Chronic Lymphocytic Leukaemia Biology, genetics and prognosis

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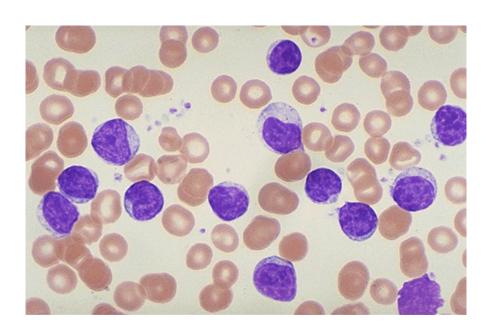




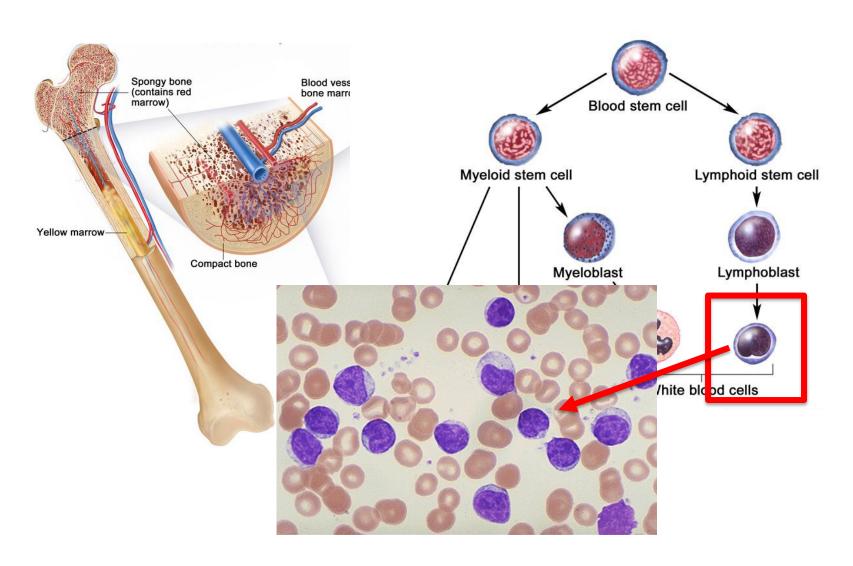


What is chronic lymphocytic leukaemia?

CLL is a low grade lymphoproliferative blood cancer caused by an accumulation of monoclonal B lymphocytes in the bone marrow, blood, lymph nodes and spleen



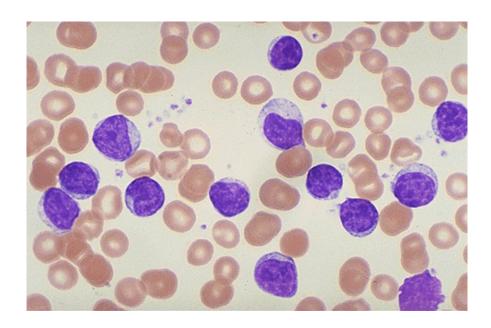
What is chronic lymphocytic leukaemia?



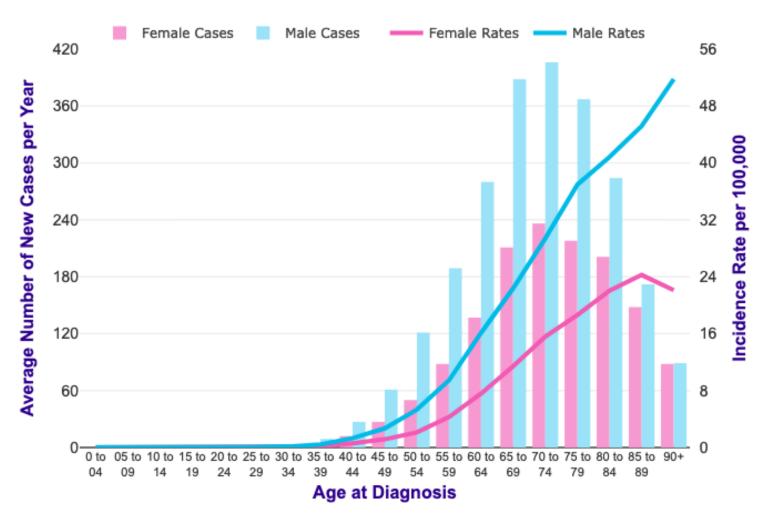
What is chronic lymphocytic leukaemia?

