

Molecular Diagnostics – Soft tissue and bone tumours

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CANCER
RESEARCH
UK

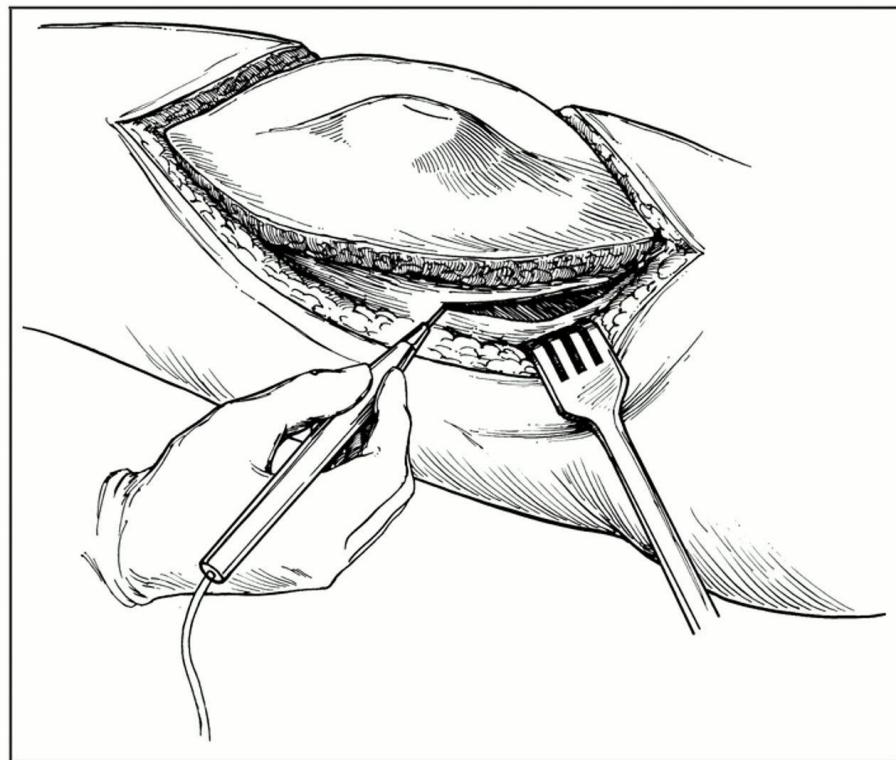
UCL
CENTRE



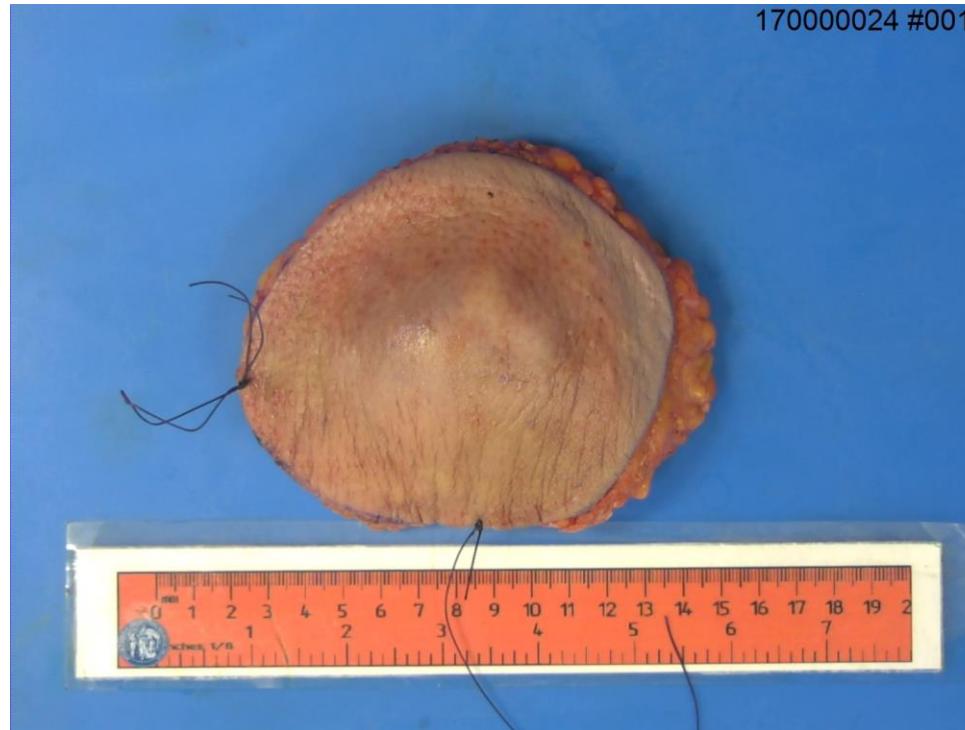
Royal National Orthopaedic Hospital 
NHS
NHS Trust

Outline

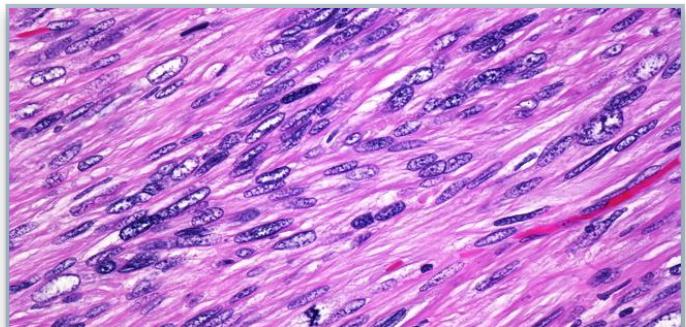
- Mesenchymal tumours– classification.
- Sarcomas and Molecular diagnostic assays (genetic).
- Recent developments in sarcoma diagnostics



Oncological Outcomes of Operative Treatment of Subcutaneous Soft-Tissue Sarcomas of the Extremities*
J Bone Joint Surg Am, 1997 Jun; 79 (6): 888 -97 .

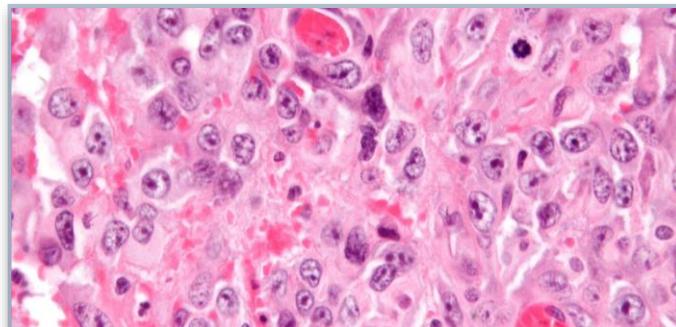


Classification of mesenchymal tumours



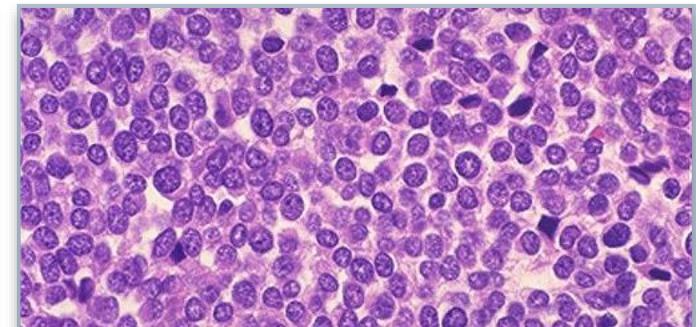
SPINDLE CELL

- Leiomyosarcoma
- Spindle cell rhabdomyosarcoma
- Fibrosarcoma
- Spindle cell sarcoma NOS



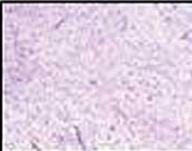
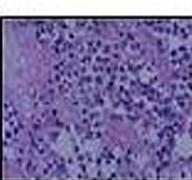
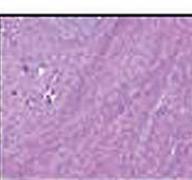
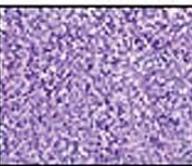
EPITHELIOID CELL

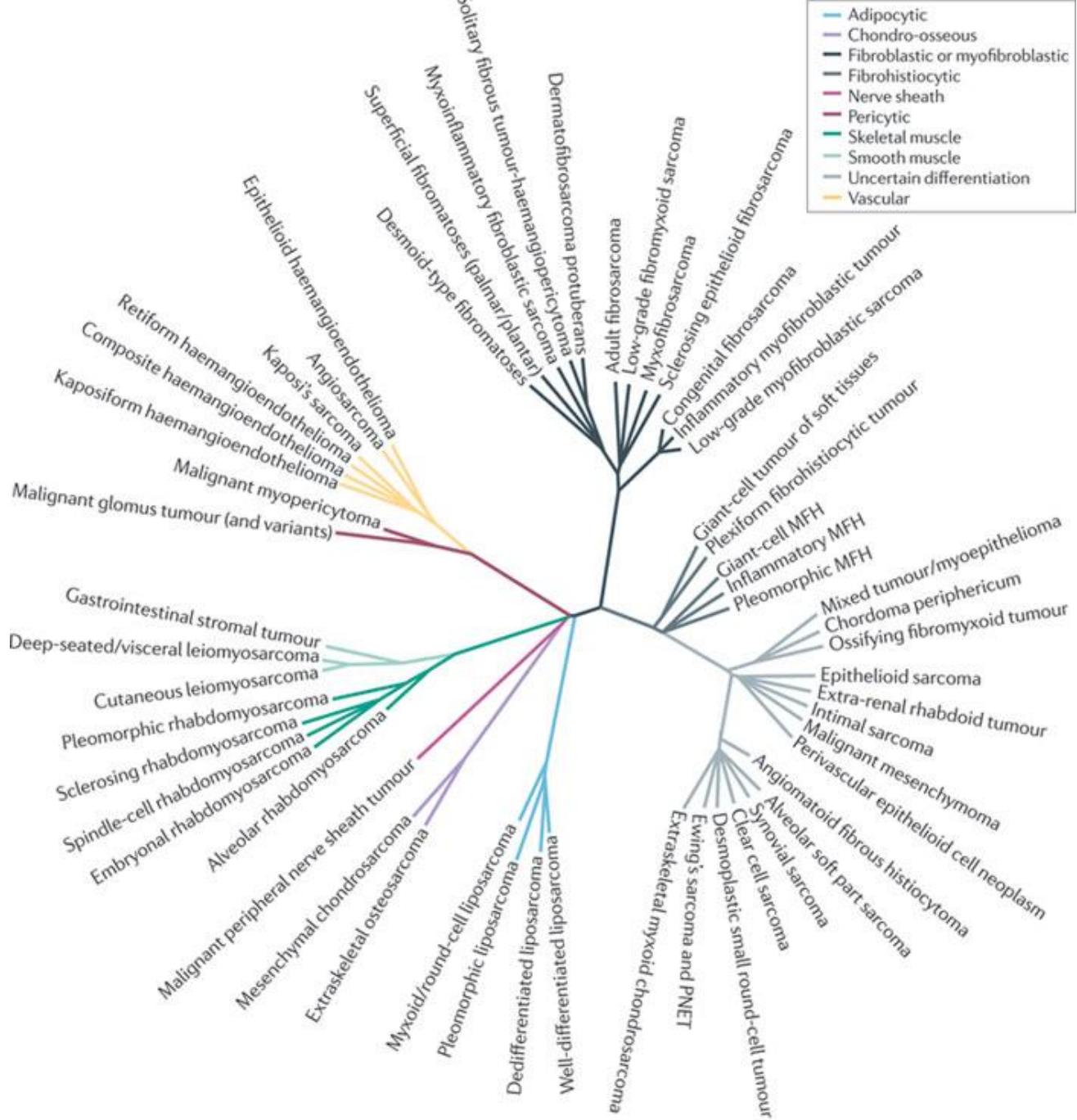
- Epithelioid sarcoma
- Epithelioid MPNST
- Epithelioid angiosarcoma
- Malignant rhabdoid tumour
- Metastatic Melanoma
- Metastatic carcinoma



ROUND CELL

- Ewing Sarcoma
- Desmoplastic round cell tumour
- Alveolar rhabdomyosarcoma
- Neuroblastoma
- Lymphoma
- Organ specific – Wilm's tumour, hepatoblastoma, pleuropulmonary blastoma

DIFFERENTIATION	Subtypes	Chromosomal traslocations	Fusion transcripts	
	ADIPOCYTIC TUMORS	<i>Lipoblastoma:</i> <i>Myxoid liposarcoma</i>	$t(7;8)(q31;q13); t(8;8)(q24;q13)$ $t(12;16)(q13;p11); t(12;22)(q13;q12)$	PLAG1-COL1A2; PLAG1-HAS2 CHOP-TLS; CHOP-EWS
	FIBROBLASTIC/ MYOFIBROBL.TUMORS	<i>Inflammatory myofibroblastic tumor</i> <i>Infantile fibrosarcoma</i> <i>Dermatofibrosarcoma protuberans/ Giant cell fibroblastoma</i>	$t(1;2)(q25;p23); t(2;19)(p23;q13); t(2;17)(p23;q23)$ $t(12;15)(p13;q25)$ $t(17;22)(q22;q13)$	TPM3-ALK; ALK-TPM4; ALK-CLTC ETV6-NTRK3 COL1A1-PDGFB
	SKELETAL MUSCLE TUMORS	<i>Alveolar rhabdomyosarcoma</i>	$t(2;13)(q35;q14); t(1;13)(p36;q14)$	PAX3-FKHR; PAX7-FKHR
	TUMORS OF UNCERTAIN DIFFERENTIATION	<i>Angiomatoid fibrous histiocytoma</i> <i>Synovial sarcoma</i> <i>Alveolar soft part sarcoma</i> <i>Clear cell sarcoma</i> <i>Extraskeletal myxoid chondrosarcoma</i> <i>Desmoplastic small round cell tumor</i>	$t(12;22)(q13;q12); t(12;16)(q13;p11)$ $t(X;18)(p11.2;q11.2)$ $t(X;17)(p11;q25)$ $t(12;22)(q13;q12)$ $t(9;22)(q22;q12); t(9;15)(q22;q21)$ $t(11;22)(p13;q12)$	SYT-SSX1/2/4 TFE3/ASPL EWS-ATF1 EWS-TEC; CHN-TFC12 EWS-WT1
	EWING SARCOMA		$t(11;22)(q24;q12); t(21;22)(q22;q12); t(17;22)(q12;q12); t(7;22)(p22;q12);$	FLI1-EWS; ERG-EWS E1AF-EWS; ETV1-EWS



Survival Rates

Figure 6b: Bone sarcoma 5-year rolling 5-year relative survival rates in males and females (UK: 1996-2010)

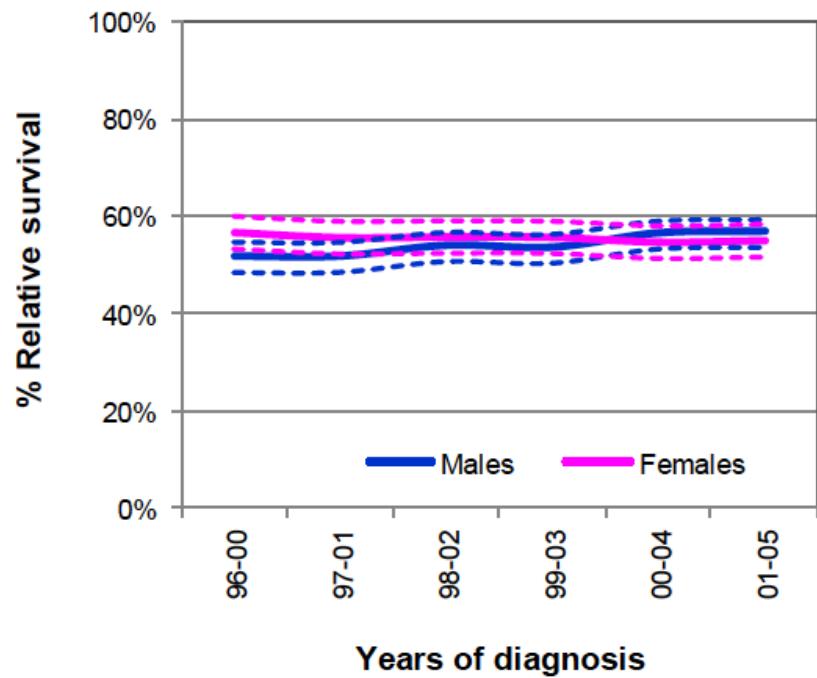
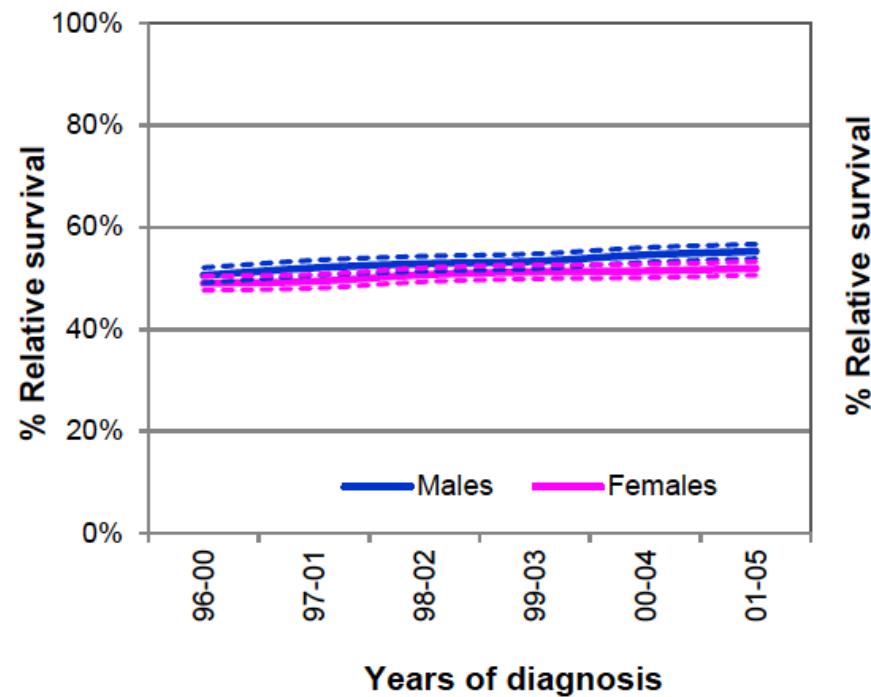
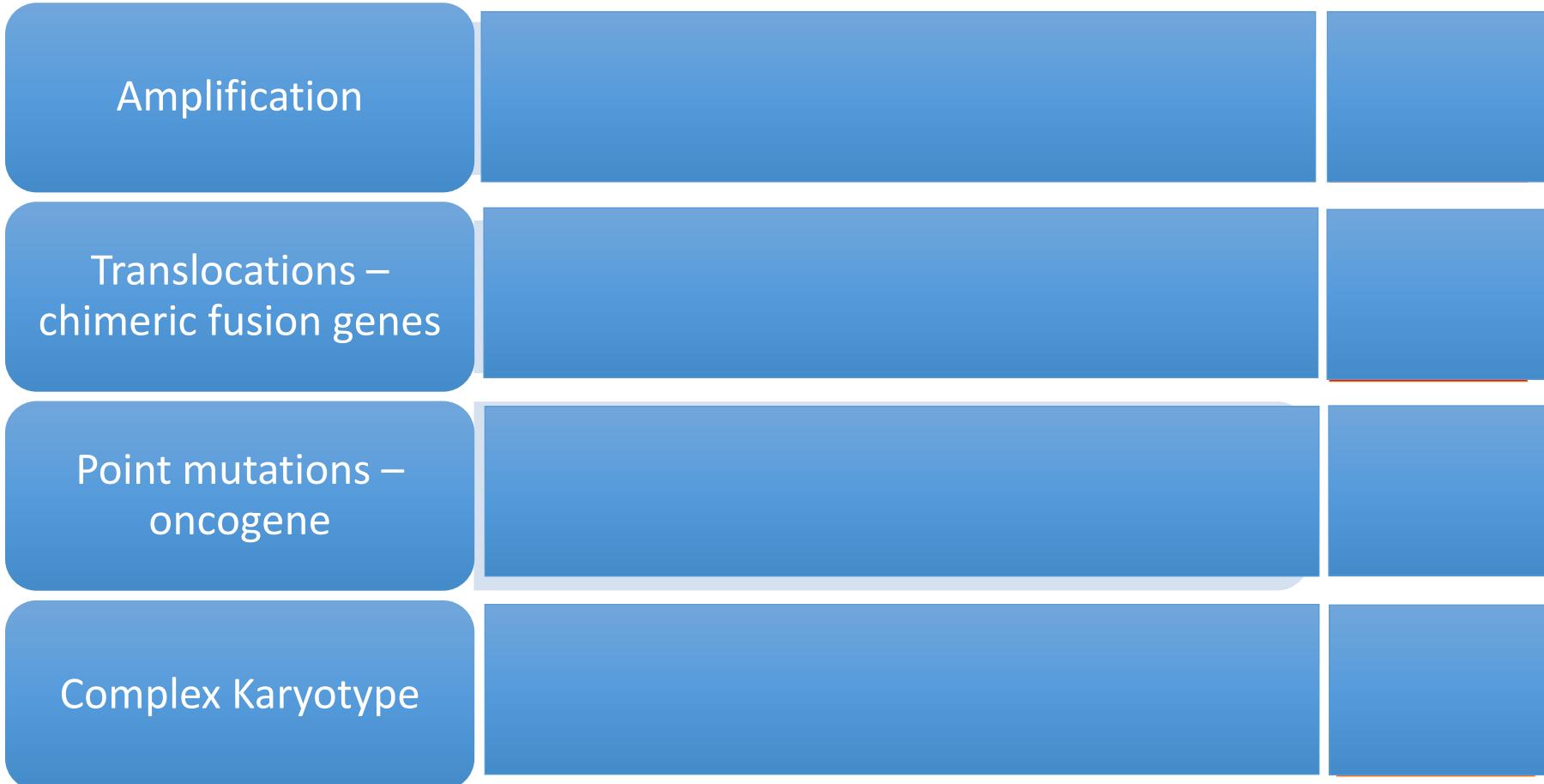


Figure 6a: Soft tissue sarcoma (excluding skin) 5-year rolling 5-year relative survival rates in males and females (UK: 1996-2010)



