

## HEMATOLOGIC ONCOLOGY TEST REQUEST FORM

BARCODE / PROTOCOL NO:

PATIENT:		PHYSICIAN:
Name and surname:		Name and surname:
Gender: F <input type="checkbox"/> M <input type="checkbox"/> Date of birth:		Phone number: E-mail:
Mother name:	Father name:	Institution:
ID number: E-mail: Phone number: Address:		Stamp:  <b>SAMPLE TAKEN AND DATE OF SAMPLE COLLECTION:</b>  Date of Receipt:                      Time: Peripheral Blood <input type="checkbox"/> Bone marrow <input type="checkbox"/> Tissue <input type="checkbox"/> Other <input type="checkbox"/>

## CLINICAL INFORMATION:

(If necessary, use new paper or attach the epicrisis printout and other necessary documents to the request sheet)

## IMPORTANT NOTE:

- Genetic tests are subject to approval. It is a legal requirement to obtain the consent of the patient and his/her guardian for those under 18.
- You can reach us for other forms at <https://www.sapiensgenetics.com/guides-forms/> or request it from our center.
- It is recommended that genetic examinations be carried out with genetic counseling before and after the test. For this purpose, you can make an appointment at our center.

**MOLECULAR GENETIC TESTS**

<input type="checkbox"/> <b>Myeloid Fusion NGS Panel (Fusion, SNV/Indel)</b> <i>SNV/Indel studied genes: ABL1, AKT3, ASXL1, BRAF, CALR, CBL, CEBPA, CREBBP, CSF3R, DCK, DNMT2, DNMT3A, ETV6, EZH2, FBXW7, FGFR2, FGFR3, FLT3, GATA1, GATA2, GNAS, IDH1, IDH2, IKZF3, JAK1, JAK2, JAK3, KDM6A, KIT, KRAS, MPL, MYD88, NOTCH1, NPM1, NRAS, PDGFRA, PH6F, PML, PTPN11, RARA, SETBP1, SF3B1, SLC29A1, SRSF2, U2AF1, WT1, XPO1</i> <i>Expression Genes Analyzed: CD274, CEBPA, CTLA4, FLT3, ID4, IRF4, IRF8, MECOM, MUC1, MYC, PDCD1, PDCD1LG2, WT1 Fusion, Splicing, Exon Skipping Genes Scanned: ABL1, BCR, CFBF, CHD1, CHIC2, CREBBP, CSF1R, ERG, ETV6, FGFR1, GLIS2, IKZF1, IKZF3, JAK2, KAT6A, KMT2A, MECOM, MKL1, MLLT4, MLLT10, MYC, MYH11, NF1, NOTCH1, NUP98, NUP214, PDCD1LG2, PDGFRA, PDGFRB, PICALM, PML, RARA, RBM15, ROS1, RUNX1, RUNX1T1, SETBP1, SETD2, TCF3, TFG</i>	<input type="checkbox"/> Irinotecan toxicity (UGT1A1) <input checked="" type="checkbox"/> JAK2 Gene Exon 12 Sequence Analysis <input type="checkbox"/> MPL Gene Exon 10 Sequence Analysis <input type="checkbox"/> B cell clonality test <input type="checkbox"/> T cell clonality test <input type="checkbox"/> Chimerism Preparation - Donor <input type="checkbox"/> Chimerism Preparation - Buyer <input type="checkbox"/> Chimerism Analysis - Recipient Tracking
