Routine Molecular Diagnosis of Cancer in Clinical Oncology



"I'm afraid you've got cows, Mr. Farnsworth."

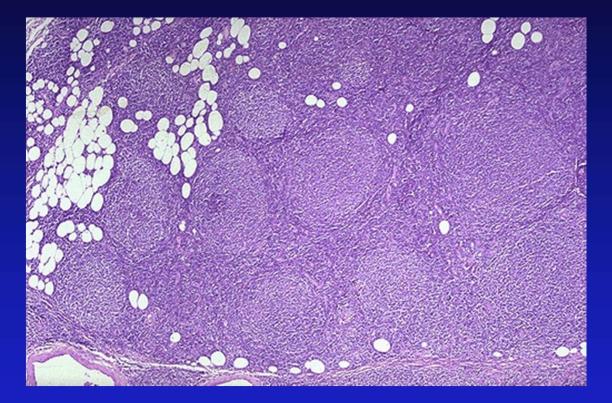
1. To provide reproducible, quantitative diagnoses for all cancer patients.

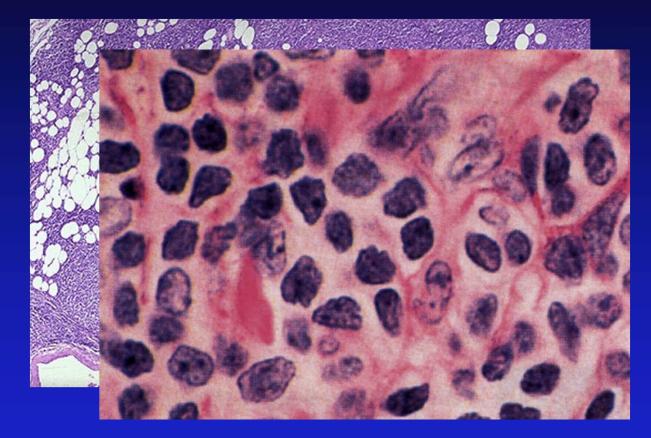
- 1. To provide reproducible, quantitative diagnoses for all cancer patients.
- 2. To clarify diagnostic distinctions that are problematic using current methods.

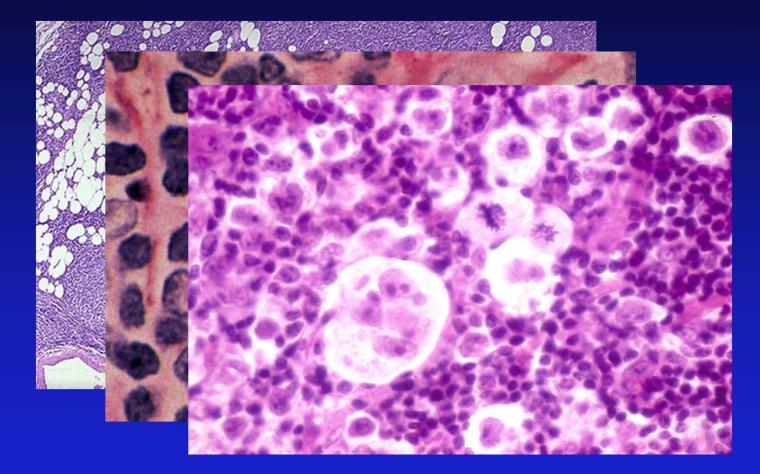
- 1. To provide reproducible, quantitative diagnoses for all cancer patients.
- 2. To clarify diagnostic distinctions that are problematic using current methods.
- 3. To deliver newly defined molecular diagnoses that influence treatment choice and/or prognosis.

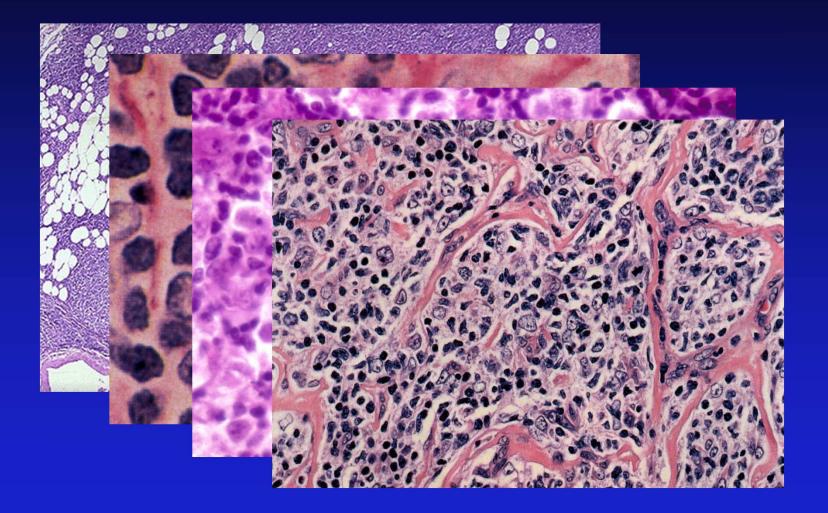
- 1. To provide reproducible, quantitative diagnoses for all cancer patients.
- 2. To clarify diagnostic distinctions that are problematic using current methods.
- 3. To deliver newly defined molecular diagnoses that influence treatment choice and/or prognosis.
- 4. To translate insights from therapeutic trials that incorporate molecular profiling.

