

## Bölüm 9

# KÜÇÜK HÜCRELİ DIŐI AKCİĐER KANSERİNDE PREDİKTİF VE PROGNOSTİK BİYOBELİRTEÇLER

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### GİRİŐ

AkciĐer kanseri, Globocan 2012 verilerine göre dünya genelinde en sık tanı alan kanser ve kansere baĐlı ölümlerin başlıca nedenidir. 2012 yılında, dünya çapında yılda tahmini 1,8 milyon yeni olgu ve 1,59 milyon ölüm kaydedilmiştir.

AkciĐer Kanseri Mutasyon Konsorsiyumunun (Lung Cancer Mutation Consortium-LCMC) katılımcı kurumları, klinisyenlerin hedefli tedavileri seçmesine ve klinik araŐtırmalara hasta almasına yardımcı olmak için 10 geni onkojenik sürücü mutasyonlar açısından analiz etmiştir (1). Moleküler analize göre küçük hücreli dışı akciĐer adenokarsinomunda en yaygın üç onkojenik sürücü mutasyon; KRAS mutasyonları, EGFR mutasyonları, ALK rearranjmanları proliferasyon ve apoptoz dahil hücre fonksiyonlarını düzenleyen sinyal iletim yolaklarında rol oynar (2). AkciĐer kanserinde son zamanlarda hem patogenezi anlama hem de tedavi yaklaşımlarını belirlemede büyük gelişmeler yaşanmıştır. Bu gelişmelerden sürücü mutasyonlar/moleküler deĐişiklikler kanserin başlangıcı ve gelişiminden sorumludurlar.

Prediktif biyobelirteç tedavinin etkinliğini öngörür, çünkü biyobelirteç ve tedavi sonuçları arasında bir ilişki vardır. Prognostik biyobelirteç tümörün doğal agresifliğini gösterir. Çünkü verilen tedaviden baĐımsız olarak hastanın sağkalımı ile ilgili bir göstergedir. Burada küçük hücreli dışı akciĐer kanserinde (KH-DAK) en sık görülen biyobelirteçlerden bahsedeceĐiz.

### EGFR MUTASYONU

Epidermal growth factor receptor (EGFR) hücre membran reseptörüdür ve kromozom 7p11.2' de lokalizedir (3). Avian erythroblastic leukemia viral (v-erb-b) oncogene (ERBB1) olarak da bilinir (3). İnsanda yetersiz ERBB sinyali bazı nörodejeneratif hastalıklarla birlikte görülürken aşırı ERBB sinyali solid tümör gelişimi ile ilişkili bulunmuştur. (3). En sık mutasyonlar tirozin kinaz doma-

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mektedir (76). Metastatik nonsquamöz KHDAK'de EGFR ve ALK negatif olan ya da bilinmeyen hastalarda PD-L1 durumuna bakılmaksızın 1. basamakta kombinasyon olarak pembrolizumab/(ya da atezolizumab/ bevacizumab)/ kemoterapi NCCN 2019 3.versiyonda kanıt 1 düzeyinde önerilmektedir ( 76,77). Squamöz KHDAK'de Pembrolizumab/karboplatin/paklitaksel (ya da nab paklitaksel) PD-L1 düzeyine bakılmaksızın NCCN 2019 3.versiyonda kanıt 1 düzeyinde önerilmektedir (78). EGFR mutasyonu olan ve ALK rearranjmanı olan hastalarda PD-L1 ekspresyon düzeyinden bağımsız olarak anti PD1 ve PD-L1 ile monoterapi 2. basamakta daha az etkilidir (79,80,81). Yine EGFR mutant hastalarda PD-L1 ekspresyonu %50 den daha fazla olanlarda dahi 1. basamakta pembrolizumab etkilidir (82).