Molecular classification of mesenchymal tumours



Atypical lipomatous tumour

Well differentiated Liposarcoma

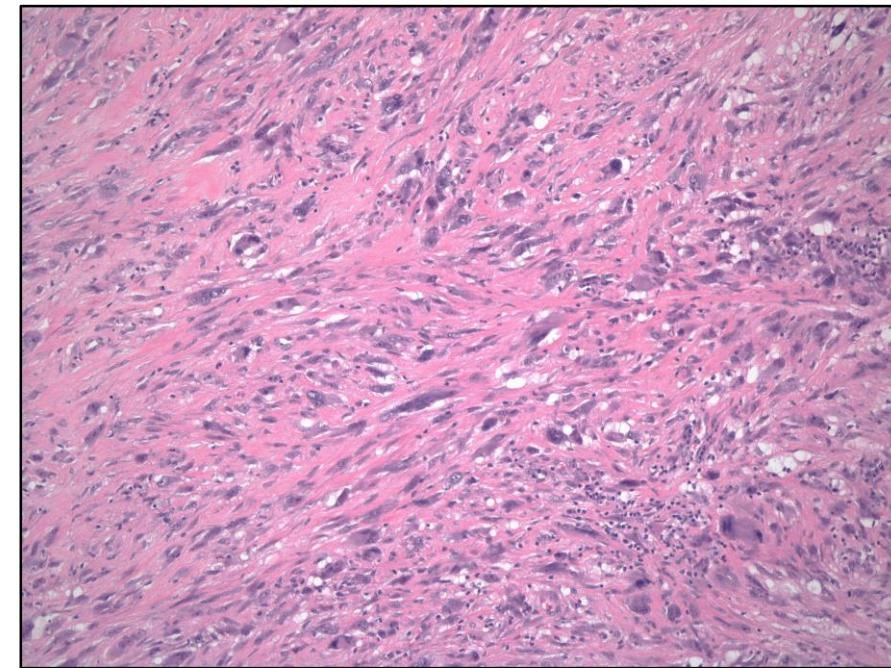
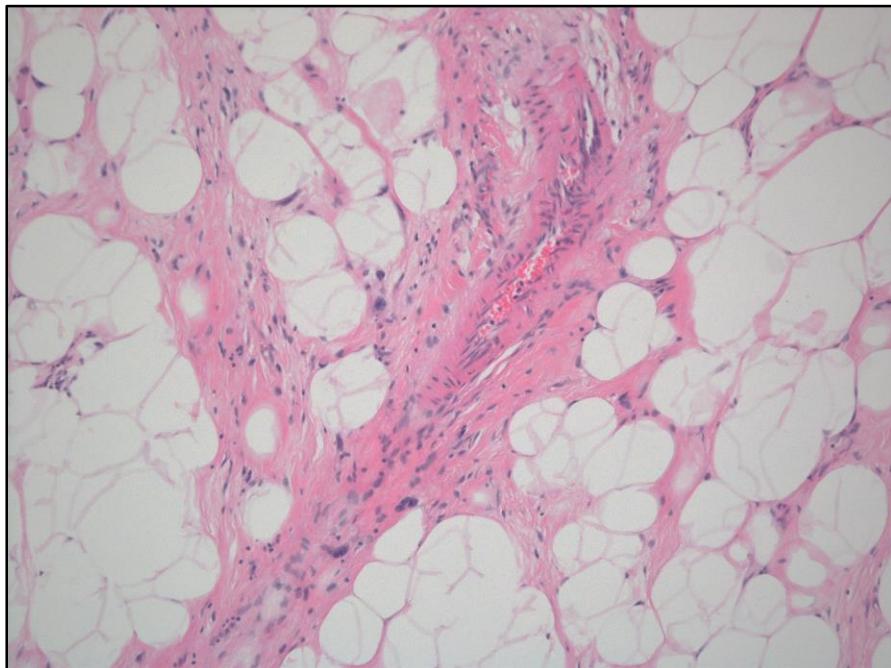
Dedifferentiated liposarcoma

Well-differentiated –

slow growing, does not metastasize, multiply recurrent,
no response to chemotherapy

Dedifferentiated–

aggressive, can metastasize, limited and transient benefit to
chemotherapy, median survival about 12 months



Gene Amplification – diagnostic tests

MDM2 amplification

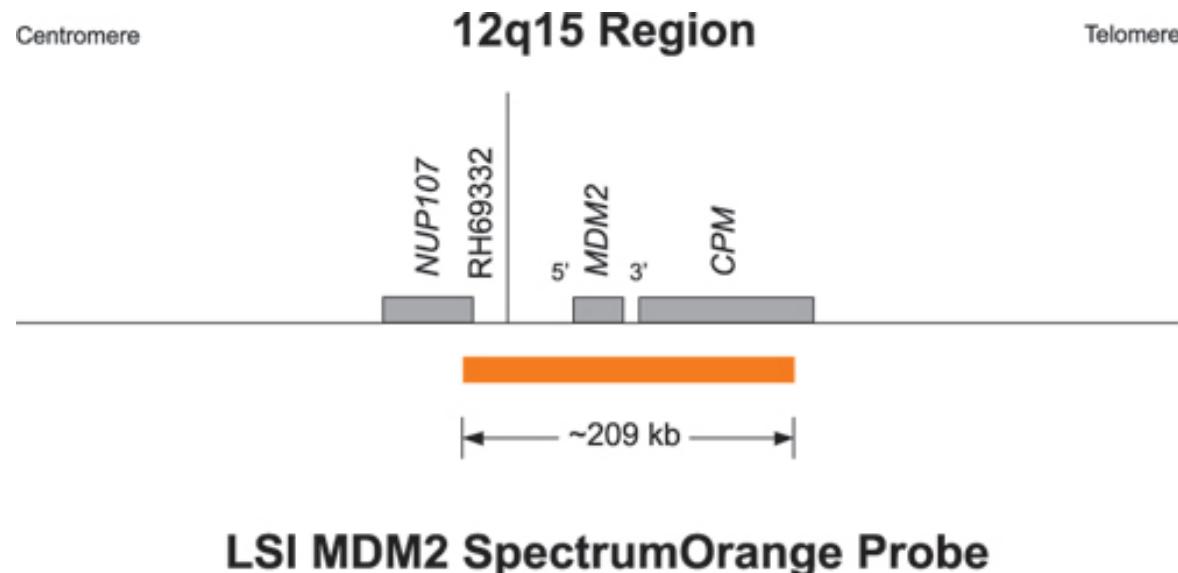
- Atypical lipomatous tumour/ dedifferentiated liposarcoma
- Low grade osteosarcoma / parosteal osteosarcoma

MDM2/CDK4 amplification – medical treatment options

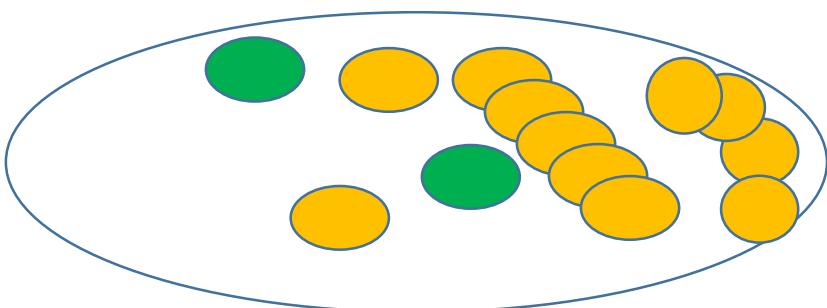
- RG7112
- Oral – inhibitor of *MDM2*
- Good response to well – dedifferentiated liposarcoma

Interphase FISH (Design for amplification)

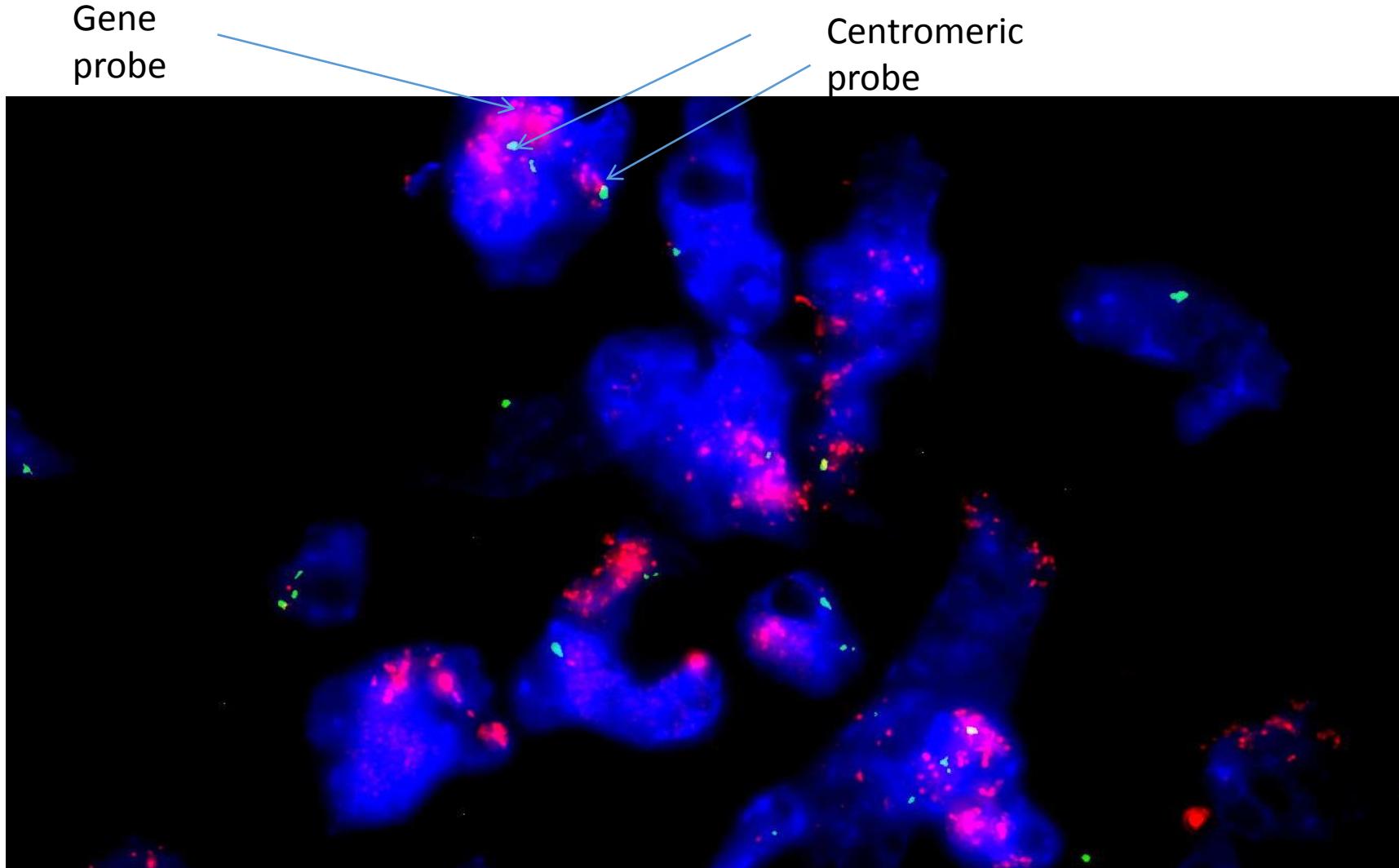
- Assay design –
- centromeric probe (enumerate the chromosomes)
- gene probe (enumerate the gene copy number)



- Addition of centromeric probe for the chromosome of interest– enables one to count number of chromosomes present.
- Diploid,
- aneuploid, polysomy
- low level copy gain,
- high level amplification

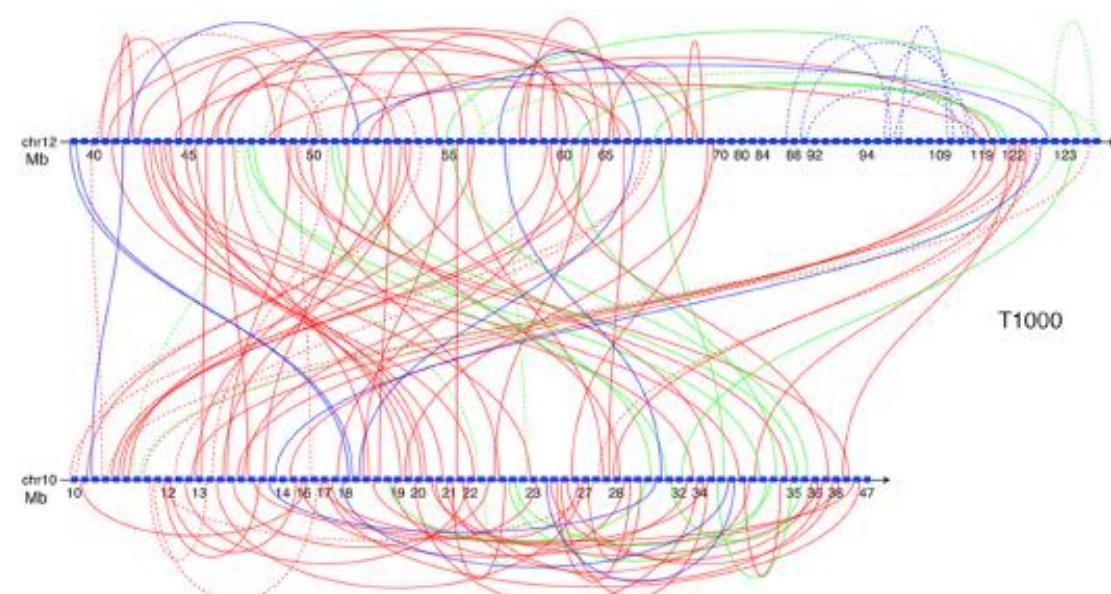
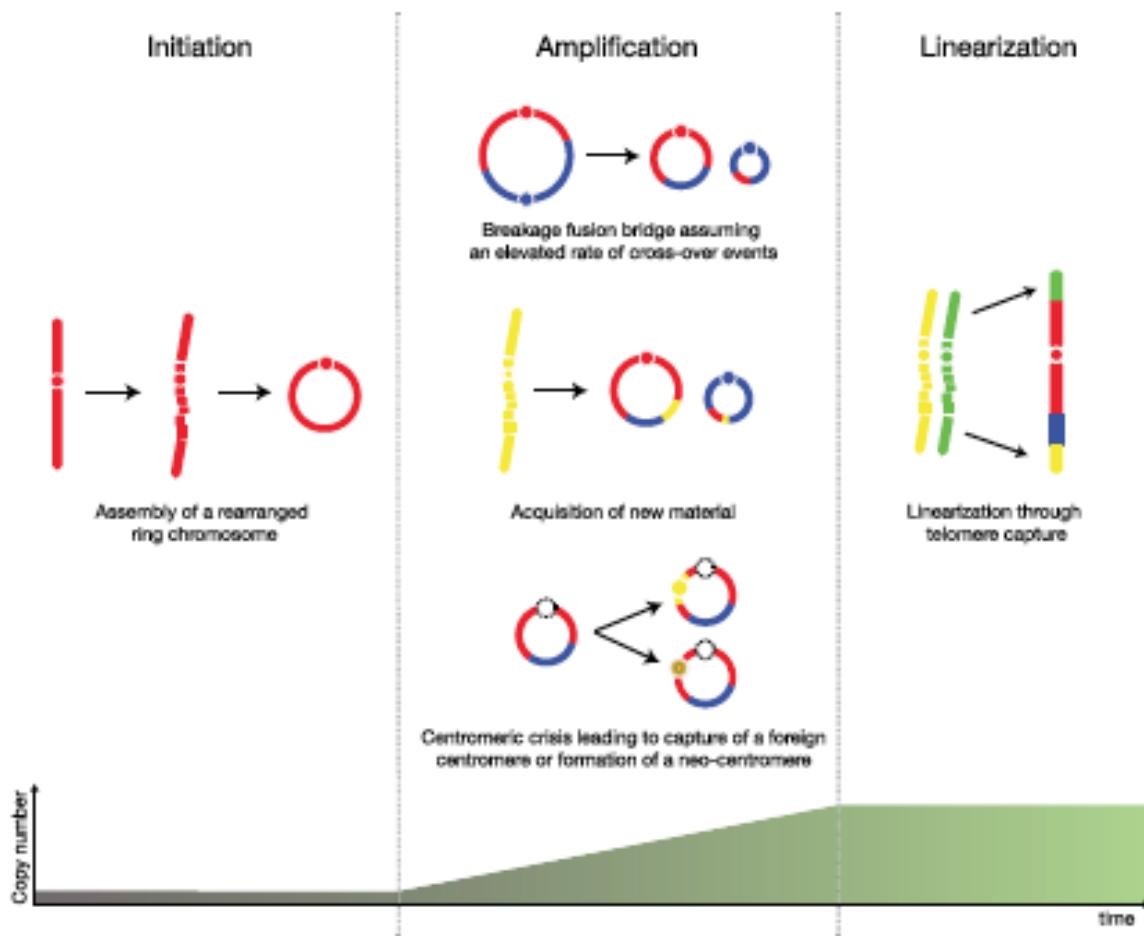


MDM2 amplification – Well differentiated liposarcoma



The Architecture and Evolution of Cancer Neochromosomes

Dale W. Garsed,^{1,8,9,15} Owen J. Marshall,^{5,15,17} Vincent D.A. Corbin,^{2,3,13,15} Arthur Hsu,^{2,15} Leon Di Stefano,² Jan Schröder,^{2,3} Jason Li,¹ Zhi-Ping Feng,^{2,3} Bo W. Kim,⁵ Mark Kowarsky,² Ben Lansdell,² Ross Brookwell,⁶ Ola Myklebost,¹⁰ Leonardo Meza-Zepeda,¹⁰ Andrew J. Holloway,¹ Florence Pedeutour,⁷ K.H. Andy Choo,⁵ Michael A. Damore,¹¹ Andrew J. Deans,¹² Anthony Papenfuss,^{2,3,4,8,13,16,*} and David M. Thomas^{1,8,14,16,*}



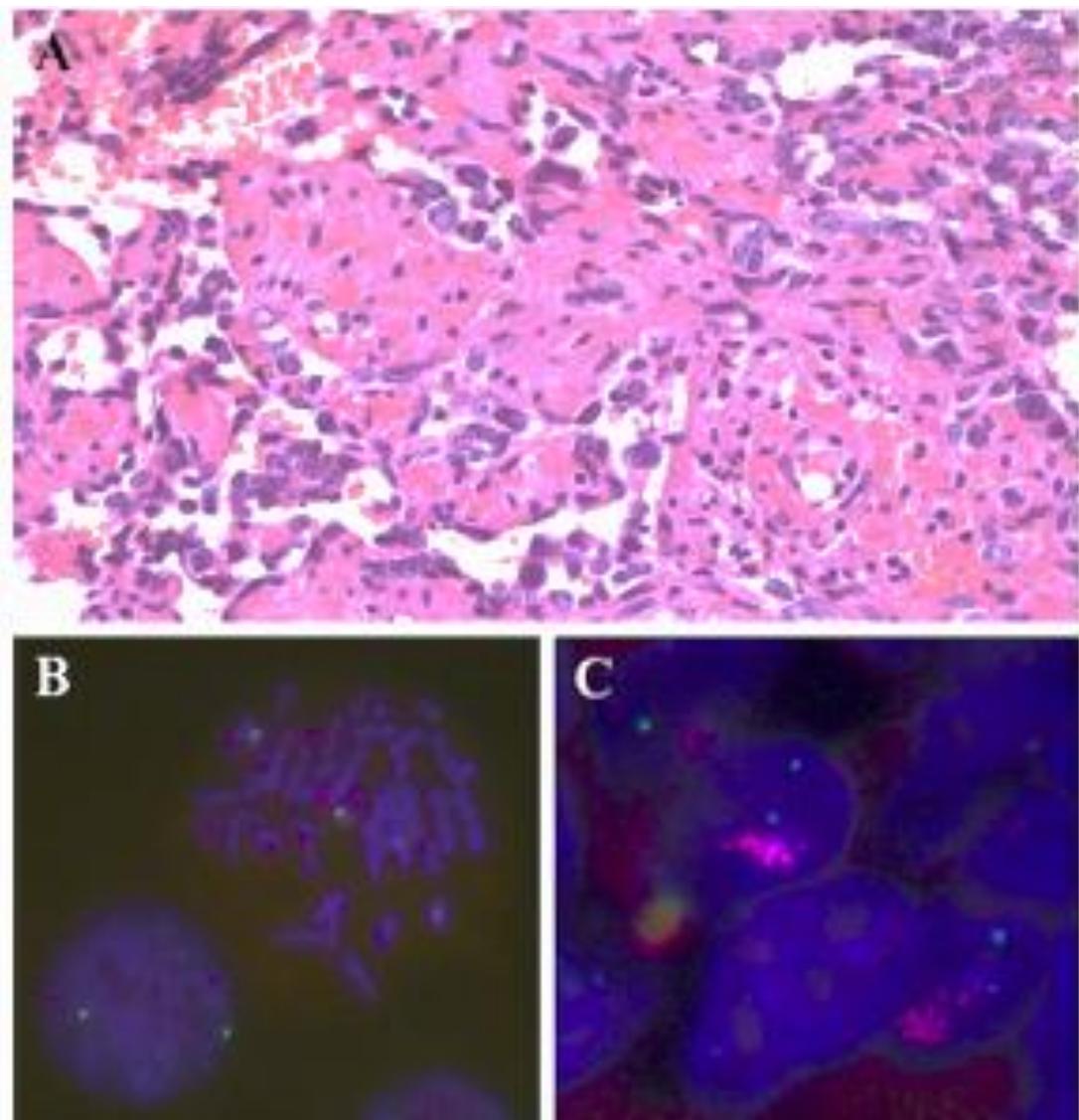
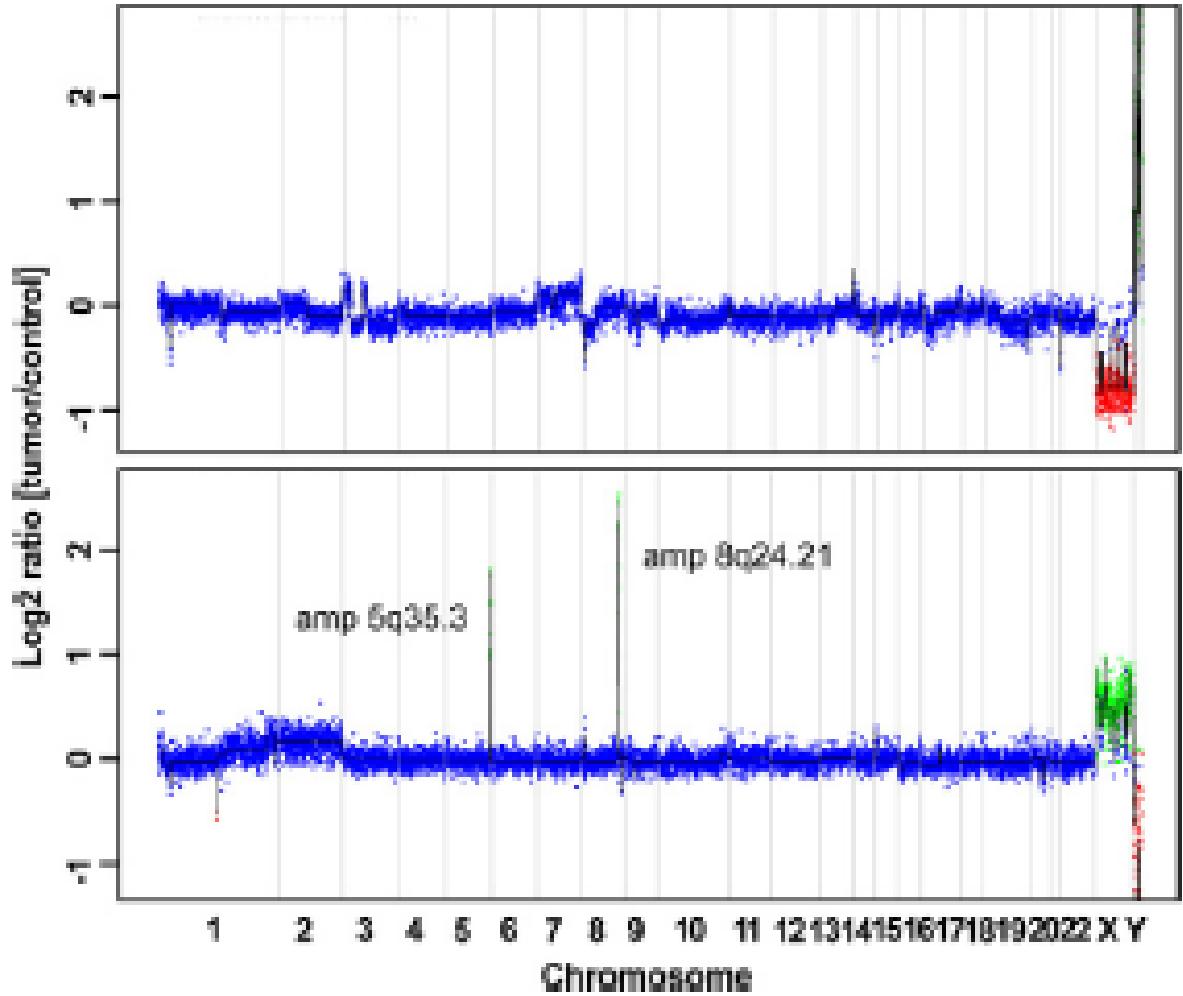
FISH as an aetiological adjunct

