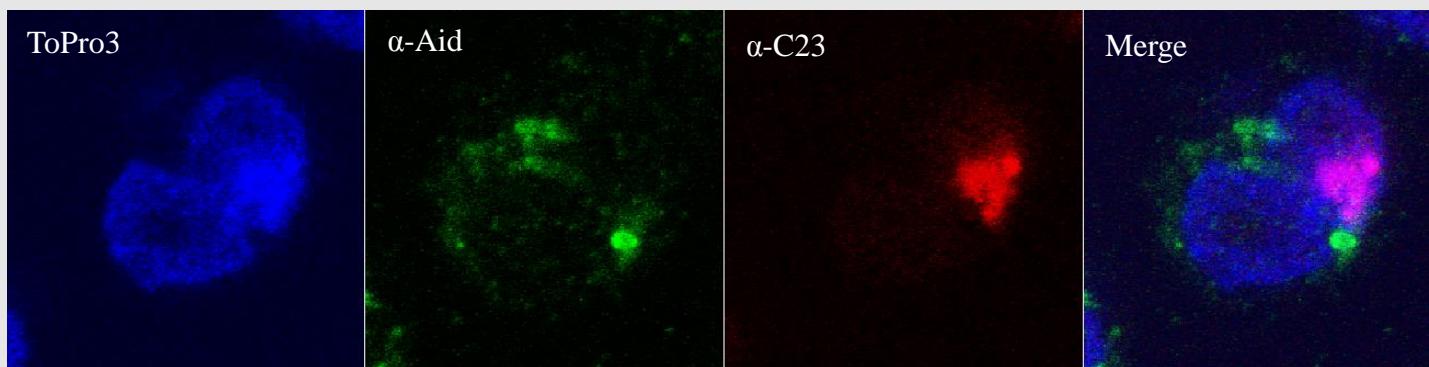
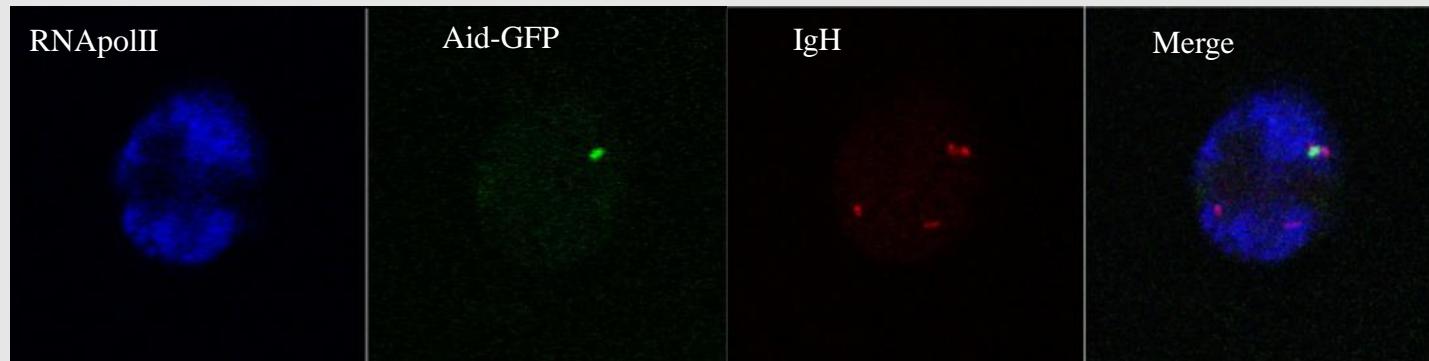
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# DAY5: SOMATIC HYPERMUTATION AND CLASS SWITCH RECOMBINATIION