Her ne kadar PD-L1 ekspresyonu optimal bir biyobelirteç olmasa da günümüzde anti PD-1 ve anti PD-L1 tedavi için uygun hasta seçiminde mevcut biyobelirteçler arasında en iyisidir. PD-L1 ekspresyonu sürekli değişken ve dinamikdir, bu nedenle pozitif bir sonuç için eşik değer belirtmek doğru olmayabilir. Yüzde 50'nin hemen altı ve üstü PD-L1 ekspresyonu olanlarda benzer sonuçlar görülebilmektedir (83). Pozitif test sonucu tanımı da hangi biyobelirteç test yöntemi kullanıldığına göre değişmektedir. Bu bağlamda tümör mutasyon yükünü (TMB) belirleyen çeşitli ölçümler araştırılmaktadır (84). TMB'nin nasıl ölçülebileceğine dair konsensus geliştirmek için uluslararası bir çaba sarf edilmektedir (85, 86, 87). Ayrıca tümör inflamasyonunun değerlendirilmesi için de çeşitli yöntemler geliştirilmektedir fakat bunların rutinimize girebilmesi için daha fazla çalışmaya ihtiyaç duyulmaktadır (29).

## SONUÇ

Son yıllarda geliştirilen moleküler testler sayesinde KHDAK'nin hem patogenezi, hem prognozu hem de tedavi hedeflerinin belirlenmesinde ilerlemeler kaydedilmiştir. Aynı zamanda bu testler tedavi direnç mekanizmalarının anlaşılmasında ve çözümünde de rol almaktadır. Şüphe yok ki, geliştirilen moleküler test yöntemleri ile gelecekte KHDAK'li birçok hastanın tümör genotipleme ile belirlenmiş spesifik tedavisi olacağı düşünülmektedir.

## KAYNAKLAR

- Sholl LM, Aisner DL, Varella-Garcia M, et al; LCMC Investigators. (2015). Multi-institutional oncogenic driver mutation analysis in lung adenocarcinoma: The Lung Cancer Mutation Consortium Experience. *J Thorac Oncol*.10(5):768-777.
- Gerber D, Gandhi L, Costa DB. (2014). Management and future directions in non-small cell lung cancer with known activating mutations. *Am Soc Clin Oncol Educ Book*. e353-365.

- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.2019 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:3236](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:3236) adresinden ulaşılmıştır.)
- Shigematsu H, Lin L, Takahashi T, et al. (2005). Clinical and biological features associated with epidermal growth factor receptor gene mutations in lung cancers. *J Natl Cancer Inst*;97(5):339-346.
- Lynch TJ, Bell DW, Sordella R, et al. (2004). Activating mutations in the epidermal growth factor receptor underlying responsiveness of non-small-cell lung cancer to gefitinib. *N Engl J Med*.;350:2129-2139
- Paez JG, Jänne PA, Lee JC, et al. (2004). EGFR mutations in lung cancer: correlation with clinical response to gefitinib therapy. *Science*.304(5676):1497-1500.
- Siegelin MD, Borczuk AC. (2014). Epidermal growth factor receptor mutations in lung adenocarcinoma. *Lab Invest*.94(2):129-137.
- Langer CJ (2013). Epidermal growth factor receptor inhibition in mutation-positive non-small-cell lung cancer: is afatinib better or simply newer? *J Clin Oncol*. 31(27):3303-6. doi: 10.1200/JCO.2013.49.8782..
- O’Kane GM, Bradbury PA, Feld R, et al. (2017). Uncommon EGFR mutations in advanced non-small cell lung cancer. *Lung Cancer*. ;109:137-144. doi: 10.1016/j.lungcan.2017.04.016. Epub 2017 Apr 27.
- Oxnard GR, Lo PC, Nishino M, et al. (2013). Natural history and molecular characteristics of lung cancers harboring EGFR exon 20 insertions. *J Thorac Oncol*.;8(2):179-84. doi: 10.1097/JTO.0b013e3182779d18.
- Finlay MR, Anderton M, Ashton S, et al. (2014). Discovery of a potent and selective EGFR inhibitor (AZD9291) of both sensitizing and T790M resistance mutations that spares the wild type form of the receptor. *J Med Chem*.;57(20):8249-67. doi: 10.1021/jm500973a.
- Rosell R, Molina MA, Costa C, et al. (2011). Pretreatment EGFR T790M mutation and BRCA1 mRNA expression in erlotinib-treated advanced non-small-cell lung cancer patients with EGFR mutations. *Clin Cancer Res*.;17(5):1160-8. doi: 10.1158/1078-0432.CCR-10-2158.
- Suda K, Mizuuchi H, Maehara Y, et al. (2012). Acquired resistance mechanisms to tyrosine kinase inhibitors in lung cancer with activating epidermal growth factor receptor mutation--diversity, ductility, and destiny. *Cancer Metastasis Rev*. Dec;31(3-4):807-14. doi: 10.1007/s10555-012-9391-7.
- Yu HA, Suzawa K, Jordan E, et al. (2018). Concurrent Alterations in EGFR-Mutant Lung Cancers Associated with Resistance to EGFR Kinase Inhibitors and Characterization of MTOR as a Mediator of Resistance. *Clin Cancer Res*. Jul 1;24(13):3108-3118. doi: 10.1158/1078-0432.CCR-17-2961.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:1097](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:1097) adresinden ulaşılmıştır.)
- Planchard D, Besse B, Groen HJM, et al. (2016). Dabrafenib plus trametinib in patients with previously treated BRAF(V600E)-mutant metastatic non-small cell lung cancer: an open-label, multicentre phase 2 trial. *Lancet Oncol*. Jul;17(7):984-993. doi: 10.1016/S1470-2045(16)30146-2.
- Paik PK, Arcila ME, Fara M, et al. (2011) Clinical characteristics of patients with lung adenocarcinomas harboring BRAF mutations. *J Clin Oncol*. May 20;29(15):2046-51. doi: 10.1200/JCO.2010.33.1280.

- Gautschi O, Milia J, Cabarrou B, et al. (2015). Targeted Therapy for Patients with BRAF-Mutant Lung Cancer: Results from the European EURAF Cohort. *J Thorac Oncol. Oct;10(10):1451-7*. doi: 10.1097/JTO.0000000000000625.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:427](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:427) adresinden ulaşılmıştır.)
- Takeuchi K, Soda M, Togashi Y, et al. (2012). RET, ROS1 and ALK fusions in lung cancer. *Nat Med; 18:378-381*.
- Shaw AT, Yeap BY, Mino-Kenudson M, et al. (2009). Clinical features and outcome of patients with non-small-cell lung cancer who harbor EML4-ALK. *J Clin Oncol.Sep 10;27(26):4247-53*. doi: 10.1200/JCO.2009.22.6993.
- Lindeman NI, Cagle PT, Beasley MB, et al. (2013). Molecular testing guideline for selection of lung cancer patients for EGFR and ALK tyrosine kinase inhibitors: guideline from the College of American Pathologists, International Association for the Study of Lung Cancer, and Association for Molecular Pathology. *J Thorac Oncol. Jul;8(7):823-59*. doi: 10.1097/JTO.0b013e318290868f.
- Ali G, Proietti A, Pelliccioni S, et al. (2014). ALK rearrangement in a large series of consecutive non-small cell lung cancers: comparison between a new immunohistochemical approach and fluorescence in situ hybridization for the screening of patients eligible for crizotinib treatment. *Arch Pathol Lab Med.138(11):1449-58*. doi: 10.5858/arpa.2013-0388-OA.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:10261](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:10261) adresinden ulaşılmıştır.)
- Shaw AT, Ou SH, Bang YJ, et al. (2014). Crizotinib in ROS1-rearranged non-small-cell lung cancer. *N Engl J Med. Nov 20;371(21):1963-71*. doi: 10.1056/NEJMoa1406766.
- Robinson DR, Wu YM, Lin SF. (2000). The protein tyrosine kinase family of the human genome. *Oncogene. Nov 20;19(49):5548-57*.
- Kim HR, Lim SM, Kim HJ, et al. (2013). The frequency and impact of ROS1 rearrangement on clinical outcomes in never smokers with lung adenocarcinoma. *Ann Oncol. Sep;24(9):2364-70*. doi: 10.1093/annonc/mdt220.
- Wu YL, Yang JC, Kim DW, et al. (2018). Phase II Study of Crizotinib in East Asian Patients With ROS1-Positive Advanced Non-Small-Cell Lung Cancer. *J Clin Oncol. May 10;36(14):1405-1411*. doi: 10.1200/JCO.2017.75.5587.
- Planchard D, Popat S, Kerr K, et al. on behalf of the ESMO Guidelines Committee, (2018). Metastatic non-small cell lung cancer: ESMO Clinical Practice Guidelines for diagnosis, treatment and follow-up. *Annals of Oncology 29* (Supplement 4): iv192–iv237, doi:10.1093/annonc/mdy275
- Lindeman N, Cagle P, Aisner D et al. (2018). Updated molecular testing guideline for the selection of lung cancer patients for treatment with targeted tyrosine kinase inhibitors: guideline from the College of American Pathologists, the International Association for the Study of Lung Cancer, and the Association for Molecular Pathology. *Arch Pathol LabMed 142: 321–346*.
- Kalemkerian GP, Narula N, Kennedy EB et al. (2018). Molecular testing guideline for the selection of patients with lung cancer for treatment with targeted tyrosine kinase inhibitors: American Society of Clinical Oncology Endorsement of the College of American Pathologists/International Association for the Study of Lung Cancer/As-