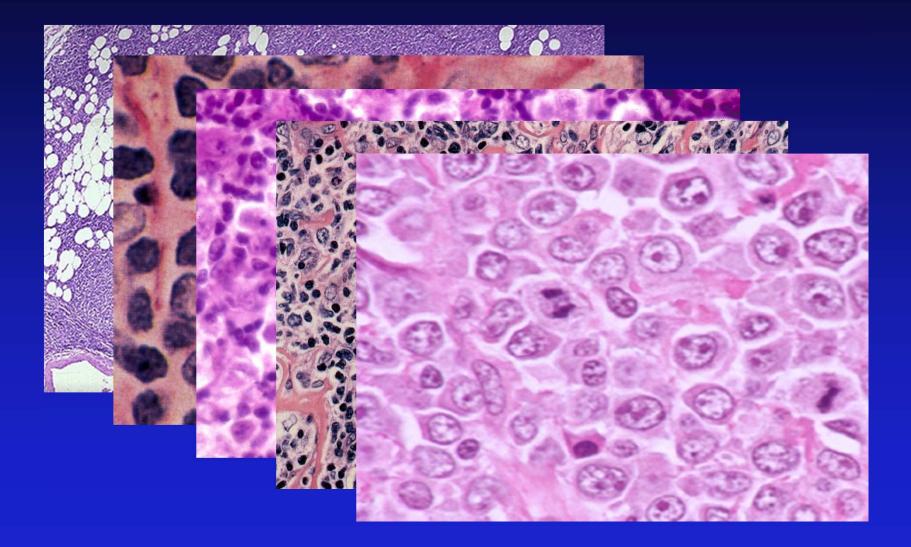
- 1. To provide reproducible, quantitative diagnoses for all cancer patients.
- 2. To clarify diagnostic distinctions that are problematic using current methods.
- 3. To deliver newly defined molecular diagnoses that influence treatment choice and/or prognosis.
- 4. To translate insights from therapeutic trials that incorporate molecular profiling.
- 5. To promote excellence in clinical science.

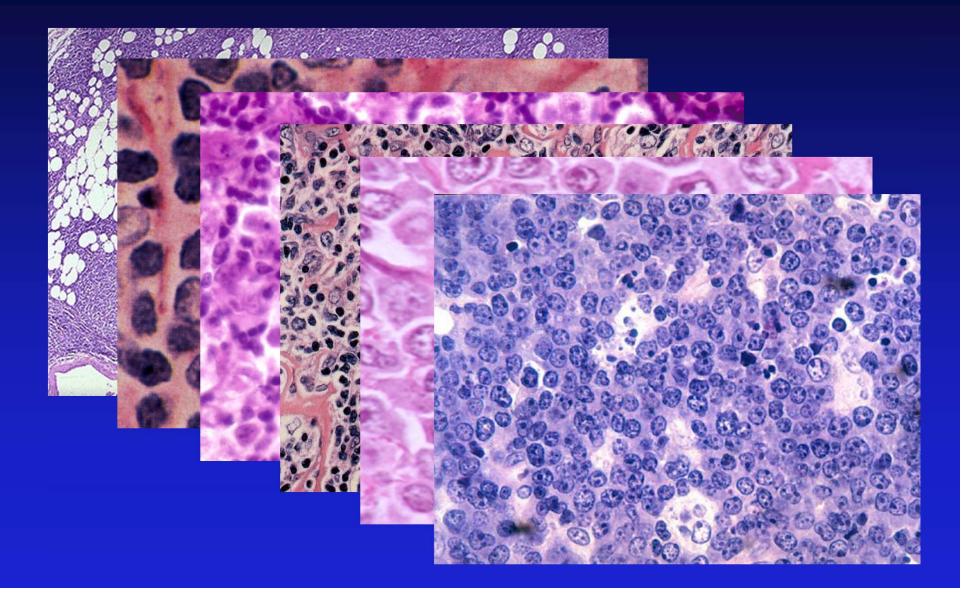












Lymphoma Subtype Diagnosis Alters Treatment Choice

Lymphoma subtype	Curable?	Therapy
Diffuse Large B Cell Lymphoma	Yes	CHOP chemotherapy + Rituximab
Primary Mediastinal B Cell Lymphoma	Yes	CHOP chemotherapy + Rituximab +/-radiation
Burkitt Lymphoma	Yes	High dose chemotherapy CNS prophylaxis
Follicular Lymphoma	No	Watchful waiting Rituximab Symptomatic chemotherapy
Mantle Cell Lymphoma	No	Watchful waiting Symptomatic chemotherapy Responsive to: Bortezomib rapamycin analogues



Goals: --Establish a molecular classification of human lymphoid malignancies. --Define molecular correlates of clinical parameters that are useful in prognosis and in the choice of optimal therapy.

Collaborating Institutions

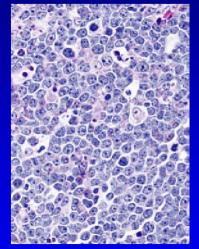
Univ. of Nebraska Medical Center British Columbia Cancer Agency Southwest Oncology Group **Cleveland Clinic** Norwegian Radium Hospital National Cancer Institute Center for Cancer Research

Univ. of Würzburg Univ. of Barcelona Univ. of Arizona Univ. of Rochester St. Bart's Hospital

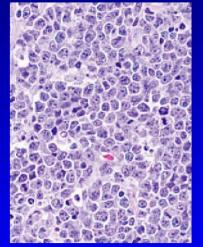
Improving the Accuracy and Reproducibility of Diagnosis Using Gene Expression Profiling

Diagnosis of Burkitt Lymphoma vs. Diffuse Large B Cell Lymphoma Alters Treatment Choice