CLL is the most frequent leukaemia in Western countries including NZ

- Incidence of 4 per 100,000 population
- Median age 70 75; M > F

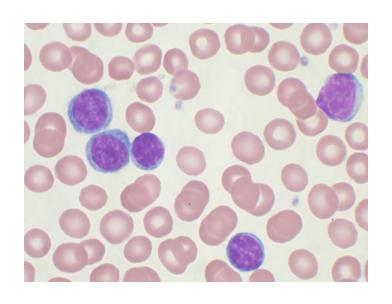


Age and Incidence of CLL

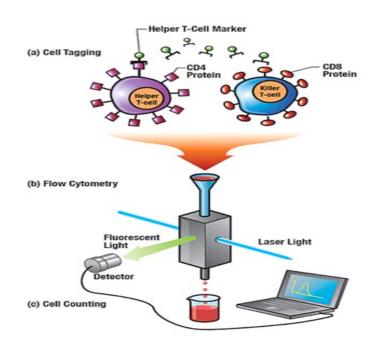


UK Cancer Registry 2015 - 17

Diagnosis of CLL



Full blood count and blood film analysis



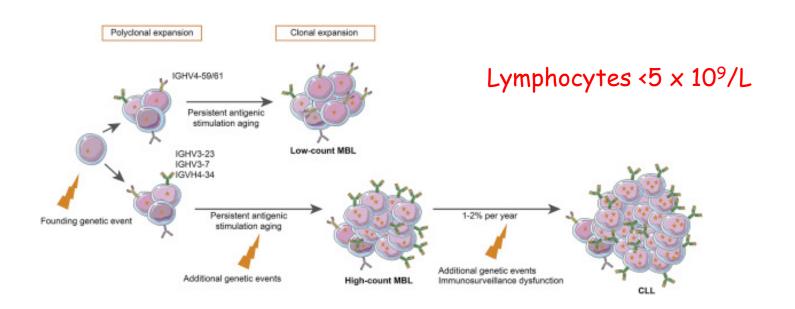
Cell marker studies "Immunophenotype"

Monoclonal B cell population with expression of CD5 and coexpression of B cell markers CD19, CD20, CD23

Diagnosis of CLL

All cases of CLL preceded by a pre-malignant phase called **Monoclonal B cell Lymphocytosis**

Present in 5% of the population age > 60, with 1-2% progressing to CLL each year



What causes CLL?

Increased understanding of the genomic landscape and the spelling mistakes / mutations in the genetic code of the CLL cells

A permissive microenvironment in the bone marrow and lymph nodes to support the CLL cells

No clearly established environmental risk factors identified to date

Increased risk of CLL in first degree relatives of patients

Is there a familial risk in CLL?

