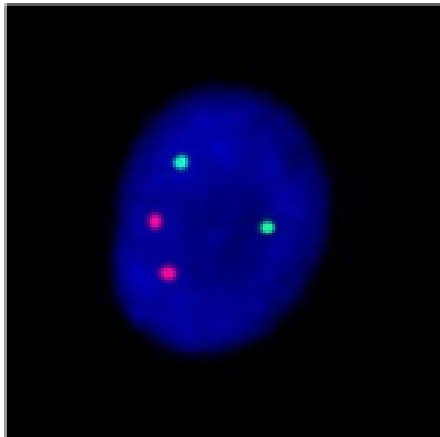


# 다발성 골수종(Multiple Myeloma)의 세포유전 검사

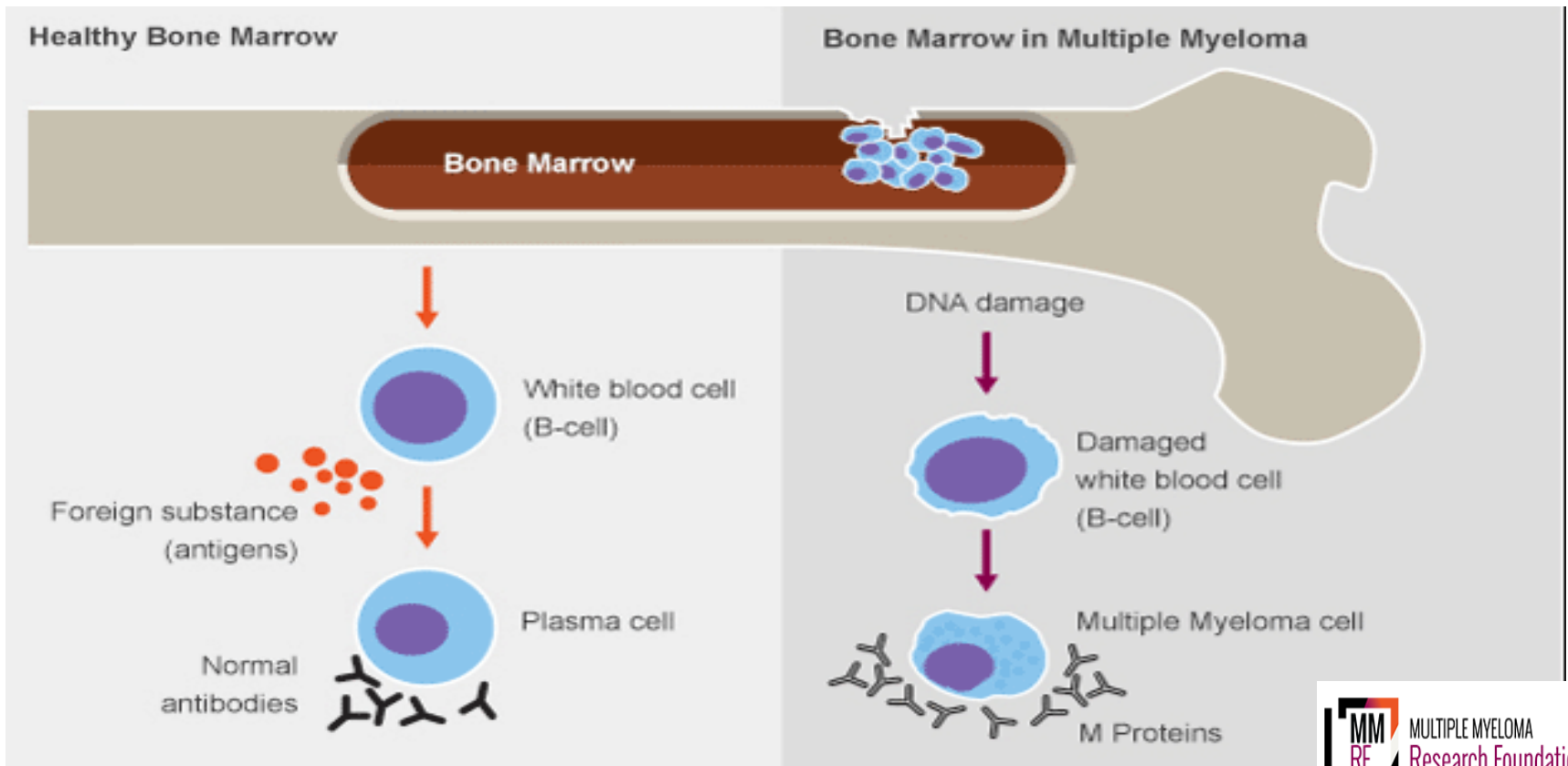


2017.6.14

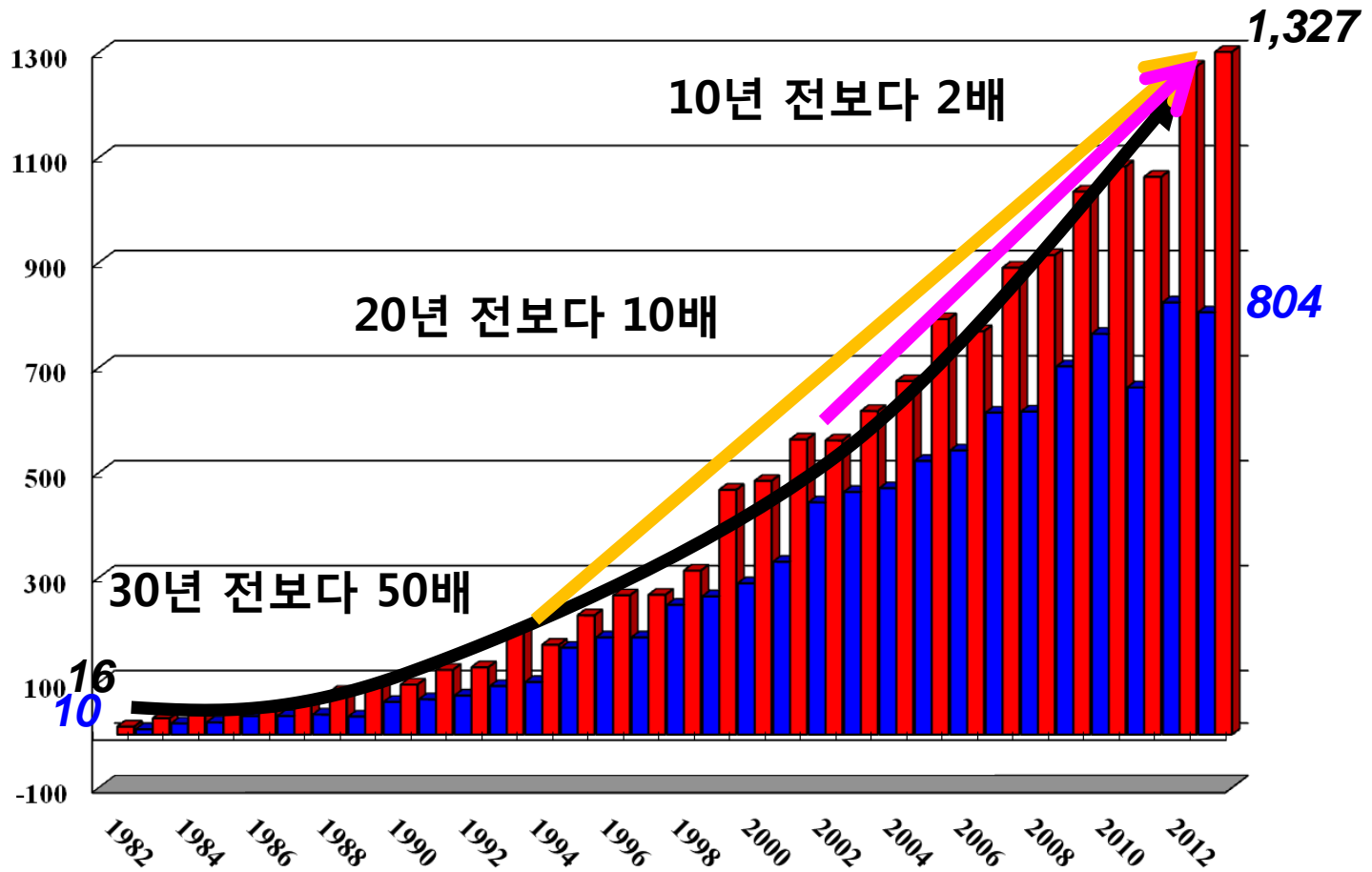
세포유전검사실  
함명희

# Multiple Myeloma?

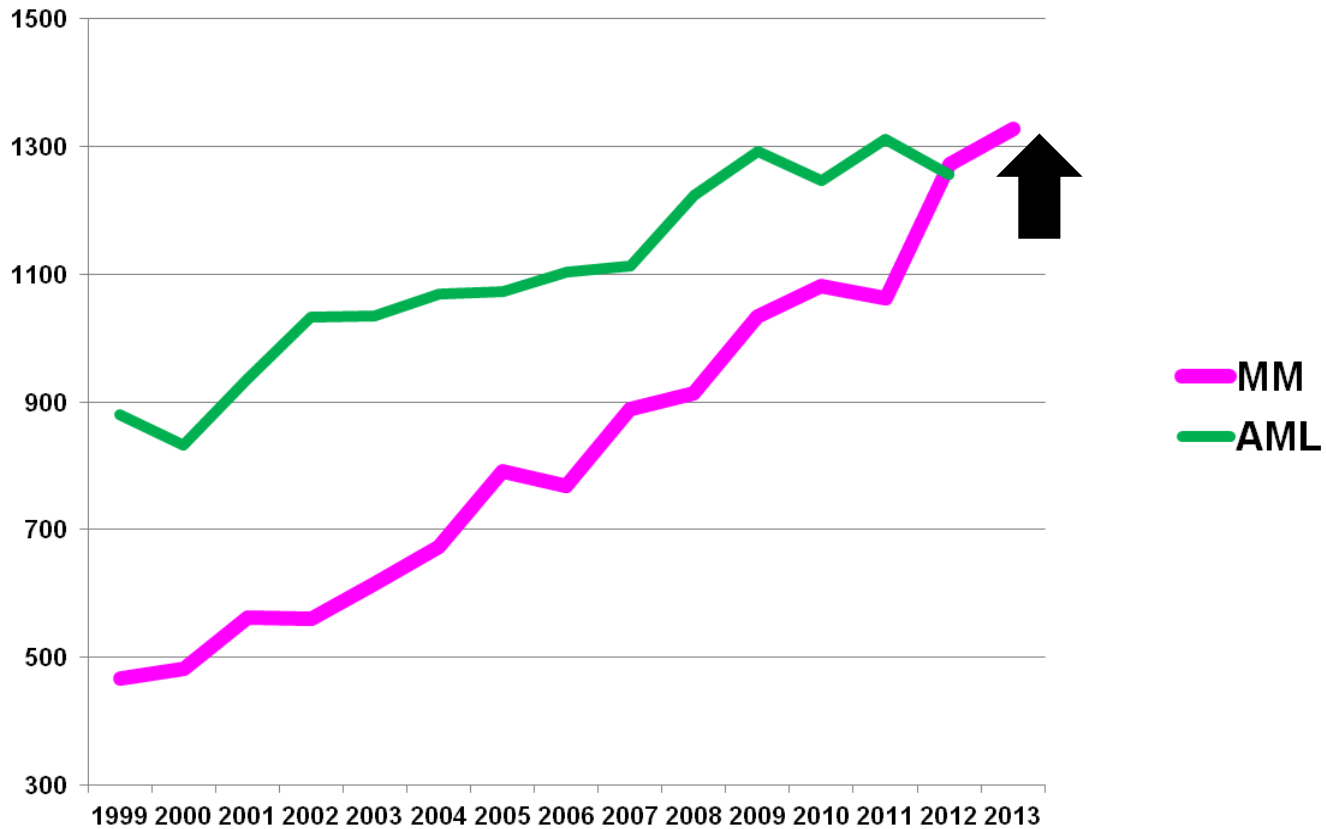
Multiple Myeloma is a malignant neoplasm of plasma cells that accumulate in bone marrow, leading to bone destruction and marrow failure.



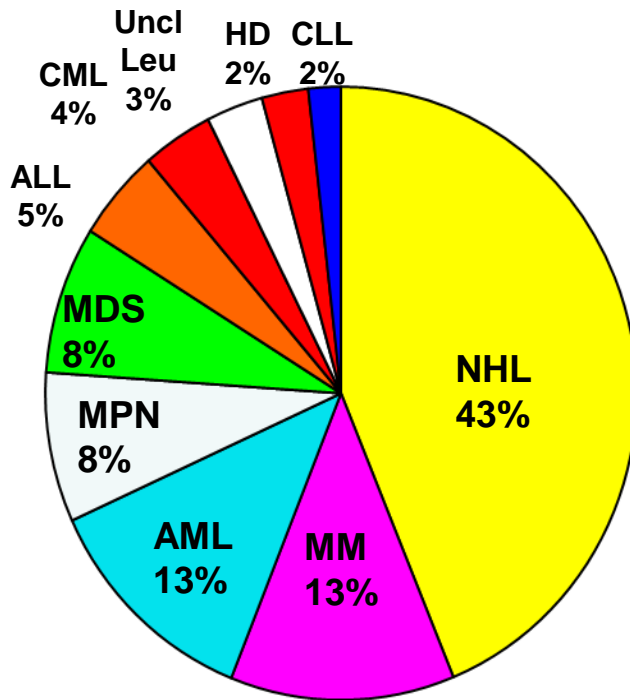
# 한국의 다발골수종 발생률/사망률



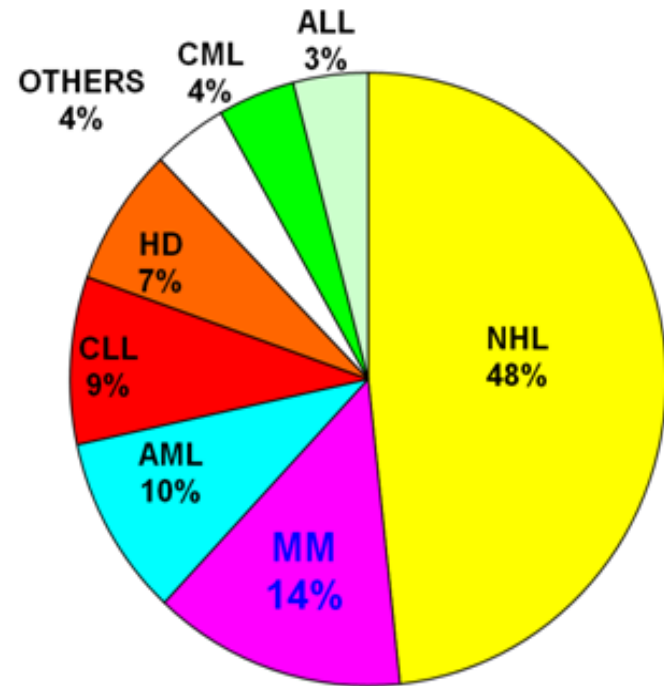
# Crude Incidence of MM and AML recent 10 years in SK



# Relative Incidence of Hematologic Malignancy in SK and US

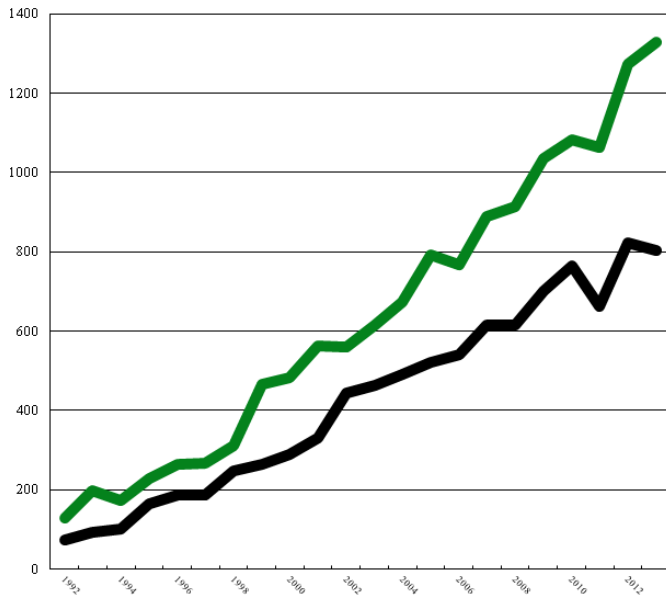


**SK 2012**

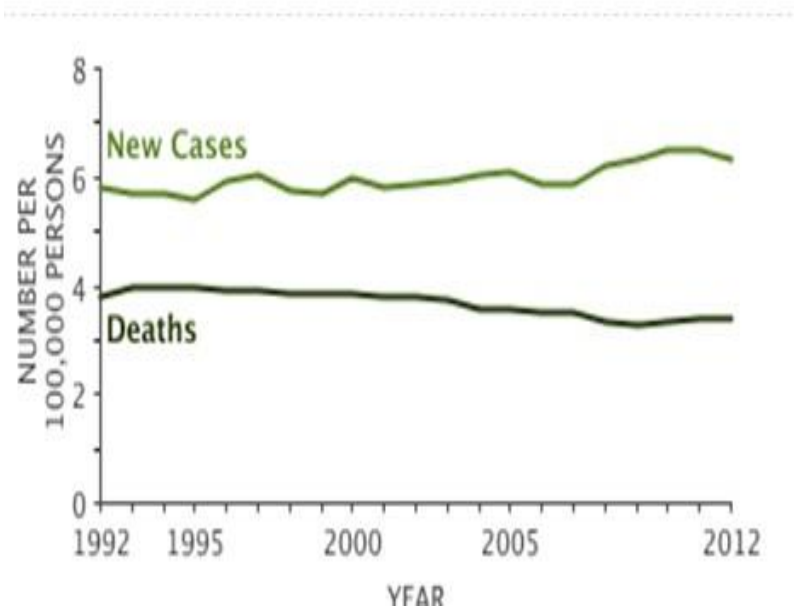


**US SEER 2010**

# 한국의 다발골수종 발생율/사망률 1992-2013



한국



미국

# Signs and symptoms



- **C**alcium elevated
- **R**enal failure
- **A**nemia
- **B**one pain
- **I**nfection

다발성 골수종의 일반적 증상

▶ 뼈의 통증과 고칼슘혈증

▶ 신부전

▶ 활력계 이상증상 (빈혈, 출혈성경향)

▶ 감염(폐렴, 요로감염)

보건복지부 국립암센터 대한식약처

# Diagnostic criteria

## NAME

Monoclonal Gammopathy of Undetermined Significance (MGUS)

Asymptomatic or Smoldering Multiple Myeloma (SMM)

Active or Symptomatic Myeloma

## DEFINITION

- Monoclonal protein present but usually < 3.0 g/dL
  - No CRAB features or other indicators of active myeloma
  - Bone marrow monoclonal plasma cells < 10%
- 
- Higher level of disease than MGUS: serum M-component can be > 3.0 g/dL and/or bone marrow plasma cells > 10%, but
  - No CRAB features or other indicators of active myeloma
- 
- Monoclonal protein present, and
  - One or more "CRAB" features and/or indicators of organ damage\*

\*Organ damage classified as "CRAB" or any other significant clinical problem linked to myeloma progression such as recurrent infections or neuropathy unrelated to treatment