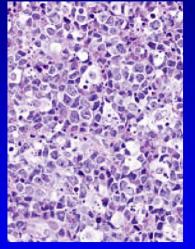
Classic Burkitt Lymphoma



Atypical Burkitt Lymphoma

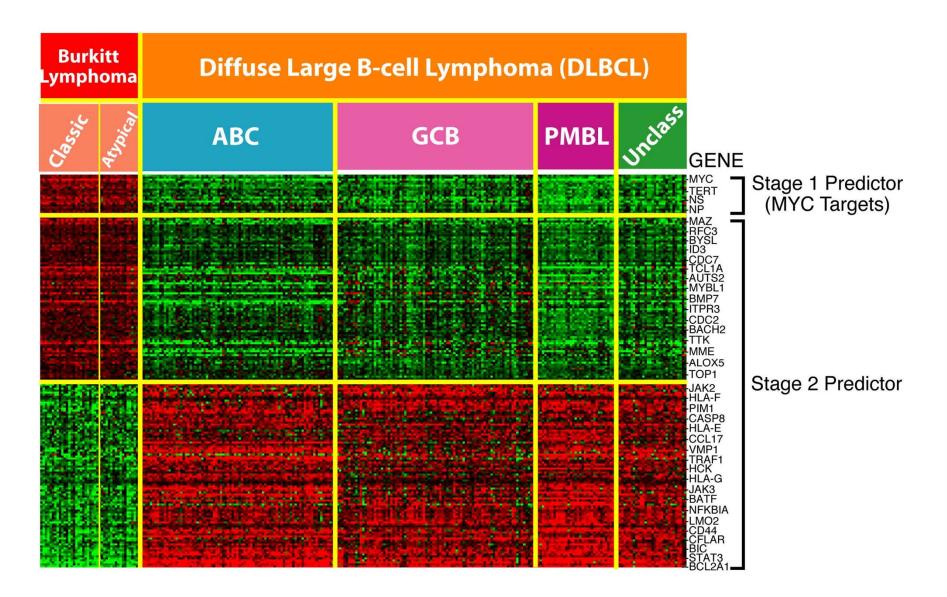


Diffuse Large B Cell Lymphoma

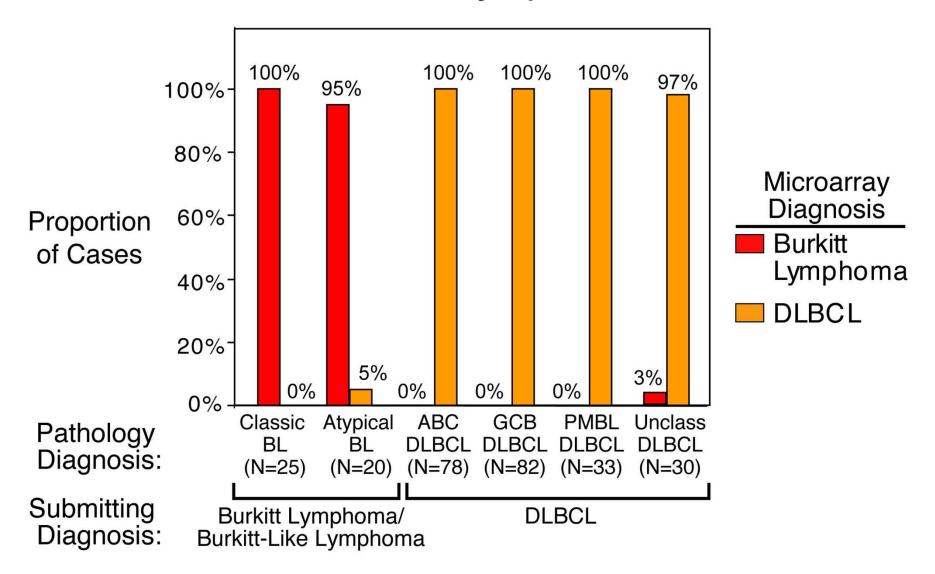


Recommended Treatment Intensive chemotherapy Intensive chemotherapy CHOP-like chemotherapy

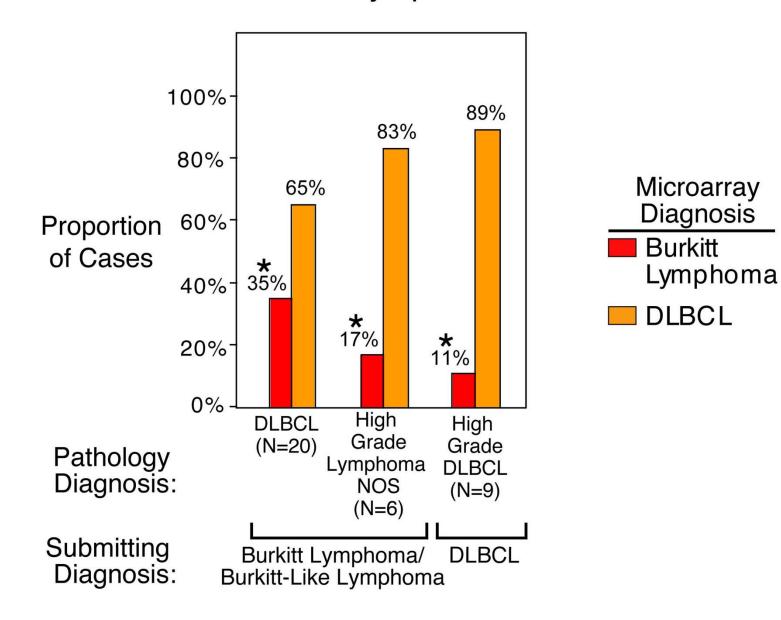
Gene Expression Differentiates Burkitt Lymphoma from all Subgroups of Diffuse Large B Cell Lymphoma



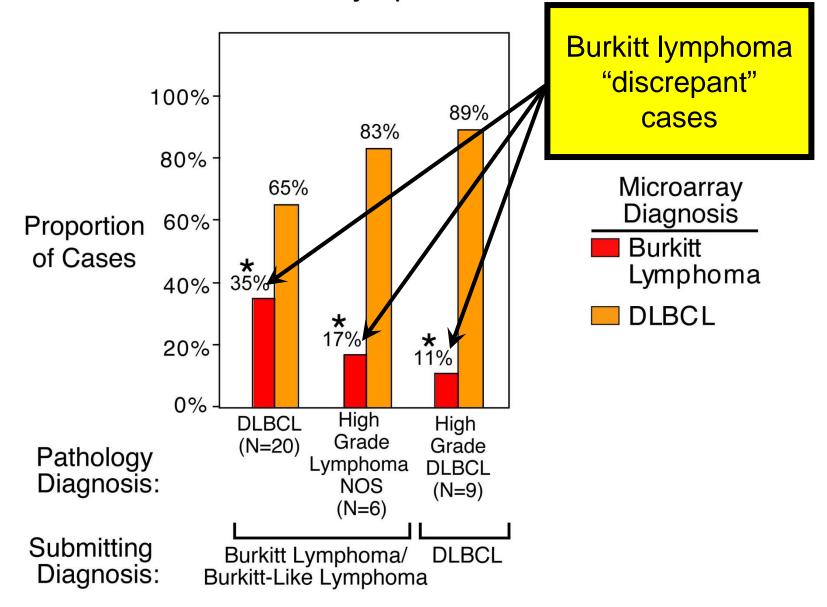
Performance of a Gene Expression-based Predictor of Burkitt Lymphoma



