

Bölüm 2

KLL'DE OLUŞAN GENETİK DEĞİŞİKLİKLER

Sedanur KARAMAN GÜLSARAN¹

GİRİŞ

Kronik lenfosittik lösemi (KLL) yaşlı nüfusun en yaygın erişkin lösemi hastalığıdır. KLL oldukça heterojen bir seyir göstermektedir. Bazı hastalar için takip yeterli iken, bazılarında hastalık hızlı seyir göstermektedir^[1]. Bu klinik heterojenite, altta yatan moleküler ve hücresel çeşitliliğe bağlıdır^[2]. KLL genetiğinin daha iyi anlaşılması, bu hastalığın prognozunu belirlemede ve tedavide önemlidir.

Floresan insitu hibridizasyon (FISH) yöntemi kullanılarak KLL hastalarının %80'ninde sitogenetik anormallikler saptanabilmektedir^[3, 4]. Konvansiyonel kar-yotip analizinde ise hastaların ancak %20-50'sinde kromozomal anomaliler saptanmaktadır; bu nedenle FISH yöntemi konvansiyonel sitogenetik yöntemle göre daha sensitiftir^[5]. KLL hastalarında FISH yöntemi kullanarak yapılan çalışmalarda del 13q14 en sık görülen genetik anormallik olup tüm hastaların yaklaşık %50'sinde bulunmaktadır. Ayrıca, hastaların yaklaşık %15-20'sinde trizomi 12 ve %10-15'inde 11q22.3 genetik mutasyonu görülmektedir. Tespit edilen diğer anormallikler ise del 6q21 ve del 17p13.1 olup, bu anormallikler hastalık sonuçları üzerinde farklı prognostik etki yaratmaktadır^[6]. Kompleks genomik değişiklik daha önce tedavi edilmemiş hastaların %15-30'unda saptanmaktadır ve hızlı hastalık progresyonu, Richter transformasyonu ile ilişkili olup bu hastalarda prognoz daha kötüdür^[7, 8].

Yapılan sitogenetik incelemeler ve FISH yöntemi ile hastalığın erken dönemlerinde tekrarlayan aberran mutasyonlar sıklıkla gözlenmezken, hastalığın seyri sırasında mutasyonların ortaya çıktığı görülmüştür^[9]. Bu nedenle tedavi değişikliği planlanırken tekrar FISH analizlerinin yapılması önerilmektedir^[10].

¹ Uzman doktor, Trakya Üniversitesi Tıp Fakültesi, sedakaraman@hotmail.com

BIRC3

Baculoviral IAP Repeat Containing 3 (BIRC3) geni, alternatif NF-kB sinyal yolunun negatif regülatörüdür. Olguların %5-8'inden daha azında görülen BIRC3 gen mutasyonu alternatif NF-kB yolunun aktivasyonuna yol açar^[63]. ATM ve BIRC3 genleri kromozom 11q bölgesinde bulunur. Tanı anında %4 BIRC3 mutasyonu saptanır iken kemoterapiye dirençli KLL'de %24'e çıkabilmektedir.

KOMPLEKS KARYOTİP VE TRANSLOKASYON

Kompleks karyotip, konvansiyonel kromozom bantlama yöntemi ile 3 veya daha fazla yapısal kromozomal anormallik varlığı olmasıdır. KLL'de kötü prognoz ile seyretmektedir. Kompleks karyotipe sahip olan hastalar normal karyotipli hastalara göre prognozu daha kötü seyretmektedir.

Translokasyonlar, KLL'de anlamlı bir alt küme oluşturmaz ve kompleks karyotipin veya trizomi 12 veya del13 gibi sitogenetik anormalliklerin bir parçası olma eğilimindedir^[78, 79]. Çok nadir görülmekle birlikte interfaz FISH ile yapılan küçük çalışmalarda KLL hastalarında %0-2 oranında t(14;18) bildirilmiştir^[80-82]. Konvansiyonel sitogenetik yöntemlerle yapılan çalışmalarda Sen ve arkadaşları^[83] 2215 KLL hastasının 2'sinde; Juliusson ve arkadaşları^[84] ise 640 KLL hastasının 3'ünde t(14;18) tespit etmiştir. KLL'deki t(14;18) moleküler yöntemlerle çalışılmış ve farklı BCL-2 değerlerine sahip olduğu, folliküler B hücreli lenfomadan farklı olduğu gösterilmiştir.