- sociation for Molecular Pathology Clinical Practice Guideline Update. *J Clin Oncol*; 36: 911–919.
- Tsao M, Hirsch F, Yatabe Y. (2016). *IASLC Atlas of ALK and ROS1 Testing in Lung Cancer*, Second Edition. Aurora, CO, USA: Editorial Rx Press.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:6407](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:6407) adresinden ulaşılmıştır.)
- Eberhard DA, Johnson BE, Amler LC, et al. (2005). Mutations in the epidermal growth factor receptor and in KRAS are predictive and prognostic indicators in patients with non-small-cell lung cancer treated with chemotherapy alone and in combination with erlotinib. *J Clin Oncol*;23(25):5900-9.
- Slebos RJ, Hruban RH, Dalesio O, et al. (1991). Relationship between K-ras oncogene activation and smoking in adenocarcinoma of the human lung. *J Natl Cancer Inst*;83(14):1024-7.
- Tsao MS, Aviel-Ronen S, Ding K, et al. (2007) Prognostic and predictive importance of p53 and RAS for adjuvant chemotherapy in non small-cell lung cancer. *J Clin Oncol*. 25(33):5240-7.
- Mitsudomi T, Steinberg SM, Oie HK, et al. (1991). Ras gene mutations in non-small cell lung cancers are associated with shortened survival irrespective of treatment intent. *Cancer Res*. 51(18):4999-5002.
- Miller VA, Riely GJ, Zakowski MF, et al. (2008). Molecular characteristics of bronchioloalveolar carcinoma and adenocarcinoma, bronchioloalveolar carcinoma subtype, predict response to erlotinib. *J Clin Oncol*.26(9):1472-8. doi: 10.1200/JCO.2007.13.0062.
- Roberts PJ, Stinchcombe TE. (2013) KRAS mutation: should we test for it, and does it matter? *J Clin Oncol*.31(8):1112-21. doi: 10.1200/JCO.2012.43.0454.
- Sholl LM, Aisner DL, Varella-Garcia M et al. (2015). Multi-institutional Oncogenic Driver Mutation Analysis in Lung Adenocarcinoma: The Lung Cancer Mutation Consortium Experience. *J Thorac Oncol*. 10(5):768-777. doi: 10.1097/JTO.0000000000000516.
- Calles A, Liao X, Sholl LM, et al. (2015).Expression of PD-1 and Its Ligands, PD-L1 and PD-L2, in Smokers and Never Smokers with KRAS-Mutant Lung Cancer. *J Thorac Oncol. Dec*;10(12):1726-35. doi: 10.1097/JTO.0000000000000687.
- Jänne PA, Shaw AT, Pereira JR, et al. (2013). Selumetinib plus docetaxel for KRAS-mutant advanced non-small-cell lung cancer: a randomised, multicentre, placebo-controlled, phase 2 study. *Lancet Oncol. Jan*;14(1):38-47. doi: 10.1016/S1470-2045(12)70489-8.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:8031](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:8031) adresinden ulaşılmıştır.)
- Farago AF, Taylor MS, Doebele RC, et al. (2018). Clinicopathologic Features of Non-Small-Cell Lung Cancer Harboring an NTRK Gene Fusion. *JCO Precis Oncol*. 2018; doi: 10.1200/PO.18.00037. Epub 2018 Jul 23.
- Drilon A, Laetsch TW, Kummar S, et al. (2018).Efficacy of Larotrectinib in TRK Fusion-Positive Cancers in Adults and Children. *N Engl J Med. Feb 22*;378(8):731-739. doi: 10.1056/NEJMoal714448.
- Drilon A, Siena S, Ou SI et al. (2017). Safety and antitumor activity of the multitargeted Pan-TRK, ROS1, and ALK inhibitor entrectinib: combined results from two phase I trials (ALKA-372-001 and STARTRK-1). *Cancer Discov*; 7: 400–409.
- Hyman DM, Laetsch TW, Kummar S et al. (2017). The efficacy of larotrectinib (LOXO-101), a selective tropomyosin receptor kinase (TRK) inhibitor, in adult and pediatric TRK fusion cancers. *J Clin Oncol* 35(18 Suppl): LBA2501.

- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:3430](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:3430) adresinden ulaşılmıştır.)
- Planchard D, Loriot Y, André F, et al. (2015). EGFR-independent mechanisms of acquired resistance to AZD9291 in EGFR T790M-positive NSCLC patients. *Annals of oncology : official journal of the European Society for Medical Oncology ;26:2073-2078*.
- Tanizaki J, Okamoto I, Okabe T, et al. (2012). Activation of HER family signaling as a mechanism of acquired resistance to ALK inhibitors in EML4-ALK-positive non-small cell lung cancer. *Clin Cancer Res;18:6219-6226*.
- Hirsch FR, Varella-Garcia M, Franklin WA, et al. (2002). Evaluation of HER-2/neu gene amplification and protein expression in non-small cell lung carcinomas. *Br J Cancer;86:1449-1456*.
- Cancer Genome Atlas Research Network. (2014). Comprehensive molecular profiling of lung adenocarcinoma. *Nature;511:543-550*.
- Kim EK, Kim KA, Lee CY, et al. (2017). The frequency and clinical impact of HER2 alterations in lung adenocarcinoma. *PLoS One;12:e0171280*.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:9967](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:9967) adresinden ulaşılmıştır.)
- Go H, Jung YJ, Kang HW, et al. (2013). Diagnostic method for the detection of KIF5B-RET transformation in lung adenocarcinoma. *Lung Cancer;82:44-50*.
- Gautschi O, Milia J, Filleron T, et al. (2017). Targeting RET in Patients With RET-Rearranged Lung Cancers: Results From the Global, Multicenter RET Registry. *J Clin Oncol;35:1403-1410*.
- Kohno T, Ichikawa H, Totoki Y, et al. (2012). KIF5B-RET fusions in lung adenocarcinoma. *Nat Med;18:375-377*.)
- Drilon A, Rekhtman N, Arcila M, et al. (2016). Cabozantinib in patients with advanced RET-rearranged non-small-cell lung cancer: an open-label, single-centre, phase 2, single-arm trial. *Lancet Oncol.17(12):1653-1660*. doi: 10.1016/S1470-2045(16)30562-9.
- Lee SH, Lee JK, Ahn MJ, et al. (2017). Vandetanib in pretreated patients with advanced non-small cell lung cancer-harboring RET rearrangement: a phase II clinical trial. *Ann Oncol. ;28(2):292-297*. doi: 10.1093/annonc/mdw559.
- HGNC (2019). *HUGO gene nomenclature committee 2019*. (25.01.19 tarihinde [https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:7029](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:7029) adresinden ulaşılmıştır.
- Paik PK, Drilon A, Fan PD, et al. (2015). Response to MET inhibitors in patients with stage IV lung adenocarcinomas harboring MET mutations causing exon 14 skipping. *Cancer Discov 5:842-849*.
- Frampton GM, Ali SM, Rosenzweig M, et al. (2015). Activation of MET via diverse exon 14 splicing alterations occurs in multiple tumor types and confers clinical sensitivity to MET inhibitors. *Cancer Discov ;5:850-859*.
- Rabeau A, Rouquette I, Vantelon JM, et al. (2017). Interest of crizotinib in a lung cancer patient with de novo amplification of MET. *Rev Mal Respir;34:57-60*.
- Engelman JA, Zejnullahu K, Mitsudomi T, et al.(2007). MET amplification leads to gefitinib resistance in lung cancer by activating ERBB3 signaling. *Science;316:1039-1043*.
- Ribas A. (2015) Releasing the Brakes on Cancer Immunotherapy. *N Engl J Med. ;373(16):1490-2*. doi: 10.1056/NEJMp1510079.
- Brahmer JR, Hammers H, Lipson EJ. (2015). Nivolumab: targeting PD-1 to bolster anti-tumor immunity. *Future Oncol.;11(9):1307-26*. doi: 10.2217/fon.15.52.