MYC – radiation induced angiosarcoma

The American Journal of Pathology, Vol. 176, No. 1, January 2010
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DOI: 10.2353/ajpath.2010.090637

Short Communication

MYC High Level Gene Amplification Is a Distinctive Feature of Angiosarcomas after Irradiation or Chronic Lymphedema



FISH – predictive and prognostic

ARTICLE

Received 12 Oct 2016 | Accepted 15 May 2017 | Published 23 Jun 2017

DOI: 10.1038/ncomms15936

OPEN

Recurrent mutation of IGF signalling genes and distinct patterns of genomic rearrangement in osteosarcoma

Sam Behjati^{1,2,3,*}, Patrick S. Tarpey^{1,*}, Kerstin Haase⁴, Hongtao Ye⁵, Matthew D. Young¹, Ludmil B. Alexandrov⁶, Sarah J. Farndon^{1,7}, Grace Collord¹, David C. Wedge⁸, Inigo Martincoren¹, Susanna L. Cooke¹, Helen Davies¹, William Mifsud⁷, Mathias Lidgren¹, Sancha Martin¹, Calli Latimer¹, Mark Maddison¹, Adam P. Butler¹, Jon W. Teague¹, Nischalan Pillay^{5,9}, Adam Shlien¹⁰, Ultan McDermott¹, P. Andrew Futreal^{1,11}, Daniel Baumhoer¹², Olga Zaikova¹³, Bodil Bjerkehagen¹³, Ola Myklebost^{13,14}, M. Fernanda Amary⁵, Roberto Tirabosco⁵, Peter Van Loo^{4,15}, Michael R. Stratton¹, Adrienne M. Flanagan^{5,9,**} & Peter J. Campbell^{1,16,**}

Cancer Medicine

ORIGINAL RESEARCH

Fibroblastic growth factor receptor 1 amplification in osteosarcoma is associated with poor response to neo-adjuvant chemotherapy

M. Fernanda Amary^{1,2}, Hongtao Ye¹, Fitim Berisha¹, Bhavisha Khatri¹, Georgina Forbes¹, Katie Lehotsky¹, Anna M. Frezza^{2,3}, Sam Behjati⁴, Patrick Tarpey⁴, Nischalan Pillay^{1,2}, Peter J. Campbell⁴, Roberto Tirabosco¹, Nadège Presneau², Sandra J. Strauss^{2,3} & Adrienne M. Flanagan^{1,2}

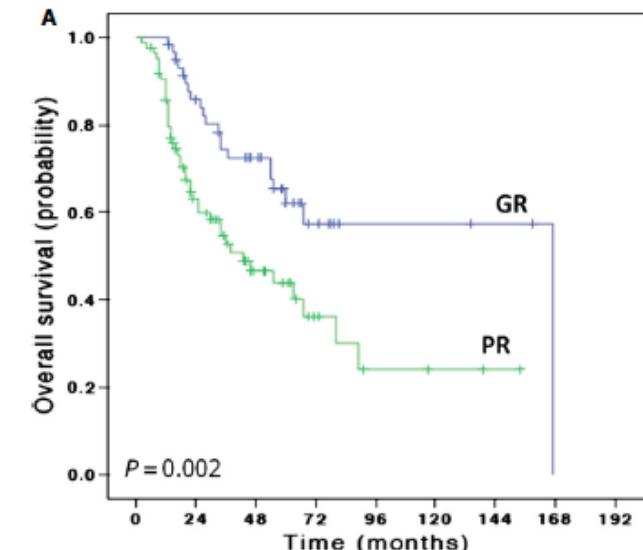
¹Histopathology, London Sarcoma Service, Royal National Orthopaedic Hospital NHS Trust, Stanmore, Middlesex, HA7 4LP, U.K.

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⁴Cancer Genome Project, Wellcome Trust Sanger Institute, Wellcome Trust Genome Campus, Hinxton, Cambridgeshire, CB10 1SA, U.K.

OSTEOSARCOMA



Gene amplification

- Diagnostic test – Atypical lipomatous tumour / Dedifferentiated liposarcoma
 - MDM2 amplification
- Predictive and prognostic utility – osteosarcoma
 - FGFR1 amplification / IGF amplification
- Biology of disease

Molecular classification of sarcomas

Amplification

Translocations –
chimeric fusion
genes

- Ewing's sarcoma – EWSR1-FLI1
- Synovial Sarcoma – SYT-SSX1

- Undifferentiated sarcoma
- Pleomorphic liposarcoma

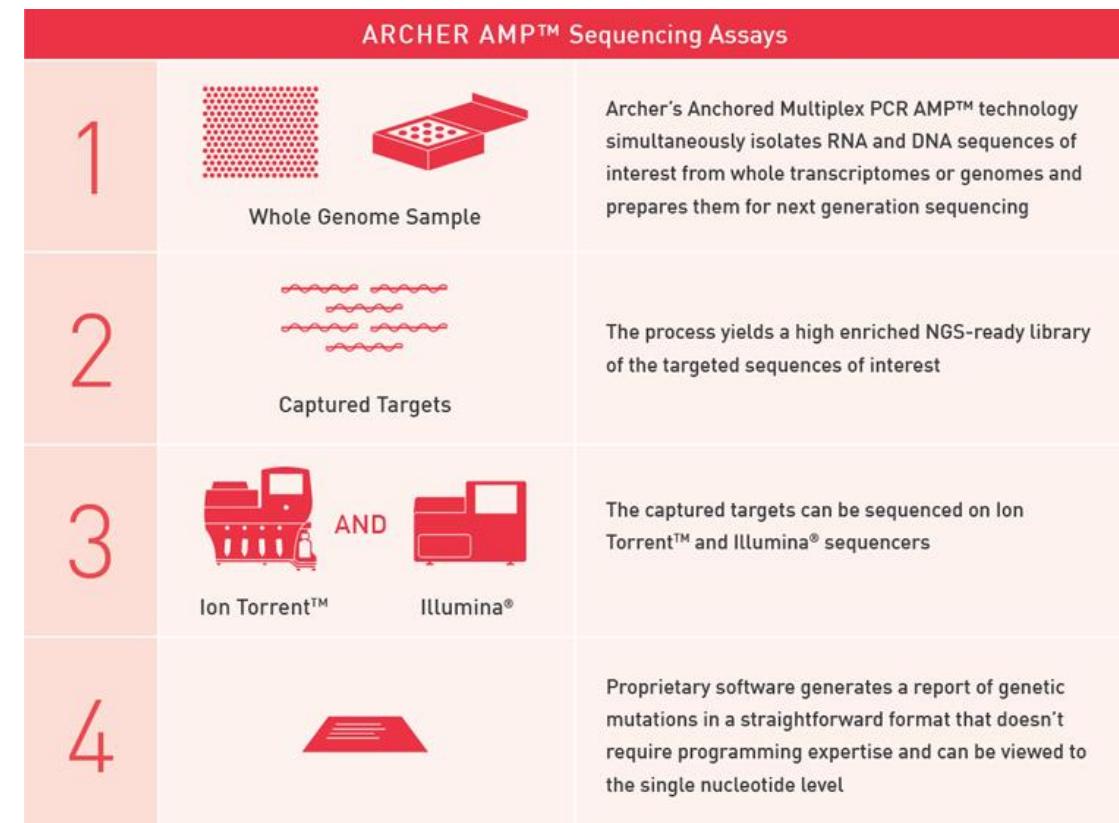
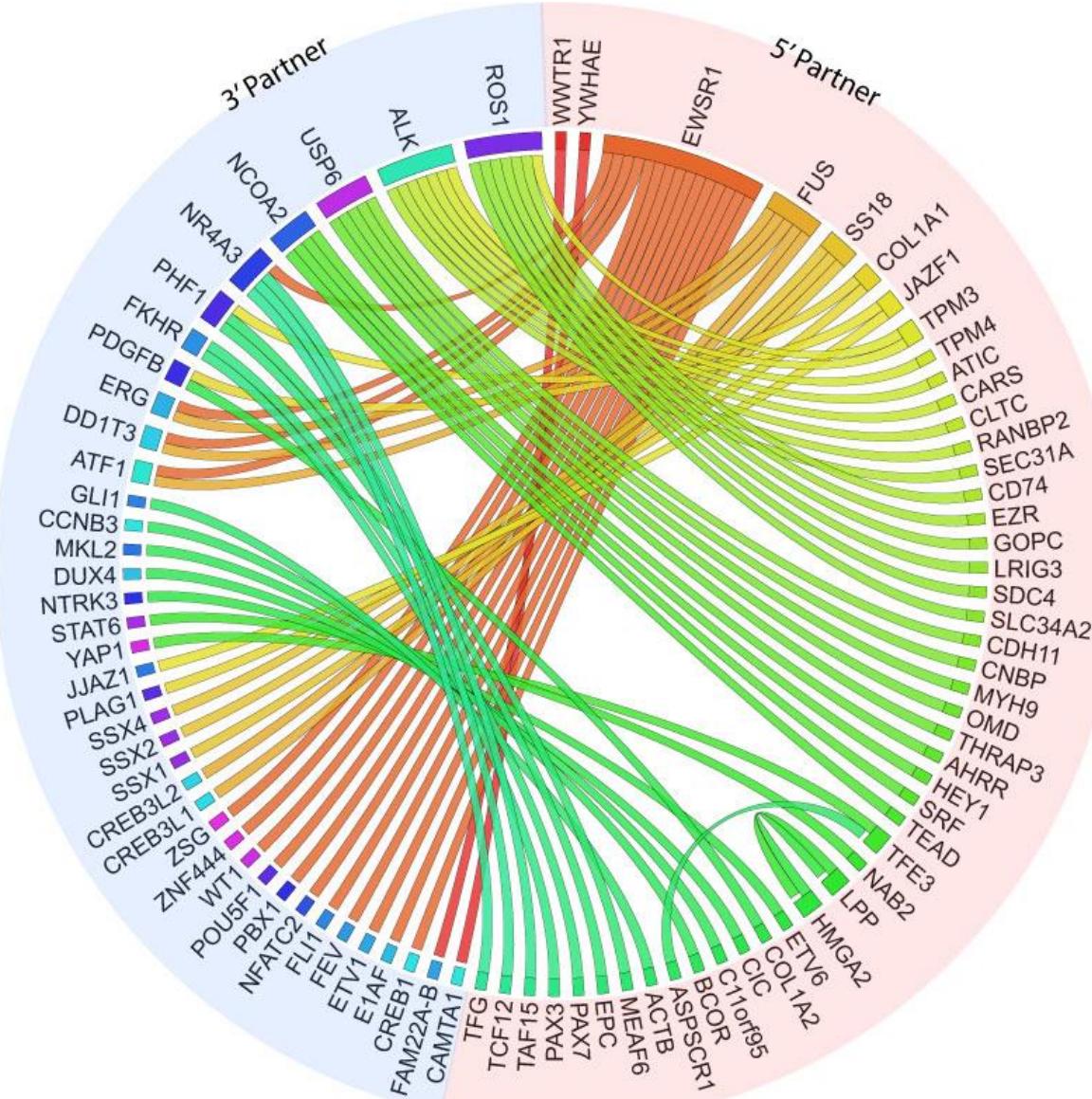
Table 3: Selected chromosomal translocations in sarcomas

Sarcoma type	Translocation	Effect of translocation
Ewing's/PNET	t(11;22) (80-85%) t(21;22) t(7;22) t(2;22) t(17;22)	EWS-FL1 translocation: some variants associated with more favourable prognosis. FL1 acts as transcription factor.
Rhabdomyosarcoma (alveolar)	t(2;13) t(1;13)	PAX-FOX01a gene fusion. Encodes chimeric transcription factor. Fusion status correlates with clinical outcome.
DFSP	t(17;22)	Fusion of collagen 1 type 1 α and PDGF β .
Synovial sarcoma	t(X;18)	Biology of fusion product not well known -?transcription co-factor.
Clear cell sarcoma	t(12;22)	EWS-ATF1
Myxoid liposarcoma	t(12;16) t(12;22)	CHOP-EWS/CHOP-TLS fusion product.
Desmoplastic small round-cell tumour	t(11;22)	EWS-WT1

Fusion gene detection – diagnostic assays

- RT-PCR
- FISH
- Immunohistochemistry – e.g. *NAB2*—*STAT6* fusion in Solitary Fibrous Tumour.
- RNA – *in situ* hybridisation (RNAscope)
- RNA-seq

Archer® FusionPlex® Sarcoma Kit

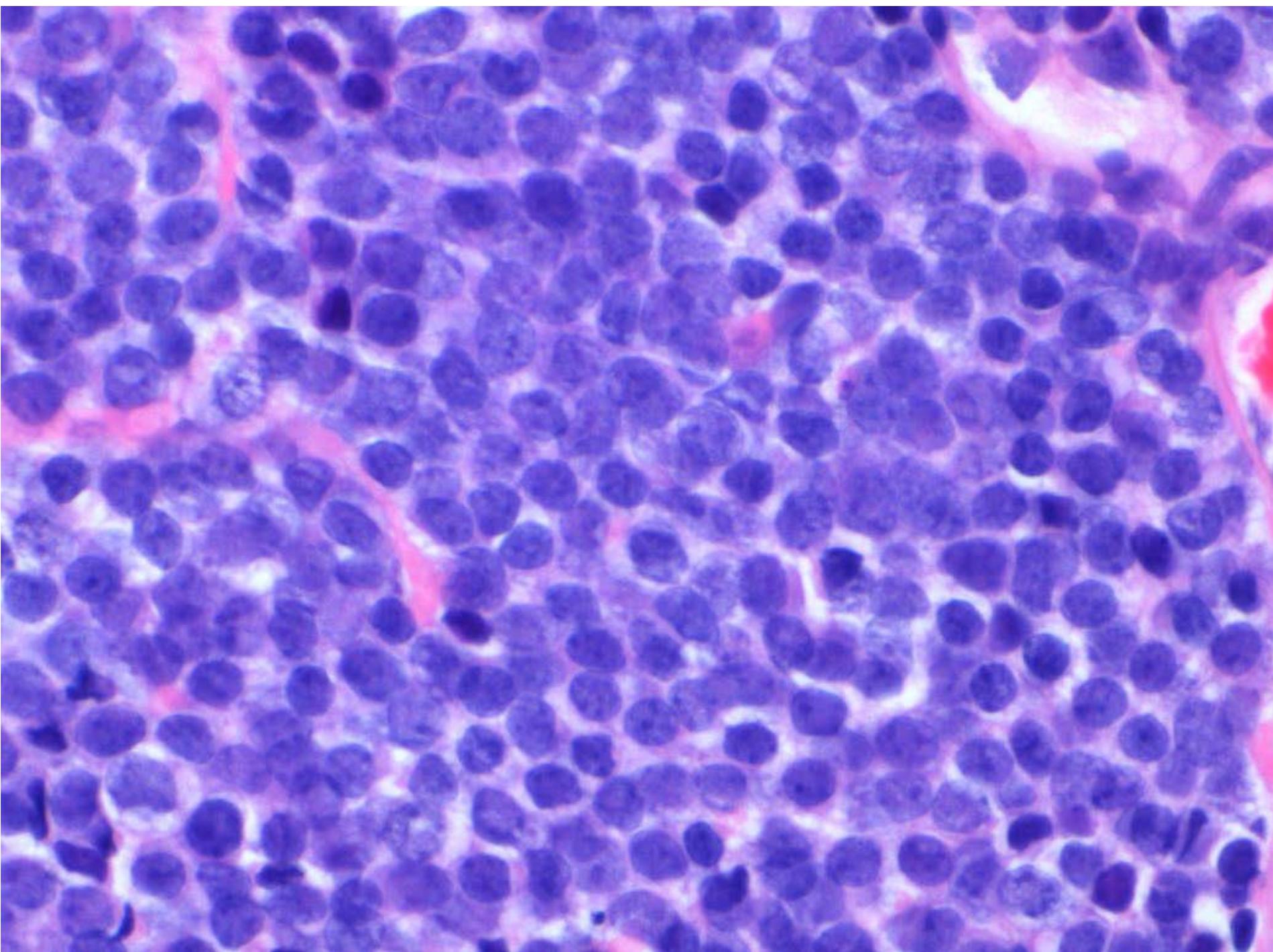




16yr old. Proximal femur fracture.

