Lymphoma (Aggressive) Panel by FISH

Indications for Ordering

• Aid in diagnosis of aggressive large B-cell lymphoma with intermediate features between Burkitt lymphoma and diffuse large B-cell lymphoma (DLBCL)
• Confirmation of suspected double hit lymphoma

Test Description

Fluorescence in situ hybridization (FISH)
• FISH probes include
  o MYC
  o IGH
  o BCL2
  o BCL6

Tests to Consider

Typical Testing Strategy
• Lymph node biopsy with morphologic and immunohistochemical evaluation
• Leukemia/lymphoma phenotyping by flow cytometry
• In aggressive B-cell lymphomas with a high proliferation index and/or with unusual morphologic, phenotypic, or clinical features
  o Tests for MYC, BCL2, and BCL6

Panel Tests

Aggressive B-Cell Lymphoma FISH Reflex, Tissue 2012710
• Formalin-fixed, paraffin-embedded (FFPE) tissue specimens
• If MYC (8q24) Gene Rearrangement by FISH is positive, then IGH-BCL2 Fusion, t(14;18) by FISH will be added
• If IGH-BCL2 Fusion, t(14;18) by FISH is negative, then BCL6 (3q27) Gene Rearrangement by FISH will be added

Lymphoma (Aggressive) Panel by FISH 2002650
• Bone marrow (BM) or whole blood specimens; other specimens may be acceptable
• FFPE and frozen specimens unacceptible

Individual Tests

MYC (8q24) Gene Rearrangement by FISH 3001300
• Detects all MYC rearrangements, including t(8;14), t(2;8), and t(8;22)

IGH-MYC (t(8;14)) by FISH 3001299
IGH-BCL2 Fusion, t(14;18) by FISH 3001298
BCL6 (3q27) Gene Rearrangement by FISH 3001311

Related Tests

Leukemia/Lymphoma Phenotyping by Flow Cytometry 2008003
• Aids in diagnosis of hematopoietic neoplasms

Chromosome FISH, Interphase 2002298
• Specific probes must be requested
  o MYC break apart, BCL2, BCL6
• Fresh tissue specimens only

Chromosome Analysis, Bone Marrow 2002292
• Diagnosis, prognosis, and monitoring of lymphoma in BM

Chromosome Analysis, Solid Tumor 2002296
• May identify additional, useful cytogenetic abnormalities in tissues that are not targeted by FISH assays

Disease Overview

Prognostic Issues

• B-cell lymphomas with two recurrent chromosomal breakpoint aberrations are referred to as double hit lymphomas and are classified as high-grade B-cell lymphomas with MYC and BCL2 and/or BCL6 (WHO 2016)
  o Usually involve MYC oncogene in association with BCL2, less often with BCL6
• Lymphomas with three translocations (usually MYC/BCL2/BCL6) are referred to as triple hit lymphomas
  o Rare
• Important to identify these lymphomas in diagnostic evaluation of morphologically aggressive lymphomas
  o They are highly resistant to standard chemotherapy
    • Poor outcome independent of regimen intensity or inclusion of rituximab
  o Individuals have shortened survival compared with those having Burkitt lymphoma or international prognostic index (IPI)-matched DLBCL
### Breakpoints Used to Identify Double Hit or Triple Hit Lymphomas

<table>
<thead>
<tr>
<th>Oncogene</th>
<th>Break apart MYC</th>
<th>BCL2</th>
<th>BCL6</th>
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<tbody>
<tr>
<td>Locus</td>
<td>8q24</td>
<td>18q21</td>
<td>3q27</td>
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<tr>
<td>Biology</td>
<td>Accelerator of cell proliferation</td>
<td>Apoptosis inhibitor</td>
<td>Transcription modifier</td>
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<tr>
<td>Cytogenetics</td>
<td>Any MYC translocation</td>
<td>BCL2/IGH – t(14;18) (q32;q21)</td>
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<td>BCL6 most commonly has a non-IGH translocation partner – BCL6 (3q27)</td>
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<td>Uncommon partner – BCL6/IGH [t(3;14)(q27;q32)]</td>
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### Specific lymphomas associated with translocation
- Burkitt lymphoma
- DLBCL
- Aggressive B-cell lymphoma not otherwise specified (NOS)
- Follicular lymphoma
- DLBCL
- Follicular lymphoma
- DLBCL
- High grade lymphomas (rare)

### Test Interpretation

#### Results
- Abnormal – t(14;18)(q32;q21) (IGH/BCL2 translocation) or other rearrangements involving BCL6 and/or MYC detected
  - Presence of two or more translocations is associated with poor prognosis in mature B-cell lymphomas
  - Single rearrangements can provide diagnostic and/or prognostic information in the appropriate context
- Normal – t(14;18)(q32;q21) (IGH/BCL2 translocation) or other rearrangements involving BCL6 or MYC not detected

#### Limitations
- Interpretation of results requires correlation with morphology and immunophenotype
- MYC and/or BCL2 overexpression can be due to other mechanisms not detected by this test
- Chromosome alterations outside probe region are not detected