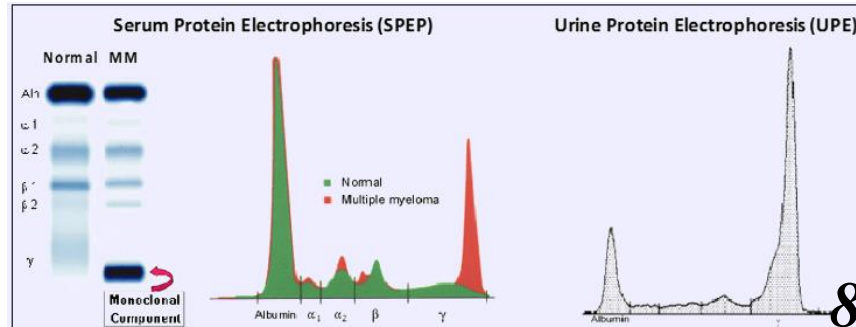
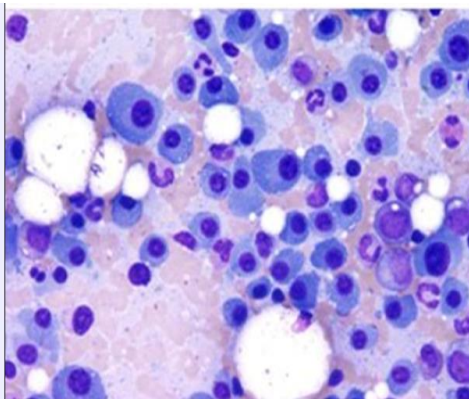
**C** - Calcium elevation (>10mg/dL)

**R** - renal dysfunction (creatinine >2mg/dl or creatinine clearance (<40ml/min)

**A** - anemia (hemoglobin <10g/dL or >2g/dL decrease from patient's normal)

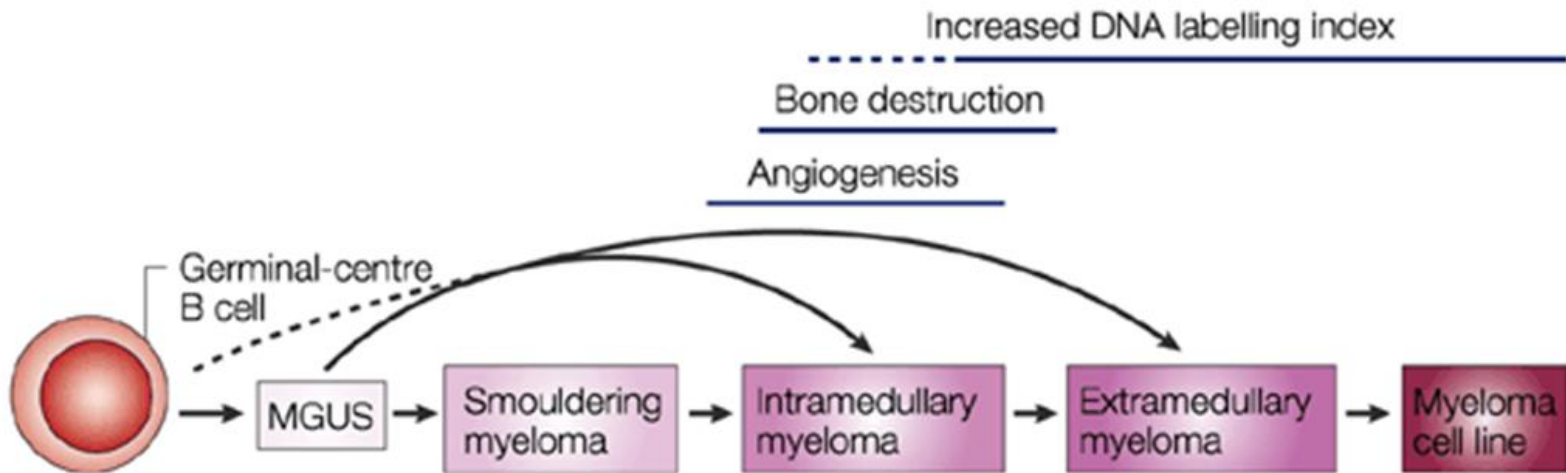
**B** - bone disease (one or more osteolytic lesions detected on skeletal radiography, WBLC CT or PET/CT)

One or more "CRAB" features or other significant problem required for diagnosis of **Symptomatic Myeloma**





# Pathogenesis



**a** Karyotypic instability

**b** Primary Ig translocations

**c** Secondary (Ig) translocations (*c-MYC*, others)

**d** 13q14 deletion/monosomy

**e** Activating mutations  
*NRAS*, *KRAS*  
*FGFR3*

**f** *TP53* mutations

# Diagnostic workup

CBC, diff-count/ blood cell morphology,  
BUN/creatinine / albumin/ electrolyte with calcium,  
Serum light chain assay/Serum quantitative Ig,  
Serum LDH/  $\beta$ 2-microglobulin

Bone marrow aspirate and  
biopsy

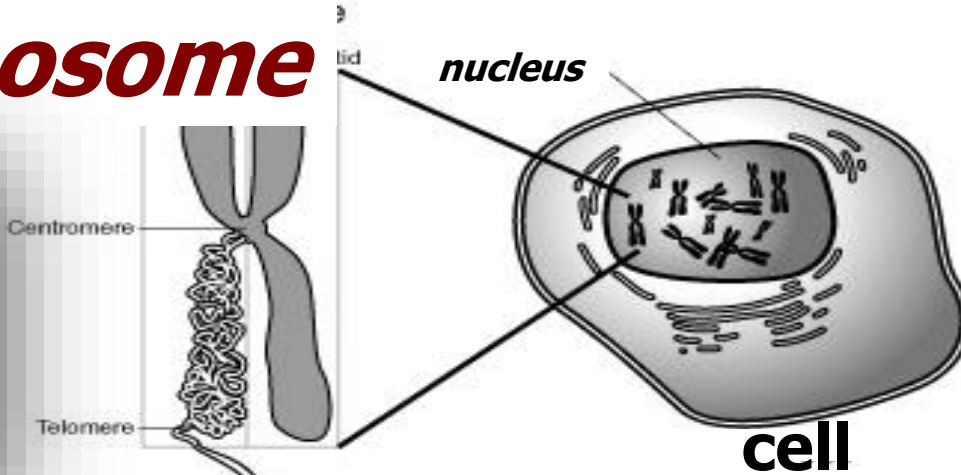
Immunohistochemistry(BM bx.) CD138, kappa/lambda