<input type="checkbox"/> <b>ALL Fusion NGS Panel (Fusion, SNV/Indel)</b> <i>SNV/Indel studied genes: ABL1, BCL2, BRAF, CREBBP, CRLF2, ETV6, EZH2, FBXW7, FLT3, IDH1, IDH2, IKZF3, IL7R, JAK1, JAK2, JAK3, KDM6A, KRAS, MPL, NOTCH1, NRAS, NT5C2, PAX5, PBX1, PTPN11, SH2B3, STAT3, STAT5B, TYK2, WT1 Genes with Expression Analysis: ABL1, AICDA, BCL2, BCL6, BLNK, CD274, CRLF2, CTLA4, DNNT, FLT3, HOXA9, HOXA10, IRF4, IRF8, LMO1, LYL1, MYC, PDCD1, PDCD1LG2, RAG1, RAG2, SOX11, TAL1, TLX1, TLX3, WT1 Fusion, Splicing, Exon Skipping Genes Scanned: ABL1, ABL2, BCL2, BCL6, BCR, CREBBP, CRLF2, CSF1R, EBF1, EPOR, ETV6, FGFR1, IKZF1, IKZF2, IKZF3, JAK2, KLF2, KMT2A, MLLT4, MYC, NF1, NOTCH1, NTRK3, NUP98, NUP214, P2RY8, PAG1, PAX5, PBX1, PDCD1LG2, PDGFRA, PDGFRB, PICALM, PTK2B, RUNX1, SEMA6A, SETD2, STIL, TAL1, TCF3, TYK2, ZCCHC7</i>	<input type="checkbox"/> Digital Droplet PCR Translocation (9;22)(q34;q11.2) <input checked="" type="checkbox"/> PCR - Translocation (9;22)(q34;q11.2) (p190) <input type="checkbox"/> PCR - Translocation (9;22)(q34;q11.2) (p210) <input type="checkbox"/> PCR - JAK2 V617F mutation <input type="checkbox"/> Digital Droplet PCR - JAK2 V617F mutation <input type="checkbox"/> PCR - FIP1L1/PDGFR fusion, del 4q12 <input type="checkbox"/> PCR -Translocation (5;12)(q33;p13)(ETV6/ACSL6) <input type="checkbox"/> PCR - Translocation (15;17)(q22;q21)(PML/RARA) <input type="checkbox"/> PCR - Translocation (8;21)(q22; q22) (RUNX1/RUNXT1) <input type="checkbox"/> PCR - CFBF-MYH11 Inversion(16)(p13;q22) <input type="checkbox"/> PCR - Translocation (1;19)(q23;p13) (E2A/PBX1) <input type="checkbox"/> PCR - Translocation (4;11)(q21;q23) (MLL/AF4) <input type="checkbox"/> PCR - Translocation (12;21)(p12;q22) <input type="checkbox"/> Thiopurine Pharmacogenetic Test (TPMT gene) <input type="checkbox"/> FLT3 Gene Exon 14, 15, 20 Sequence Analysis <input type="checkbox"/> GATA1 Gene Exon 2 Sequence Analysis <input type="checkbox"/> IDH1 Gene Exon 4 Sequence Analysis
<input type="checkbox"/> <b>Lymphoma Fusion NGS Panel (Fusion, SNV/Indel)</b> <i>SNV/Indel: AKT3, BAX, BCL2, BIRC3, BRAF, BTK, CARD11, CCND1, CD79B, CREBBP, DNMT3A, ETV6, EZH2, FBXW7, IDH1, IDH2, JAK1, JAK3, KRAS, MYD88, NOTCH1, NOTCH2, NRAS, PLCG1, PLCG2, RHOA, SF3B1, STAT3, STAT5B, STAT6, WT1, CD44, CDC25A, CDKN2A, CDKN2B, CECPD, CEPE, CEPPG, CREB3L2, CTLA4, CYB5R2, DENND3, DLEU1, DNMT3B, DNNT, E2F2, ENTDP1, EXOC2, FAM216A, FOXP1, FUT8, IL16, IRF4, IRF8, ITPKB, 