Radiological Differential diagnosis:

- Infection
- Langerhan's cell histiocytosis
- Osteosarcoma
- Ewing sarcoma



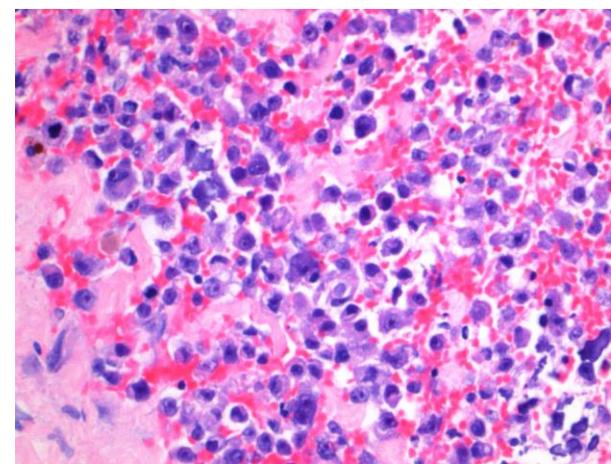
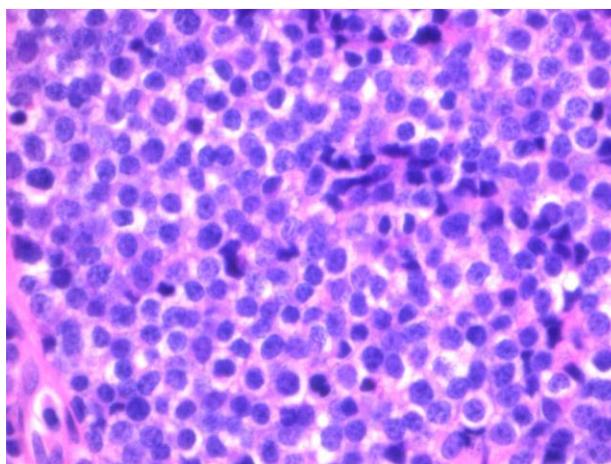
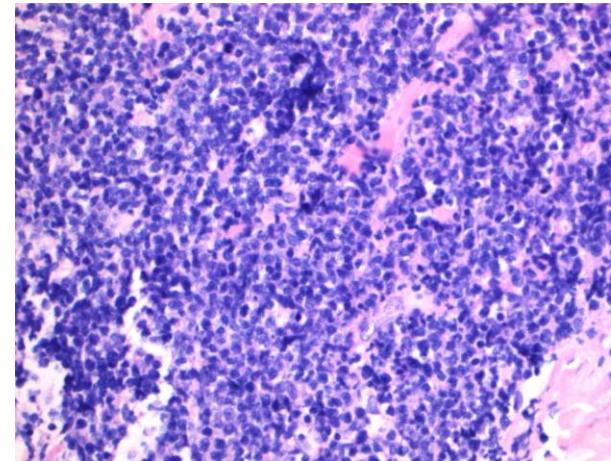
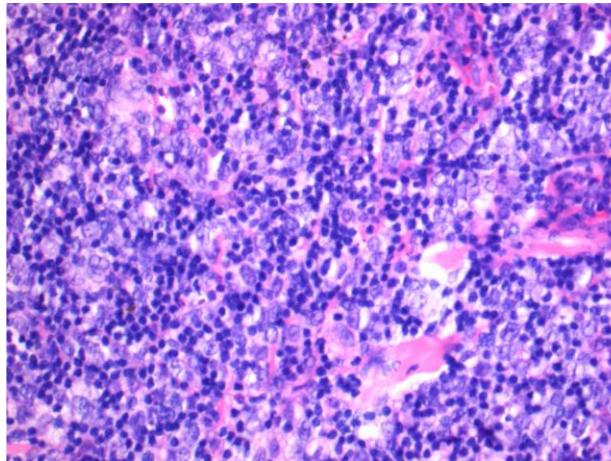
Differential Diagnosis

- Ewing's sarcoma
- Ewing's-like Round Cell Sarcomas
- Small Cell Osteosarcoma
- Lymphoma
- Alveolar Rhabdomyosarcoma
- Desmoplastic Small Round Cell Tumour
- Round Cell Liposarcoma (HG Myxoid LPS, WHO 2013)
- Mesenchymal Chondrosarcoma
- Neuroblastoma
- Metastatic Melanoma

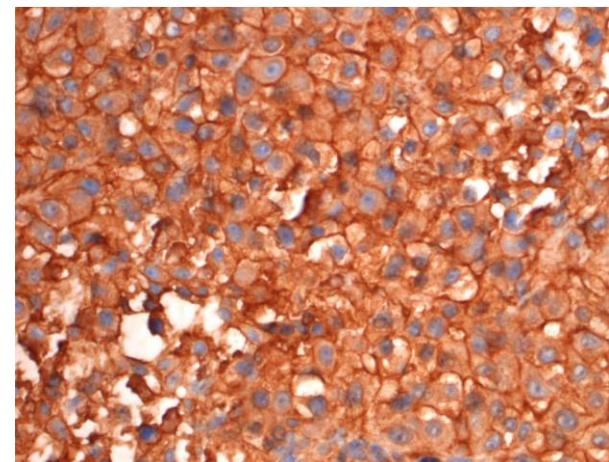
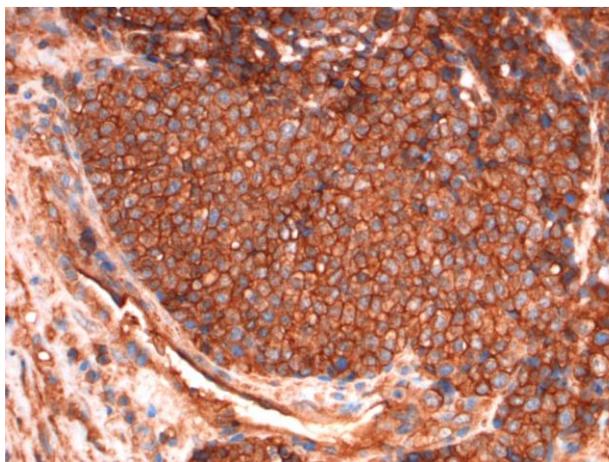
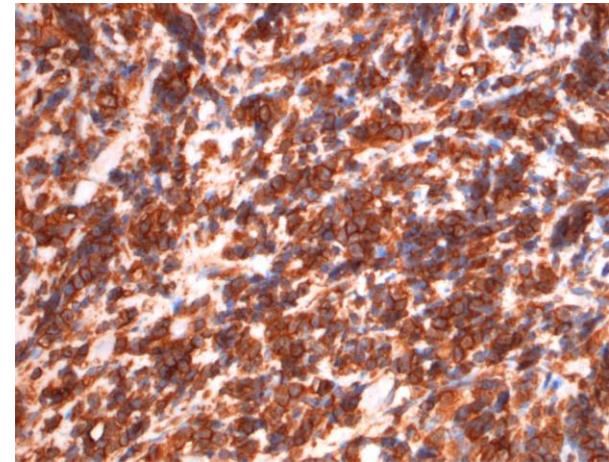
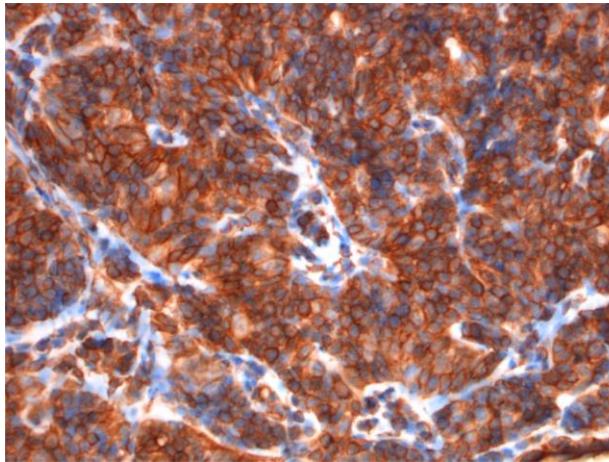
Immunohistochemistry

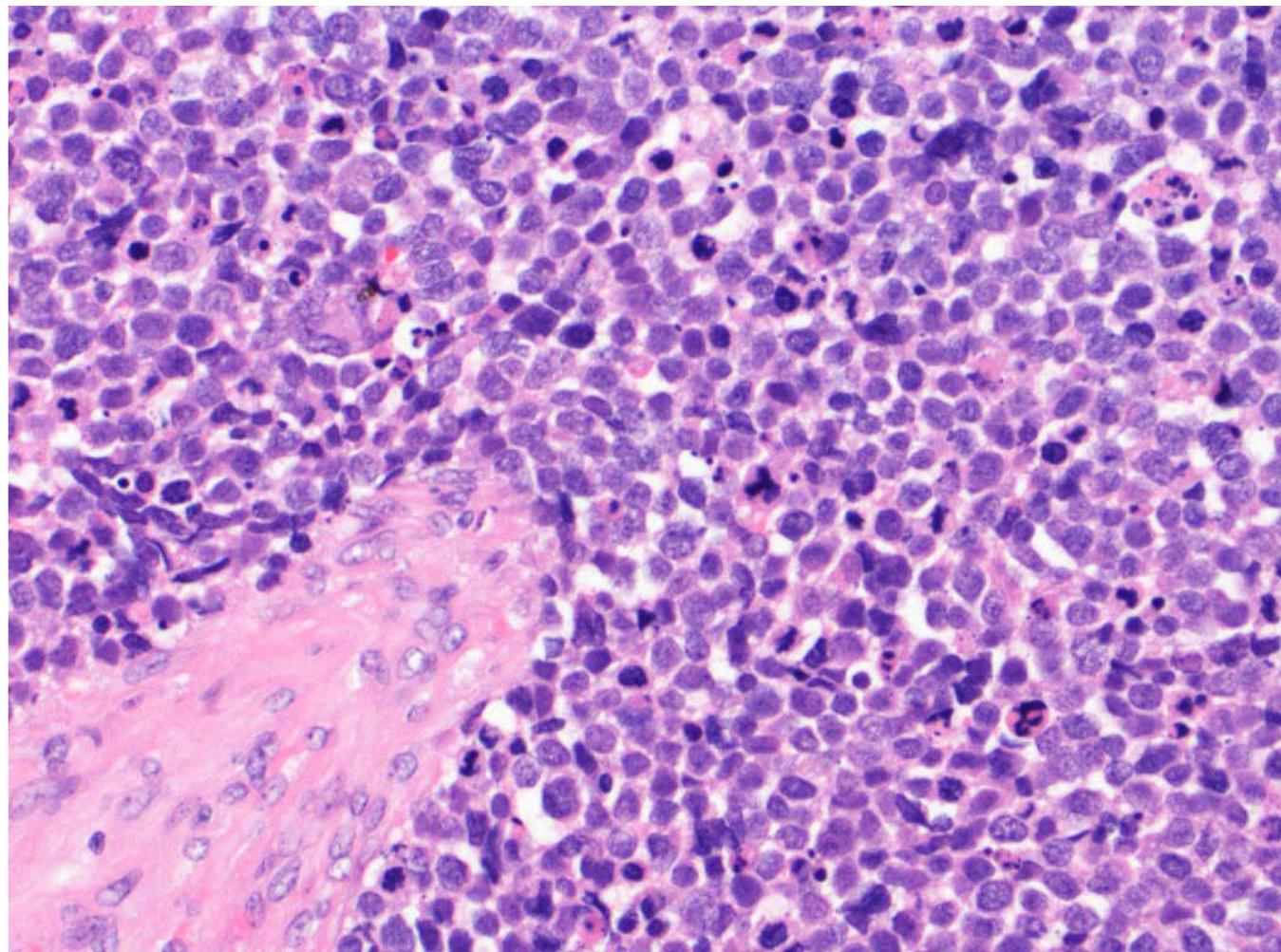
- CD99
- CD45
- TdT
- S100
- Desmin
- Pan cytokeratin

The value of CD99



The value of CD99





The common immunophenotype

- CD99
- CD45
- TdT
- S100
- Desmin
- Pankeratin

+++

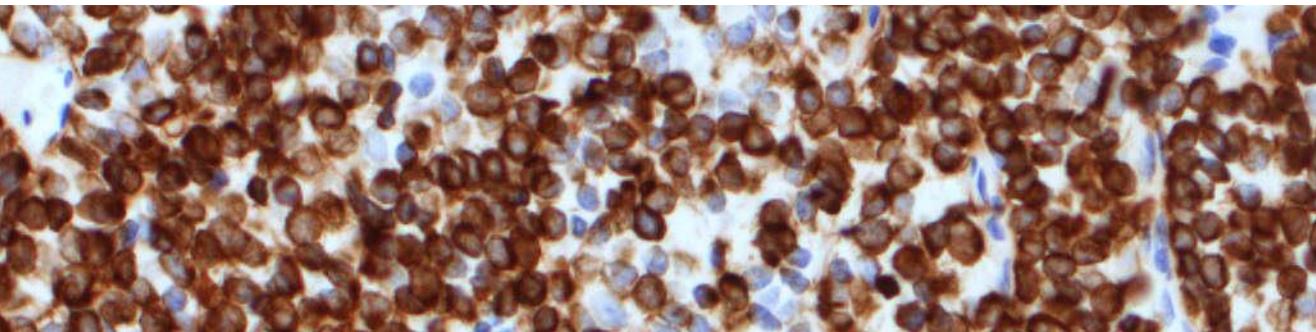
Neg.

Neg.

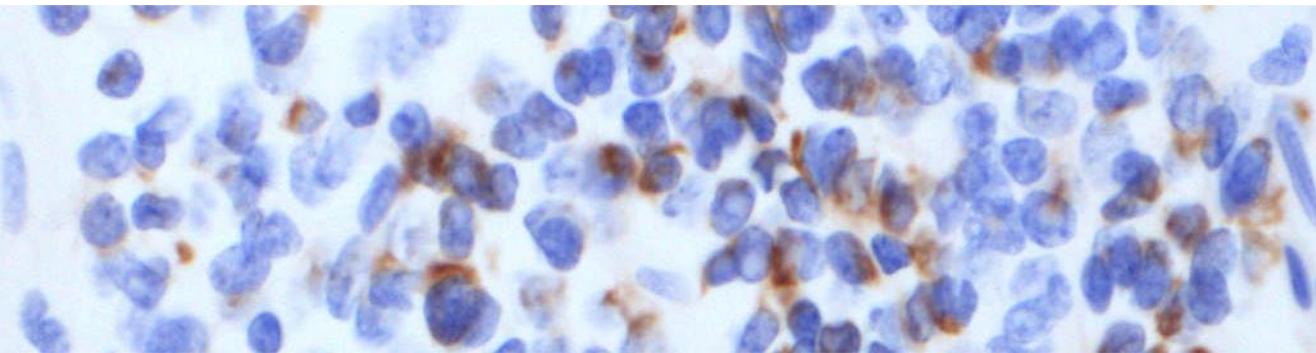
Neg.

Neg.

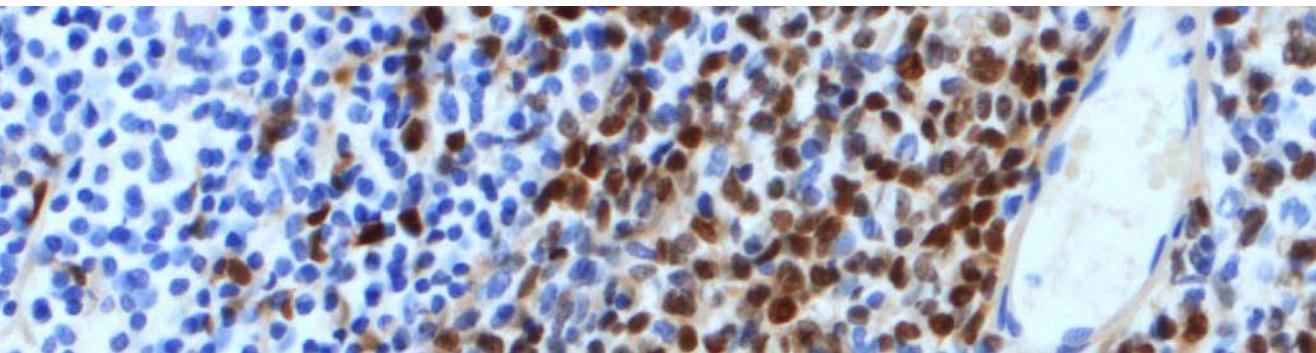
Neg.



MNF116
(~10%)



CK20



S100

The uncommon immunophenotype

Molecular confirmation needed

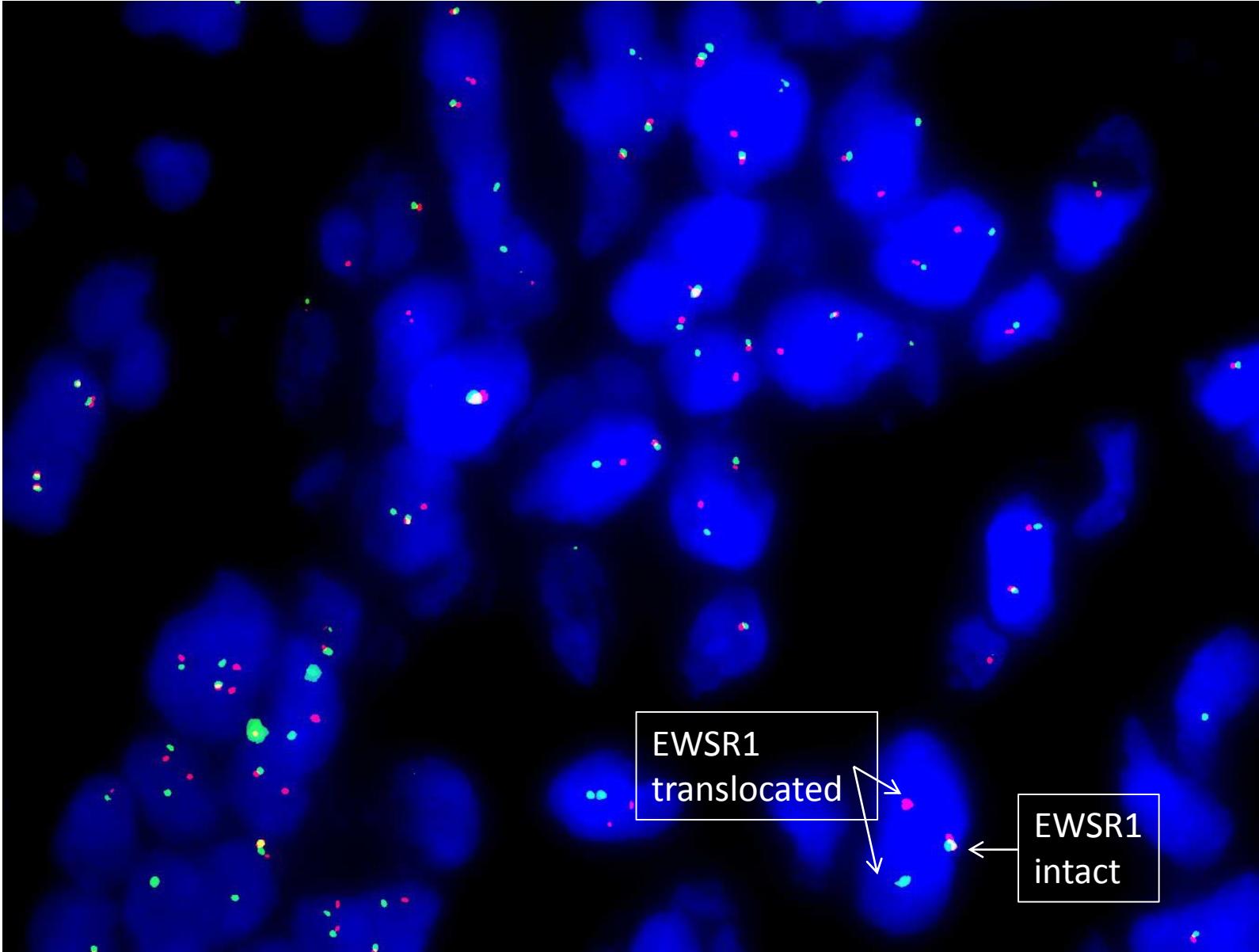
Break-apart FISH

- Fusion detection: break-apart probe corresponding to the most commonly rearranged partner.
- E.g. FISH for Ewing sarcoma
- EWSR1 – FLI1

