

Patient Name: Norman, George Patient DOB / Sex: 09/19/1994 / M

Accession / CaseNo: 430071 / NTP20-001848

# Molecular Genetics NGS Lymphoma Profile For FL

Client 8090 Oncology Institute

800 N Brand ave Glendale, CA 91205 Phone: (818) 500-7887 Fax: (818) 500-7800 Patient Name: Norman, George Patient DOB / Sex: 09/19/1994 / M Specimen Type: **Paraffin Tissue** 

Specimen ID: 79-44 MRN: 3006788 Ordering Physician(s): Vasillis Margaritis, MD Accession / CaseNo: 0067889 / TIM03-04379

Collection Date: 12/13/2022

Received Date: 12/14/2022 09:07:00 AM PDT Report Date: 12/23/2022 08:07:00 AM PDT

### **Results Summary**

XXX	<u>1</u> Clinically Significant Translocation Detected	T(14;18) (q32;q21)				
Q	Additional Studies	BCL 2 overexpression: DETECTED				
	Pertinent Negatives NO abnormalitites detected in the following genes: Cycline D1					
Laborate Anti-						

#### Interpretation

- BCL 2 gene overexpression: Detected. Dependent on lymphoma type, BCL mutations are often associated with a poor prognosis aggressive disease. Clinicopathologic correlation recommended.

#### **Profile Results Detail**

Molecular Testing Detail								
Gene name	Alteration	Chromosome Change	Chromosome Change	Consequence	Expression level (%)	Read Depth		
BCL2	Translocation	T(14;18)	Q32;q21	Apoptosis inhibition	90	124		
Cycline D1	Not detected	N/A	N/A	N/A	0	0		

Biomarker/Assay	Results		
BCL2 gene analysis	Overexpression T(14;18) q(32;21)		

## **Test Description & Methodology**

#### **Test Description**

The NGS Lymphoma Profile for Follicular lymphoma uses next-generation sequencing (NGS) as listed below. Test orders include summary interpretation of all results to help guide treatment decisions.

#### Clinical Significance

The NGS Lymphoma Profile for FL is useful to excision biopsy of lymph nodes that are indeterminate or suspicious on cytology. Presence of mutations or gene rearrangements as detected by FISH predicts malignancy with varying degrees of specificity depending on the gene alteration and histological subtype. BCL2 rearrangement/overexpression is associated with poor prognosis follicular lymphoma

<sup>§</sup> See full list of genes tested in Biomarkers Evaluated section at end of report.