Discrepancies Between Molecular and Pathological Diagnoses of Burkitt Lymphoma



Discrepancies Between Molecular and Pathological Diagnoses of Burkitt Lymphoma

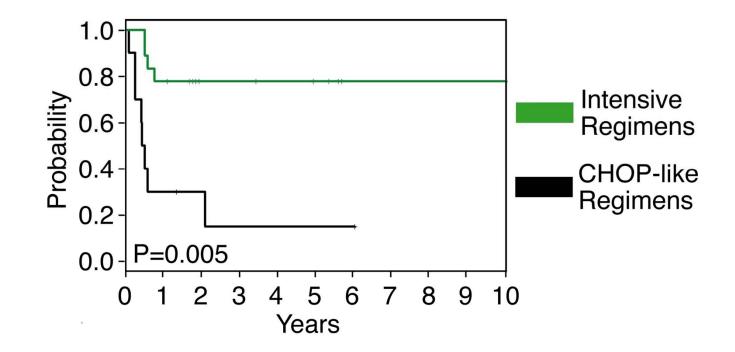


Effect of Treatment on Overall Survival in Burkitt Lymphoma

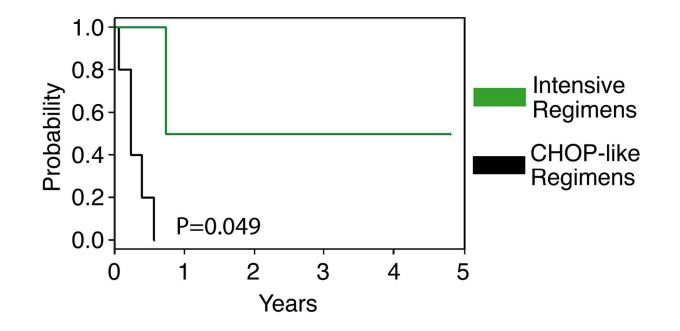
Classification of Treatments

CHOP-Like CHOP CNOP

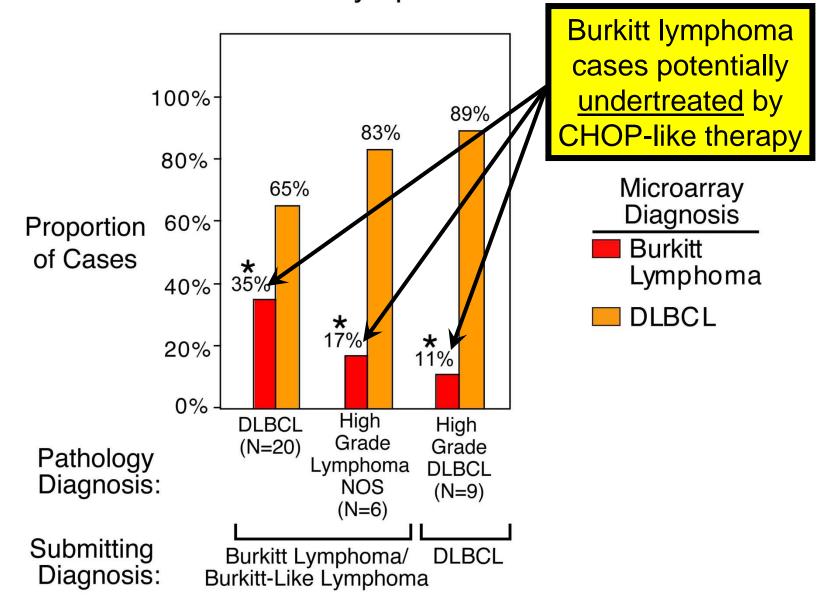
Intensive BFM CODOX-M/IVAC Regimens requiring autologous stem cell rescue High Cure Rates in Burkitt Lymphoma Treated With Intensive Regimens But Not CHOP-like Regimens



Burkitt Lymphoma Discrepant Cases are Not Curable With CHOP-like Regimens



Discrepancies Between Molecular and Pathological Diagnoses of Burkitt Lymphoma

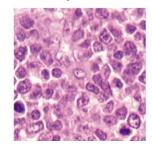


Conclusions

- Burkitt lymphoma has a distinct molecular profile that can reliably distinguish it from all forms of diffuse large B-cell lymphoma.
- Current means used for the diagnosis of Burkitt lymphoma disagree with the molecular diagnosis of Burkitt lymphoma in 17% of cases.
- The distinction between Burkitt lymphoma and DLBCL is critical because of significant differences in treatment.

=>Molecular diagnosis of Burkitt lymphoma will improve patient outcome.