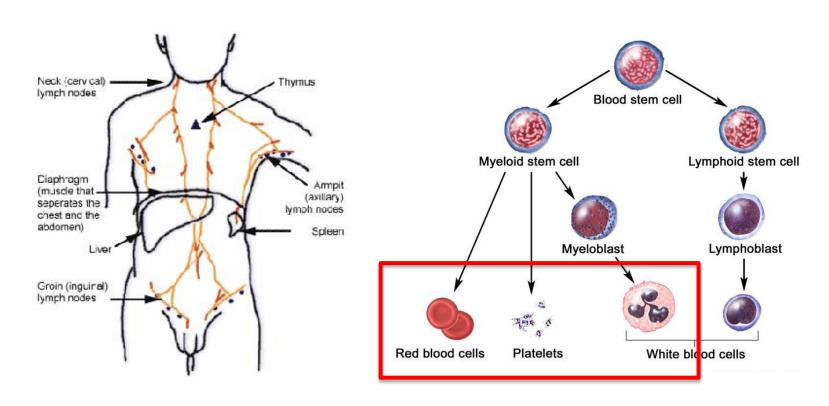
No single predisposition gene identified unlike breast and colon cancer

Likely to be interaction of multiple genes and variations within those genes

Risk for relatives remains very low

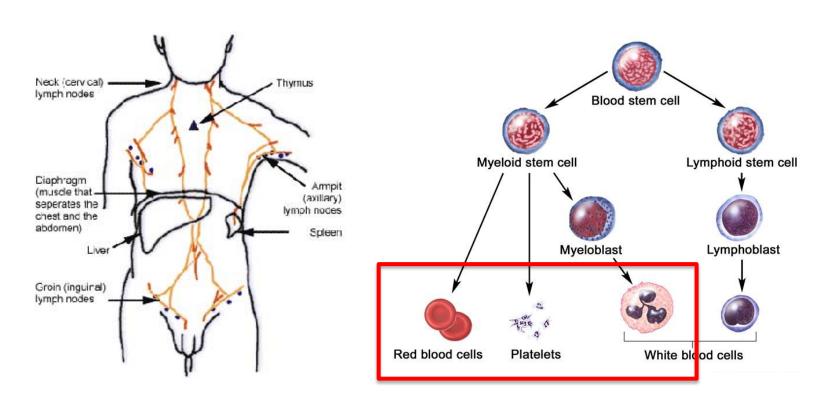


Presentation of CLL



Raised lymphocyte count in the peripheral blood Suppression of the normal blood counts (in red box) Enlarged lymph nodes and / or spleen

Presentation of CLL



70% of patients early stage of disease at presentation with no anaemia or thrombocytopenia and no significantly enlarged nodes

Predicting outcome in CLL

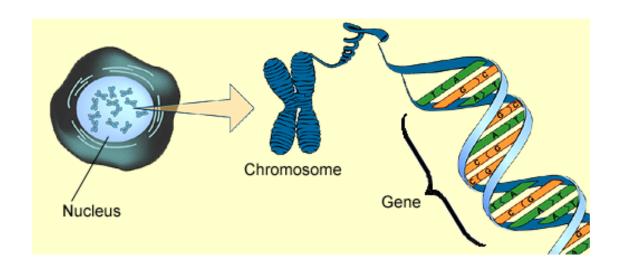
Recent discoveries in genetic alterations in CLL

The importance of the immunoglobulin gene mutation status

Incorporating these findings into new prognostic scores for patients

The increasing role of measuring minimal residual disease

Personalized Haematology



Chronic Lymphocytic Leukaemia
Improved understanding of the mutations / spelling
mistakes that occur in the genes of the CLL cells

Treatment decisions made based on the different mutations found in each case of CLL