1, LIMD1, LMO2, LRMP, LZTS1, MAL, MAML3, MME, MUC1, MYBL1, MYC, NEK6, NFKB1, NME1, PAICS, PDCD1, PDCD1LG2, PIM1, PIM2, PPAT, PRKAR2B, PTPN1, PYCR1, RAB29, RAG1, RAG2, RANBP1, S1PR2, SERPINA9, SH3PB5, STRBP, TNFRSF13B, TNFSF4, WT1 Fusion, Splicing, Exon Skipping: AKT3, ALK, BCL2, BCL6, BCR, BIRC3, CFBF, CCND1, CDK6, CHIC2, CIITA, DEK, DUSP22, EIF4A1, ETV6, JAK2, KMT2A, MALT1, MKL1, MLF1, MLLT10, MYC, NFKB2, NOTCH1, P2RY8, PDCD1LG2, PDGFRA, PRDM16, STIL, TCF3, TP63</i>	<input type="checkbox"/> IDH2 Gene Exon 4 Sequence Analysis <input type="checkbox"/> Perforin Gene Sequence Analysis <input type="checkbox"/> STX11 (Syntaxin 11) Gene Sequence Analysis <input type="checkbox"/> Sequence analysis of the Munc 13-4 (UNC13D) Gene <input type="checkbox"/> CALR Gene Exon 9 Sequence Analysis <input type="checkbox"/> CEBPA Gene Sequence Analysis <input type="checkbox"/> CSF3R Gene Sequence Analysis <input type="checkbox"/> ASXL1 Gene Exon 12 Sequence Analysis <input type="checkbox"/> Kostmann (HAX1 Gene Sequence Analysis) <input type="checkbox"/> NPM1 Mutation Analysis (MutA, MutB, MutD) <input type="checkbox"/> Shwachman-Diamond SBDS Sequence Analysis <input type="checkbox"/> PDGFRB Gene Exon 12-20 Sequence Analysis
<input type="checkbox"/> <b>Ph-like Leukemia Fusion NGS Panel (Fusion, SNV/Indel)</b> <i>SNV/Indel studied genes: ABL1, BCL2, BRAF, CREBBP, CRLF2, ETV6, EZH2, FBXW7, FLT3, IDH1, IDH2, IKZF3, IL7R, JAK1, JAK2, JAK3, KDM6A, KRAS, MPL, NOTCH1, NRAS, NT5C2, PAX5, PBX1, PTPN11, SH2B3, STAT3, STAT5B, TYK2, WT1 Genes with Expression Analysis: ABL1, AICDA, BCL2, BCL6, BLNK, CD274, CRLF2, CTLA4, DNNT, FLT3, HOXA9, HOXA10, IRF4, IRF8, LMO1, LYL1, MYC, PDCD1, PDCD1LG2, RAG1, RAG2, SOX11, TAL1, TLX1, TLX3, WT1 Fusion, Splicing, Exon Skipping Genes Scanned: ABL1, ABL2, BCL2, BCL6, BCR, CREBBP, CRLF2, CSF1R, EBF1, EPOR, ETV6, FGFR1, IKZF1, IKZF2, IKZF3, JAK2, KLF2, KMT2A, MLLT4, MYC, NF1, NOTCH1, NTRK3, NUP98, NUP214, P2RY8, PAG1, PAX5, PBX1, PDCD1LG2, PDGFRA, PDGFRB, PICALM, PTK2B, RUNX1, SEMA6A, SETD2, STIL, TAL1, TCF3, TYK2, ZCCHC7</i>	
<input type="checkbox"/> <b>BCR IGH-V Somatic Mutation NGS Panel</b>	
<input type="checkbox"/> <b>TCR NGS Panel</b>	
<input type="checkbox"/> <b>Myeloid &amp; Lymphoid NGS Panel (SNV/Indel)</b>	
<input type="checkbox"/> <b>Imatinib Resistance</b>	
<input type="checkbox"/> <i>G250E, Y253H, E255K, T315I, F317L, M351T mutations</i>	