Immunophenotyping CD19,CD38,CD45,CD56,CD138

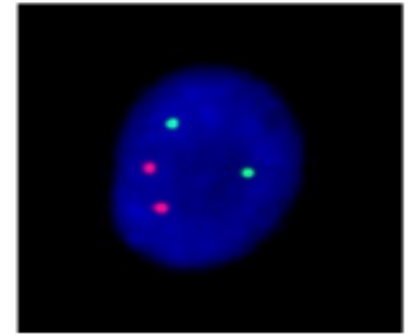
**Cytogenetic analysis**

**FISH** 1q gain, CDKN2A deletion,  
RB1 deletion, p53, IGH, t(4;14),  
t(14;16)

# ***Chromosome***



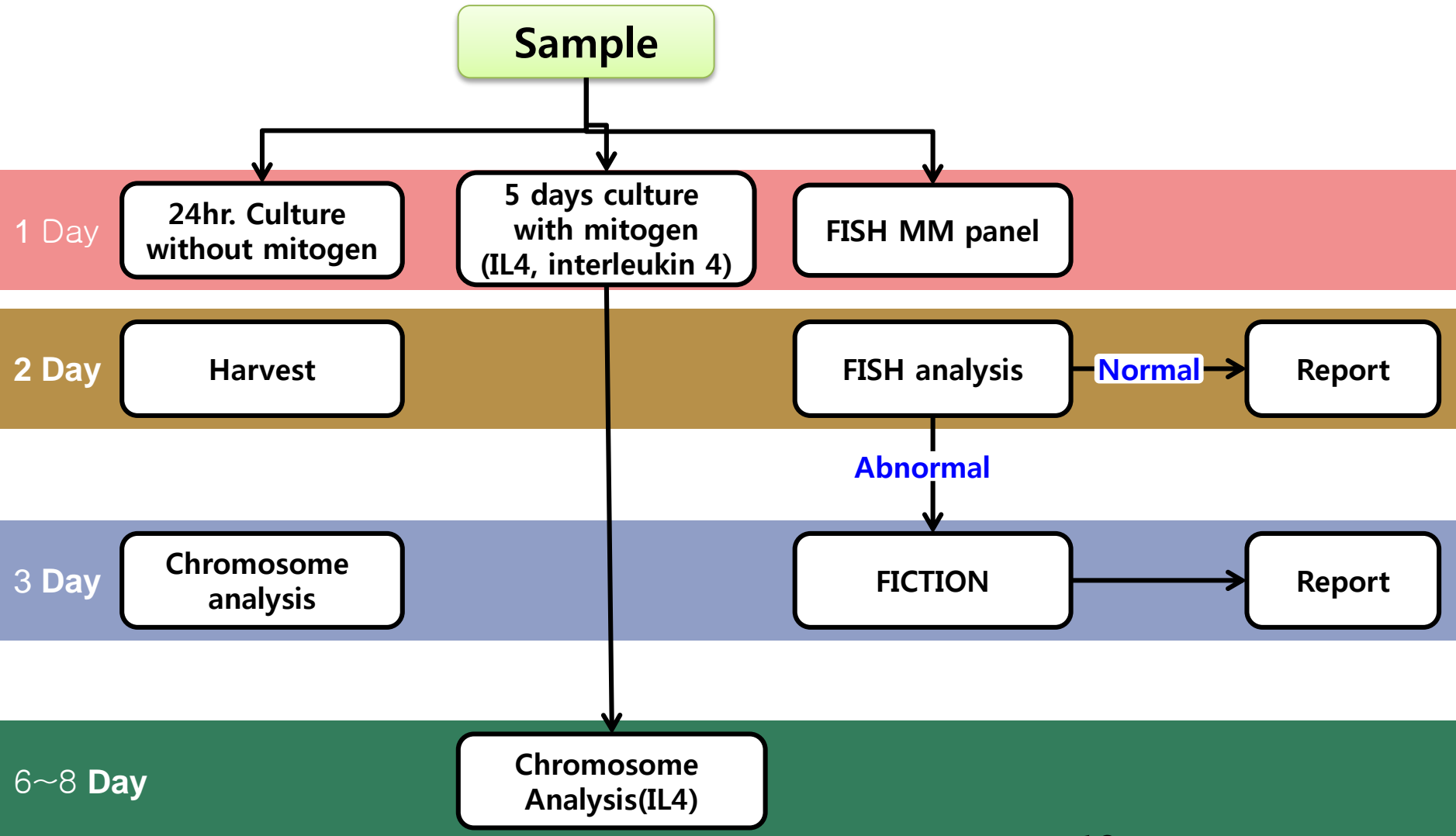
# ***FISH***



# ***DNA study***



# Work flow

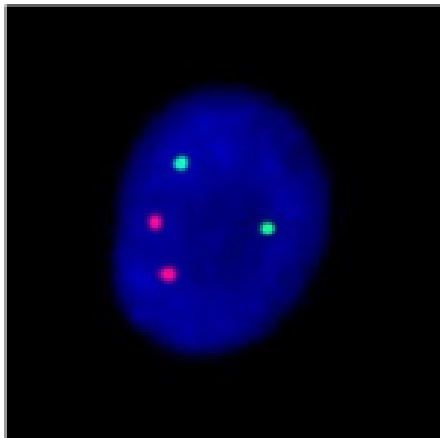




# of cytogenetics & FISH

<b>Chromosome analysis</b>	Long culture period(2~7 day)
	Low proliferation rate of plasma cells in vitro
	Abnormal clone is not detected until disease has reached an advanced stage
	Poor chromosome banding quality
	Sensitivity N/20 metaphase
<b>FISH</b>	Circumvents the need for cell division(1~2 day)
	Abnormalities by artifact
	Available for the detection of submicroscopic alterations
	for the clarification of complex alterations and also for MRD purposes
	Sensitivity N/200~300 interphase

# Complementary G-banding vs. FISH



**iFISH**



**G-banding**

# Chromosome aberrations in MM

- Marked karyotypic instability (significant molecular heterogeneity )

Nonhyperdiploidy	~half	
Recurrent translocations with IGH gene(14q32)	11q13(CCND1)	16%
	4p16(FGFR3)	15%
	16q23(MAF)	5%
	6p21(CCND3)	3%
	20q12(MAFB)	2%

Hyperdiploidy	~remaining half	
Multiple trisomies usually do not have coexistent 14q32 translocations.	Odd no. chromosome 3, 5, 7, 9, 11, 15, 19, 21	

# Chromosome aberrations in MM

## Hypodiploidy (<46)

- Loss of 13, specifically 13q14.3 and/or
- Loss of 17, specifically 17p13.1(*TP53*)
- Includes structural abnormalities chr. 1,4,6,14,16,20
- Loss of 8, 17, Y, X
- 1p loss, 1q gain, 4q loss, 6q loss, 20 loss
- Rearrangement 14q32, 16q
- 70~90 chromosome, double content
- Adverse prognosis



# Chromosome aberrations in MM

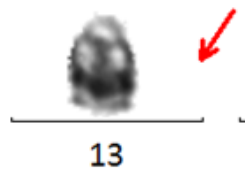
## Hyperdiploidy

- Gain of 3, 5, 7, 9, 11, 15, 19, 21
- standard risk category as long as no deletion 13q or 17p
- the most common translocation: t(11;14)

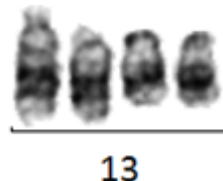
# Chromosome aberrations in MM

## Deletion of 13q/loss of 13

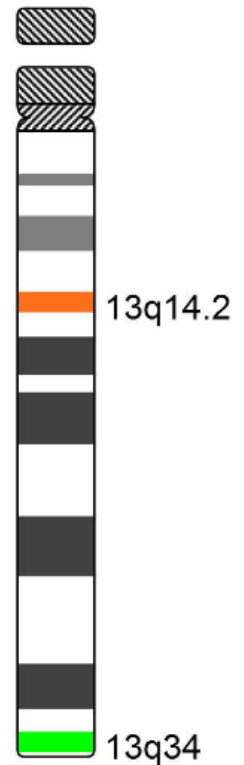
- Most frequent (~50% of abnormal karyotype)
- Interstitial deletion of 13q involving 13q14.2(RB1) or 13q14.3(D13S319); **cryptic**, only FISH
- Deletion with conventional cytogenetics(10~20%)
- Poor outcome, high relapse rate



**-13**



**del(13)(q14q22)**





ELSEVIER

Cancer Genetics and Cytogenetics 168 (2006) 124–132

CANCER GENETICS  
AND  
CYTOGENETICS

## Identification of 13q deletion, trisomy 1q, and IgH rearrangement as the most frequent chromosomal changes found in Korean patients with multiple myeloma

Soo-Mee Bang<sup>a,1,2</sup>, Young Ree Kim<sup>b,1</sup>, Han Ik Cho<sup>c</sup>, Hyun Sook Chi<sup>d</sup>, Eul-Ju Seo<sup>d</sup>,  
Chan Jeoung Park<sup>d</sup>, Soo Jin Yoo<sup>d</sup>, Hee Chan Kim<sup>e</sup>, Hong Gu Chun<sup>e</sup>, Hyun Chung Min<sup>f,g</sup>,  
Bo Ra Oh<sup>f,g</sup>, Tae Young Kim<sup>f,g</sup>, Jae Hoon Lee<sup>a</sup>, Dong Soon Lee<sup>c,f,g,\*</sup>

<sup>a</sup>Department of Internal Medicine, Gachon Medical School, Gil Medical Center, 1198 Guwol-dong, Namdong-gu, Incheon 405-760, Korea

<sup>b</sup>Department of Laboratory Medicine, Jeju National University College of Medicine, 66 Jeju-daehakro, Jeju 690-756, Korea

<sup>c</sup>Department of Laboratory Medicine, Seoul National University College of Medicine, 28 Yeongeon-dong, Jongno-gu, Seoul 110-744, Korea

<sup>d</sup>Department of Laboratory Medicine, College of Medicine, University of Ulsan and Asan Medical Center,  
388-1 Pungnab2-dong, Songpa-gu, Seoul 138-736, Korea

<sup>e</sup>Department of Biomedical Engineering, <sup>f</sup>Cancer Research Institute, Seoul National University College of Medicine, Seoul,  
28 Yeongeon-dong, Jongno-gu, Seoul 110-744, Korea

<sup>g</sup>National Research Laboratory for Molecular Cell Imaging, Seoul National University College of Medicine,  
28 Yeongeon-dong, Jongno-gu, Seoul 110-744, Korea

Received 19 October 2005; received in revised form 6 February 2006; accepted 15 February 2006

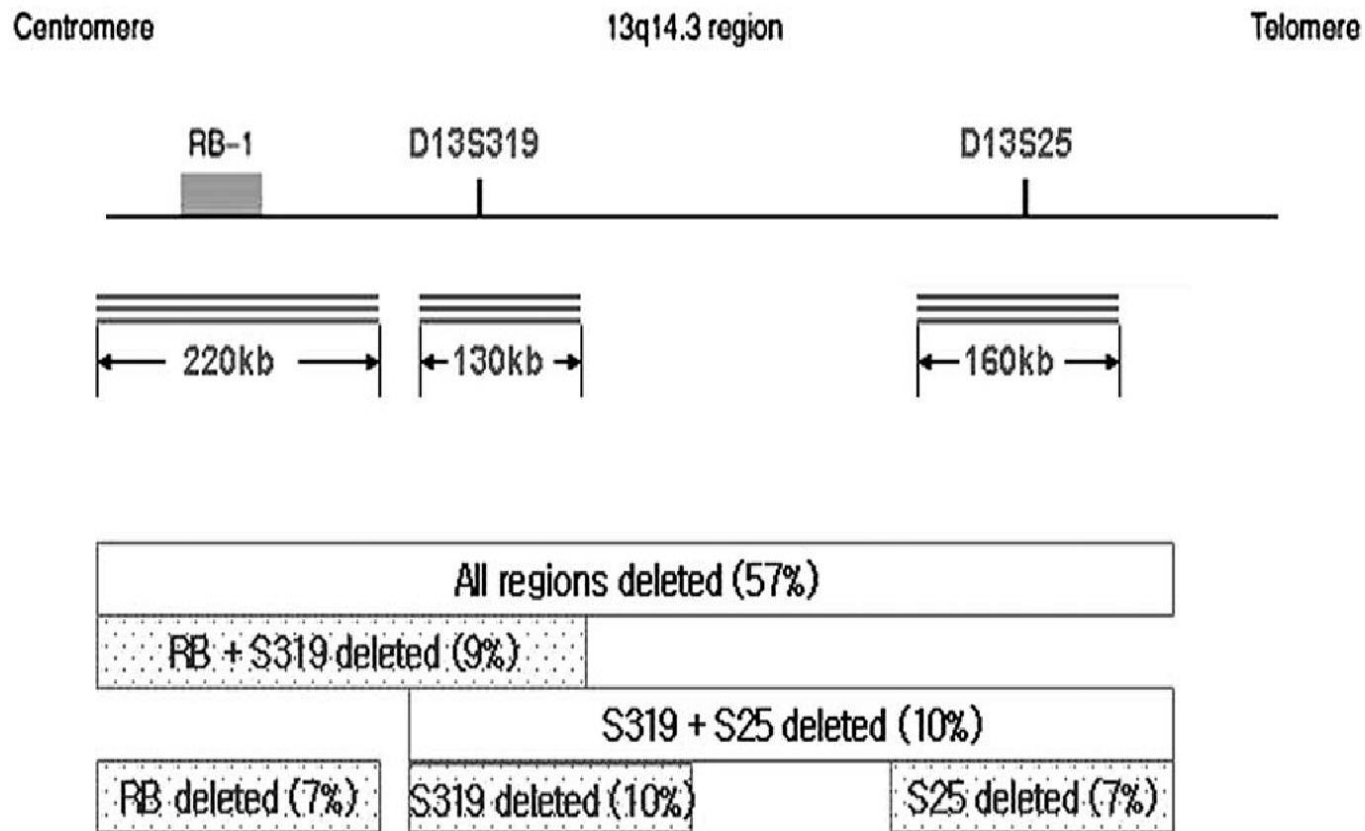
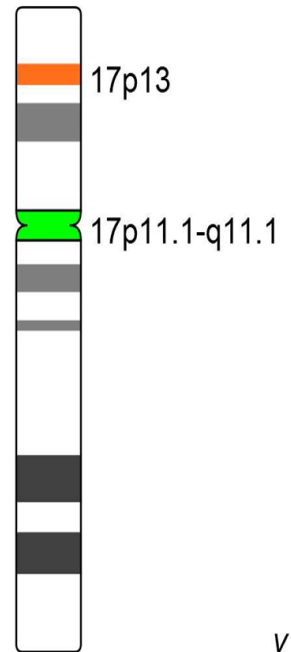