FISH studies in CLL

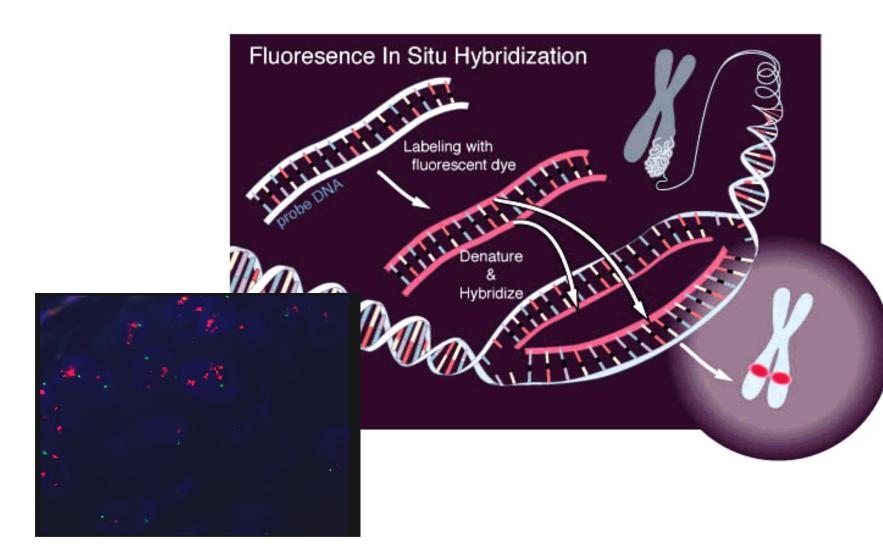
FISH is an acronym for ...

Fluorescence In Situ Hybridization



This technique exploits the ability of a fluorescent labelled DNA molecule to bind specifically to DNA

FISH studies in CLL



FISH studies in CLL

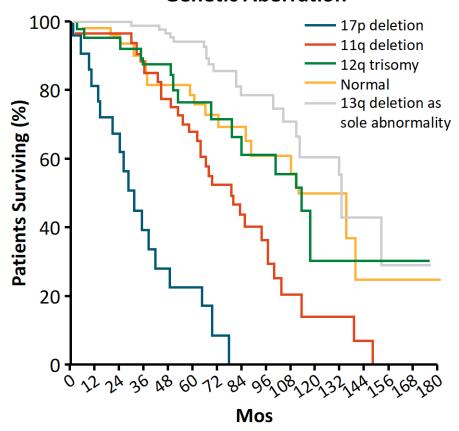
Four chromosomes analyzed:

- Del 13q
 55%
- Trisomy 12 20%
- Del 11q
 25%

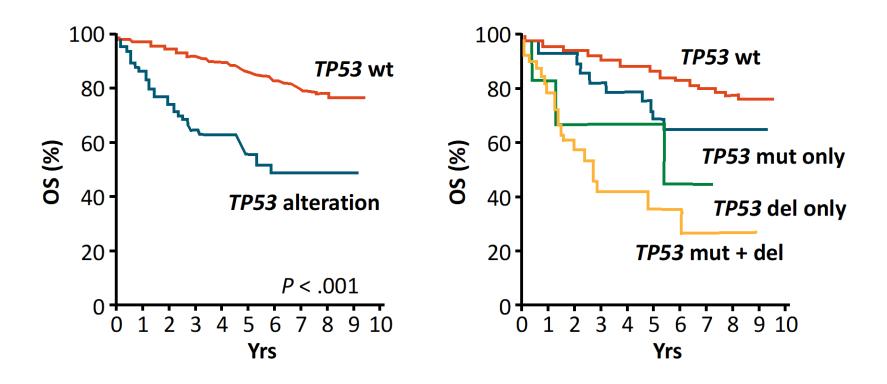


TP53 gene Add sequencing gene if FISH negative

Probability of OS From Diagnosis, by Genetic Aberration



Impact of TP53 mutations

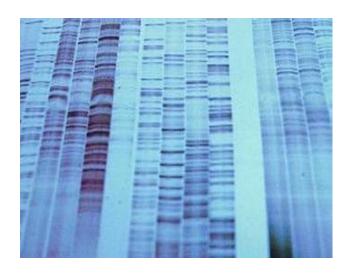


Poor response to conventional therapy eg FCR Good responses to novel agents Ibrutinib and Venetoclax Venetoclax funded by Pharmac