**CYTOGENETICS - CHROMOSOME TESTS**

- |   |   |
|---|---|
| <input type="checkbox"/> Chromosome analysis in bone marrow aspiration material | <input type="checkbox"/> Chromosome analysis from peripheral blood (Leukemia) |
| <input type="checkbox"/> DEB Test (Fanconi aplastic anemia)                     | <input type="checkbox"/> Mitomycin C Test                                     |

**MOLECULAR CYTOGENETICS - FISH TESTS**

- |   |  |
|---|--|
| <input type="checkbox"/> FISH- 17p13.1 (TP53) deletion                                    | <input type="checkbox"/> FISH - 4q12 rearrangements (FIP1LI/PDGFRB fusion)     |
| <input type="checkbox"/> FISH - 5q32 (PDGFRB) rearrangements                              | <input type="checkbox"/> FISH - 5q31 deletion (EGR1)                           |
| <input type="checkbox"/> FISH - 10q24 (TLX1) rearrangements                               | <input type="checkbox"/> Fish - Monosomy / Trisomy 8                           |
| <input type="checkbox"/> FISH - 11q22.3 (ATM) deletion                                    | <input type="checkbox"/> FISH - Monosomy/Trisomy 5                             |
| <input type="checkbox"/> FISH - 11q23 (MLL) rearrangements                                | <input type="checkbox"/> FISH - Monosomy/Trisomy 7                             |
| <input type="checkbox"/> FISH - 13q14.3 (D13S319)deletion                                 | <input type="checkbox"/> FISH - 8q24 (MYC) rearrangements                      |
| <input type="checkbox"/> FISH - 13q14.3 (RB1) deletion                                    | <input type="checkbox"/> FISH - 17q21 (RARA) rearrangements                    |
| <input type="checkbox"/> FISH - 1q21 (CKS1B)/1q32-36 (CDKN2C) Amplification/Deletion      | <input type="checkbox"/> FISH - SS18 remasters                                 |
| <input type="checkbox"/> FISH - 20q12 (D20S108) deletion                                  | <input type="checkbox"/> FISH - Translocation (1;19)(q23;p13) (TCF3/PBX1)      |
| <input type="checkbox"/> FISH - 13q34 deletion  | <input type="checkbox"/> FISH - Translocation (11;14)(q13;q32)(IGH/CCND1)      |
| <input type="checkbox"/> FISH - 14q11.2 (TCR) rearrangements                              | <input type="checkbox"/> FISH - Translocation (11;18)(q21;q21) (API2/MALT1)    |
| <input type="checkbox"/> FISH - 14q32 deletion  | <input type="checkbox"/> FISH - Translocation (11;19)(q23;p13.1) (MLL/ENL)     |
| <input type="checkbox"/> FISH - 5q35.1 TLX3 rearrangements                                | <input type="checkbox"/> FISH - Translocation (12;21)(p12;q22)(ETV6/RUNX1)     |
| <input type="checkbox"/> FISH - 6q23 (MYB) deletion                                       | <input type="checkbox"/> FISH - Translocation (14;16)(q32;q23) (MAF/IGH)       |
| <input type="checkbox"/> FISH - 7q22/del 7q35 (MDS) deletion                              | <input type="checkbox"/> FISH - Translocation (14;18)(q32;q21)(IGH/BCL2)       |
| <input type="checkbox"/> FISH - 20q13.2 (LSI ZNF217)deletion                              | <input type="checkbox"/> FISH - Translocation (14;20)(q32;q12)(IGH/MAFB)       |
| <input type="checkbox"/> FISH - 5q33-q34 (CSF1R) deletion                                 | <input type="checkbox"/> FISH - Translocation (15;17)(q22;q21)(PML/RARA)       |
| <input type="checkbox"/> FISH - 7q31 deletion   | <input type="checkbox"/> FISH - Translocation (4;11)(q21;q23) (MLL/AFF1)       |
| <input type="checkbox"/> FISH - 7q34 (TCRB) rearrangements                                | <input type="checkbox"/> FISH - Translocation (4;14)(p16;q32)(IGH/FGFR3)       |
| <input type="checkbox"/> FISH - 9p21 (CDKN2A) deletion                                    | <input type="checkbox"/> FISH - Translocation (6;14)(p21;q32)(CCND3/IGH)       |
| <input type="checkbox"/> FISH - 9q34 (ASS1) deletion                                      | <input type="checkbox"/> FISH - Translocation (6;9)(p22;q34) (DEK/NUP214)      |
| <input type="checkbox"/> FISH - 18q21 (BCL2) rearrangements                               | <input type="checkbox"/> FISH - Translocation (8;14)(q24; q32)(MYC/IGH)        |
| <input type="checkbox"/> FISH - 3q27 (BCL6) rearrangements                                | <input type="checkbox"/> FISH - Translocation (8;21)(q22; q22) (RUNX1/RUNX1T1) |
| <input type="checkbox"/> FISH - 11q13 (CCND1) rearrangements                              | <input type="checkbox"/> FISH - Translocation (9;11)(p22;q23) (AF9/MLL)        |
| <input type="checkbox"/> FISH - DDIT3 rearrangements                                      | <input type="checkbox"/> FISH - Translocation (9;22)(q34;q11.2)(BCR/ABL)       |
| <input type="checkbox"/> FISH - EWSR1 rearrangements                                      | <input type="checkbox"/> FISH - Trisomy 10                                     |
| <input type="checkbox"/> FISH - Translocation (14;18)(q32;q21) (IGH/MALT1)                | <input type="checkbox"/> FISH - Trisomy 11                                     |
| <input type="checkbox"/> FISH - Inversion (16)(p13;q22) / Translocation (16;16)(p13;q22)  | <input type="checkbox"/> FISH - Trisomy 12                                     |
| <input type="checkbox"/> FISH - Inversion (3)(q21;q26.2) / Translocation(3;3)(q21;q26.2)  | <input type="checkbox"/> FISH - Trisomy 17                                     |
| <input type="checkbox"/> FISH - Chimerism Analysis (Transplantation from Different Sexes) | <input type="checkbox"/> FISH - Trisomy 4                                      |
| <input type="checkbox"/> FISH - 2p12 (IGK) rearrangements                                 | <input type="checkbox"/> FISH - Trisomy 6                                      |
| <input type="checkbox"/> FISH - 22q11 IGL rearrangements                                  | <input type="checkbox"/> FISH - C-MET Amplifications                           |

\*For tests and research projects not listed here, please contact us.  
international@sapiens.com.tr