Defining New Molecular Subgroups of Cancer by Gene Expression Profiling



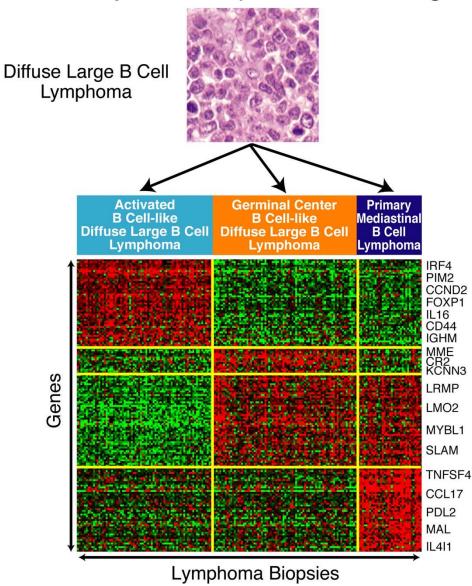
Diffuse large B cell lymphoma

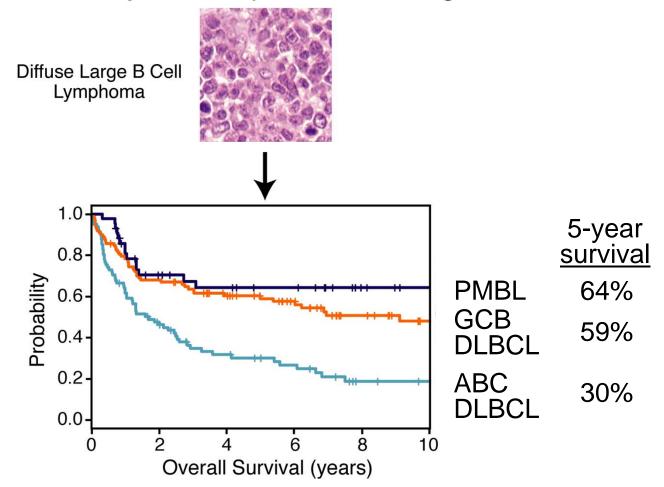
40% of Non-Hodgkin lymphomas

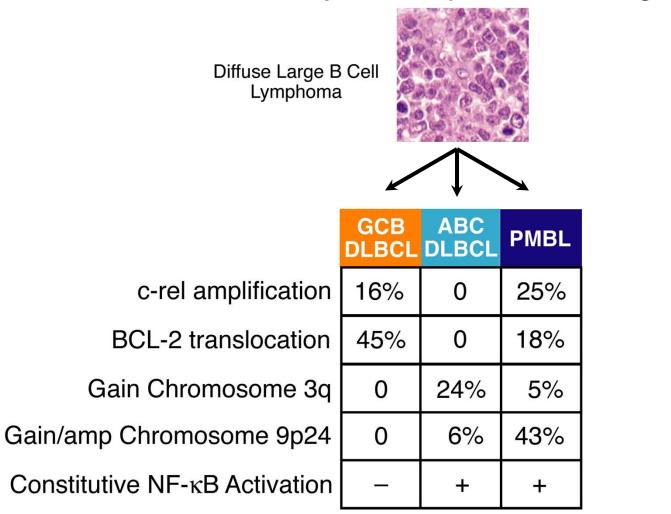
~23,000 new diagnoses/yr

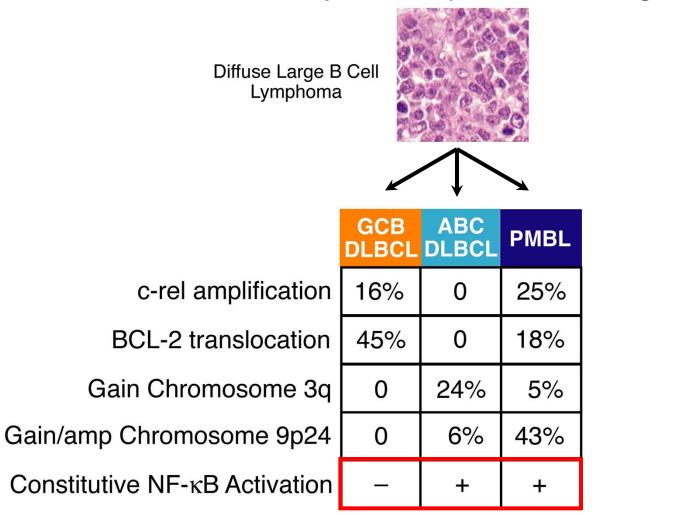
~40% cure rate

~10,000 deaths/yr

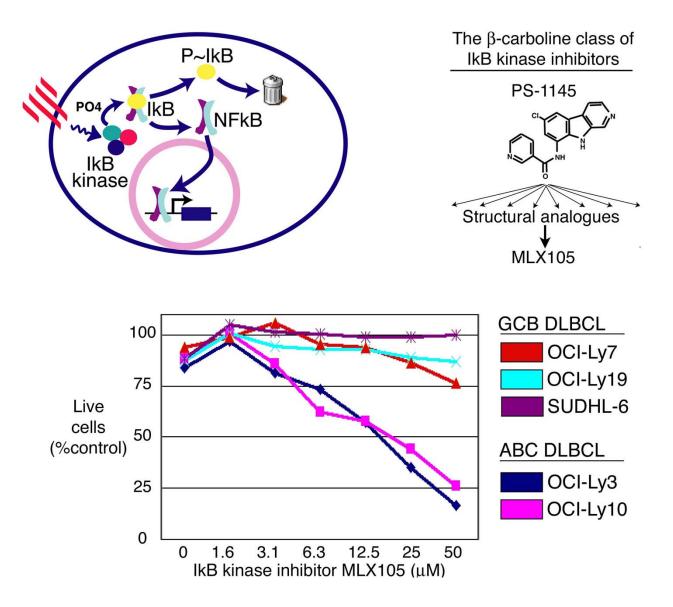






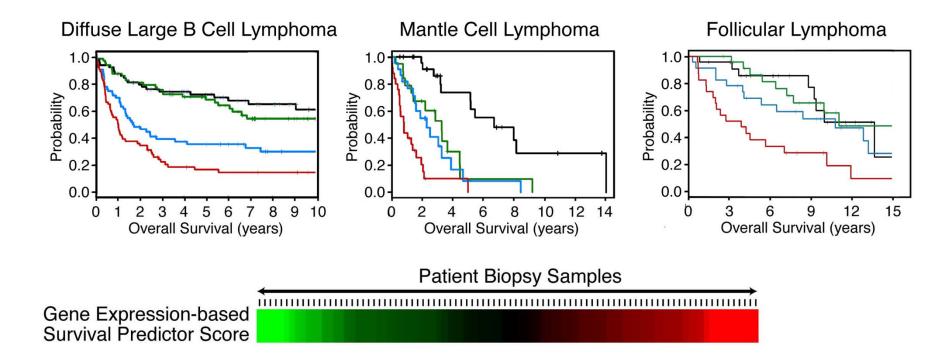


Validation of the NF-kB Pathway as a Therapeutic Target in Activated B Cell-like Diffuse Large B Cell Lymphoma

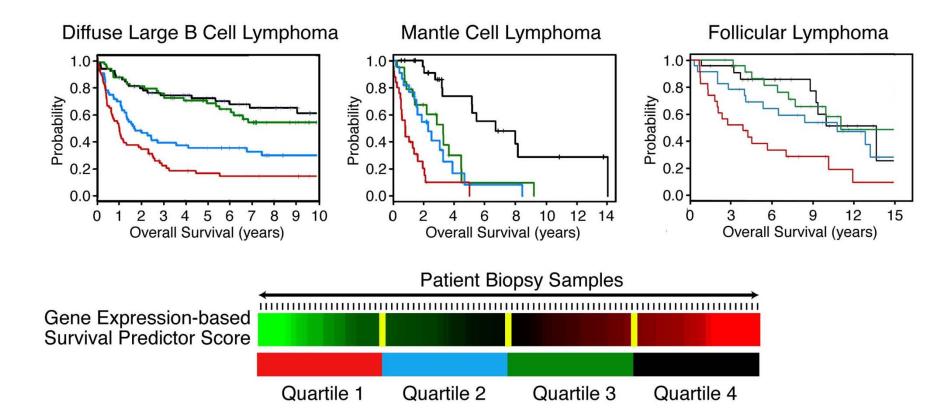


Molecular Predictors of Outcome In Cancer Using Gene Expression Profiling

Survival Prediction Based on the Gene Expression Profile of the Diagnostic Biopsy



Survival Prediction Based on the Gene Expression Profile of the Diagnostic Biopsy



Why Can Gene Expression Profiling Predict Outcome in Cancer?