SONUÇ

KLL heterojen seyir gösteren bir hastalık olup, hastalığın tanı anından takip eden süreçte yeni genetik anomaliler, mutasyonlar gelişebilmektedir. Tanı anında genomik olarak risk değerlendirmesi, hasta yönetimi için yararlı olmaktadır^[85]. Hedefe yönelik tedavilerin, konvansiyonel tedavilerin önüne geçtiği dönemde, genetik bilgi hastalığın tanısında ve tedavisini planlamak için klinisyenlere yol gösterici olmaktadır.

Anahtar Kelimeler: KLL, genetik

KAYNAKÇA

1. Chiorazzi N, Rai KR and Ferrarini M. Chronic lymphocytic leukemia. *New England Journal of Medicine*, 2005. 352(8): p. 804-815.
2. Chiorazzi N and Ferrarini M. Cellular origin (s) of chronic lymphocytic leukemia: cautionary notes and additional considerations and possibilities. *Blood*, 2011. 117(6): p. 1781-1791.
3. Döhner H, Stilgenbauer S, Benner A, et al. Genomic aberrations and survival in chronic lymphocytic leukemia. *New England Journal of Medicine*, 2000. 343(26): p. 1910-1916.
4. Puente XS, Pinyol M, Quesada V, et al. Whole-genome sequencing identifies recurrent mutations in chronic lymphocytic leukaemia. *Nature*, 2011. 475(7354): p. 101.