- Brahmer J, Reckamp KL, Baas P, et al. (2015). Nivolumab versus Docetaxel in Advanced Squamous-Cell Non-Small-Cell Lung Cancer. *N Engl J Med.*;373(2):123-35. doi: 10.1056/NEJMoa1504627.
- Antonia SJ, Villegas A, Daniel D, et al. (2017). Durvalumab after Chemoradiotherapy in Stage III Non-Small-Cell Lung Cancer. *N Engl J Med.*;377(20):1919-1929. doi: 10.1056/NEJMoa1709937.
- Fehrenbacher L, Spira A, Ballinger M, et al. (2016). Atezolizumab versus docetaxel for patients with previously treated non-small-cell lung cancer (POPLAR): a multicentre, open-label, phase 2 randomised controlled trial. *Lancet.*;387(10030):1837-46. doi: 10.1016/S0140-6736(16)00587-0.
- Kerr KM, Hirsch FR. (2016). Programmed death ligand-1 immunohistochemistry: friend or foe? *Arch Pathol Lab Med* ; 140: 326–331.
- Hirsch FR, McElhinny A, Stanforth D et al. (2017). PD-L1 immunohistochemistry assays for lung cancer: results from phase 1 of the blueprint PD-L1 IHC assay comparison project. *J Thorac Oncol* 12: 208–222.
- Ratcliffe MJ, Sharpe A, Midha A et al. (2017). Agreement between programmed cell death ligand-1 diagnostic assays across multiple protein expression cutoffs in non-small cell lung cancer. *Clin Cancer Res*; 23: 3585–3591.
- Rimm DL, Han G, Taube JM et al. (2017). A prospective, multi-institutional, pathologist-based assessment of 4 immunohistochemistry assays for PD-L1 expression in non-small cell lung cancer. *JAMA Oncol*; 3: 1051–1058.
- Adam J, Le Stang N, Rouquette I et al. (2018). Multicenter French harmonization study for PD-L1 IHC testing in non-small cell lung cancer. *Ann. Oncol*; 29: 953–958.
- Reck M, Rodriguez-Abreu D, Robinson AG et al. (2016). Pembrolizumab versus chemotherapy for PD-L1-positive non-small-cell lung cancer. *N Engl J Med*; 375: 1823–1833.
- Gandhi L, Rodríguez-Abreu D, Gadgeel S, et al. (2018). Pembrolizumab plus Chemotherapy in Metastatic Non-Small-Cell Lung Cancer. *N Engl J Med.*;378(22):2078-2092. doi: 10.1056/NEJMoa1801005.
- Socinski MA, Jotte RM, Cappuzzo F, et al. (2018). Atezolizumab for First-Line Treatment of Metastatic Nonsquamous NSCLC. *N Engl J Med.*;378(24):2288-2301. doi: 10.1056/NEJMoa1716948.
- Paz-Ares L, Luft A, Vicente D, et al. (2018). Pembrolizumab plus Chemotherapy for Squamous Non-Small-Cell Lung Cancer. *N Engl J Med.*;379(21):2040-2051. doi: 10.1056/NEJMoa1810865.
- Rittmeyer A, Barlesi F, Waterkamp D, et al. (2017). Atezolizumab versus docetaxel in patients with previously treated non-small-cell lung cancer (OAK): a phase 3, open-label, multicentre randomised controlled trial. *Lancet.* 2017 Jan 21;389(10066):255-265. doi: 10.1016/S0140-6736(16)32517-X.
- Herbst RS, Baas P, Kim DW, et al. (2016). Pembrolizumab versus docetaxel for previously treated, PD-L1-positive, advanced non-small-cell lung cancer (KEYNOTE-010): a randomised controlled trial. *Lancet.*;387(10027):1540-50. doi: 10.1016/S0140-6736(15)01281-7.
- Gainor JF, Shaw AT, Sequist LV, et al. (2016). EGFR Mutations and ALK Rearrangements Are Associated with Low Response Rates to PD-1 Pathway Blockade in Non-Small Cell Lung Cancer: A Retrospective Analysis. *Clin Cancer Res.*;22(18):4585-93. doi: 10.1158/1078-0432.CCR-15-3101.