EWSR1 BREAK APART DESIGN – DUAL COLOUR



LSI EWSR1 Dual Color, Break Apart Rearrangement Probe



EWSR1
translocated

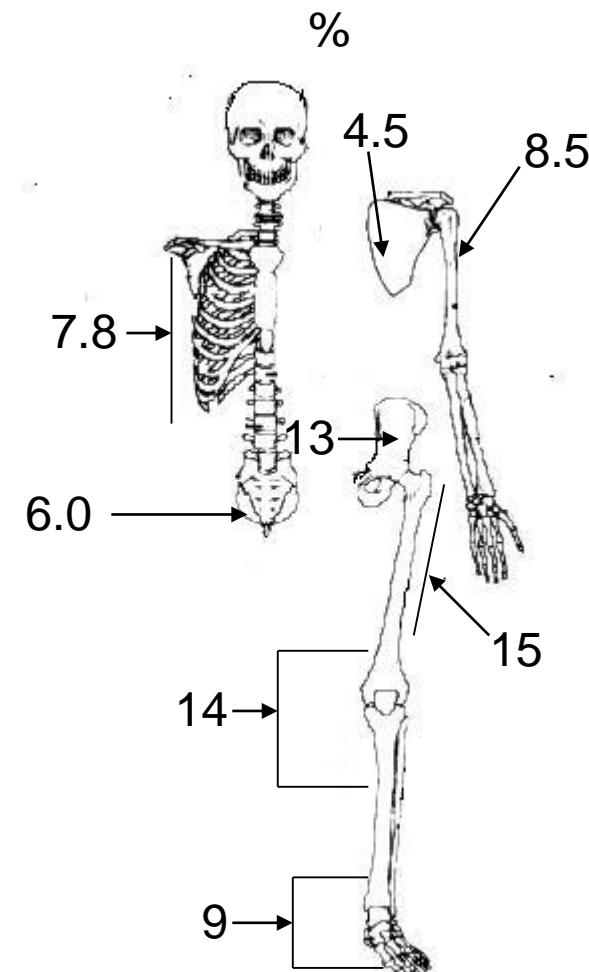
EWSR1
intact

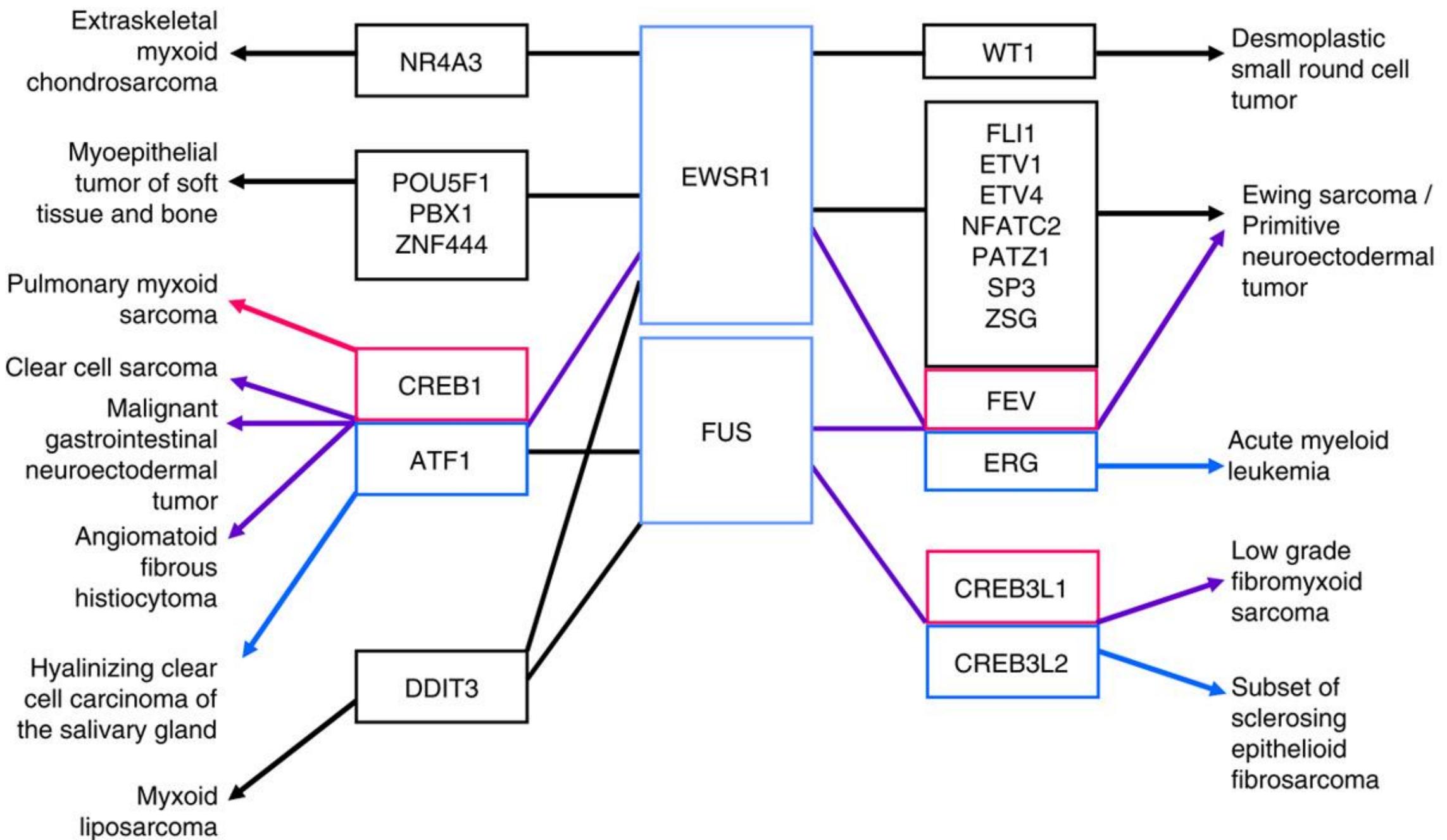


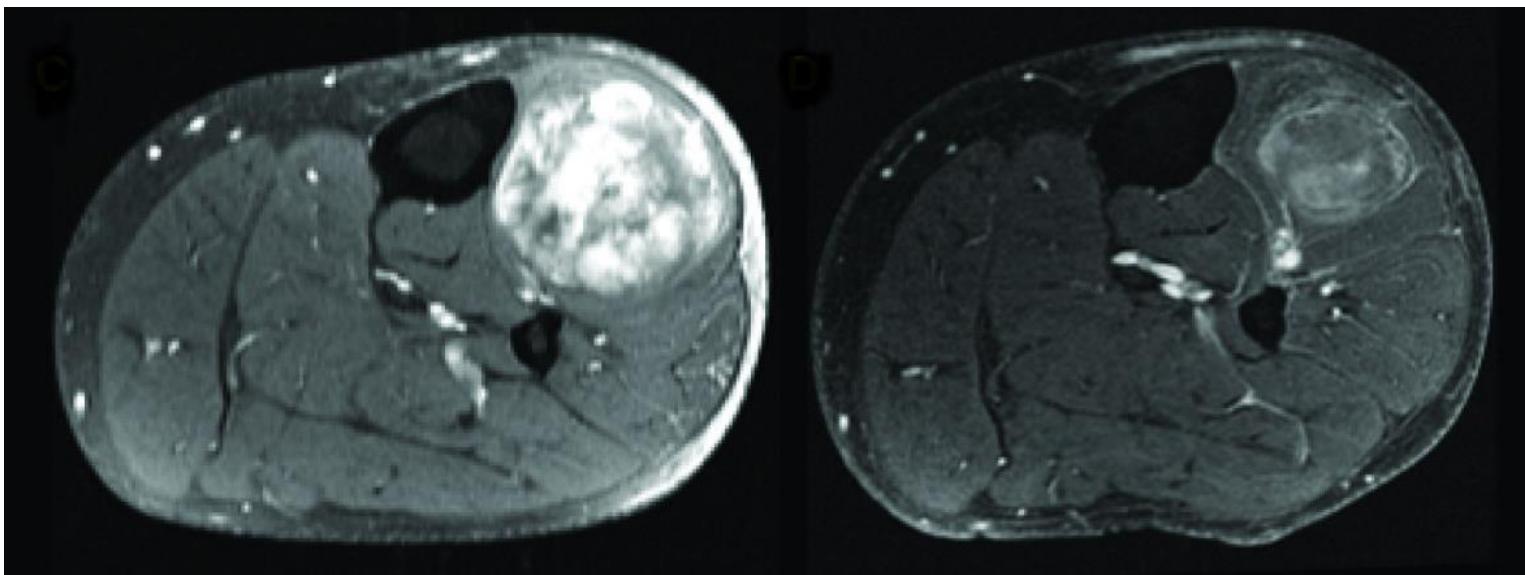
Codman's triangle

EWING'S SARCOMA

- * Peak incidence : II decade
- * Male predilection
- * Pain is the main symptom
- * 70% bone; 30% soft tissue
- * *EWSR1* gene rearrangement (~95%)
- * Chemosensitive tumour

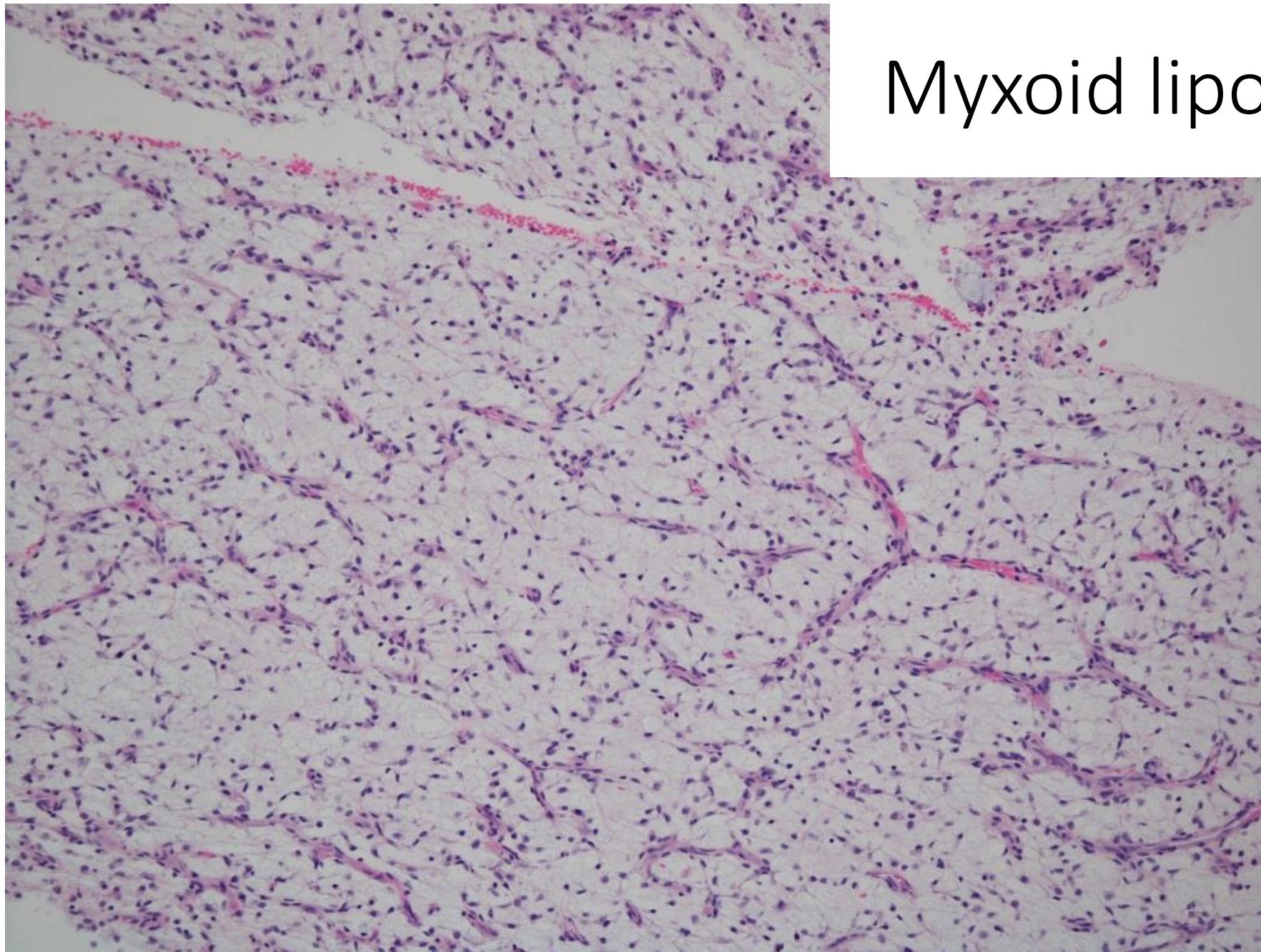






45yr old male – lump in
the leg

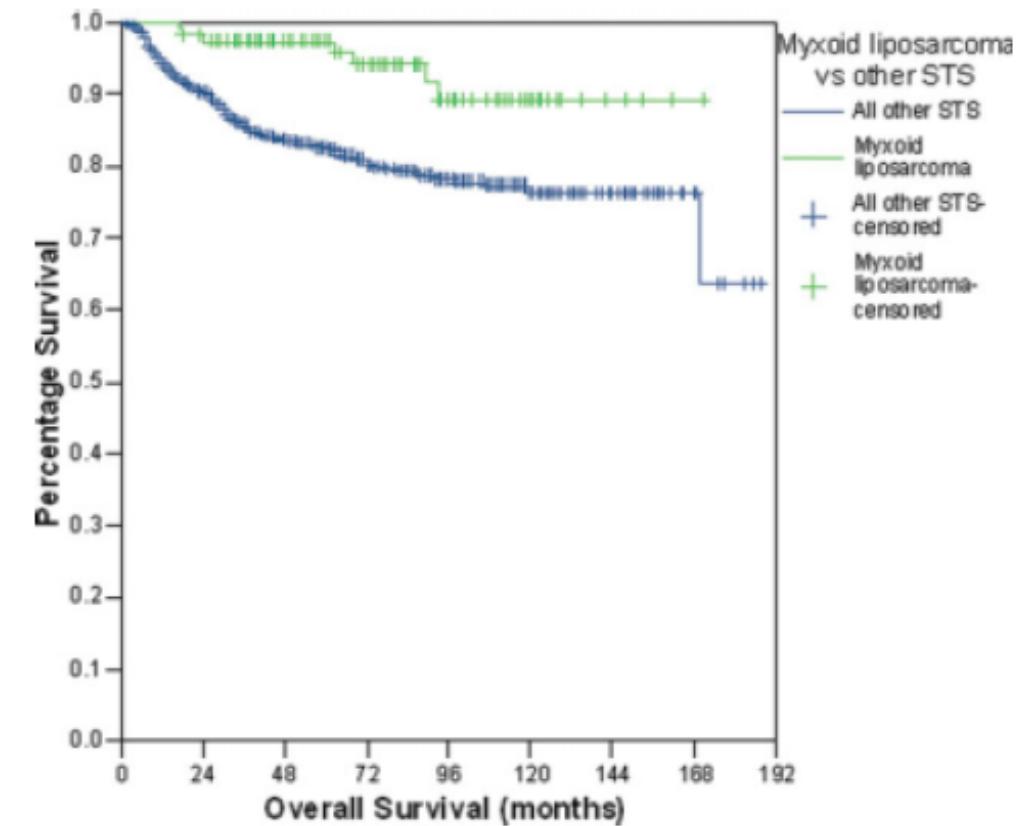
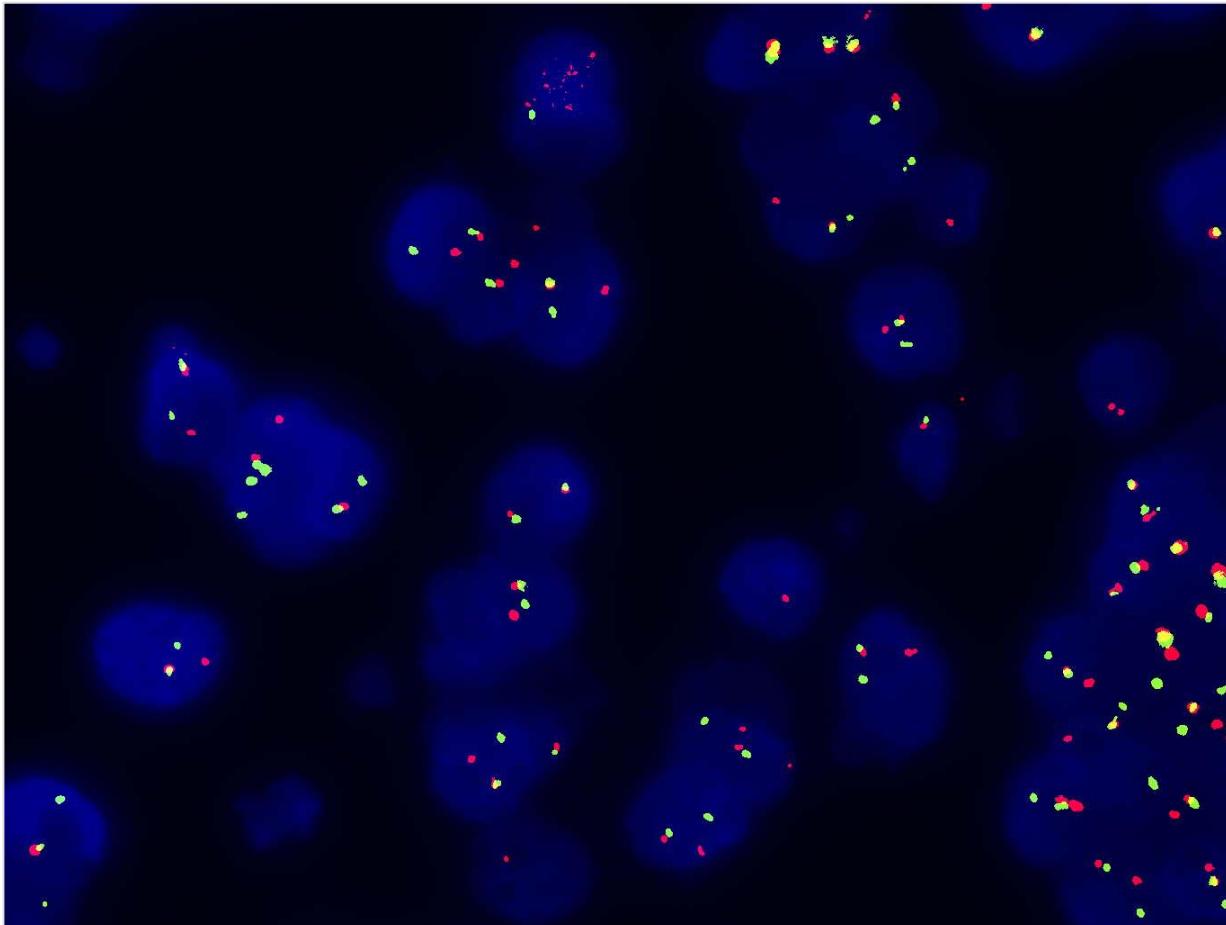
Myxoid liposarcoma



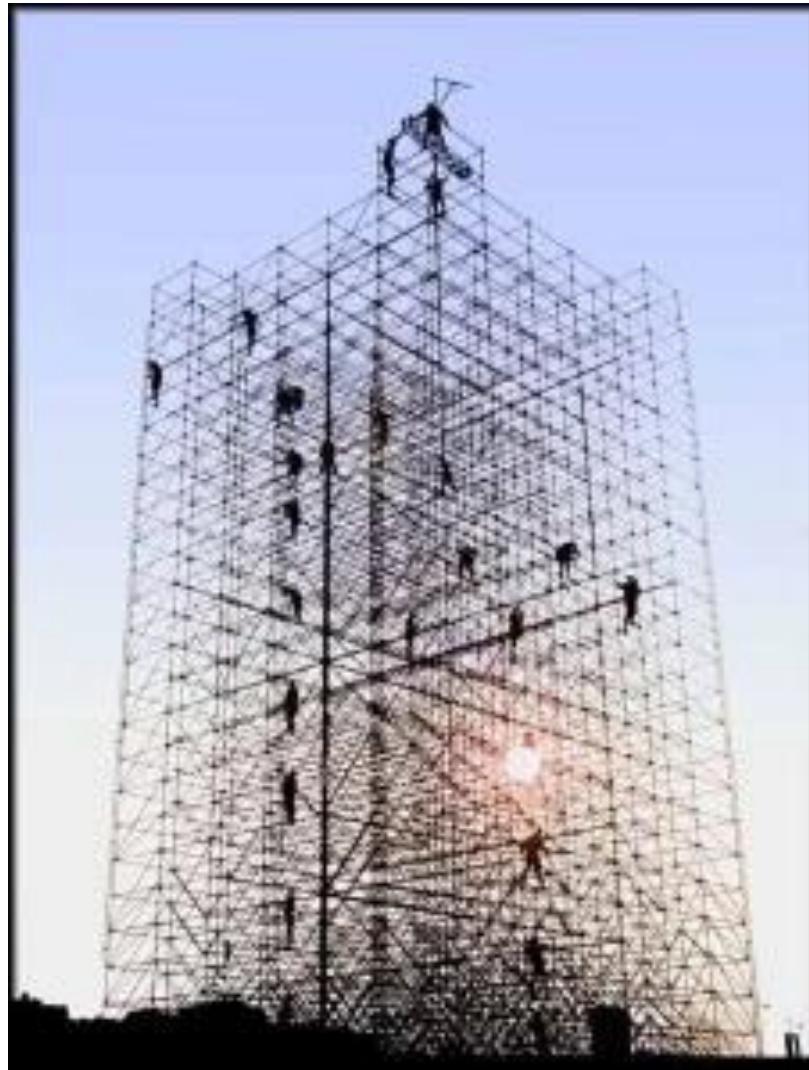
Myxoid/round cell liposarcoma

Break apart FISH – *DDIT3*

EWSR1-DDIT3 / FUS-DDIT3

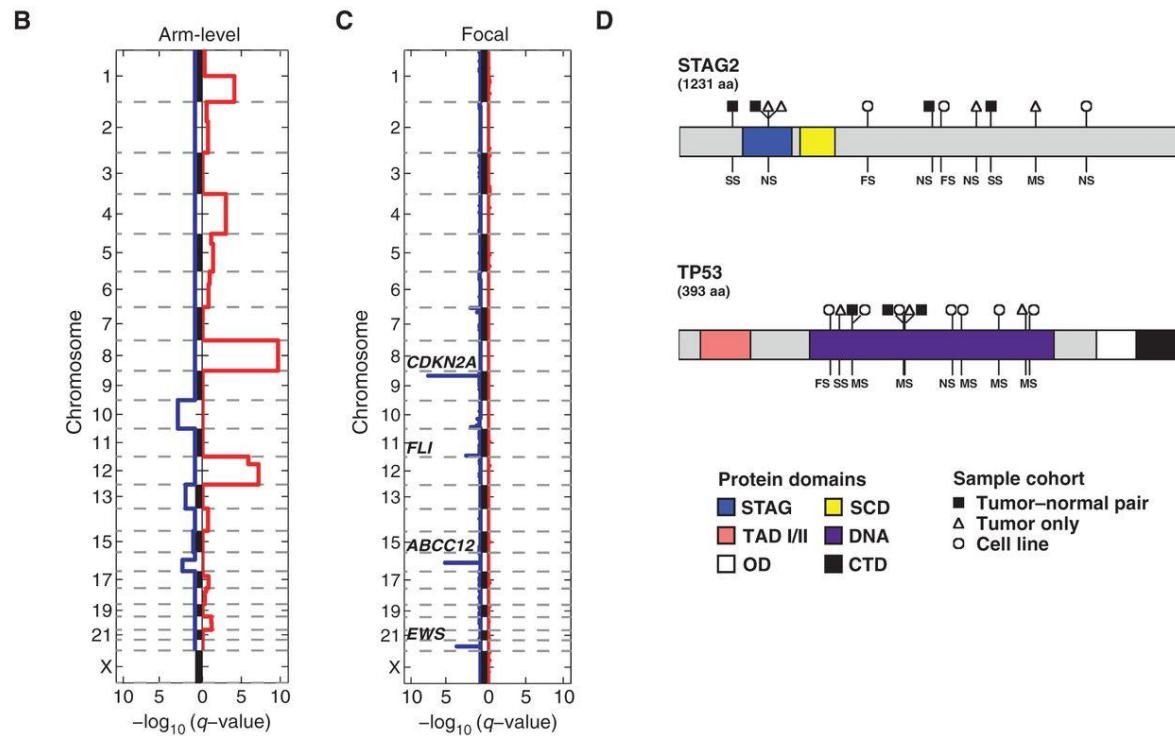
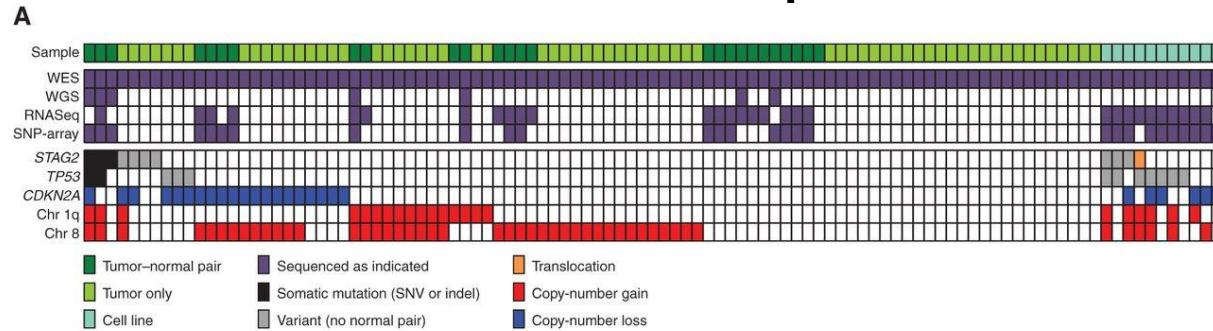