5. Glassman AB and Hayes KJ. The value of fluorescence in situ hybridization in the diagnosis and prognosis of chronic lymphocytic leukemia. *Cancer genetics and cytogenetics*, 2005. 158(1): p. 88-91.
6. Malek S. Molecular biomarkers in chronic lymphocytic leukemia, in *Advances in Chronic Lymphocytic Leukemia*. 2013, Springer. p. 193-214.
7. Ouillette P, Fossum S, Parkin B, et al. Aggressive chronic lymphocytic leukemia with elevated genomic complexity is associated with multiple gene defects in the response to DNA double-strand breaks. *Clinical Cancer Research*, 2010. 16(3): p. 835-847.
8. Kujawski L, Ouillette P, Erba H, et al. Genomic complexity identifies patients with aggressive chronic lymphocytic leukemia. *Blood*, 2008. 112(5): p. 1993-2003.
9. Shanafelt TD, Witzig TE, Fink SR, et al. Prospective evaluation of clonal evolution during long-term follow-up of patients with untreated early-stage chronic lymphocytic leukemia. *Journal of Clinical Oncology*, 2006. 24(28): p. 4634-4641.
10. Hallek M, Cheson BD, Catovsky D, et al. Guidelines for the diagnosis and treatment of chronic lymphocytic leukemia: a report from the International Workshop on Chronic Lymphocytic Leukemia updating the National Cancer Institute-Working Group 1996 guidelines. *Blood*, 2008. 111(12): p. 5446-5456.
11. Fitchett M, Griffiths MJ, Oscier DG, et al. Chromosome abnormalities involving band 13q14 in hematologic malignancies. *Cancer genetics and cytogenetics*, 1987. 24(1): p. 143-150.
12. Juliusson G, Oscier DG, Fitchett M, et al. Prognostic subgroups in B-cell chronic lymphocytic leukemia defined by specific chromosomal abnormalities. *New England Journal of Medicine*, 1990. 323(11): p. 720-724.
13. Peterson LC, Lindquist LL, Church S, et al. Frequent clonal abnormalities of chromosome band 13q 14 in b-cell chronic lymphocytic leukemia: Multiple clones, subclones, and nonclonal alterations in 82 midwestern patients. *Genes, Chromosomes and Cancer*, 1992. 4(4): p. 273-280.
14. Zech L and Mellstedt H. Chromosome 13 new marker for B-cell chronic lymphocytic leukemia. *Hereditas*, 1988. 108(1): p. 77-84.
15. Stilgenbauer S, Lichter P, and Döhner H. Genetic features of B-cell chronic lymphocytic leukemia. *Reviews in clinical and experimental hematology*, 2000. 4(1): p. 48-72.
16. Van Dyke DL, Shanafelt TD, Call TG, et al. A comprehensive evaluation of the prognostic significance of 13q deletions in patients with B-chronic lymphocytic leukaemia. *British journal of haematology*, 2010. 148(4): p. 544-550.
17. Van Dyke DL, Werner L, Rassenti LZ, et al. The Dohner fluorescence in situ hybridization prognostic classification of chronic lymphocytic leukaemia (CLL): the CLL Research Consortium experience. *British journal of haematology*, 2016. 173(1): p. 105-113.
18. Garg R, Wierda W, Ferrajoli A, et al. The prognostic difference of monoallelic versus biallelic deletion of 13q in chronic lymphocytic leukemia. *Cancer*, 2012. 118(14): p. 3531-3537.
19. Sparkes RS, Sparkes MC, Wilson MG, et al. Regional assignment of genes for human esterase D and retinoblastoma to chromosome band 13q14. *Science*, 1980. 208(4447): p. 1042-1044.
20. Weinberg RA. The retinoblastoma protein and cell cycle control. *cell*, 1995. 81(3): p. 323-330.
21. Liu Y, D Grandér D, Einhorn S, et al. Retinoblastoma gene deletions in B-cell chronic lymphocytic leukemia. *Genes, Chromosomes and Cancer*, 1992. 4(3): p. 250-256.
22. Neubauer A, Kant E, Rochlitz C, et al. Altered expression of the retinoblastoma susceptibility gene in chronic lymphocytic leukaemia. *British journal of haematology*, 1993. 85(3): p. 498-503.
23. Garcia-Marco JA, Caldas C, Price CM, et al. Frequent somatic deletion of the 13q12. 3 locus encompassing BRCA2 in chronic lymphocytic leukemia. *Blood*, 1996. 88(5): p. 1568-1575.
24. Kaur P. *Chronic Lymphocytic Leukemia*. 2018: Springer.
25. Gaidano G, Ballerini P, Gong JZ, et al. p53 mutations in human lymphoid malignancies: association with Burkitt lymphoma and chronic lymphocytic leukemia. *Proceedings of the National Academy of Sciences*, 1991. 88(12): p. 5413-5417.
26. El Roubi S, Thomas A, Costin D, et al. p53 gene mutation in B-cell chronic lymphocytic leukemia is associated with drug resistance and is independent of MDR1/MDR3 gene expression. *Blood*, 1993. 82(11): p. 3452-3459.