- Lisberg A, Cummings A, Goldman JW, et al. (2018). A Phase II Study of Pembrolizumab in EGFR-Mutant, PD-L1+, Tyrosine Kinase Inhibitor Naïve Patients With Advanced NSCLC. *J Thorac Oncol*. Aug;13(8):1138-1145. doi: 10.1016/j.jtho.2018.03.035. Epub 2018 Jun 1.
- Kerr KM, Nicolson MC. (2015) Non-Small Cell Lung Cancer, PD-L1, and the Pathologist. *Arch Pathol Lab Med*.;140(3):249-54. doi: 10.5858/arpa.2015-0303-SA.
- Hellmann MD, Ciuleanu T-E, Pluzanski A et al. (2018). Nivolumab plus ipilimumab in lung cancer with a high tumor mutational burden. *N Engl J Med*; 378: 2093–2104.].
- Rizvi NA, Hellmann MD, Snyder A et al. (2015). Cancer immunology. Mutational landscape determines sensitivity to PD-1 blockade in nonsmall cell lung cancer. *Science*; 348: 124–128.
- Carbone DP, Reck M, Paz-Ares L et al. (2017) First-line nivolumab in stage IV or recurrent non-small-cell lung cancer. *N Engl J Med* ; 376:2415–2426.
- Rizvi H, Sanchez-Vega F, La K et al. (2018). Molecular determinants of response to anti programmed cell death (PD)-1 and anti-programmed death-ligand (PD-L)-ligand 1 blockade in patients with non-small-cell lung cancer profiled with targeted next-generation sequencing. *J Clin Oncol*; 36: 633–641.