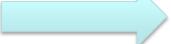
Progress in Genomic Technology

1990

2020



200 million fold



600 bases per day

120,000,000,000 bases per day

270 000 years per human genome

1 day per human genome

Progress in Genomic Technology

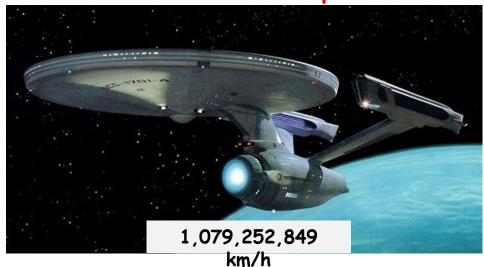


10 fold increase in speed

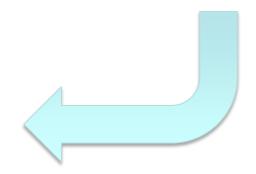




40 million fold increase in speed

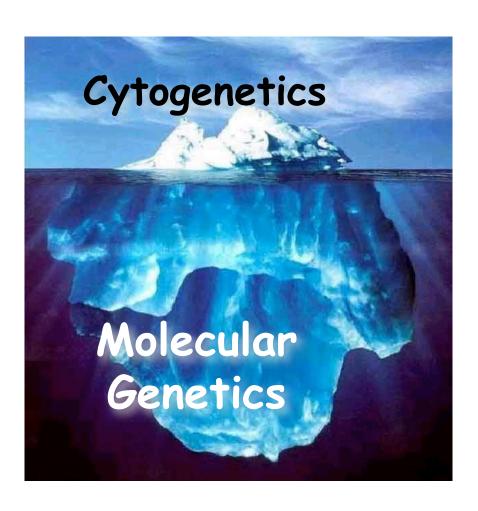




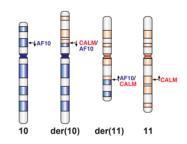


4 million fold increase in speed