27. Fenaux P, Preudhomme C, Lai JL, et al. Mutations of the p53 gene in B-cell chronic lymphocytic leukemia: a report on 39 cases with cytogenetic analysis. *Leukemia*, 1992. 6(4): p. 246-250.
28. Gaidano G, Newcomb EW, Gong JZ, et al. Analysis of alterations of oncogenes and tumor suppressor genes in chronic lymphocytic leukemia. *The American journal of pathology*, 1994. 144(6): p. 1312.
29. Dohner H, Fischer K, Bentz M, et al. p53 gene deletion predicts for poor survival and non-response to therapy with purine analogs in chronic B-cell leukemias. *Blood*, 1995. 85(6): p. 1580-1589.
30. Hoffman R, Benz Jr EJ, Silberstein LE, et al. Hematology: basic principles and practice. 2013: *Elsevier Health Sciences*.
31. Cordone I, Masi S, Mauro FR, et al. p53 expression in B-cell chronic lymphocytic leukemia: a marker of disease progression and poor prognosis. *Blood*, 1998. 91(11): p. 4342-4349.
32. Grever MR, Lucas DM, Dewald GW, et al. Comprehensive assessment of genetic and molecular features predicting outcome in patients with chronic lymphocytic leukemia: results from the US Intergroup Phase III Trial E2997. *Journal of Clinical Oncology*, 2007. 25(7): p. 799-804.
33. Lozanski G, Heerema NA, Flinn IW, et al. Alemtuzumab is an effective therapy for chronic lymphocytic leukemia with p53 mutations and deletions. *Blood*, 2004. 103(9): p. 3278-3281.
34. Tam CS, Shanafelt TD, Wierda WG, et al. De novo deletion 17p13. 1 chronic lymphocytic leukemia shows significant clinical heterogeneity: the MD Anderson and Mayo Clinic experience. *Blood*, 2009. 114(5): p. 957-964.
35. Döhner H, Stilgenbauer S, James MR, et al. 11q deletions identify a new subset of B-cell chronic lymphocytic leukemia characterized by extensive nodal involvement and inferior prognosis. *Blood*, 1997. 89(7): p. 2516-2522.
36. Stilgenbauer S, Liebisch P, James MR et al. Molecular cytogenetic delineation of a novel critical genomic region in chromosome bands 11q22. 3-923.1 in lymphoproliferative disorders. *Proceedings of the National Academy of Sciences*, 1996. 93(21): p. 11837-11841.
37. Austen B, Skowronska A, Baker C, et al. Mutation status of the residual ATM allele is an important determinant of the cellular response to chemotherapy and survival in patients with chronic lymphocytic leukemia containing an 11q deletion. *Journal of Clinical Oncology*, 2007. 25(34): p. 5448-5457.
38. Skowronska A, Parker A, Ahmed G, et al. Biallelic ATM inactivation significantly reduces survival in patients treated on the United Kingdom Leukemia Research Fund Chronic Lymphocytic Leukemia 4 trial. *Journal of Clinical Oncology*, 2012. 30(36): p. 4524-4532.
39. Tsujimoto Y, Yunis J, Onorato-Showe L, et al. Molecular cloning of the chromosomal breakpoint of B-cell lymphomas and leukemias with the t (11; 14) chromosome translocation. *Science*, 1984. 224(4656): p. 1403-1406.
40. Gahrton G, Robert KH, Friberg K, et al. Nonrandom chromosomal aberrations in chronic lymphocytic leukemia revealed by polyclonal B-cell-mitogen stimulation. *Blood*, 1980. 56(4): p. 640-647.
41. Strati P, Abruzzo LV, Wierda WG, et al. Second cancers and Richter transformation are the leading causes of death in patients with trisomy 12 chronic lymphocytic leukemia. *Clinical Lymphoma Myeloma and Leukemia*, 2015. 15(7): p. 420-427.
42. Fegan C., et al. Karyotypic evolution in CLL: identification of a new sub-group of patients with deletions of 11q and advanced or progressive disease. *Leukemia*, 1995. 9(12): p. 2003-2008.
43. Garcia-Marco J., et al. Correlation of trisomy 12 with proliferating cells by combined immunocytochemistry and fluorescence in situ hybridization in chronic lymphocytic leukemia. *Leukemia*, 1996. 10(11): p. 1705-1711.
44. Auer RL, Bienz N, Neilson J, et al. The sequential analysis of trisomy 12 in B-cell chronic lymphocytic leukaemia. *British journal of haematology*, 1999. 104(4): p. 742-744.
45. Juliusson G. and Gahrton G. Chromosome aberrations in B-cell chronic lymphocytic leukemia: pathogenetic and clinical implications. *Cancer genetics and cytogenetics*, 1990. 45(2): p. 143-160.

