

T-Cell Acute Lymphoblastic Leukemia/Lymphoma (ALL), FISH, Adult, Varies

Overview

Useful For

<u>Detecting, at diagnosis, recurrent common chromosome abnormalities associated with T-cell acute lymphoblastic leukemia/lymphoma (T-ALL) in adult patients using a laboratory-designated probe set algorithm</u>

As an adjunct to conventional chromosome studies in adult patients with T-ALL

Evaluating specimens in which chromosome studies are unsuccessful

This test should not be used to screen for residual T-ALL

Reflex Tests

Test Id	Reporting Name	Available Separately	Always Performed
TALAB	Probe, Each Additional	No, (Bill Only)	No
	(TALAF)		

Testing Algorithm

This test includes a charge for the probe application, analysis, and professional interpretation of results for 10 probe sets (20 individual fluorescence in situ hybridization [FISH] probes). Additional charges will be incurred for all reflex or additional probe sets performed. Analysis charges will be incurred based on the number of cells analyzed per probe set. If no cells are available for analysis, no analysis charges will be incurred.

This test is performed as panel testing only using the following analysis algorithm.

If the patient clinically relapses, a conventional chromosome study is useful to identify cytogenetic changes in the neoplastic clone or the possible emergence of a new therapy-related myeloid clone.

The **diagnostic** adult T-cell acute lymphoblastic leukemia (ALL) FISH panel includes testing for the following abnormalities using the FISH probes listed:

t(9p24.1;var) or JAK2 rearrangement, JAK2 break-apart probe set

ABL1 amplification or t(9;22)(q34;q11.2), ABL1/BCR probe set

t(11q23;var) or KMT2A rearrangement, KMT2A break-apart probe set

1p33 rearrangement or STIL deletion, TAL1/STIL probe set

t(5;14)(q35;q32) or *TLX3::BCL11B* fusion, TLX3/BCL11B probe set

t(7q34;var) or TRB rearrangement, TRB break-apart probe set

t(14q11.2;var) or TRA rearrangement, TRA break-apart probe set

t(10;11)(p12;q14) or MLLT10::PICALM fusion, MLLT10/PICALM fusion probe set

t(9q34;var) or ABL1 rearrangement, ABL1 break-apart probe set

t(5q32;var) or PDGFRB rearrangement, PDGFRB break-apart probe set



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Appropriate ancillary probes may be performed at consultant discretion to render comprehensive assessment. Any additional probes used will have the results included within the final report and will be performed at an additional charge. In the following situations, additional (reflex) testing may be performed at the laboratory's discretion and may be influenced by available karyotype results or other FISH testing.

When a TRB rearrangement is identified, testing in an attempt to identify the translocation partner may be performed. Probes include identification of t(7;10)(q34;q24) TRB::TLX1 fusion or t(7;11)(q34;p13) TRB::LMO2 fusion.

When a *KMT2A* rearrangement is identified, testing with 1 or more dual-fusion (D-FISH) probe sets may be performed in an attempt to identify the translocation partner for the following abnormalities:

t(4;11)(q21;q23) or KMT2A::AFF1 fusion, AFF1/KMT2A probe set

t(6;11)(q27;q23) or KMT2A::AFDN ;fusion, AFDN/KMT2A probe set

t(9;11)(p22;q23) or KMT2A::MLLT3 fusion, MLLT3/KMT2A probe set

t(10;11)(p12;q23) or KMT2A::MLLT10 fusion, MLLT10/KMT2A probe set

t(11;19)(q23;p13.1) or KMT2A::MLLT1 fusion, KMT2A/ELL probe set

t(11;19)(q23;p13.3) or KMT2A::ELL fusion, KMT2A/MLLT1 probe set

When a TRA rearrangement is identified, testing in an attempt to identify the translocation partner may be performed. Probes include identification of t(10;14)(q24;q11.2) TRA::TLX1 fusion or t(11;14)(p13;q11.2) TRA::LMO2 fusion.