Genetics of leukaemia



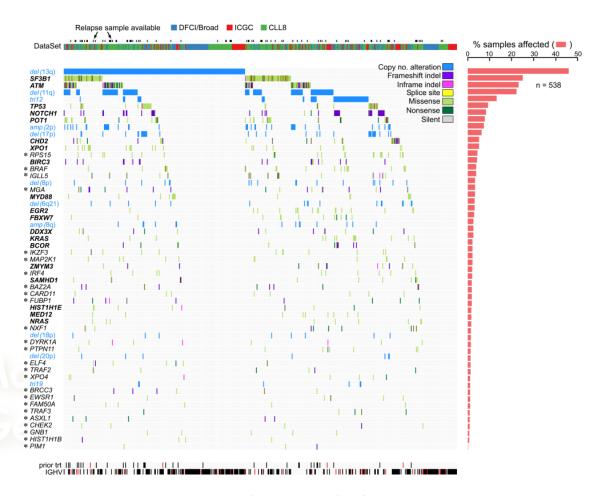
Translocations



Gene Sequencing



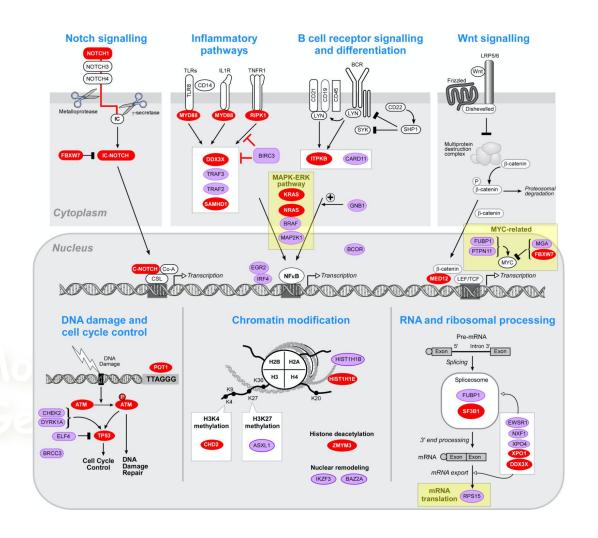
Genetics of CLL



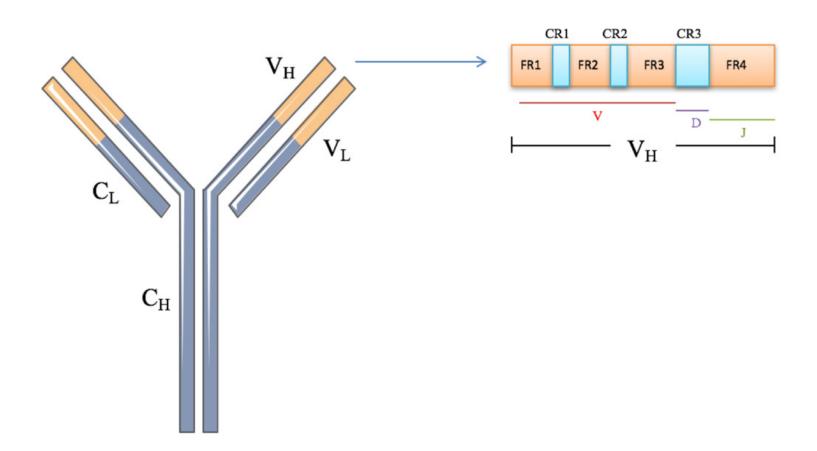
538 patients with CLL whole exome sequencing 44 recurrently mutated genes

Landau et al Nature 2015

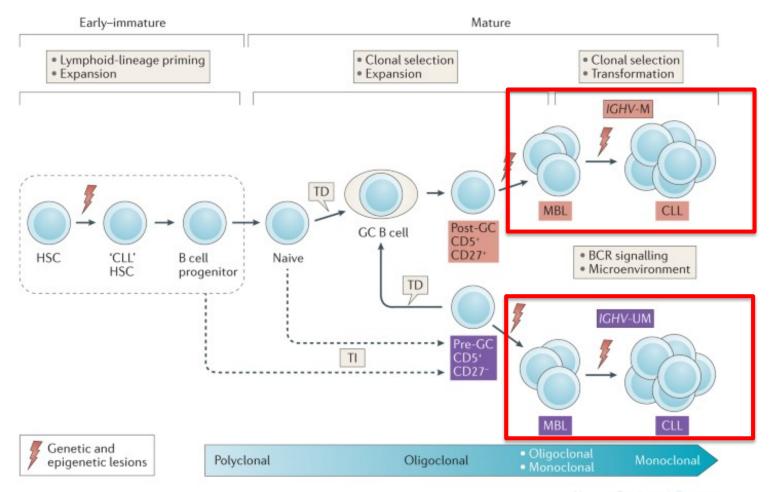
Genetics of CLL and Treatment



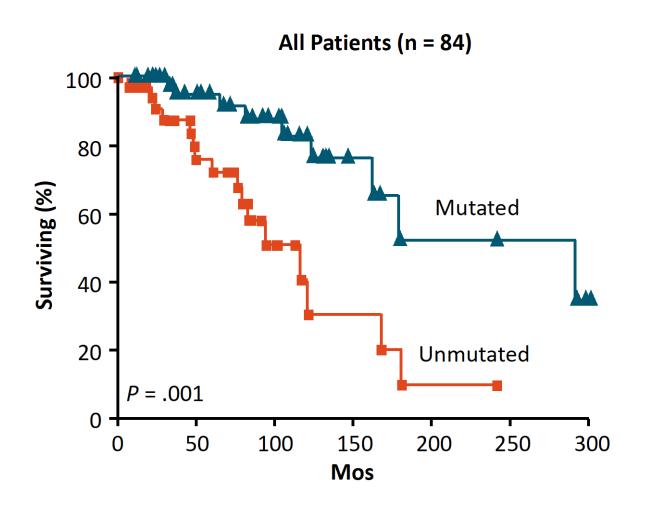
Impact of IGVH Mutation Status



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Impact of IGVH Mutation Status