International Cancer Genome consortium/ TCGA

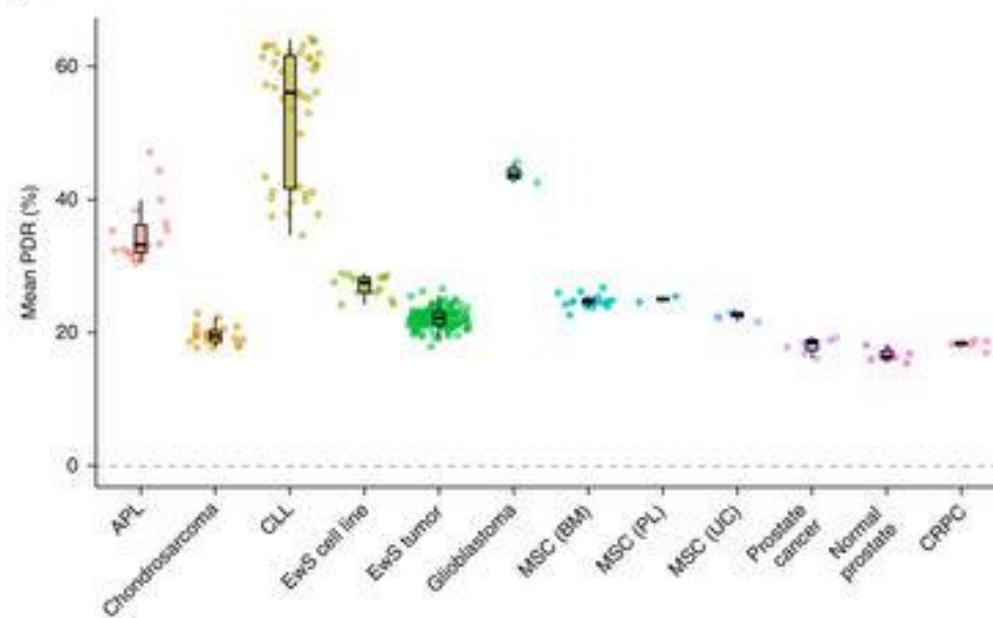
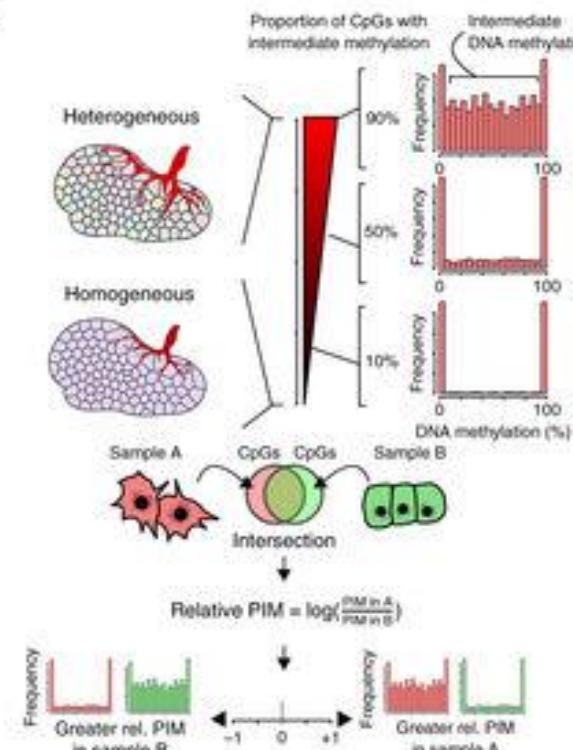


- Bone tumours:
 1. Chondrosarcoma: *IDH1/IDH2*
 2. Giant cell tumours of bone – *H3F3A* (G34W)
 3. Chondroblastoma: *H3F3B* (K36M)
 4. Osteosarcoma: IGF signalling
 5. Chordoma:
 6. Ewing's sarcoma – *STAG2, TP53, CDKN2A*

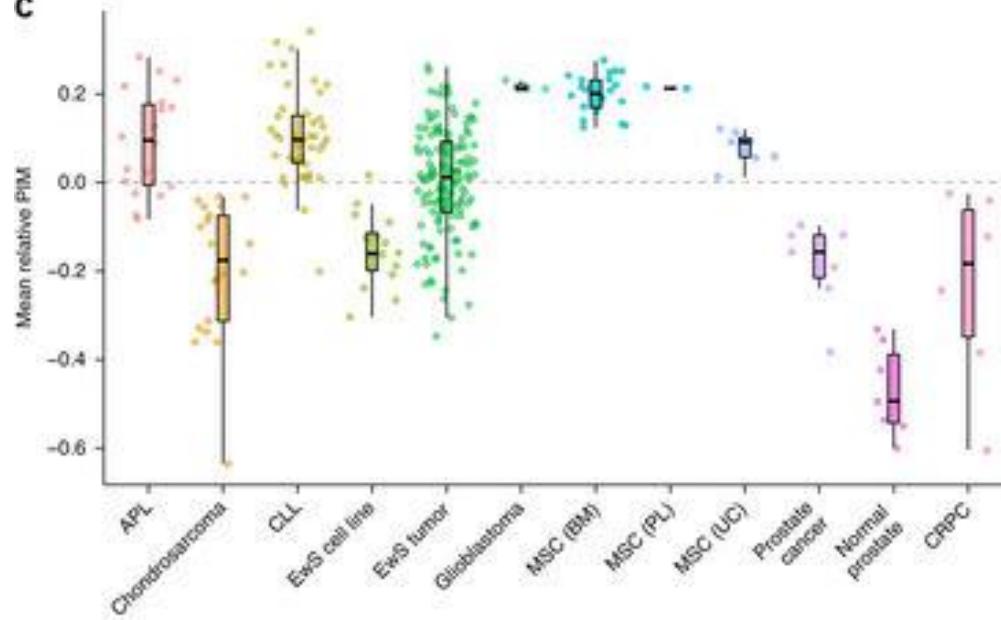
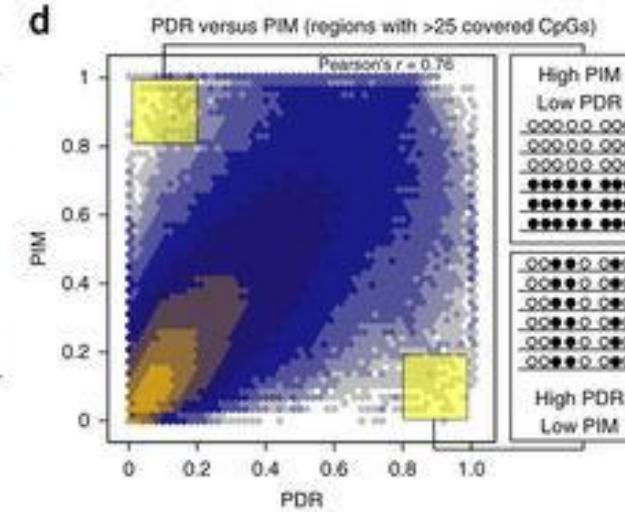
Genomic landscape of Ewing sarcoma



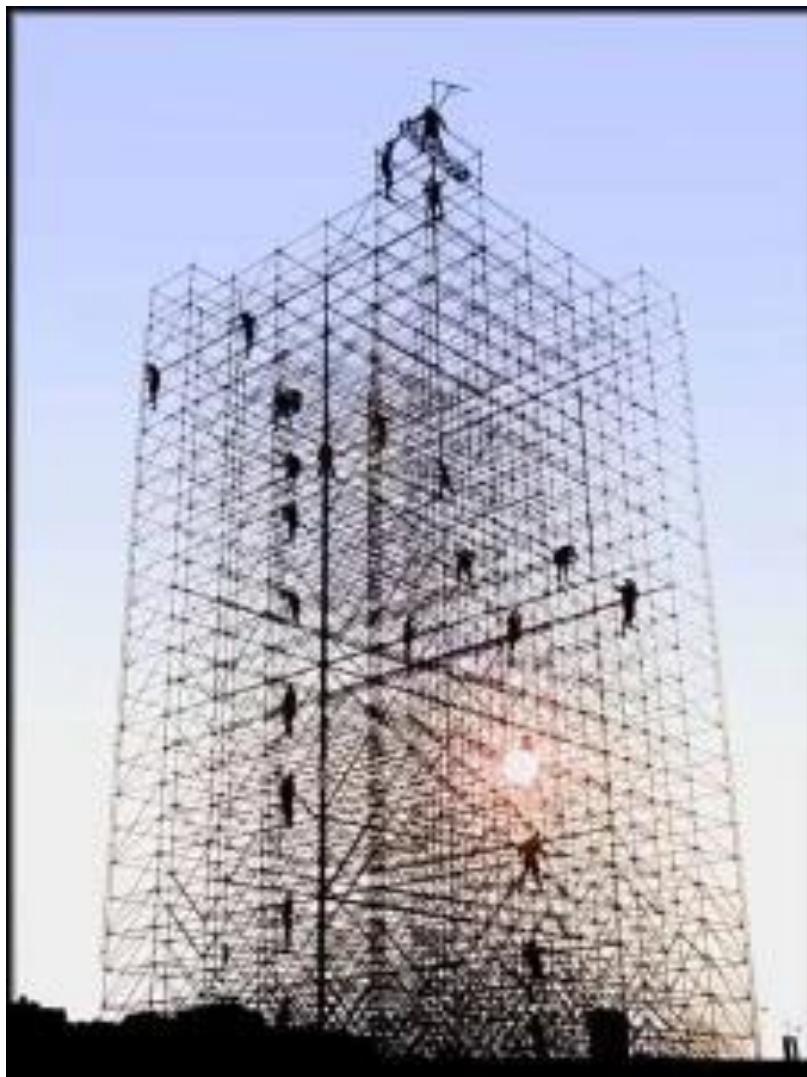
Cancer Discovery
Nov 2014, Vol 4,
Issue 1

a**b**

DNA methylation heterogeneity defines a disease spectrum in Ewing sarcoma

c**d**

The post TCGA, ICGC era



Output from genome sequencing studies

- Biological processes implicated in cancer development.
- Tumour heterogeneity.
- Evolution of metastasis.
- Mutational processes involved in carcinogenesis.
- Identification of drug targets.
- Outcome and response to therapy.
- Identification of new cancer genes.

Dramatic increase in our understanding of genomic events that characterise cancer but...

- 1) Clinical implementation of this knowledge to inform decision making is a major challenge.
- 1) Do not fully understand the interaction between molecular therapeutic agents and the genetic mutations they target...

[Recurrent BRAF Gene Fusions in a Subset of Pediatric Spindle Cell Sarcomas: Expanding the Genetic Spectrum of Tumors With Overlapping Features With Infantile Fibrosarcoma.](#)

Kao YC, Fletcher CDM, Alaggio R, Wexler L, Zhang L, Sung YS, Orhan D, Chang WC, Swanson D, Dickson BC, Antonescu CR.

Am J Surg Pathol. 2017 Sep 4. doi: 10.1097/PAS.0000000000000938. [Epub ahead of print]

[Recurrent BRAF Gene Rearrangements in Myxoinflammatory Fibroblastic Sarcomas, but Not Hemosiderotic Fibrolipomatous Tumors.](#)

Kao YC, Ranucci V, Zhang L, Sung YS, Athanasian EA, Swanson D, Dickson BC, Antonescu CR.

Am J Surg Pathol. 2017 Jul 7. doi: 10.1097/PAS.0000000000000899. [Epub ahead of print]

[Expanding the molecular signature of ossifying fibromyxoid tumors with two novel gene fusions: CREBBP-BCORL1 and KDM2A-WWTR1.](#)

Kao YC, Sung YS, Zhang L, Chen CL, Huang SC, Antonescu CR.

Genes Chromosomes Cancer. 2017 Jan;56(1):42-50. doi: 10.1002/gcc.22400. Epub 2016 Aug 26.

[Myopericytomatosis: Clinicopathologic Analysis of 11 Cases With Molecular Identification of Recurrent PDGFRB Alterations in Myopericytomatosis and Myopericytoma.](#)

Hung YP, Fletcher CDM.

Am J Surg Pathol. 2017 Aug;41(8):1034-1044. doi: 10.1097/PAS.0000000000000862.

PRC2 is recurrently inactivated through *EED* or *SUZ12* loss in malignant peripheral nerve sheath tumors

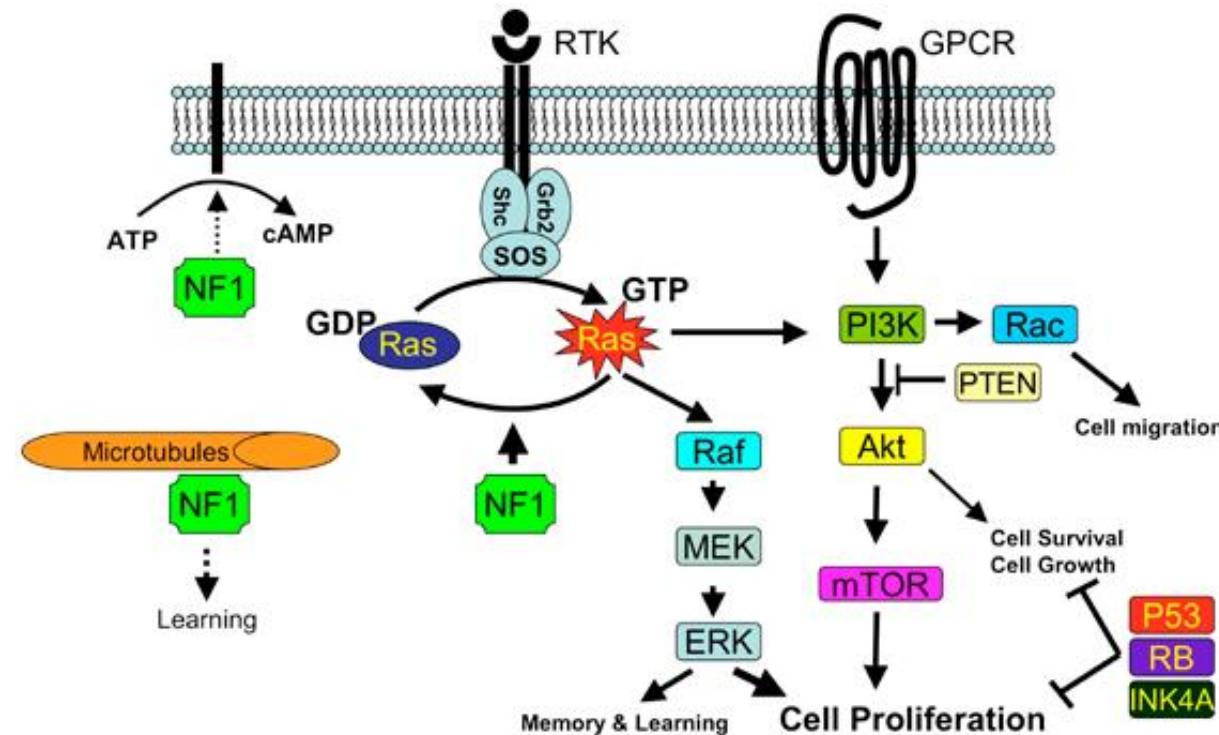
William Lee^{1,2,17}, Sewit Teckie^{2,3,17}, Thomas Wiesner^{3,17}, Leili Ran^{3,17}, Carlos N Prieto Granada⁴, Mingyan Lin⁵, Sinan Zhu³, Zhen Cao³, Yupu Liang³, Andrea Sboner^{6–8}, William D Tap^{9,10}, Jonathan A Fletcher¹¹, Kety H Huberman¹², Li-Xuan Qin¹³, Agnes Viale¹², Samuel Singer¹⁴, Deyou Zheng^{5,15,16}, Michael F Berger^{3,4}, Yu Chen^{3,9,10}, Cristina R Antonescu⁴ & Ping Chi^{3,9,10}

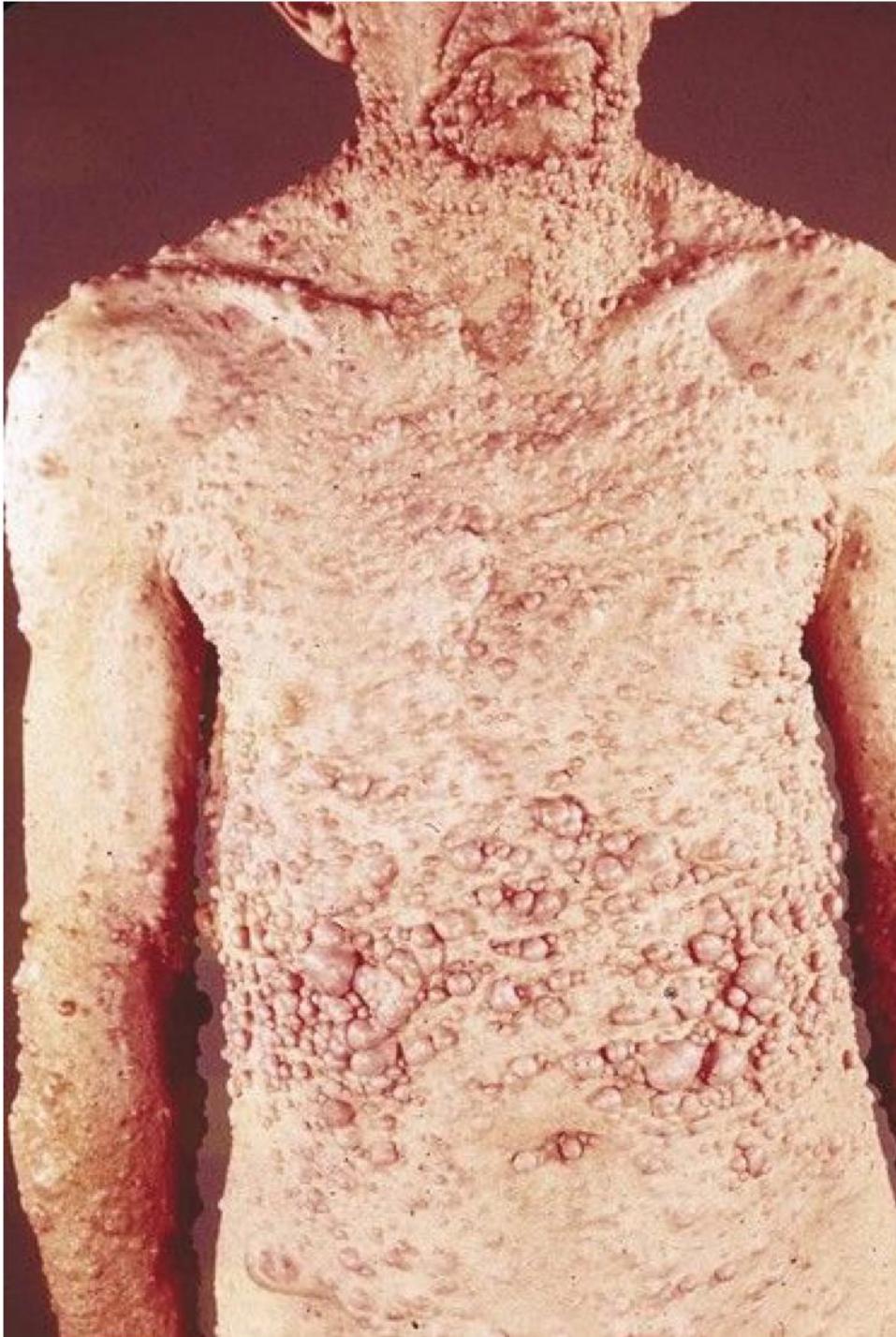
Somatic mutations of *SUZ12* in malignant peripheral nerve sheath tumors

Ming Zhang^{1,2}, Yuxuan Wang^{1,2}, Sian Jones³, Mark Sausen³, Kevin McMahon^{1,2}, Rajni Sharma⁴, Qing Wang^{1,2}, Allan J Belzberg⁵, Kaisorn Chaichana⁵, Gary L Gallia⁵, Ziya L Gokaslan⁵, Greg J Riggins⁵, Jean-Paul Wolinsky⁵, Laura D Wood⁴, Elizabeth A Montgomery⁴, Ralph H Hruban⁴, Kenneth W Kinzler^{1,2}, Nickolas Papadopoulos^{1,2}, Bert Vogelstein^{1,2} & Chetan Bettegowda^{1,2,5}

Neurofibromatosis Type I

- Common genetic disease.
- 1 in 3500 people. AD with high penetrance.
- NF1 – deletions, insertions, splice site mutations, mis-sense, non-sense mutations