For more information see Acute Leukemias of Ambiguous Lineage Testing Algorithm.

Special Instructions

• Acute Leukemias of Ambiguous Lineage Testing Algorithm

Method Name

Fluorescence In Situ Hybridization (FISH)

NY State Available

Yes

Specimen

Specimen Type

Varies

Ordering Guidance

This test is only performed on specimens from patients with T-cell acute lymphoblastic leukemia/lymphoma (T-ALL) who are 31 years and older.

This test is intended for instances when the entire T-ALL fluorescence in situ hybridization (FISH) panel is needed for an **adult** patient.



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This test **should NOT be used** to screen for residual T-ALL. at follow-up, or if the patient clinically relapses, conventional cytogenetic studies (CHRBM / Chromosome Analysis, Hematologic Disorders, Bone Marrow) are useful to identify cytogenetic changes in the neoplastic clone or the possible emergence of a new therapy-related myeloid clone. Additionally, targeted T-ALL FISH probes can be evaluated based on the abnormalities identified in the diagnostic study.

If targeted T-cell ALL FISH probes are preferred, order TALMF / T-Cell Acute Lymphoblastic Leukemia/Lymphoma (ALL), Specified FISH, Varies.

If this test is ordered on a patient 30 years or younger, this test will be canceled and automatically reordered by the laboratory as TALFP / Pediatric T-Lymphoblastic Leukemia/Lymphoma panel, FISH, Varies.

If this test is ordered and the laboratory is informed that the patient is on a Children's Oncology Group (COG) protocol, this test will be canceled and automatically reordered by the laboratory as COGTF / T-Cell Acute Lymphoblastic Leukemia/Lymphoma (ALL), Children's Oncology Group Enrollment Testing, FISH, Varies.

If BALAF / B-Cell Acute Lymphoblastic Leukemia/Lymphoma (ALL), Adult, FISH, Varies is ordered concurrently with this test, the laboratory may cancel TALAF and automatically reorder as TALMF / T-Cell Acute Lymphoblastic Leukemia/Lymphoma (ALL), Specified FISH, Varies with the following FISH probes: TLX3/BCL11B, break-apart TRB, break-apart TRA, MLLT10/PICALM, and TAL1/STIL. If an abnormality is identified that would result in reflex testing in TALAF, the same reflex testing will be performed in the TALMF. This cancelation is necessary to avoid duplicate testing. Probes for break-apart PDGFRB, break-apart JAK2, ABL1/BCR, break-apart ABL1, and break-apart MLL will still be performed as part of the adult B-ALL FISH panel.

For testing paraffin-embedded tissue samples from patients with T-cell lymphoblastic leukemia/lymphoma, order TLBLF / T-Lymphoblastic Leukemia/Lymphoma, FISH, Tissue. If a paraffin-embedded tissue sample is submitted for this test, testing will be canceled and TLBLF will be added and performed as the appropriate test.

Additional Testing Requirements

At diagnosis, conventional cytogenetic studies (CHRBM / Chromosome Analysis, Hematologic Disorders, Bone Marrow) and this fluorescence in situ hybridization panel should be performed. If there is limited specimen available, only this test will be performed.

Shipping Instructions

Advise Express Mail or equivalent if not on courier service.

Necessary Information

- **1.** A reason for testing must be provided. If this information is not provided, an appropriate indication for testing may be entered by Mayo Clinic Laboratories.
- 2. A flow cytometry and/or a bone marrow pathology report should be submitted with each specimen. The laboratory will not reject testing if this information is not provided, but appropriate testing and interpretation may be compromised or delayed.