Fig. 2. The sensitivity of three different probes for 13q deletion. A total of 58 patients had at least one locus deleted on 13q. Among them, 33 patients (57%) proved to be deleted at all three loci. Six patients lost D13S25 plus D13S319 (10%), and five patients lost *RB* plus D13S319 (9%). Six patients lost only D13S319 (10%), and four patients each lost *RB* (7%) or D13S25 (7%).

# Chromosome aberrations in MM

## Deletion of 17p

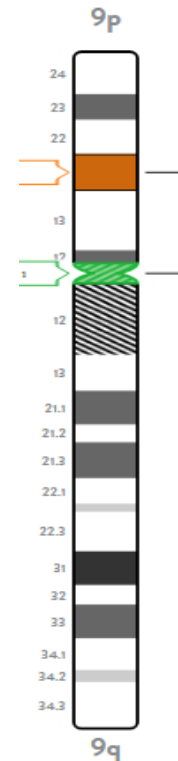
- Deletion of TP53 (~10%)
- Tumor suppressor gene
- Adverse prognostic outcome
- Occur as secondary events during disease progression.



# Chromosome aberrations in MM

## Deletion of CDKN2A

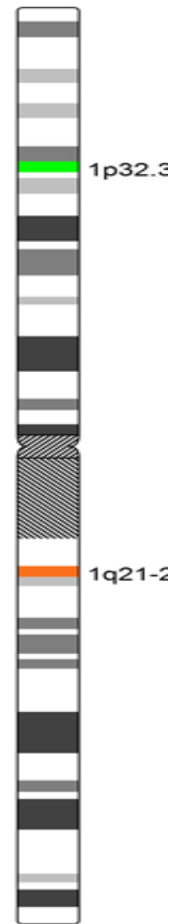
- Deletions of 9p13 or methylation
- If cryptic deletion, only FISH
- Tumor suppressor gene



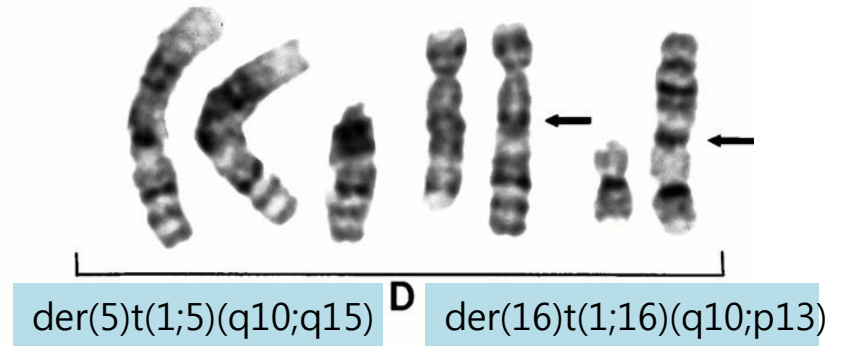
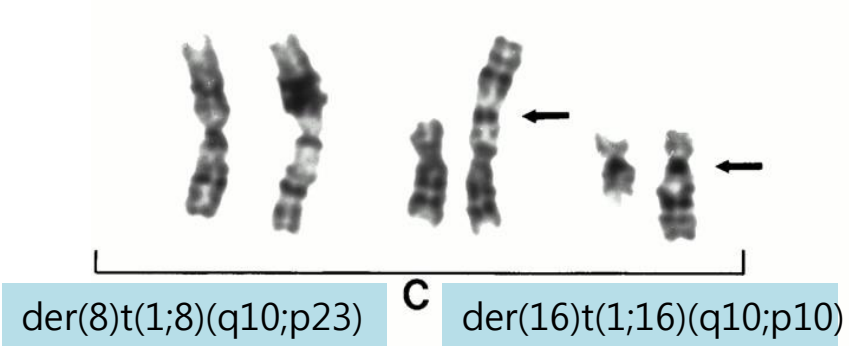
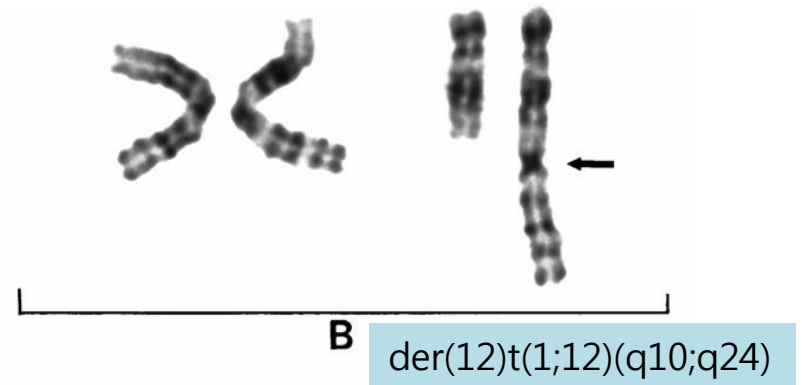
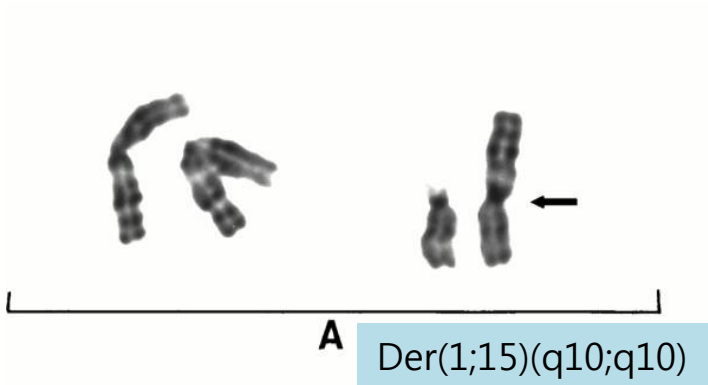
# Chromosome aberrations in MM

## Chromosome 1 Abnormalities

- Deletions of 1p: 1p12~1p31
- Gains of 1q: 1q21,  
the second most frequent(40%~70%)
- Translocations involving either arm:  
t(1;15)(q10;q10), der(1;16)(q10;p10),  
der(1;19)(q10;p10), i(1)(q10)
- Unfavorable prognosis



# Chromosome 1 Abnormalities

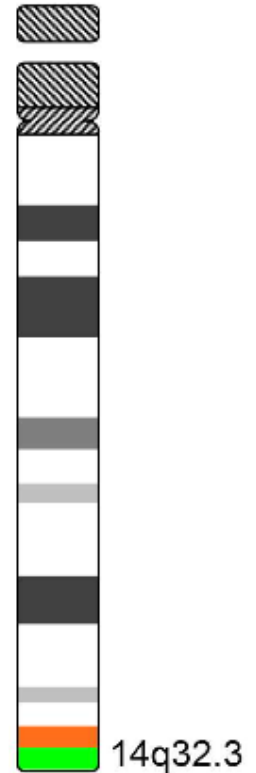




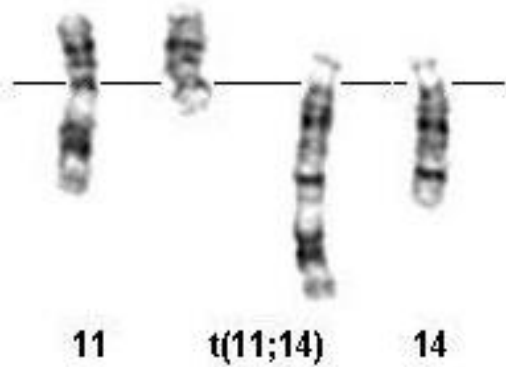
# Chromosome aberrations in MM

## IGH @ Rearrangement

- ~40% of patients with MM.
- Translocation of oncogenes into this region may lead to their increased expression, contributing to disease initiation or disease progression and therapeutic resistance.
- t(11;14): 20~25%, transforms to an aggressive phenotype after acquiring a secondary genetic "hit," CDKN2A inactivation by promoter methylation.
- t(4;14): 15%, cryptic, tend to be very frequent in hypodiploid karyotype, high-risk prognostic category
- t(14;16): 5~7%, cryptic, tend to be very frequent in hypodiploid karyotype



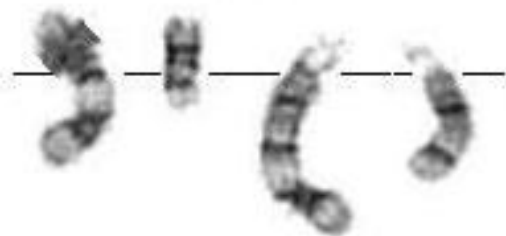
# IGH @ Rearrangement



**t(11;14)**



**t(14;16)**



**t(6;14)**

# Cytogenetic prognostic group in M.M

## Unfavorable risk

t(4;14) or t(14;16)