Within a current diagnostic category, gene expression profiling can identify:

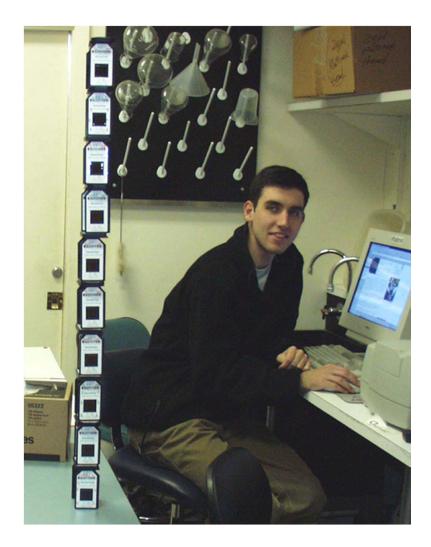
 Heterogeneity in cell of origin.
Heterogeneity in oncogenic pathways.
Heterogeneity in common cellular functions e.g. proliferation, survival, cell-cell interactions.

This heterogeneity is present in the tumor at the time of diagnosis.

Routine Molecular Diagnosis of Cancer in Clinical Oncology:

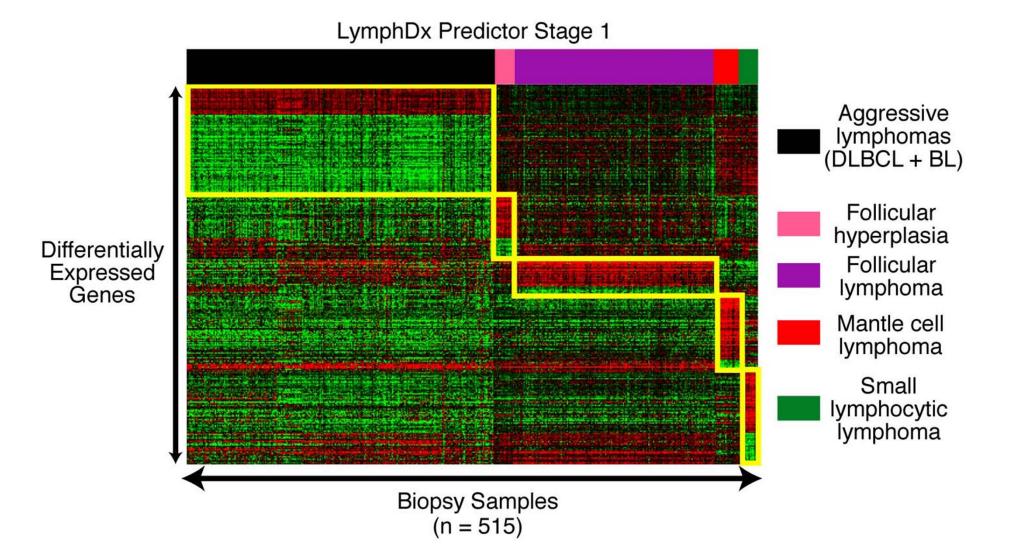
Development of a Lymphoma Diagnostic Microarray

The LymphDx Project Team

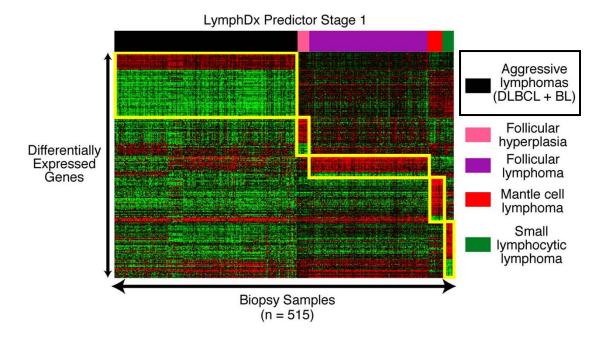


- Metabolism Branch, CCR, NCI
 - George Wright
 - Sandeep Dave
 - Bruce Tan
 - Andreas Rosenwald
 - Michael Chiorazzi
 - Hong Zhao
 - Liming Yang
 - Louis Staudt
- Members of the LLMPP
- Affymetrix
 - John Palma
 - Janet Warrington

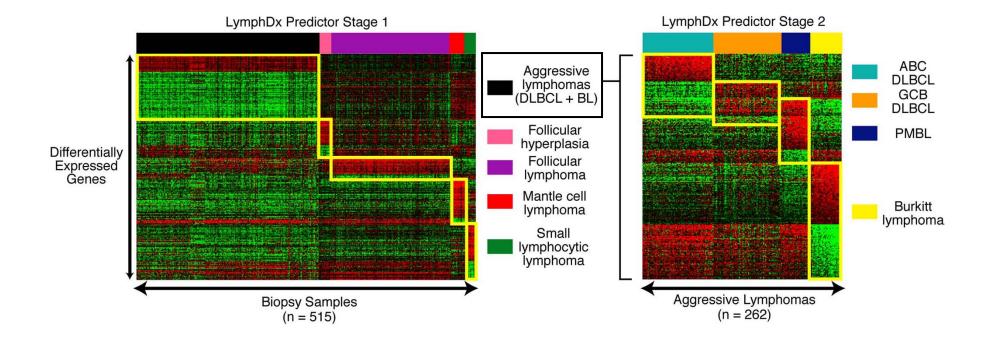
Gene Expression-based Diagnosis of Lymphoma