46. Han T, Ozer H, Sadamori N, et al. Prognostic importance of cytogenetic abnormalities in patients with chronic lymphocytic leukemia. *New England Journal of Medicine*, 1984. 310(5): p. 288-292.
47. Pittman S and Catovsky D. Prognostic significance of chromosome abnormalities in chronic lymphocytic leukaemia. *British journal of haematology*, 1984. 58(4): p. 649-660.
48. Stilgenbauer S, Bullinger L, Lichter P, et al. Genetics of chronic lymphocytic leukemia: genomic aberrations and V [subH] gene mutations status in pathogenesis and clinical course. *Leukemia* (08876924), 2002. 16(6).
49. Montserrat E. Classical and new prognostic factors in chronic lymphocytic leukemia: where to now? *The Hematology Journal*, 2002. 3(1): p. 7-9.
50. Döhner H. Prognostic implications of findings from cytogenetics and molecular genetics. *Chronic Lymphocytic Leukemia: Case-Based session Hematology*, 2001: p. 141-5.
51. Naylor M and Capra JD. Mutational status of Ig VH genes provides clinically valuable information in B-cell chronic lymphocytic leukemia. *Blood*, 1999. 94(6): p. 1837-1839.
52. Soysal T. Kronik Lenfositik Lösemi: Güncel Yaklaşımlar. XXIX. *Ulusal Hematoloji Kongresi*, 2002, Antalya.
53. Hamblin TJ, Davis Z, Gardiner A, et al. Unmutated Ig VH genes are associated with a more aggressive form of chronic lymphocytic leukemia. *Blood*, 1999. 94(6): p. 1848-1854.
54. Damle RN, Wasil T, Fais F, et al. Ig V Gene Mutation Status and CD38 Expression As Novel Prognostic Indicators in Chronic Lymphocytic Leukemia: Presented in part at the 40th Annual Meeting of The American Society of Hematology, held in Miami Beach, FL, December 4-8, 1998. *Blood*, 1999. 94(6): p. 1840-1847.
55. Kröber A, Seiler T, Benner A, et al. VH mutation status, CD38 expression level, genomic aberrations, and survival in chronic lymphocytic leukemia. *Blood*, 2002. 100(4): p. 1410-1416.
56. Hamblin TJ, Orchard JA, Ibbotson RE, et al. CD38 expression and immunoglobulin variable region mutations are independent prognostic variables in chronic lymphocytic leukemia, but CD38 expression may vary during the course of the disease. *Blood*, 2002. 99(3): p. 1023-1029.
57. Oscier DG, Gardiner AC, Mould SJ, et al. Multivariate analysis of prognostic factors in CLL: clinical stage, IGVH gene mutational status, and loss or mutation of the p53 gene are independent prognostic factors. *Blood*, 2002. 100(4): p. 1177-1184.
58. Wan Y and Wu CJ, SF3B1 mutations in chronic lymphocytic leukemia. *Blood*, 2013. 121(23): p. 4627-4634.
59. Wang L, Lawrence MS, Wan Y, et al. SF3B1 and other novel cancer genes in chronic lymphocytic leukemia. *New England Journal of Medicine*, 2011. 365(26): p. 2497-2506.
60. Landau DA, Carter SL, Stojanov P, et al. Evolution and impact of subclonal mutations in chronic lymphocytic leukemia. *Cell*, 2013. 152(4): p. 714-726.
61. Fabbri G, Rasi S, Rossi D, et al. Analysis of the chronic lymphocytic leukemia coding genome: role of NOTCH1 mutational activation. *Journal of Experimental Medicine*, 2011. 208(7): p. 1389-1401.
62. Quesada V, Conde L, Villamor N, et al. Exome sequencing identifies recurrent mutations of the splicing factor SF3B1 gene in chronic lymphocytic leukemia. *Nature genetics*, 2012. 44(1): p. 47.
63. Rossi D, Fangazio M, Rasi S, et al. Disruption of BIRC3 associates with fludarabine chemorefractoriness in TP53 wild-type chronic lymphocytic leukemia. *Blood*, 2012. 119(12): p. 2854-2862.
64. Landau DA and Wu CJ. Chronic lymphocytic leukemia: molecular heterogeneity revealed by high-throughput genomics. *Genome medicine*, 2013. 5(5): p. 47.
65. Rosati E., et al. Constitutively activated Notch signaling is involved in survival and apoptosis resistance of B-CLL cells. *Blood*, 2009. 113(4): p. 856-865.
66. Rossi D, Rasi S, Fabbri G, et al. Mutations of NOTCH1 are an independent predictor of survival in chronic lymphocytic leukemia. *Blood*, 2012. 119(2): p. 521-529.
67. Balatti V, Bottoni A, Palamarchuk A, et al. NOTCH1 mutations in CLL associated with trisomy 12. *Blood*, 2012. 119(2): p. 329-331.

