

Distinct NSCLC *EGFR* Variants in a Family With Li-Fraumeni Syndrome: Case Report



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ABSTRACT

Introduction: Heritable lung cancer may occur in the context of germline *TP53* mutations (Li-Fraumeni syndrome). Limited cases of intrafamily tumor genomic characteristics have been reported.

Main concerns, Important Clinical Findings, Primary Diagnoses, Interventions, Outcomes: A 40-year-old woman with no smoking history or known environmental exposure risk was incidentally found to have stage II (T2N1) NSCLC harboring an *EGFR* exon 19 p.Glu746_Ala750 deletion. Family history was notable for an identical twin sister with colorectal cancer (diagnosed at age 31 y) and a mother with stage I NSCLC harboring an *EGFR* exon 21 c.2573T>G (p.Leu858Arg) mutation (diagnosed at age 69 y). Genetic testing revealed a germline *TP53* c.542G>A (p.Arg181His) mutation in the patient, her mother, and her sister, consistent with Li-Fraumeni syndrome. No germline *EGFR* mutations were detected.

Conclusion: Shared germline *TP53* mutations may be associated with distinct NSCLC somatic *EGFR* mutations within families with Li-Fraumeni syndrome. Further understanding of the association between genetic cancer syndromes and lung cancer risk may improve early lung cancer detection in populations not otherwise meeting screening eligibility.

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Introduction

Lung cancer almost always occurs sporadically. More than 80% of cases are attributable to environmental exposure, most often tobacco but also radon, asbestos, and other carcinogens. Although the identification of somatic driver alterations in oncogenes such as *EGFR* or *ALK* provides a mechanistic description for lung cancer in individuals without obvious environmental causes, why some individuals develop these genomic events remains unknown. In rare cases, heritable lung cancer may occur in the context of a pathogenic germline variant.^{1,2} Here, we report lung cancer cases harboring distinct somatic *EGFR* variants occurring within family members found to have a germline *TP53* mutation consistent with Li-Fraumeni syndrome.

Case Presentation

A 40-year-old woman with no smoking history, no concerning environmental exposures, and no

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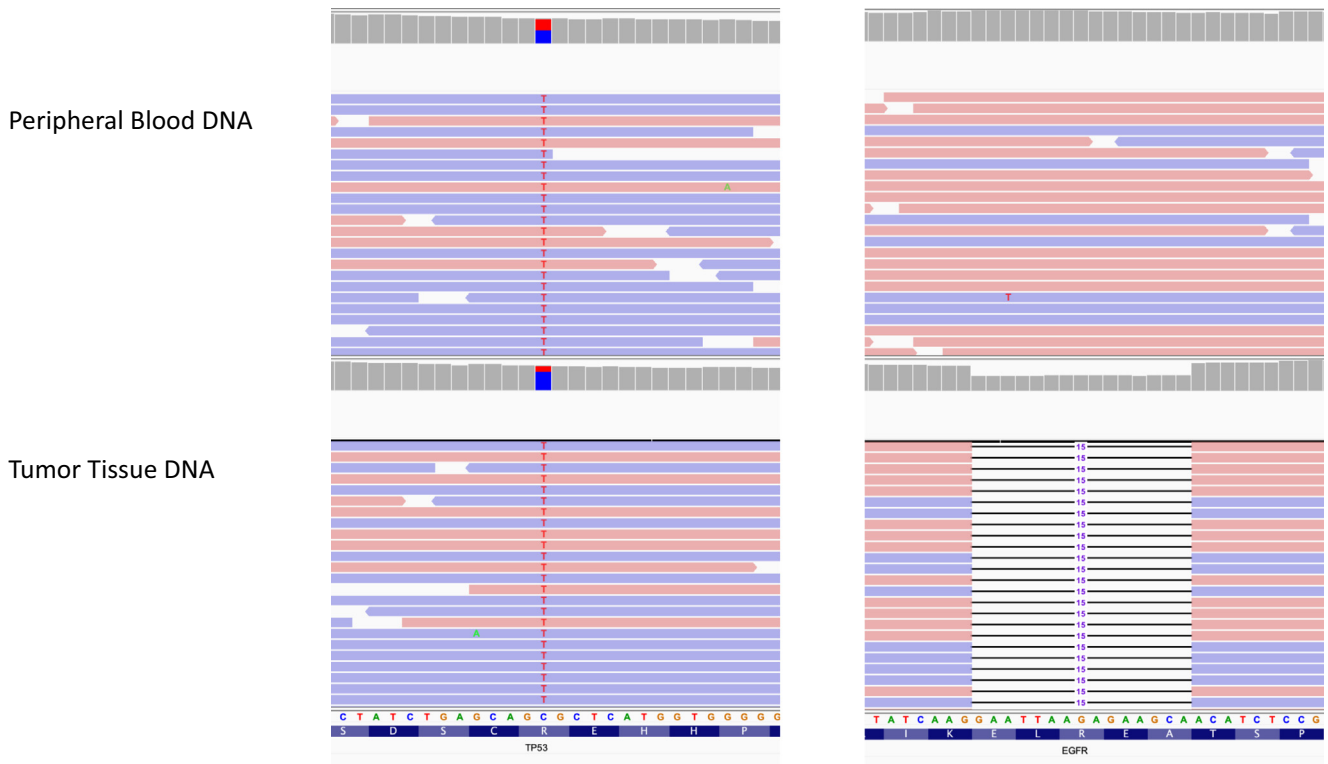