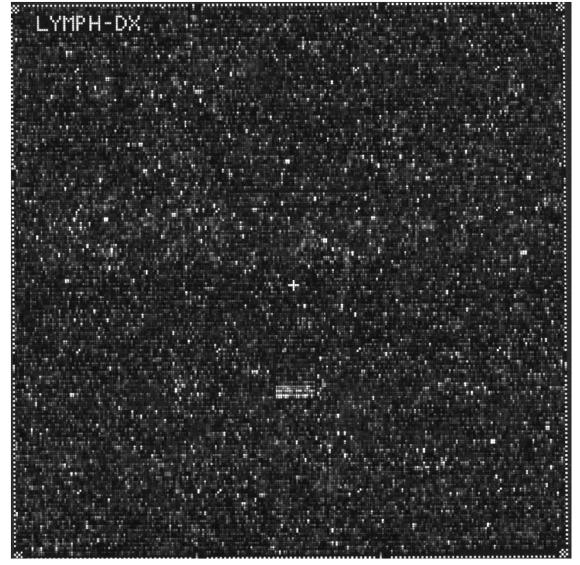
Gene Expression-based Diagnosis of Lymphoma



Gene Expression-based Diagnosis of Lymphoma



The LymphDx Custom DNA Microarray

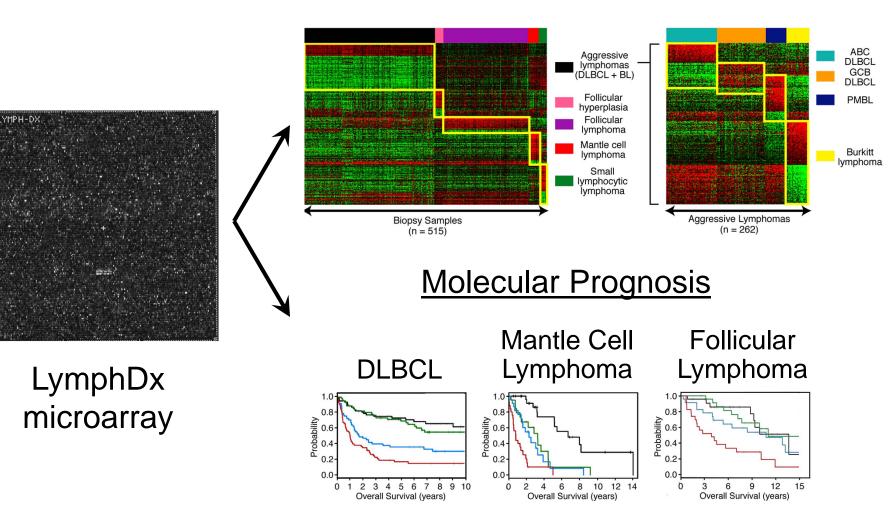


~2643 human genes:

Lymphoma diagnostic genes Lymphoma prognostic genes EBV, HHV8, HTLV1 viral genes Genes encoding all human kinases cytokines chemokines cytokine receptors chemokine receptors

Invariantly expressed control genes

LymphDx: The One-stop Shopping Approach to Lymphoma Diagnosis



Molecular Diagnosis



GOAL: Implementation of a gene expression-based molecular diagnosis of lymphoma in routine clinical practice.

Collaborating LLMPP Institutions

Univ. of Nebraska Medical CenterUniv. of WürzburgBritish Columbia Cancer CenterUniv. of BarcelonaSouthwest Oncology GroupUniv. of ArizonaCleveland ClinicUniv. of RochesterNorwegian Radium HospitalSt. Bart's HospitalNational Cancer Institute Center for Cancer Research



PHASE 1

Affymetrix whole genome profiles of retrospectively ascertained lymph node biopsies (non-Hodgkin and Hodgkin lymphoma, other cancers, benign conditions) (n= 2000)

Design custom diagnostic microarray



PHASE 1

Affymetrix whole genome profiles of retrospectively ascertained lymph node biopsies (non-Hodgkin and Hodgkin lymphoma, other cancers, benign conditions) (n= 2000)

Design custom diagnostic microarray

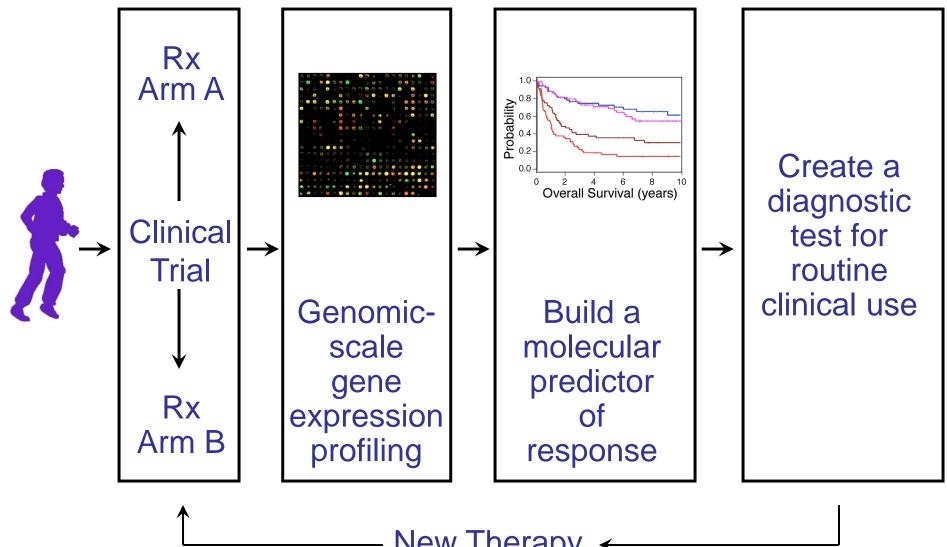
PHASE 2

Diagnostic microarray profiles of prospectively ascertained lymph node biopsies (n= 2000)