68. Oscier DG, Rose-Zerilli MJ, Winkelmann N, et al. The clinical significance of NOTCH1 and SF3B1 mutations in the UK LRF CLL4 trial. *Blood*, 2013. 121(3): p. 468-475.
69. Villamor N, Conde L, Martínez-Trilloset A, et al. NOTCH1 mutations identify a genetic subgroup of chronic lymphocytic leukemia patients with high risk of transformation and poor outcome. *Leukemia*, 2013. 27(5): p. 1100.
70. De Keersmaecker K, Michaux L, Bosly A, et al. Rearrangement of NOTCH1 or BCL3 can independently trigger progression of CLL. *Blood*, 2012. 119(16): p. 3864-3866.
71. Malcovati L, Papaemmanuil E, Bowen DT, et al. Clinical significance of SF3B1 mutations in myelodysplastic syndromes and myelodysplastic/myeloproliferative neoplasms. *Blood*, 2011. 118(24): p. 6239-6246.
72. Koboldt DC, Fulton RS, McLellan MD, et al. Comprehensive molecular portraits of human breast tumours. *Nature*, 2012. 490(7418): p. 61-70.
73. Rossi D, Brusca A, Spina V, et al. Mutations of the SF3B1 splicing factor in chronic lymphocytic leukemia: association with progression and fludarabine-refractoriness. *Blood*, 2011. 118(26): p. 6904-6908.
74. O'Neill LA and Bowie AG. The family of five: TIR-domain-containing adaptors in Toll-like receptor signalling. *Nature Reviews Immunology*, 2007. 7(5): p. 353.
75. Ngo VN, Young RM, Schmitz R, et al. Oncogenically active MYD88 mutations in human lymphoma. *Nature*, 2011. 470(7332): p. 115.
76. Zenz T, Mertens D, Küppers R, et al. From pathogenesis to treatment of chronic lymphocytic leukaemia. *Nature Reviews Cancer*, 2010. 10(1): p. 37.
77. Zenz T, Mertens D and Stilgenbauer S. Biological diversity and risk-adapted treatment of chronic lymphocytic leukemia. 2010, *Haematologica*.
78. Miller CR, Stephens D, Ruppert AS, et al. Jumping translocations, a novel finding in chronic lymphocytic leukaemia. *British journal of haematology*, 2015. 170(2): p. 200-207.
79. De Braekeleer M, Tous C, Guéganic N, et al. Immunoglobulin gene translocations in chronic lymphocytic leukemia: a report of 35 patients and review of the literature. *Molecular and clinical oncology*, 2016. 4(5): p. 682-694.
80. Yabumoto K, Ohno H, Doi S, et al. Involvement of the BCL3 gene in two patients with chronic lymphocytic leukemia. *International journal of hematology*, 1994. 59(3): p. 211-218.
81. Dyer MJS and Oscier D. The configuration of the immunoglobulin genes in B cell chronic lymphocytic leukemia. *Leukemia*, 2002. 16(6): p. 973.
82. Aoun P, Blair HE, Smith LM, et al. Fluorescence in situ hybridization detection of cytogenetic abnormalities in B-cell chronic lymphocytic leukemia/small lymphocytic lymphoma. *Leukemia & lymphoma*, 2004. 45(8): p. 1595-1603.
83. Sen F, Lai R and Albitar M. Chronic lymphocytic leukemia with t (14; 18) and trisomy 12: report of 2 cases and review of the literature. *Archives of pathology & laboratory medicine*, 2002. 126(12): p. 1543-1546.
84. Juliusson G and Gahrton G. 5 Cytogenetics in CLL and related disorders. *Baillière's clinical haematology*, 1993. 6(4): p. 821-848.
85. Cheson BD, Bennett JM, Grever M, et al. National Cancer Institute-sponsored Working Group guidelines for chronic lymphocytic leukemia: revised guidelines for diagnosis and treatment. *Blood*, 1996. 87(12): p. 4990-4997.