Specimen Required

Submit only 1 of the following specimens:



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Preferred

Specimen Type: Bone marrow

Container/Tube:

Preferred: Yellow top (ACD)

Acceptable: Green top (sodium heparin) or lavender top (EDTA)

Specimen Volume: 2 to 3 mL **Collection Instructions:**

1. It is preferable to send the first aspirate from the bone marrow collection.

2. Invert several times to mix bone marrow.

3. Send bone marrow specimen in original tube. Do not aliquot.

Acceptable

Specimen Type: Whole blood

Container/Tube:

Preferred: Yellow top (ACD)

Acceptable: Green top (sodium heparin) or lavender top (EDTA)

Specimen Volume: 6 mL **Collection Instructions:**

1. Invert several times to mix blood.

2. Send whole blood specimen in original tube. Do not aliquot.

Forms

If not ordering electronically, complete, print, and send a <u>Hematopathology/Cytogenetics Test Request</u> (T726) with the specimen.

Specimen Minimum Volume

Bone marrow: 1 mL; Whole blood: 2 mL

Reject Due To

All specimens will be evaluated at Mayo Clinic Laboratories for test suitability.

Specimen Stability Information

Specimen Type	Temperature	Time	Special Container
Varies	Ambient (preferred)		
	Refrigerated		

Clinical & Interpretive

Clinical Information

T-cell acute lymphoblastic leukemia (T-ALL) accounts for approximately 25% of cases of adult lymphoblastic leukemia. An abnormal karyotype is found in 50% to 70% of T-ALL cases, although many of the classic abnormalities are "cryptic" by conventional chromosome studies and must be identified by fluorescence in situ hybridization (FISH) studies and are



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associated with various prognoses. One predictive marker, amplification of the *ABL1* gene region, has been identified in 5% of T-ALL, and these patients may be responsive to targeted tyrosine kinase inhibitors.

A summary of the characteristic chromosome abnormalities identified in T-ALL is listed in the following table.

Table. Common Chromosome Abnormalities in T-cell Acute Lymphoblastic/Leukemia

Cytogenetic change	Genes involved
del(1p33)	TAL1/STIL
t(5;14)(q35;q32)	TLX3::BCL11B
t(5q32;var)	PDGFRB
t(10;11)(p13;q14)	PICALM::MLLT10
Episomal amplification	ABL1
t(9p24.1;var)	JAK2
t(9q34;var)	ABL1
t(11q23;var)	KMT2A
t(4;11)(q21;q23)	KMT2A::AFF1
t(6;11)(q27;q23)	KMT2A:: <i>AFDN</i>
t(9;11)(p21.3;q23)	KMT2A::MLLT3
t(10;11)(p13;q23)	KMT2A::MLLT10
t(11;19)(q23;p13.1)	KMT2A::ELL
t(11;19)(q23;p13.3)	KMT2A::MLLT1
t(7q34;var)	TRB
t(6;7)(q23;q34)	TRB::MYB
t(7;10)(q34;q24)	TRB ::TLX1
t(7;11)(q34;p15)	TRB ::LMO1
t(7;11)(q34;p13)	TRB ::LMO2
t(14q11.2;var)	TRA
t(8;14)(q24.21;q11.2)	TRA::MYC
t(10;14)(q24;q11.2)	TLX1::TRA
t(11;14)(p15;q11.2)	LMO1::TRA
t(11;14)(p13;q11.2)	LMO2 ::TRA
del(17p)	TP53
Complex karyotype (> or =4 abnormalities)	

Reference Values

An interpretive report will be provided.

Interpretation

A neoplastic clone is detected when the percent of cells with an abnormality exceeds the normal reference range for any



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given probe set.

The absence of an abnormal clone does not rule out the presence of a neoplastic disorder.

Cautions

This test is not approved by the US Food and Drug Administration, and it is best used as an adjunct to clinical and pathologic information.

Fluorescence in situ hybridization (FISH) is not a substitute for conventional chromosome studies because the latter detects chromosome abnormalities associated with other hematological disorders that would go undetected in a targeted T-cell acute lymphoblastic leukemia/lymphoma FISH panel test.

Bone marrow is the preferred specimen type for this FISH test. If bone marrow is not available, a blood specimen may be used if there are circulating malignant cells in the blood specimen (as verified by a hematopathologist).

If no FISH signals are observed post-hybridization, the case will be released indicating a lack of FISH results.