Generate data for regulatory approval

Evolving Molecular Diagnosis To Match Changes in Cancer Treatment

The Molecular Diagnosis Cycle



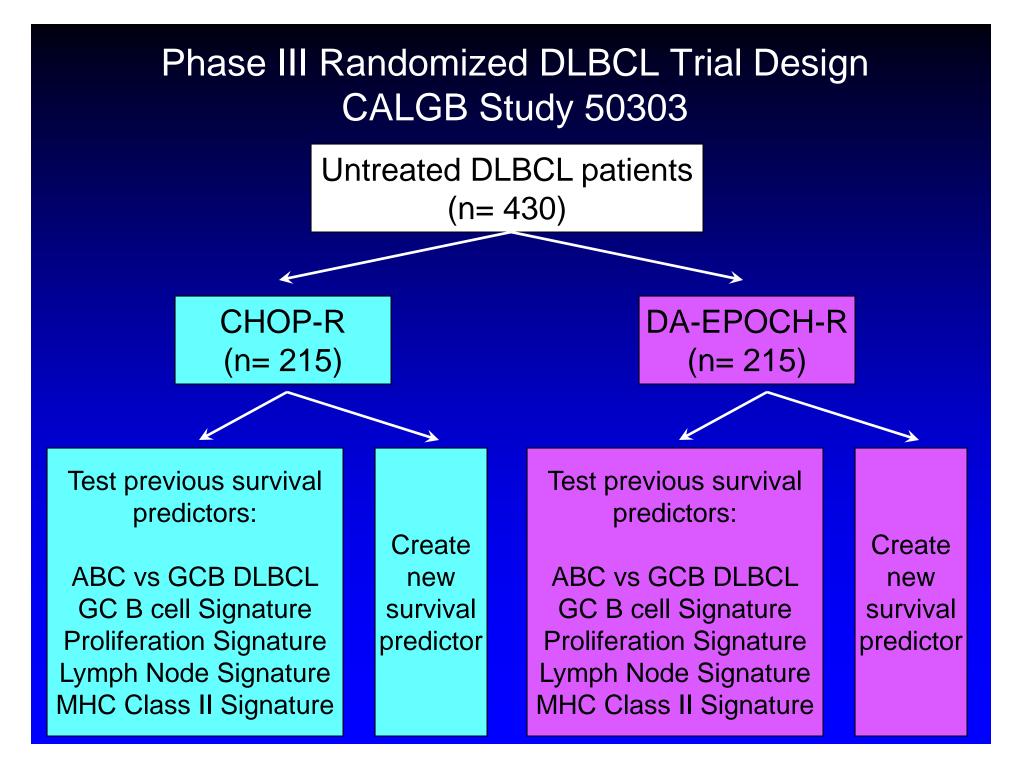
New Therapy +

Phase III Randomized Clinical Trial of CHOP-Rituximab vs. Dose-adjusted EPOCH-Rituximab with Gene Expression Profiling Analysis in Untreated Diffuse Large B cell Lymphoma

> CALGB Study 50303 Opened for accrual: May 2005

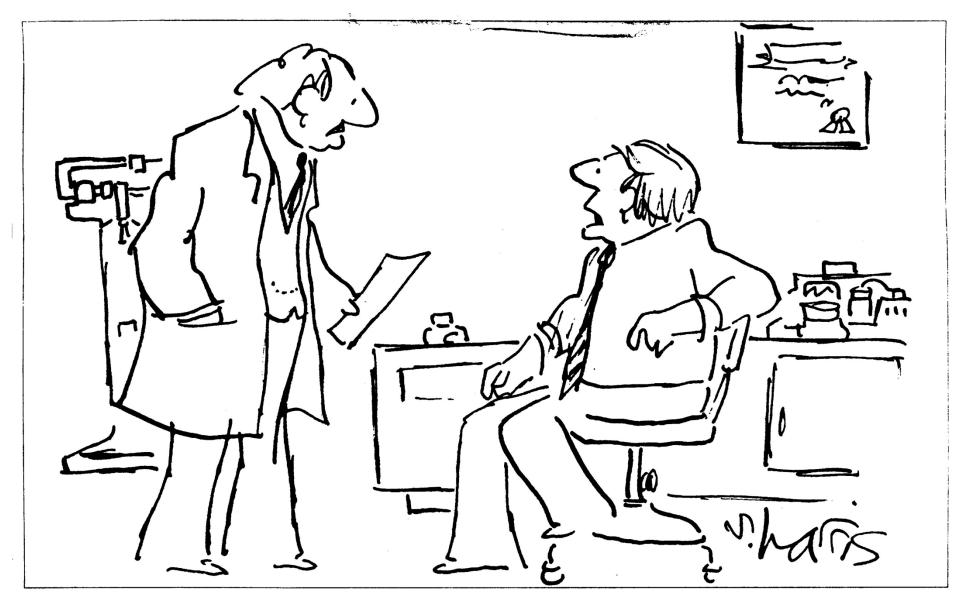
> > Study Chairs:

 Wyndham H. Wilson, Center for Cancer Research, NCI Bruce Cheson, CALGB Lymphoma Committee
Andrew D. Zelenetz, CALGB Lymphoma Committee
Richard Fisher, Chair, SWOG Lymphoma Committee
Louis M. Staudt, Center for Cancer Research, NCI

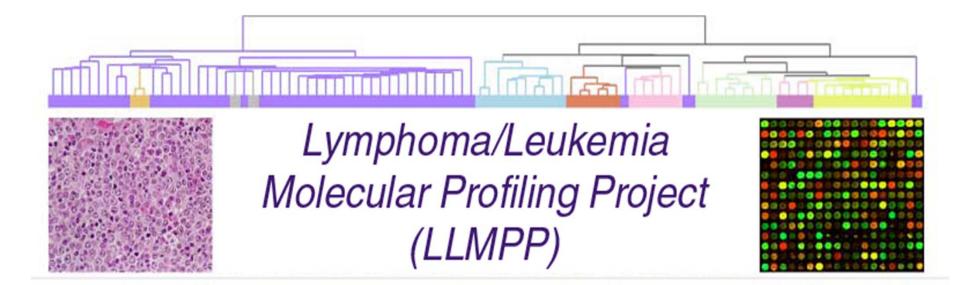


Why We Need Gene Expression Profiling for All Cancer Patients

- 1. To provide reproducible, quantitative diagnoses for all cancer patients.
- 2. To clarify diagnostic distinctions that are problematic using current methods.
- 3. To deliver newly defined molecular diagnoses that influence treatment choice and/or prognosis.
- 4. To translate insights from therapeutic trials that incorporate molecular profiling.
- 5. To promote excellence in clinical science.



"I feel fine, but I thought there may be something amiss on the molecular level"



Collaborating Institution Univ. of Nebraska Medical Center British Columbia Cancer Agency Southwest Oncology Group Univ. of Würzburg Univ. of Barcelona Norwegian Radium Hospital St. Bart's Hospital, London

Center for Cancer Research, National Cancer Institute John Chan Randy Gascoyne Rich Fisher Konrad Muller-Hermelink Elias Campo Erlend Smeland Andrew Lister

Lou Staudt Sandeep Dave George Wright