Hamblin et al Blood 1999

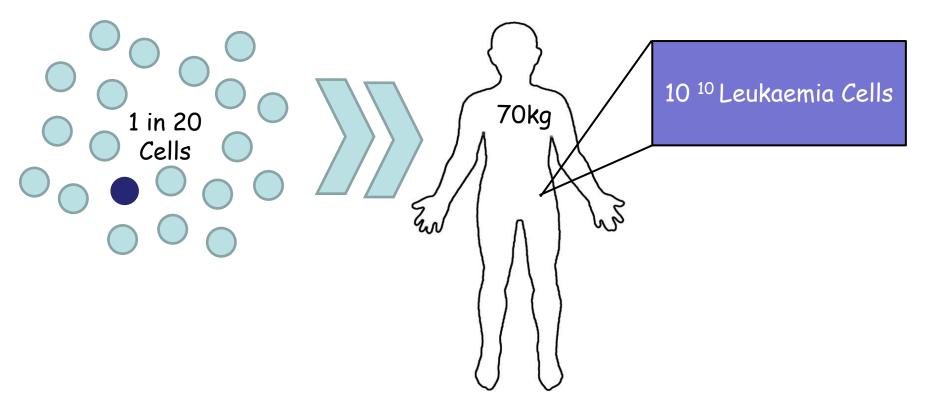
Integration of biologic markers into clinical staging

Prognostic factor	Points
Del17p on FISH or <i>TP53</i> mutation	4
Unmutated IGHV genes	2
Serum β2 microglobulin >3.5 mg/L	2
Rai stage I–IV	1
Age >65 years	1

Cumulative CLL- IPI score	Risk category	5-year TFS ^a
0–1	Low risk	78%
2–3	Intermediate risk	54%
4–6	High risk	32%
7–10	Very high risk	0%