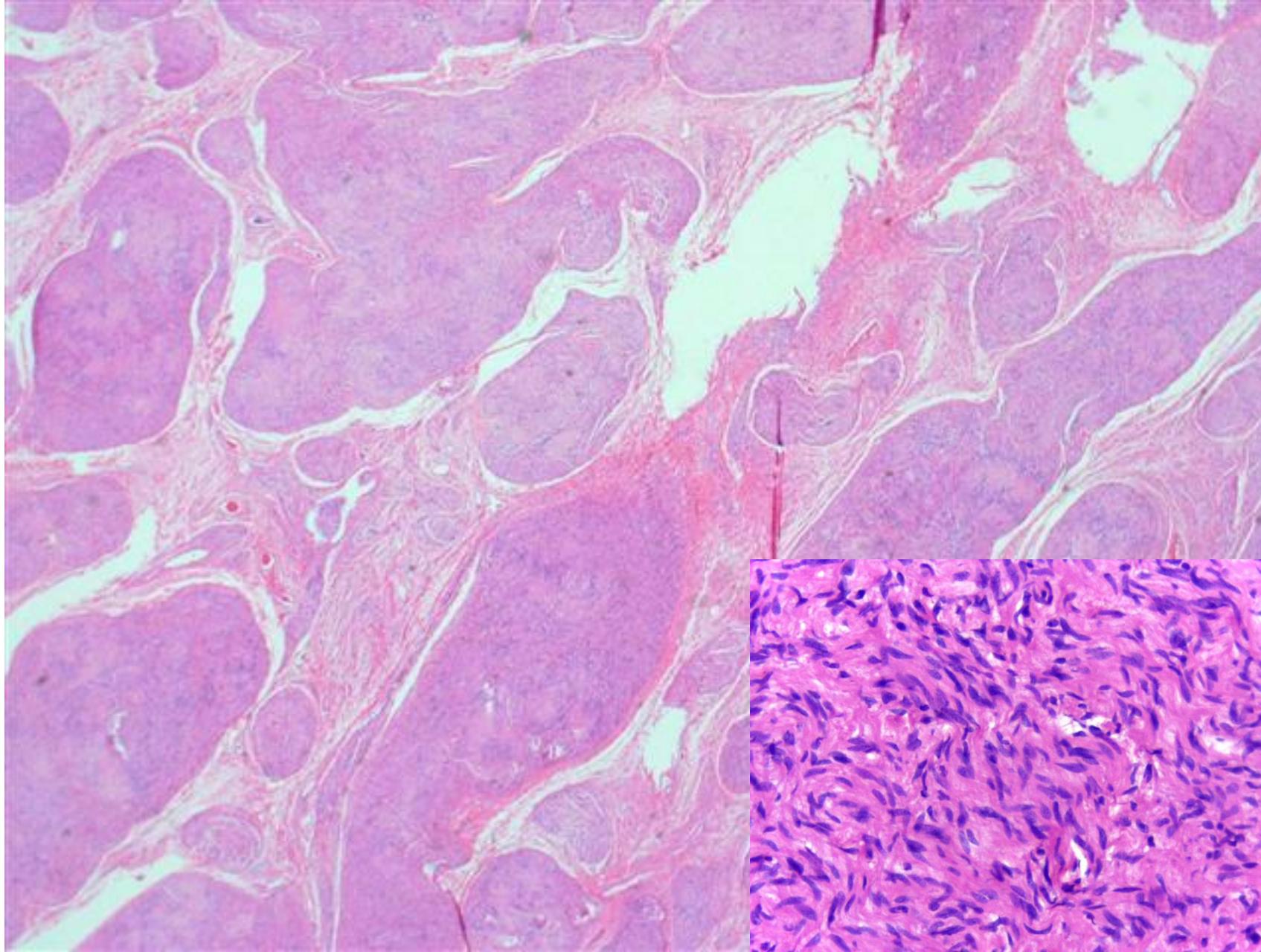




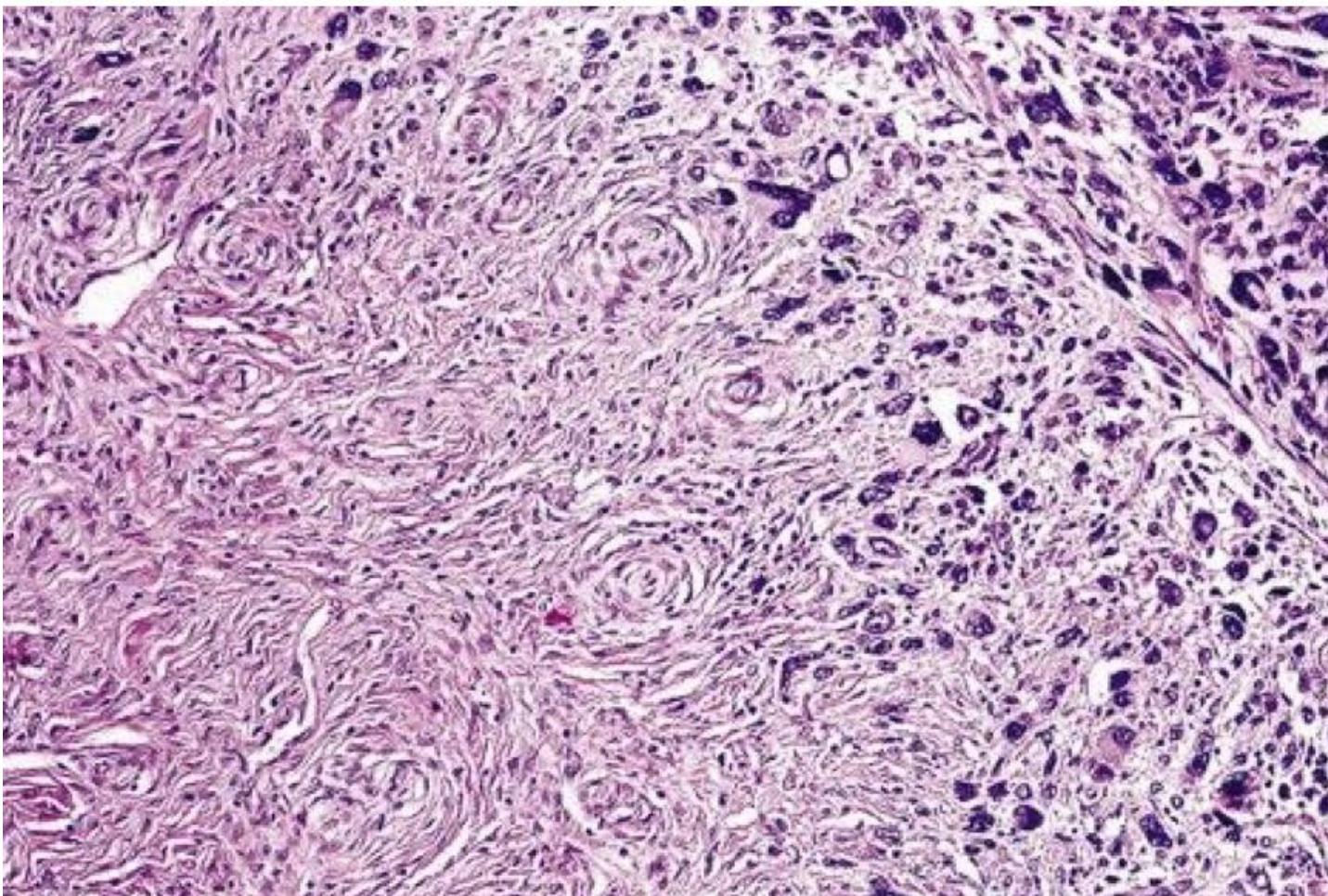
Enzinger and Weiss's Soft Tissue Tumors, 6th Edition

PLEXIFORM NEUROFIBROMA

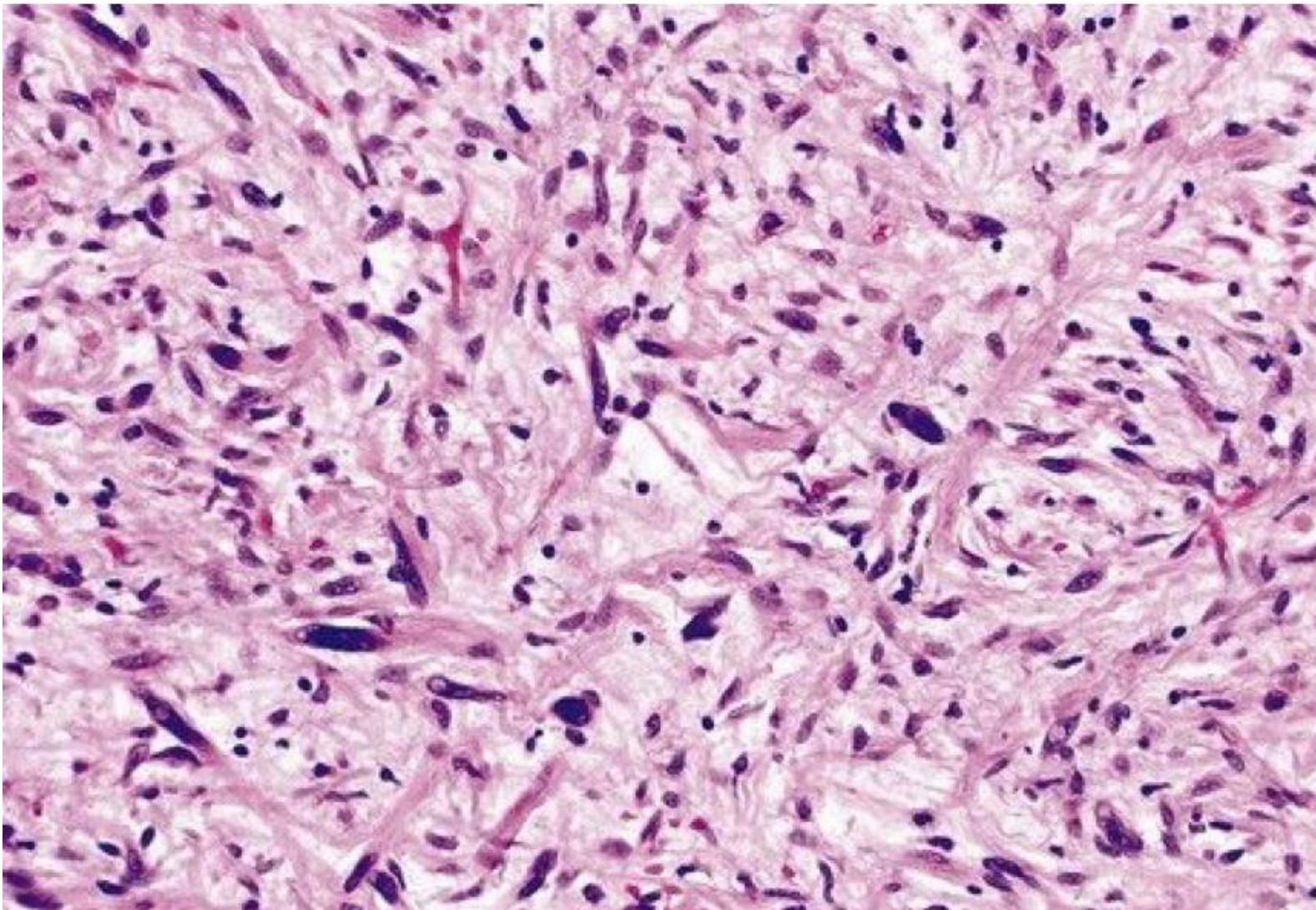




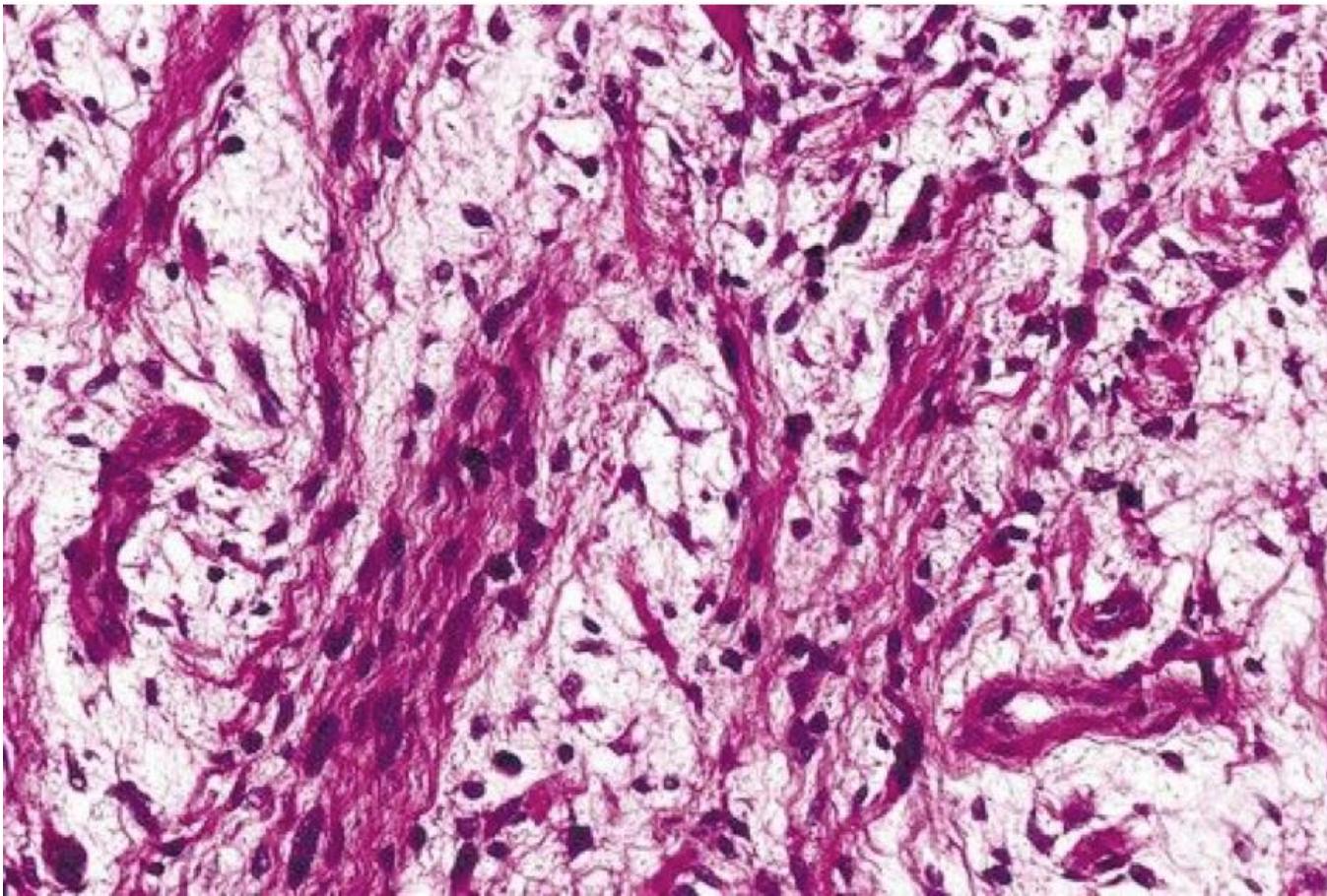
- Distinction between a neurofibroma with atypical features and MPNST Grade 1 is one of the most difficult – histological continuum.



“Atypical” neurofibroma

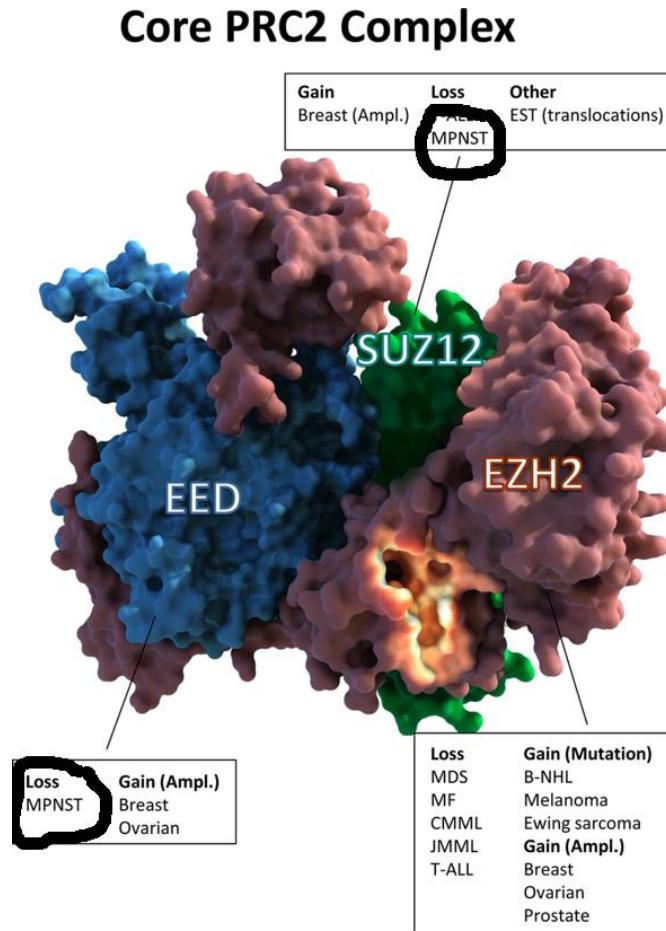


Low grade MPNST



Enzinger and Weiss's Soft Tissue Tumors, 6th Edition

MPNST have mutations in the PRC2 complex



- H3K27me3
- Transcriptional repression
-

Loss of H3K27me3 Expression Is a Highly Sensitive Marker for Sporadic and Radiation-induced MPNST

Carlos N. Prieto-Granada, MD,*† Thomas Wiesner, PhD,‡ Jane L. Messina, MD,† Achim A. Jungbluth, MD,* Ping Chi, MD, PhD,‡§|| and Cristina R. Antonescu, MD*

Am J Surg Pathol • Volume 40, Number 4, April 2016

TABLE 2. H3K27me3 Monoclonal Antibody IHC Results of the Different Entities Included in the MPNST Differential Diagnosis and Miscellaneous Tumors

Diagnosis	H3K27me3 IHC Loss/Total Cases
Cutaneous melanoma	
Pure desmoplastic melanoma	0/37
Mixed desmoplastic melanoma	0/11
Spindle cell melanoma	0/5
Synovial sarcoma (MF, BF, and PD)	0/113
GIST	
KIT/PDGFR α mutant	0/109
SDHB-deficient WT pediatric and adult	0/13
WT dedifferentiated GIST	0/1
Liposarcoma	
Well differentiated	0/31
Dedifferentiated	0/44
Ossifying fibromyxoid tumor	0/6
Soft tissue myoepithelial carcinomas	0/6
MFS	0/63

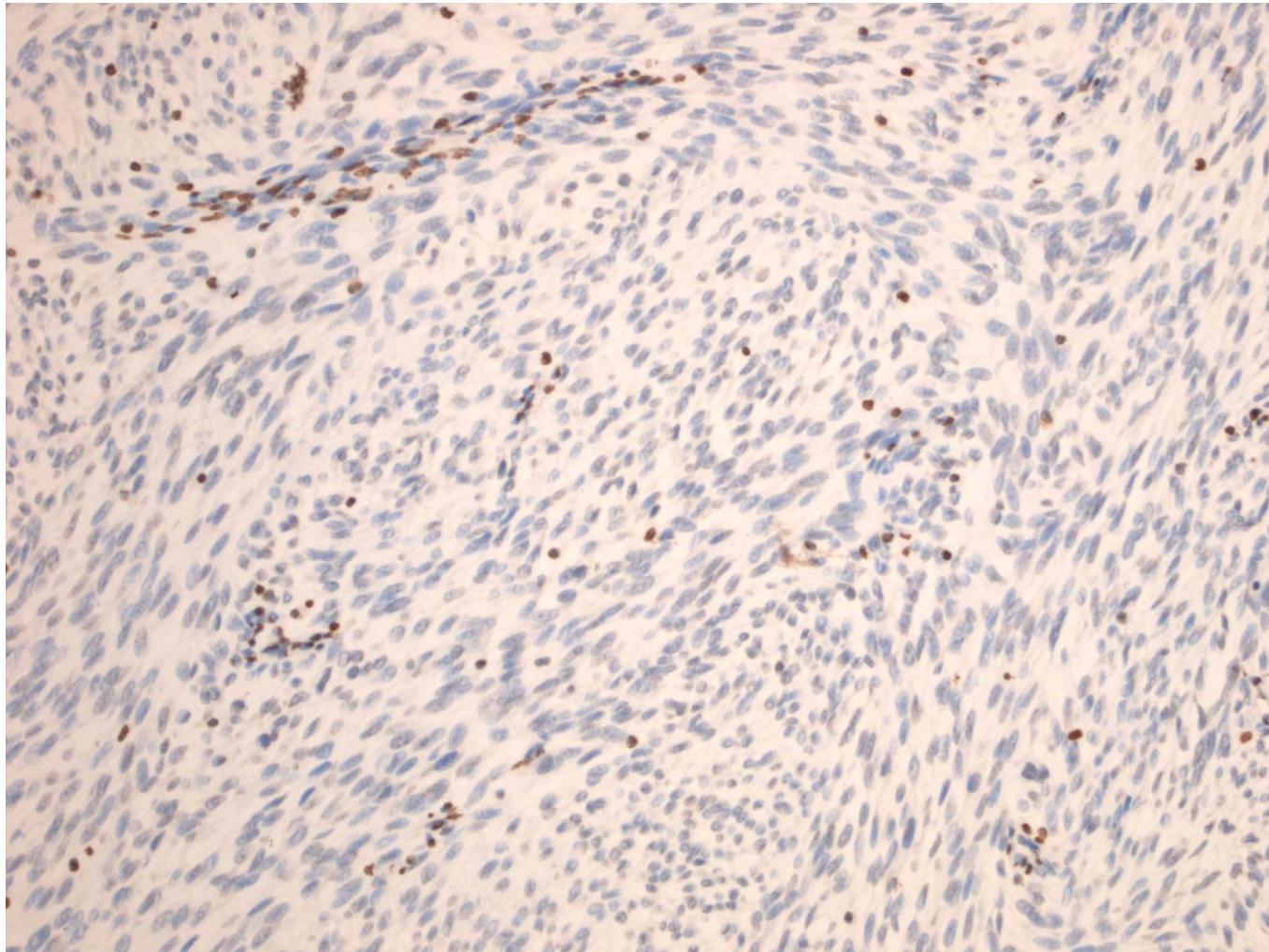
H3K27Me3 loss
in >90% of
MPNST

In the case of GIST, KIT and PDGFR α were WT.

BF indicates biphasic; MF, monophasic; PD, poorly differentiated.