Deletion 17p13 (p53)

13q deletion or aneuploidy

Hypodiploidy

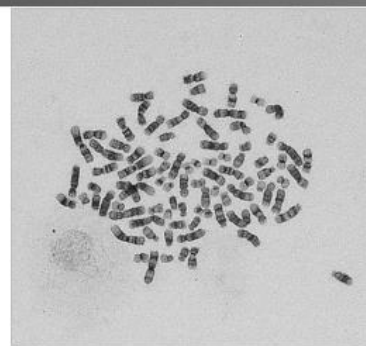
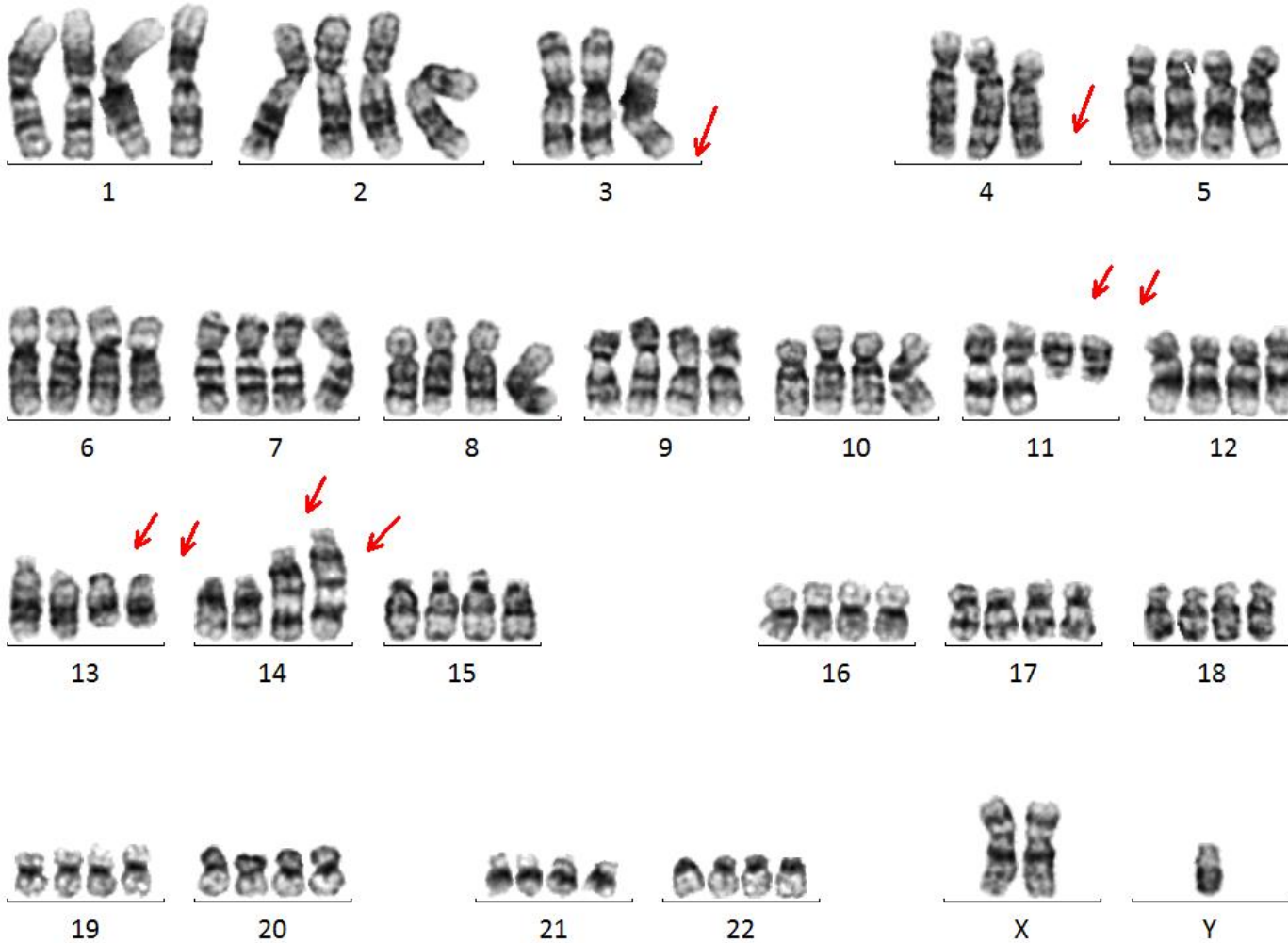
complex abnormalities

## Favorable risk

t(11;14)

Absence of unfavorable risk genetics

Hyperdiploidy



Assign

Rotate 180°/90°

Rotate X

Shift

Clean

Reduce

Magnify

Staining

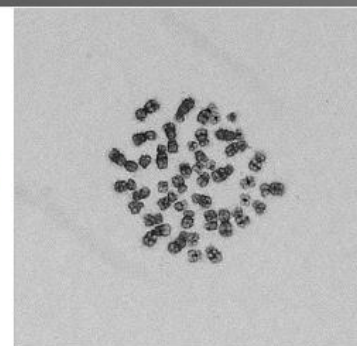
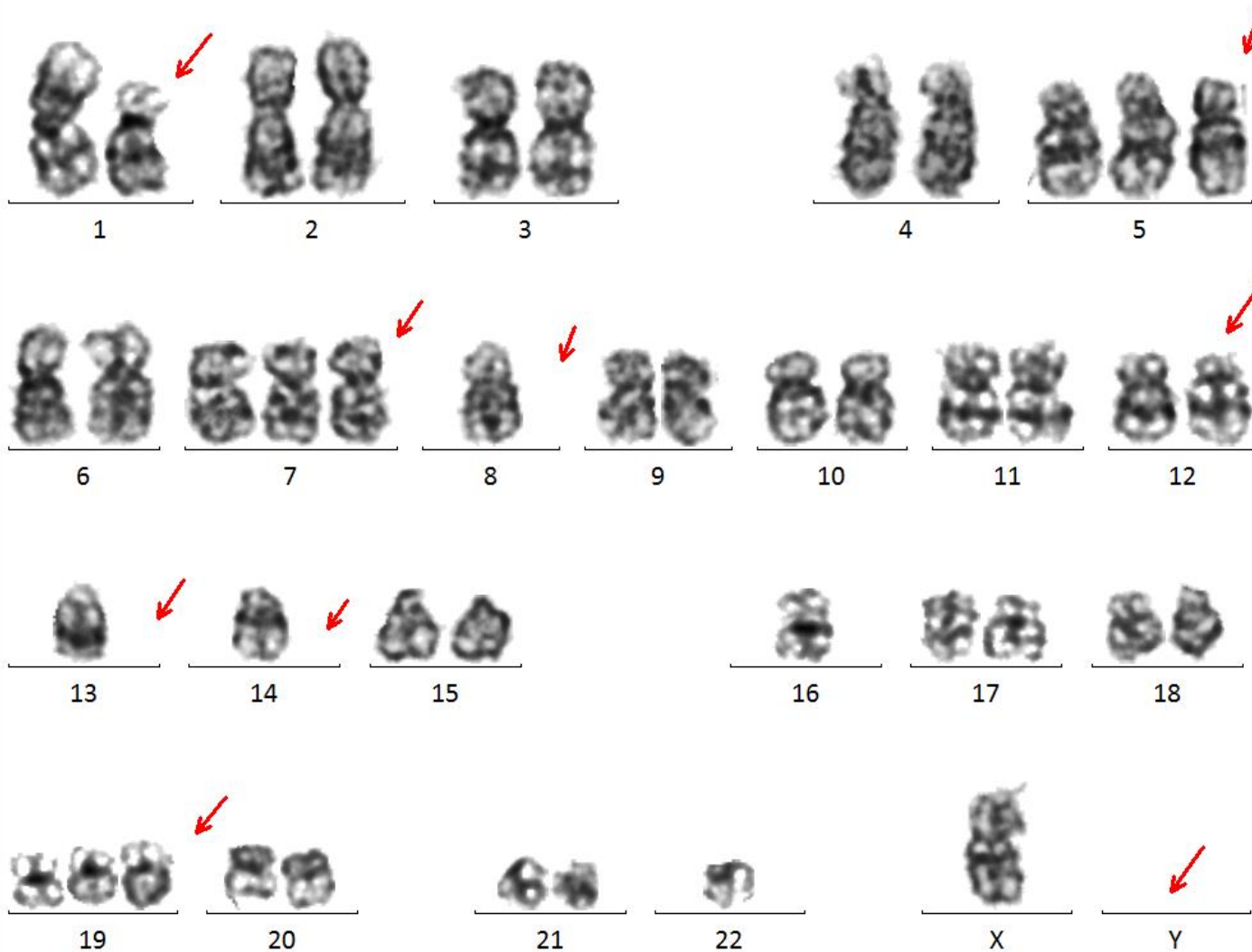
Annotate