Clinical Reference

- 1. WHO Classification of Tumours Editorial Board, eds. Haematolymphoid tumours. 5th ed. IARC Press; 2024. WHO Classification of Tumours, Volume 11
- 2. Gesk S, Martin-Subero JI, Harder L, et al. Molecular cytogenetic detection of chromosomal breakpoints in T-cell receptor gene loci. Leukemia. 2003;17(4):738-745
- 3. Chin M, Mugishima H, Takamura M, et al. Hemophagocytic syndrome and hepatosplenic (gamma)(delta) T-cell lymphoma with isochromosome 7q and 8 trisomy. J Pediatr Hematol Oncol. 2004;26(6):375-378
- 4. Graux C, Cools J, Michaux L, Vandenberghe, P, Hagemeijer A. Cytogenetics and molecular genetics of T-cell acute lymphoblastic leukemia: from thymocyte to lymphoblast. Leukemia. 2006;20:1496-1510
- 5. Cayuela JM, Madani A, Sanhes L, Stern MH, Sigaux F. Multiple tumor-suppressor gene 1 inactivation is the most frequent genetic alteration in T-cell acute lymphoblastic leukemia. Blood. 1996;87:2180-2186
- 6. Hayette S, Tigaud I, Maguer-Satta V, et al. Recurrent involvement of the *MLL* gene in adult T-lineage acute lymphoblastic leukemia. Blood. 2002;99:4647-4649
- 7. Graux C, Cools J, Melotte C, et al. Fusion of *NUP214* to *ABL1* on amplified episomes in T-cell acute lymphoblastic leukemia. Nat Genet. 2004;36:1084-1089

Performance

Method Description

This test is performed using commercially available and laboratory-developed probes. Rearrangements involving *TAL1/STIL, PDGFRB, TRB, JAK2, ABL1, KMT2A*, and *TRA* are detected using a dual-color break-apart (BAP) strategy probe sets. Dual-color, dual-fusion fluorescence in situ hybridization (D-FISH) strategy probe sets are used to detect t(5;14), t(9;22), t(10;11), and in reflex testing when rearrangements of *KMT2A, TRB*, or *TRA* genes are detected. Amplification of the *ABL1* gene region is detected using a D-FISH probe strategy. For enumeration and BAP strategy probe sets, 100 interphase nuclei are scored; 200 interphase nuclei are scored when D-FISH probes are used. All results are expressed as



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the percent abnormal nuclei.(Unpublished Mayo method)

PDF Report

No

Day(s) Performed

Monday through Friday

Report Available

7 to 10 days

Specimen Retention Time

4 weeks

Performing Laboratory Location

Mayo Clinic Laboratories - Rochester Main Campus

Fees & Codes

Fees

- Authorized users can sign in to <u>Test Prices</u> for detailed fee information.
- Clients without access to Test Prices can contact <u>Customer Service</u> 24 hours a day, seven days a week.
- Prospective clients should contact their account representative. For assistance, contact <u>Customer Service</u>.

Test Classification

This test was developed and its performance characteristics determined by Mayo Clinic in a manner consistent with CLIA requirements. It has not been cleared or approved by the US Food and Drug Administration.

CPT Code Information

88271x20, 88275x10, 88291x1-FISH Probe, Analysis, Interpretation; 10 probe sets 88271x2, 88275x1-FISH Probe, Analysis; each additional probe set (if appropriate)

LOINC® Information

Test ID	Test Order Name	Order LOINC® Value
TALAF	Adult ALL (T-cell), FISH	101663-3

Result ID	Test Result Name	Result LOINC® Value
609558	Result Summary	50397-9
609559	Interpretation	69965-2
609560	Result Table	93356-4
609561	Result	62356-1
GC071	Reason for Referral	42349-1



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GC072	Specimen	31208-2
609562	Source	31208-2
609563	Method	85069-3
609564	Additional Information	48767-8
609565	Disclaimer	62364-5
609566	Released By	18771-6