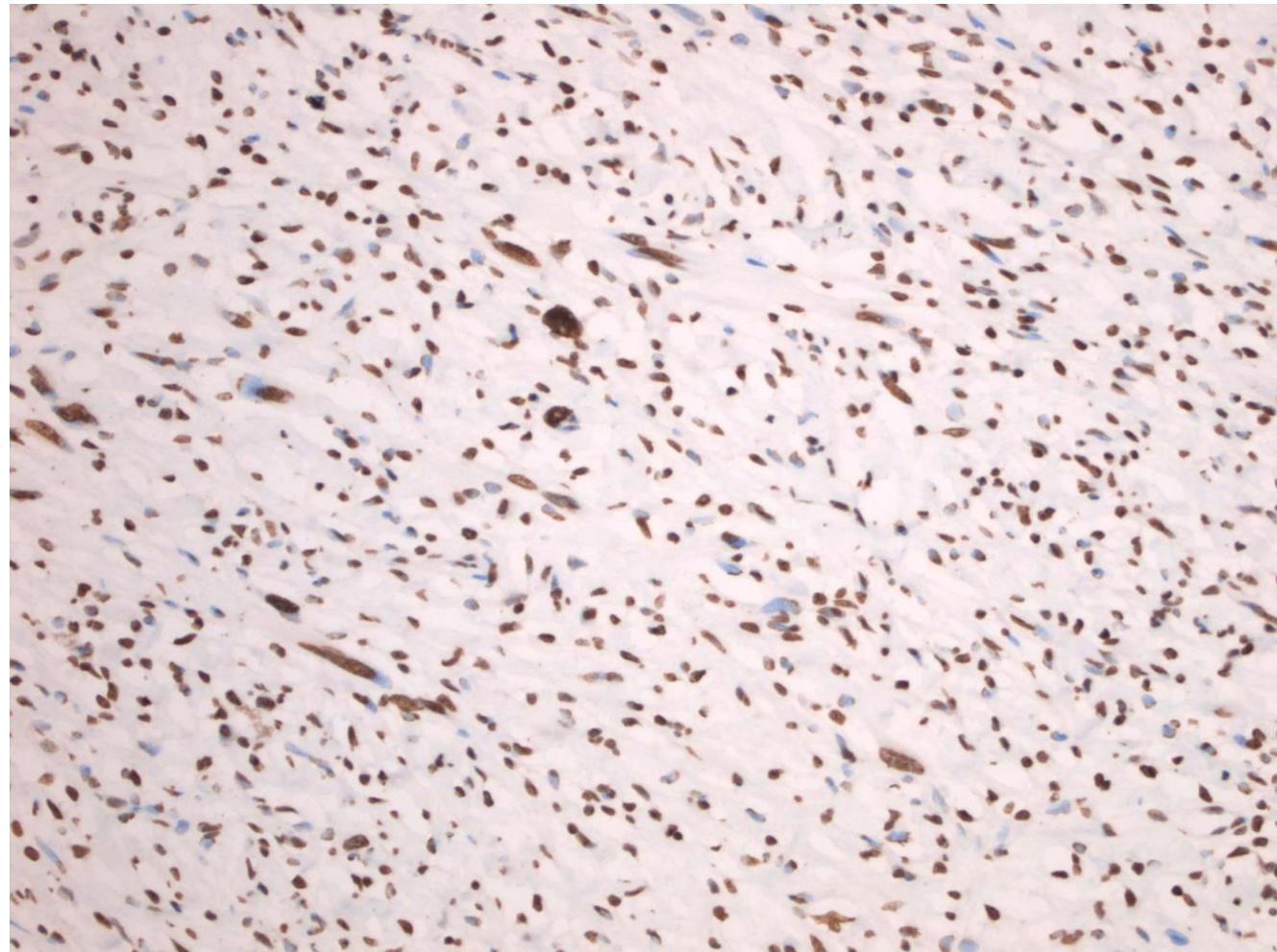
MPNST Grade 3

H3K27me3 -
immunohistochemistry



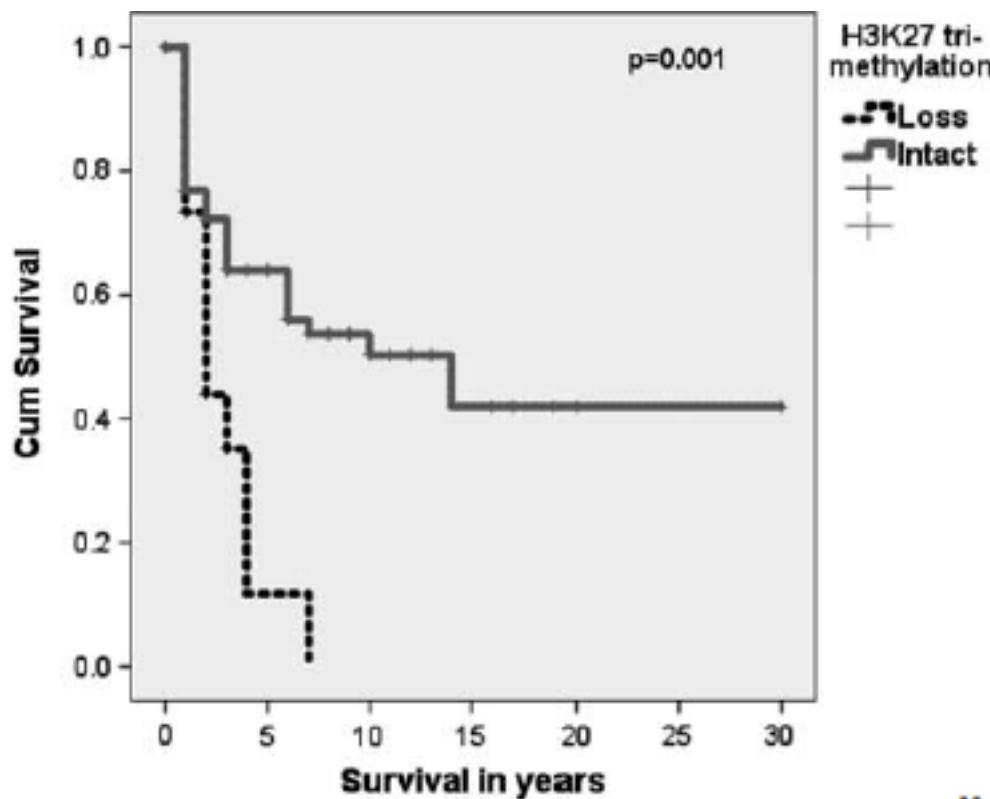
Atypical Neurofibroma

H3K27me3 -
immunohistochemistry



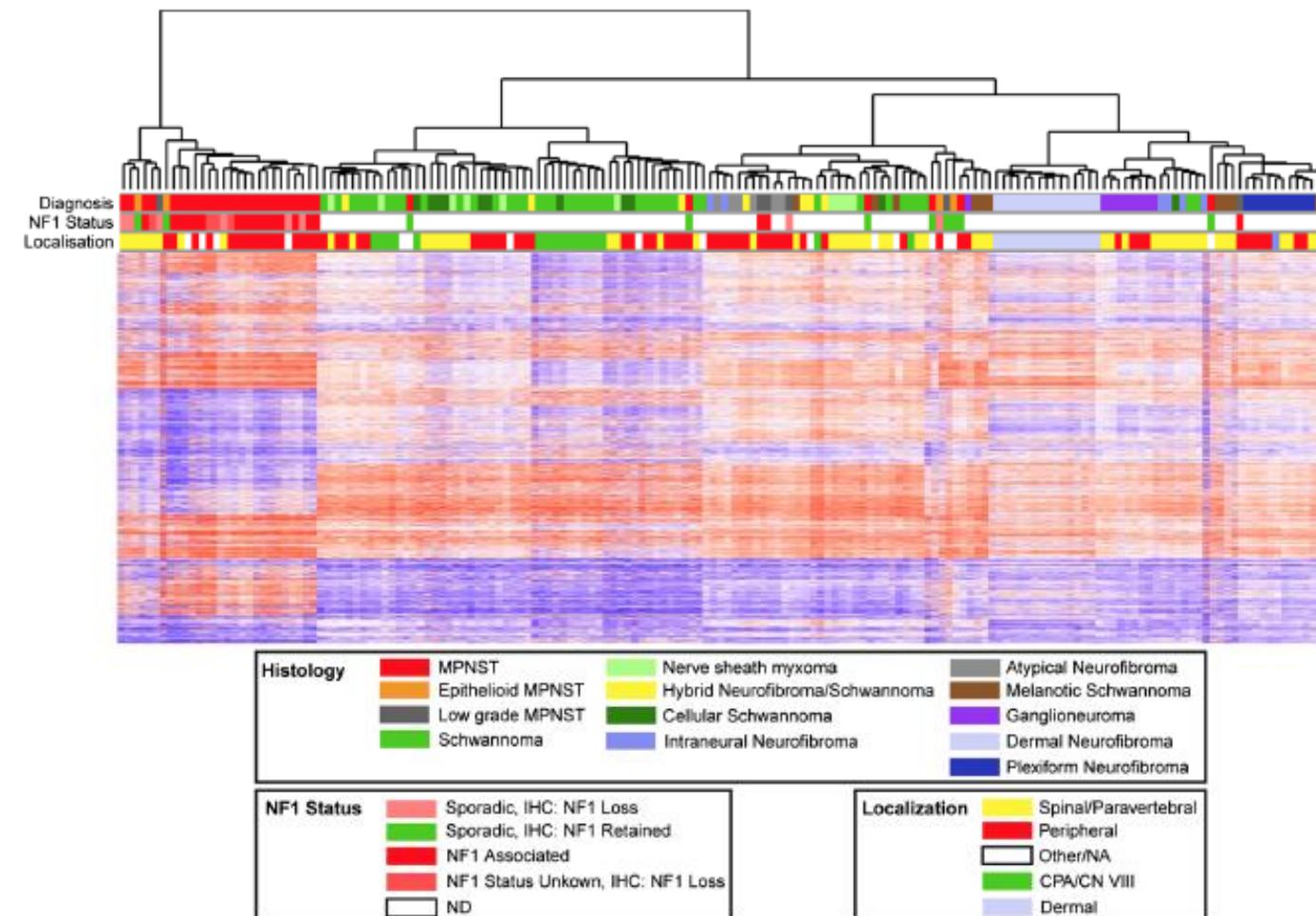
H3K27me3

- Prognostic utility



Methylation-based classification of benign and malignant peripheral nerve sheath tumors

Manuel Röhrich^{1,2} · Christian Koelsche^{1,2} · Daniel Schrimp^{1,2} · David Capper^{1,2} ·
Felix Sahn^{1,2} · Annekathrin Kratz^{1,2} · Jana Reuss² · Volker Hovestadt³ · David T. W. Jones⁴ ·
Melanie Bewerunge-Hudler⁵ · Albert Becker⁶ · Joachim Wels⁷ · Christian Mawrin⁸ ·
Michel Mittelbronn^{9,10} · Arlie Perry¹¹ · Victor-Felix Mautner¹² · Gunhild Mechtersheimer¹³ ·
Christian Hartmann¹⁴ · Ali Fuat Okuducu¹⁵ · Mirko Arp¹⁶ · Marcel Seitz-Rosenhagen¹⁶ ·
Daniel Hägggl¹⁶ · Stefanie Helm¹⁷ · Werner Paulus¹⁷ · Jens Schitzenhelm¹⁸ · Rezvan Ahmadi¹⁹ ·
Christel Herold-Mende¹⁹ · Andreas Unterberg¹⁹ · Stefan M. Pfister^{4,20} · Andreas von Deimling^{1,2} ·
David E. Reuss^{1,2}



MPNST

Machine learning and diagnostics

Train / Test Split		
Sarcoma Types	Train	Test
Synovial Sarcoma	7	3
Undifferentiated Sarcoma	35	15
Dedifferentiated Liposarcoma	44	19
Leiomyosarcoma(lms)	72	31
MPNST	6	3
Myxifibrosarcoma	15	7
Total	179	78

4.6 Summary of Classifier performances

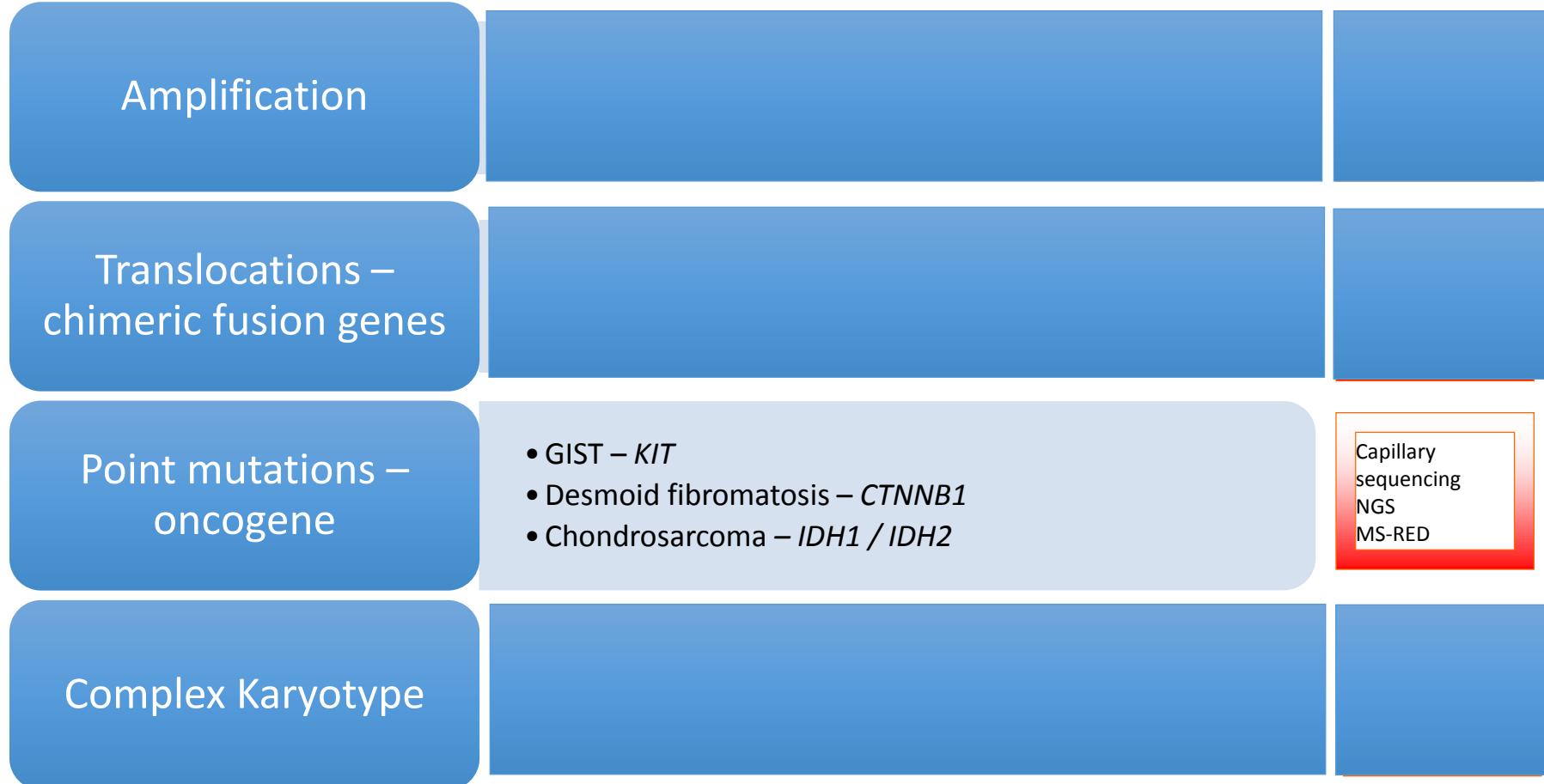
Type of Feature Reduction	Principal Component Feature Extraction	HighVariance Feature Selection	Chi Square	Chi Square	High Dimensional Data			McDeOon		
Type of Classifier	Logistic Regression	Logistic Regression	Logistic Regression	Support Vector Classifier	Support Vector Classifier	XGBoost	Random Forest	Support Vector Classifier	XGBoost	Random Forest
Classifier Performance Comparison	61.5	66.66	70.51	74.35	78.79	71.79	69.23	72.5	71.5	77.5

Figure 4.14: Summary of classification performance

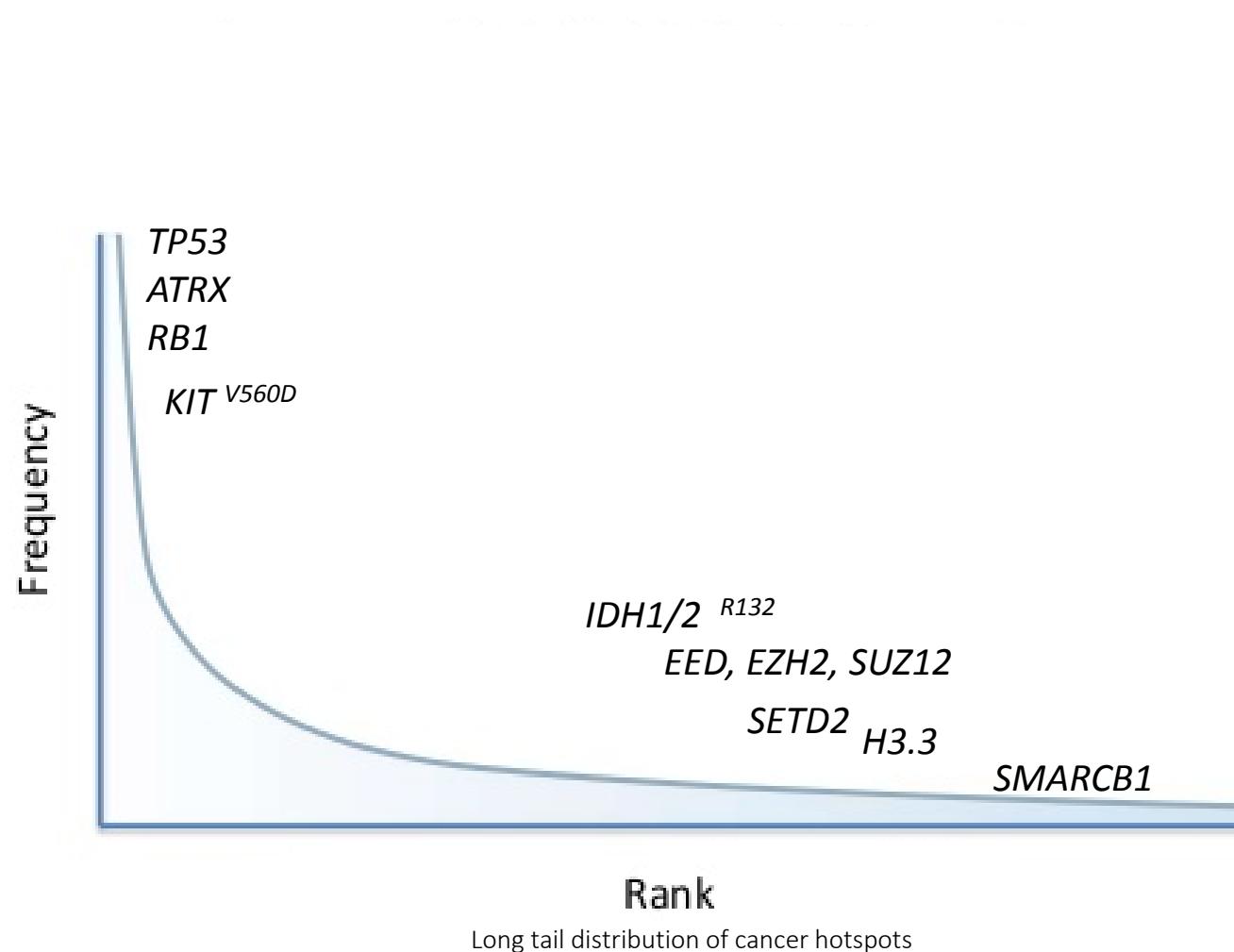
UCL MSc study – computer science

Manmohan Nair
Chris Steele

Molecular classification of mesenchymal tumours



Sarcoma hotspot mutations



Applications of molecular techniques

- Lipomatous tumours – ALT, Myxoid liposarcoma
- Small round blue cell tumours – Ewing sarcoma
- MPNST
- FISH
- Next generation sequencing
- Methylation profiling

Pathologist's role – beyond single gene mutations

- Era of immuno-oncology – need for biomarkers of response to checkpoint inhibition.
- - PD1/PDL1 expression, CTLA4
- Mutational burden

Chalmes et al. *Genome Medicine* (2017) 9:34
DOI 10.1186/s13073-017-0424-2

Genome Medicine

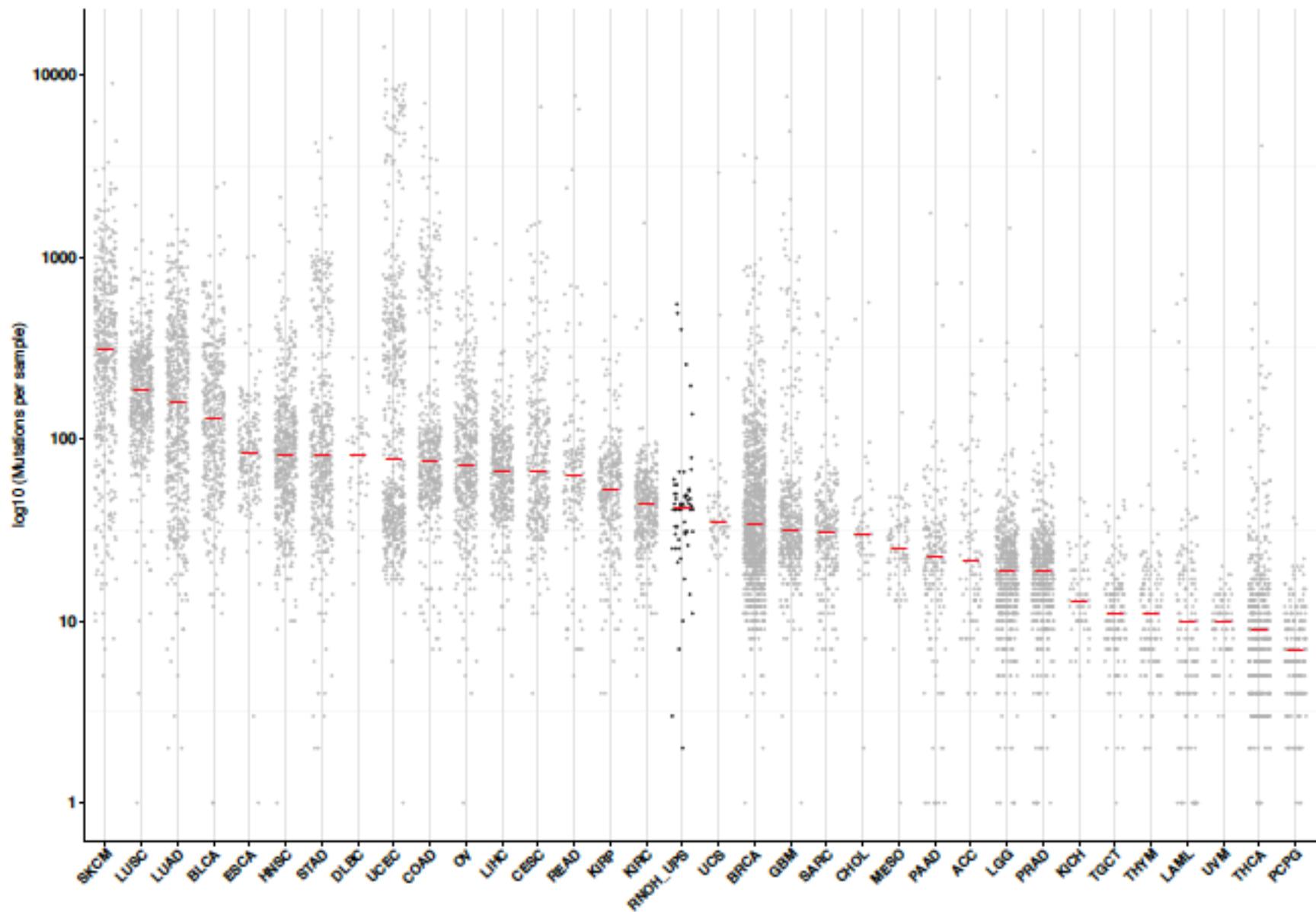
RESEARCH

Open Access



Analysis of 100,000 human cancer genomes reveals the landscape of tumor mutational burden

Mutational burden



Thanks for your attention.