DM17-1182 ◀ 234 ▶ ◀ A ▶ 89<4n>,XXY,-Y,-3,-4,t(11;14)(q13;q32)x2,del(13)(q14q22)x2 89

i142~A -6793/19635 CID:262

MSD2016 1705

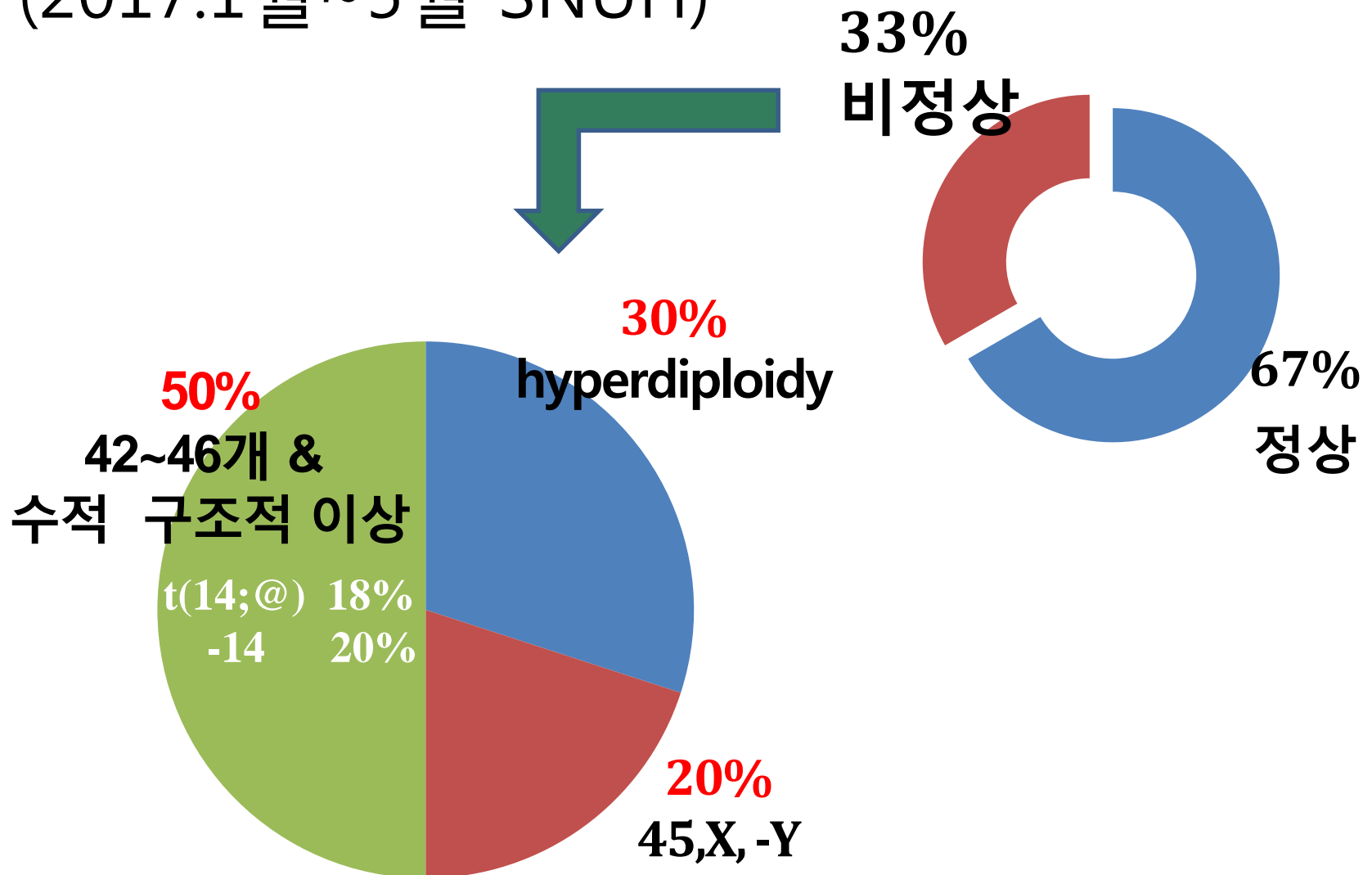
Adm GBand



- Assign
- Rotate 180°/ 90°
- Rotate X
- Shift
- Clean
- Reduce
- Magnify
- Staining

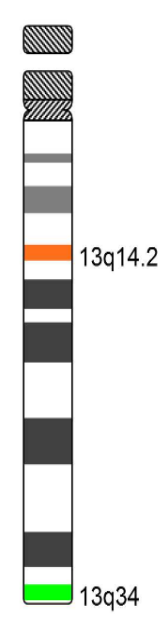
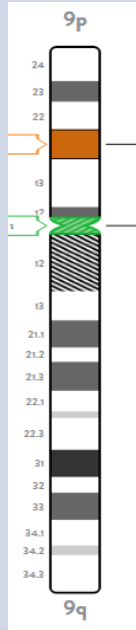
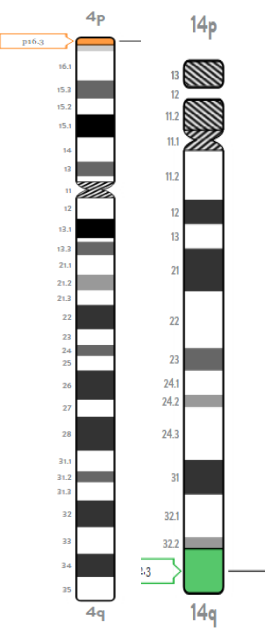
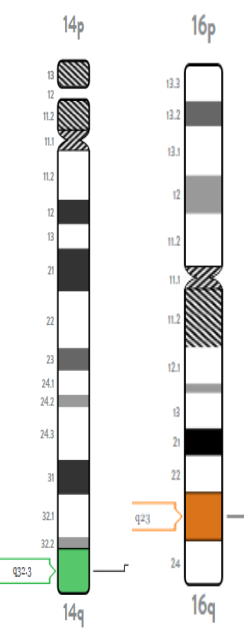
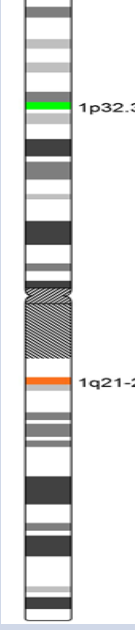
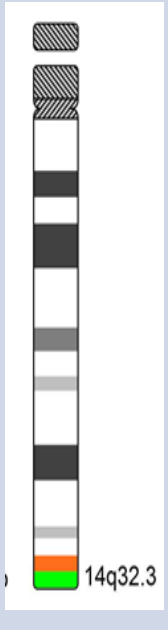
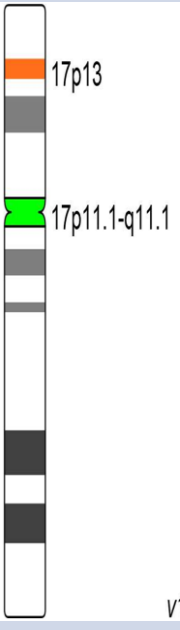
Annotate

# M.M 염색체검사 양성률 (2017.1월~5월 SNUH)

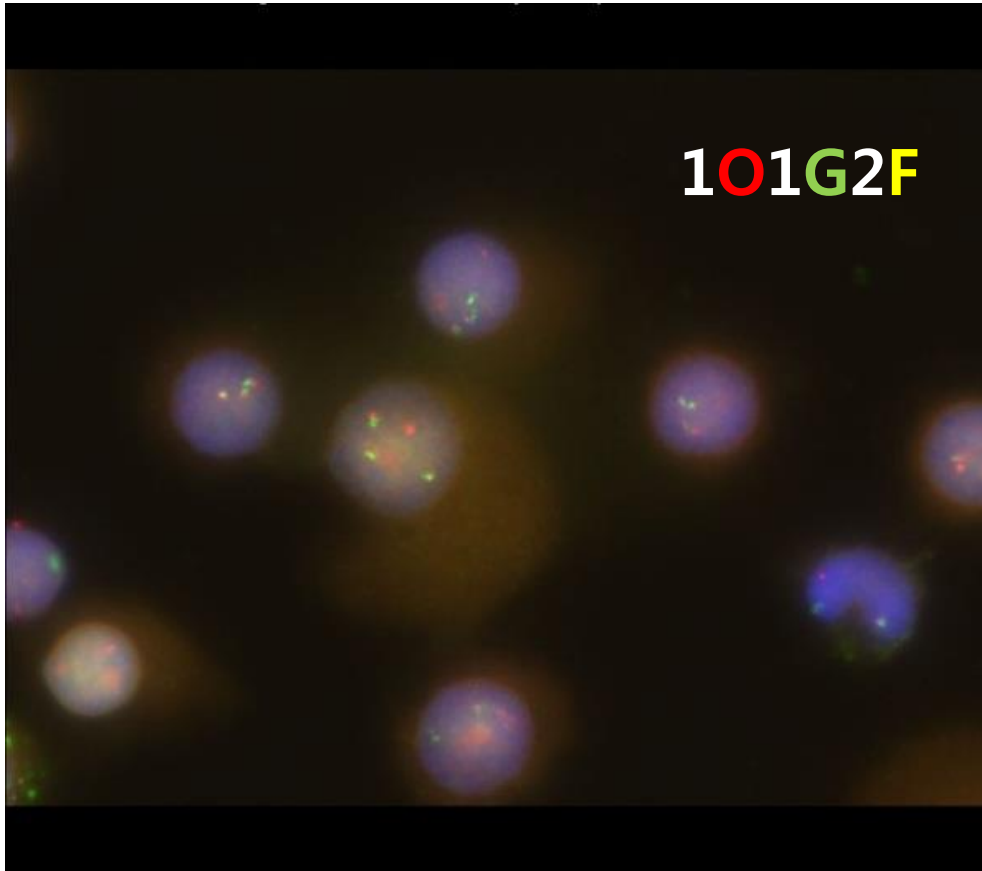


# M.M panel-A(Diagnosis); 7 item

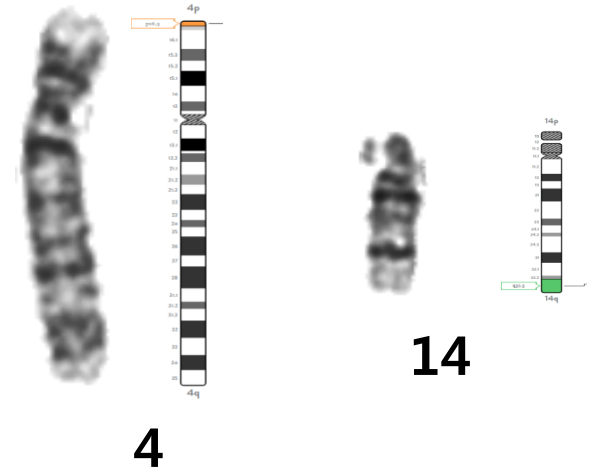
## M.M panel-B(IMWG risk); 4 item

PROBE	D13S319 A B	CDKN2A A	T(4;14) A B	T(14;16) A B	1P/1Q A	IGH A	P53 A B
Locus & color							
Vendor	Meta systems	Abbott	Abbott	Abbott	Meta systems	Meta systems	Meta systems