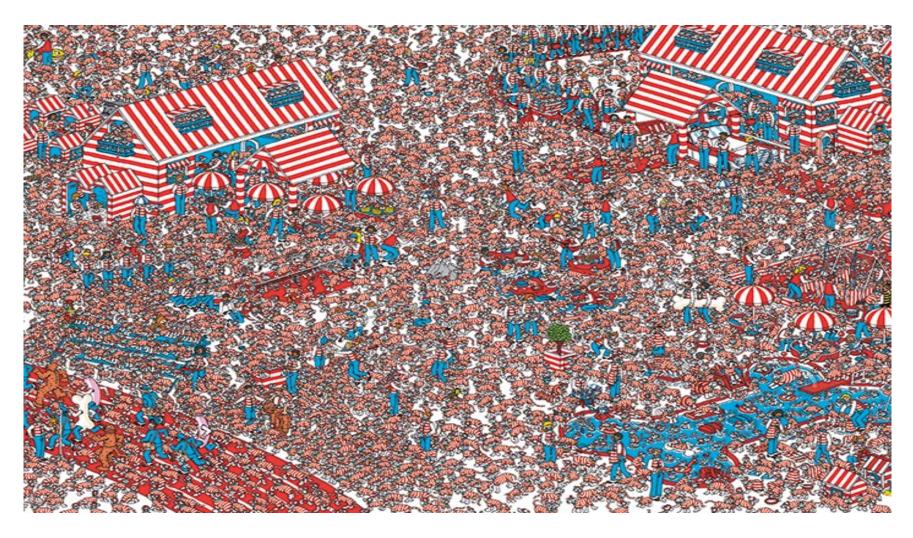
CLL-IPI Lancet Oncology 2016

Minimal residual leukaemia

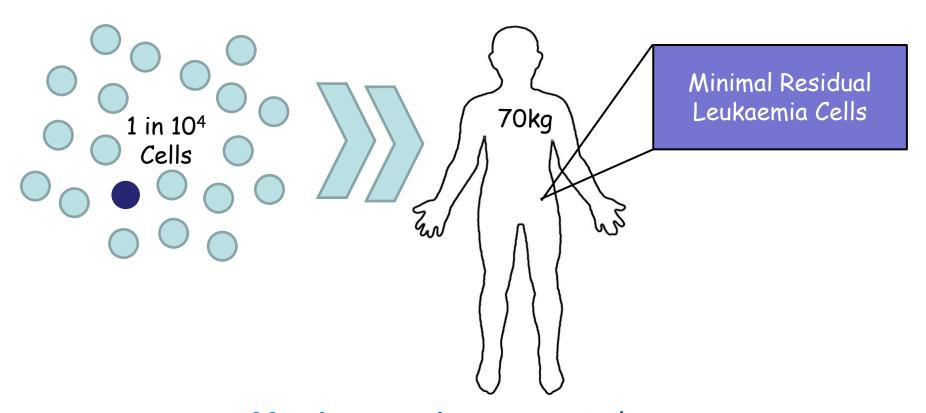


Remission is defined normal blood count, normal bone marrow and resolution of lymphadenopathy

Detection of Minimal Residual Disease (MRD)

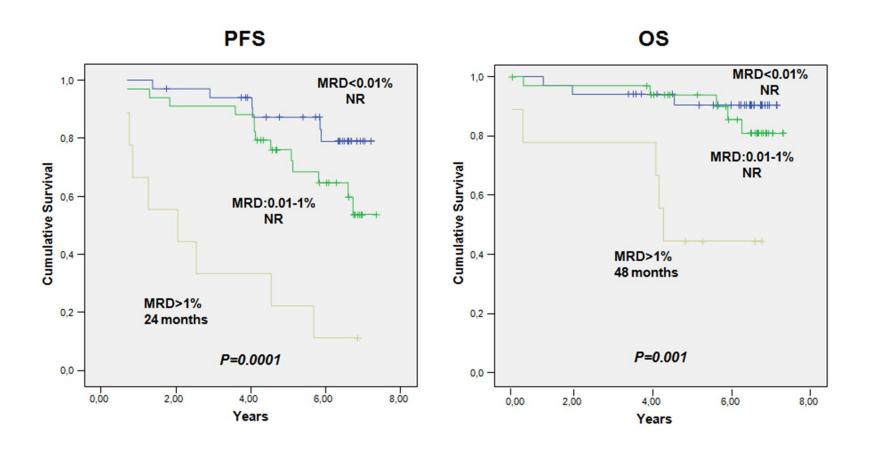


Detection of MRD in CLL by Flow Cytometry



Monitor patients post therapy Improved outcomes if MRD negative

Prognostic Impact of MRD in CLL



Biology of CLL Conclusions

Significant advances in understanding the genetics and biology of CLL

- Impact of TP53 status
- Impact of IGVH mutation status
- Emerging data from genomic sequencing studies

Moving into contemporary practice

Emerging role of MRD monitoring

Thank you

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Niloofar Zandvakili

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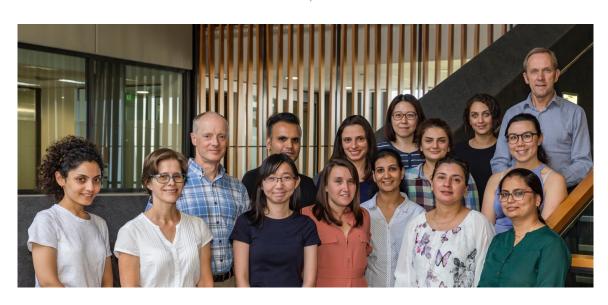
Lachlan Macdonald

Jessica Chase

Christina Walker Chloé Morin



our mission is to care, our vision is to cure





The Family of Marijana Kumerich