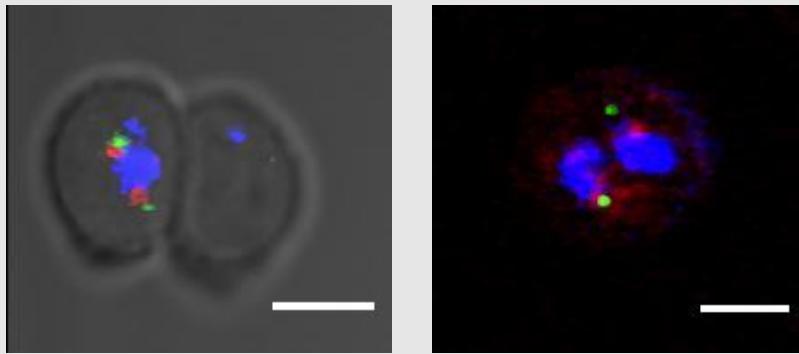


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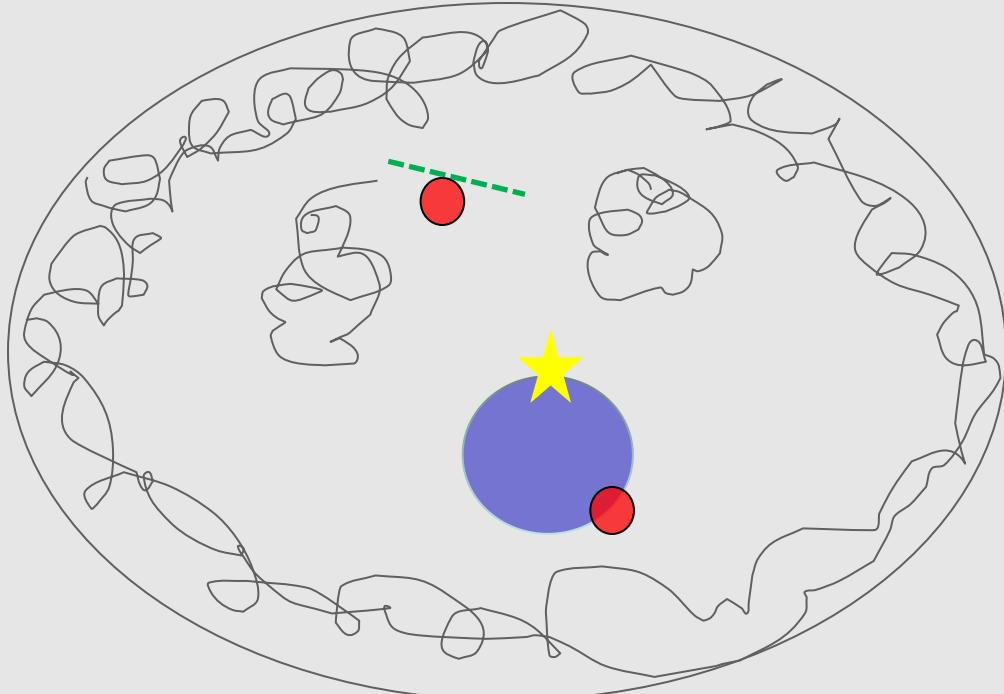
# TRANSCRIPTIONALLY-ACTIVE IgH ALLELE AND AID ARE LOCATED CLOSE TO THE NUCLEOLUS



# THE $\gamma$ H2AX FOCI SPECIFICALLY COLOCALIZE WITH IgH AND THE NUCLEOLUS IN DAY<sub>5</sub> B-LYMPHOCYTES



NUCLEOLUS (Anti-B23)  $\gamma$ H2AX IgH



# Equipe "Chromatine, Développement et Cancer"

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- **Marc Lipinski**, DR CNRS
- **Yegor Vassetzky**, DR CNRS
- **Chrystèle Bilhou-Nabera**, MCU-PH
- **Andrei Pichugin**, postdoctorant
- **Tatiana Tsfasman**, postdoctorante
- **Yara Bou Saada**, doctorante UPS
- **Diana Markozashvili**, doctorante (cotutelle Russie)
- **Ilya Sklyar**, doctorant (cotutelle Russie)
- **Rawan El-Amine**, doctorante (cotutelle Liban)
- **Carla Dib**, M2 UPS



- **Vincent Ribrag**, IGR, Villejuif
- **Sergey Razin**, IBG, Moscow
- **Olga Iarovaya**, IBG, Moscow
- **M.-N. Prioleau**, IJM, Paris
- **S. Bury-Moné**, ENS Cachan