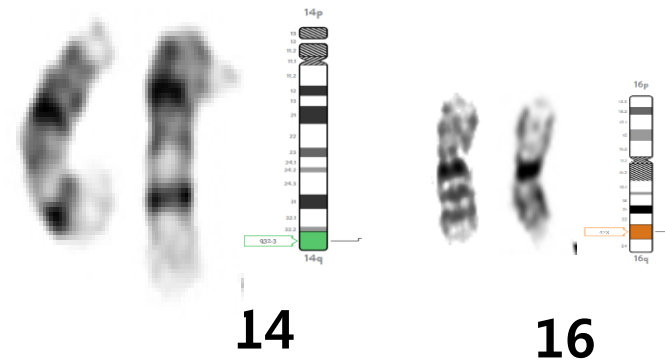
# Cryptic translocation



**t(4;14)(p16;q32)**

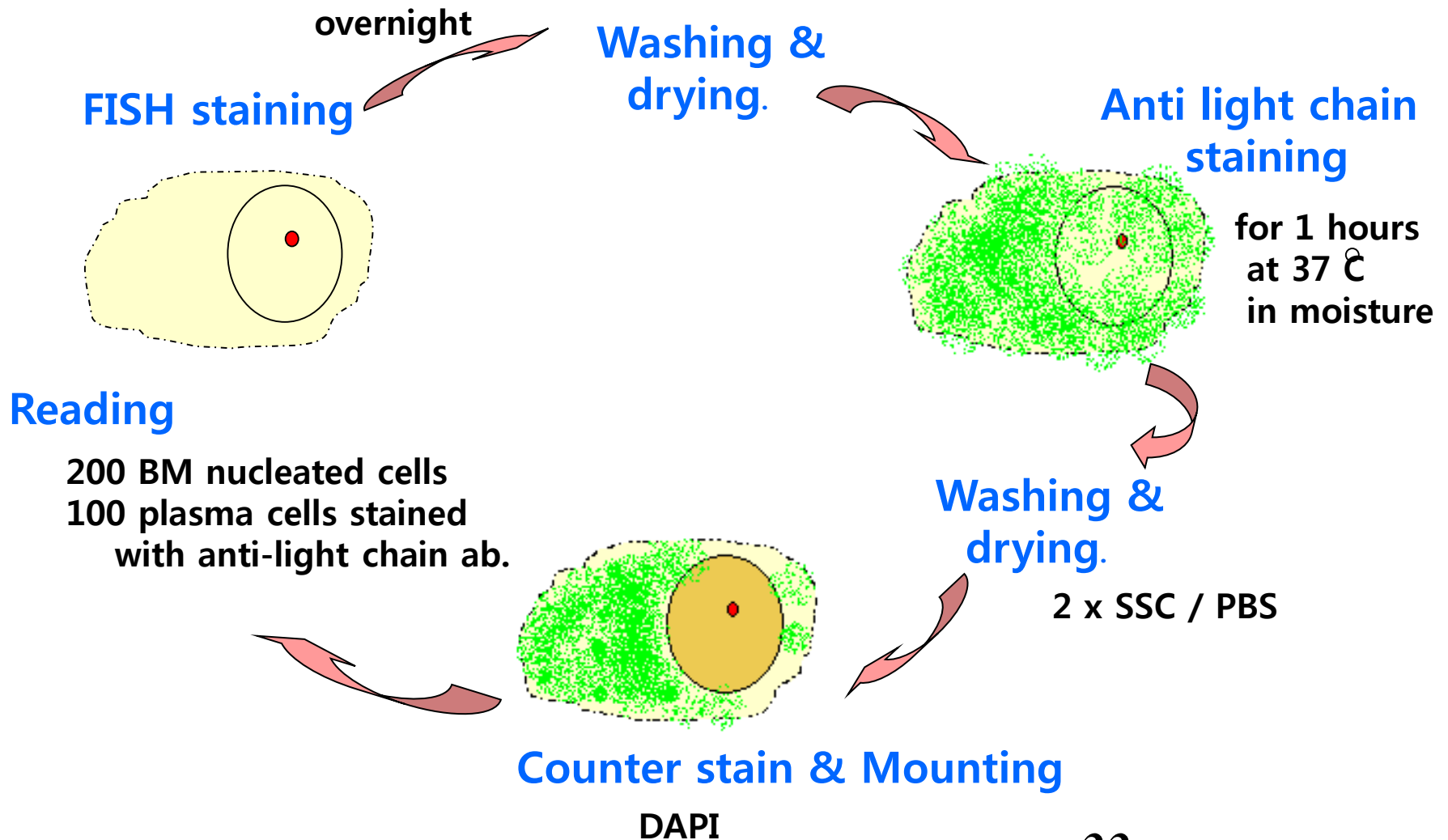


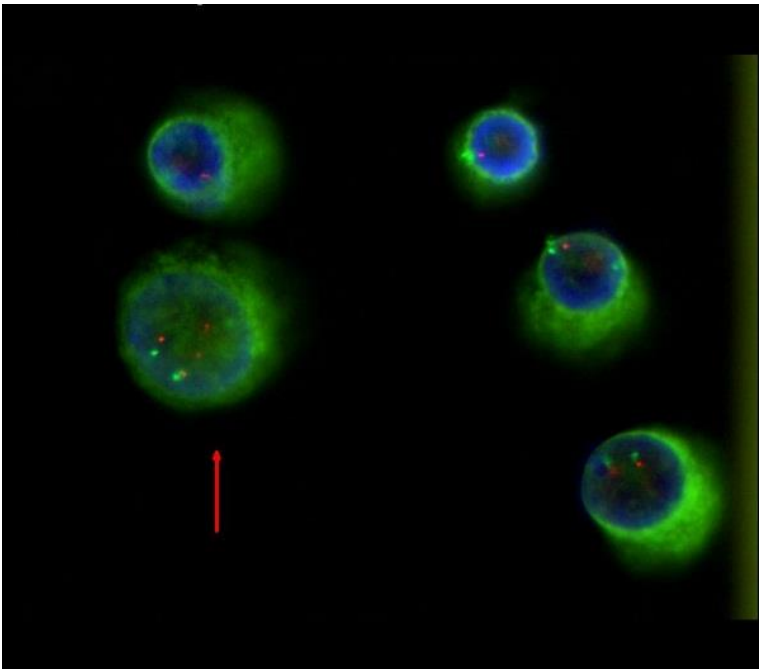
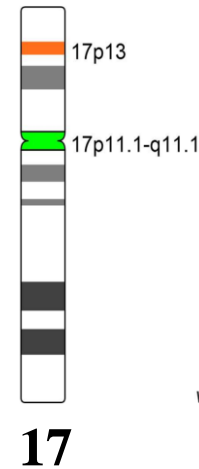
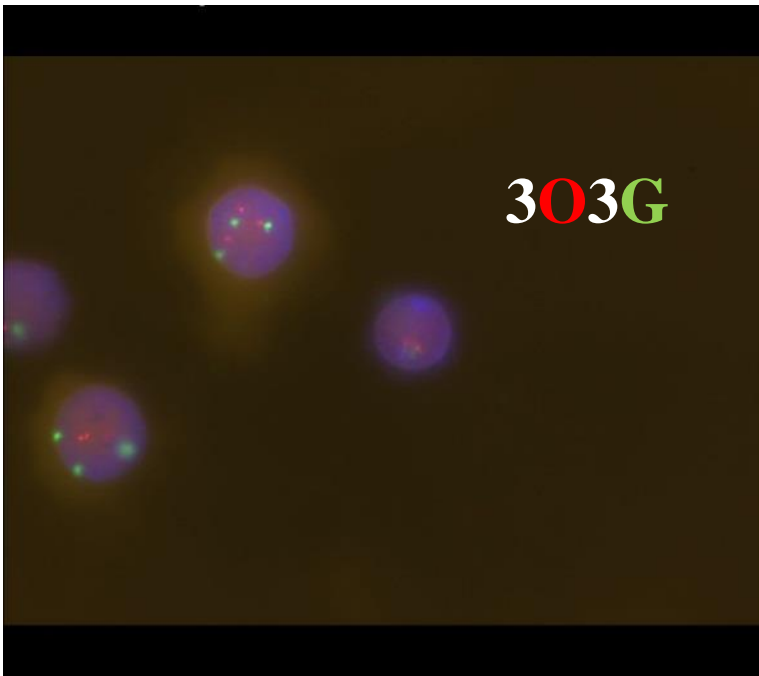
**t(14;16)(q32;q23)**





# Secondary immunohistochemical stain (rabbit to human kappa & lambda antibody-FITC)





**Plasma cell sorting;  
Immunomagnetic bead or  
FICTION (Fluorescence  
Immunophenotyping and  
interphase Cytogenetics as a Tool  
for the Investigation Of  
Neoplasms)**

# Treatment

- Chemotherapy--기본
- 자가조혈모세포이식--65세 이하
- 동종조혈모세포이식—노령 ,전신상태불량
- 방사선치료--증상완화목적
- 뼈질환치료제—골절방지
- 수혈
- 투석--신기능 악화

# Prognosis

- 치료저항성이 생겨 예후는 좋지 못함
- 항암화학요법이 도입되기 전에는 6개월 정도의 평균
- 현재는 항암치료 만으로도 2~3년의 생존기간을 보이고
- 자가조혈모세포이식을 한 경우는 5년
- 최근 도입된 신약들이 점점 광범위하게 도입됨에 따라 더 긴 생존기간을 보일 것으로 생각되며
- 국내에서는 보험 보험급여 문제로 효과 좋은 신약을 쓰기 힘들다.