Figure 1. *TP53* and *EGFR* variants detected in the proband. The *EGFR* variant was only detected in the tumor tissue but not in the peripheral blood. The *TP53* variant was detected in both tissue and peripheral blood, revealing germline inheritance. Images from Integrated Genomics Viewer software.

cardiopulmonary symptoms underwent total body computed tomography (CT) as part of an initial annual executive medical assessment. These imaging studies revealed a right upper lobe mass. Result of a CT-guided biopsy revealed adenocarcinoma consistent with lung primary. No distant disease was found on brain magnetic resonance imaging (MRI) and positron emission tomography-CT. The patient underwent right upper lobectomy and regional lymph node dissection, with pathologic evaluation identifying a stage II (T2N1M0) lung adenocarcinoma harboring an *EGFR* exon 19 p.Glu746_Ala750 deletion (Fig. 1).

Review of the patient's family history noted multiple relatives with cancer (Fig. 2). Specifically, her mother, who had a distant and minimal smoking history, had been diagnosed with having a stage 1B RUL NSCLC harboring an *EGFR* exon 21 c.2573T>G (p.Leu858Arg) mutation at age 69 years (Fig. 1). At age 31 years, an identical twin sister had been diagnosed with having invasive moderately differentiated adenocarcinoma of the sigmoid colon with well-differentiated neuroendocrine tumor features, *KRAS* wild type, *NRAS* wild type, and *BRAF* wild type, with immunohistochemistry for mismatch repair proteins intact. A maternal aunt had breast cancer diagnosed at age 65 years, and her son (first cousin to the proband) was diagnosed with having

prostate cancer at age 55 years. On the basis of the occurrence of multiple cancers including young-onset cases, the patient and family members were referred for genetic counseling. Germline genetic testing by a 91-gene pan-cancer panel (see [Supplementary Materials](#)) in the proband revealed a heterozygous likely pathogenic germline *TP53* variant c.542G>A (p.Arg181His), which was subsequently confirmed in her mother and her twin sister, consistent with Li-Fraumeni syndrome. No germline *EGFR* mutations were detected.

Because of concern for heightened leukemogenic potential of cytotoxic agents in the setting of a germline *TP53* mutation,³ adjuvant chemotherapy was not recommended. Approximately 6 weeks after surgical resection, the patient started adjuvant osimertinib.

Furthermore, cancer screening was implemented for these family members on the basis of published *TP53* variant guidelines. Specific testing—designed to limit radiation exposure given concerns for heightened carcinogenic potential—includes complete physical examination every 6 months, clinical breast examination every 6 months with breast MRI every 12 months, annual brain MRI, annual whole-body MRI, annual dermatologic examination, colonoscopy and upper endoscopy every 2 to 5 years, and ultrasonography of the abdomen and pelvis every 12 months.⁴

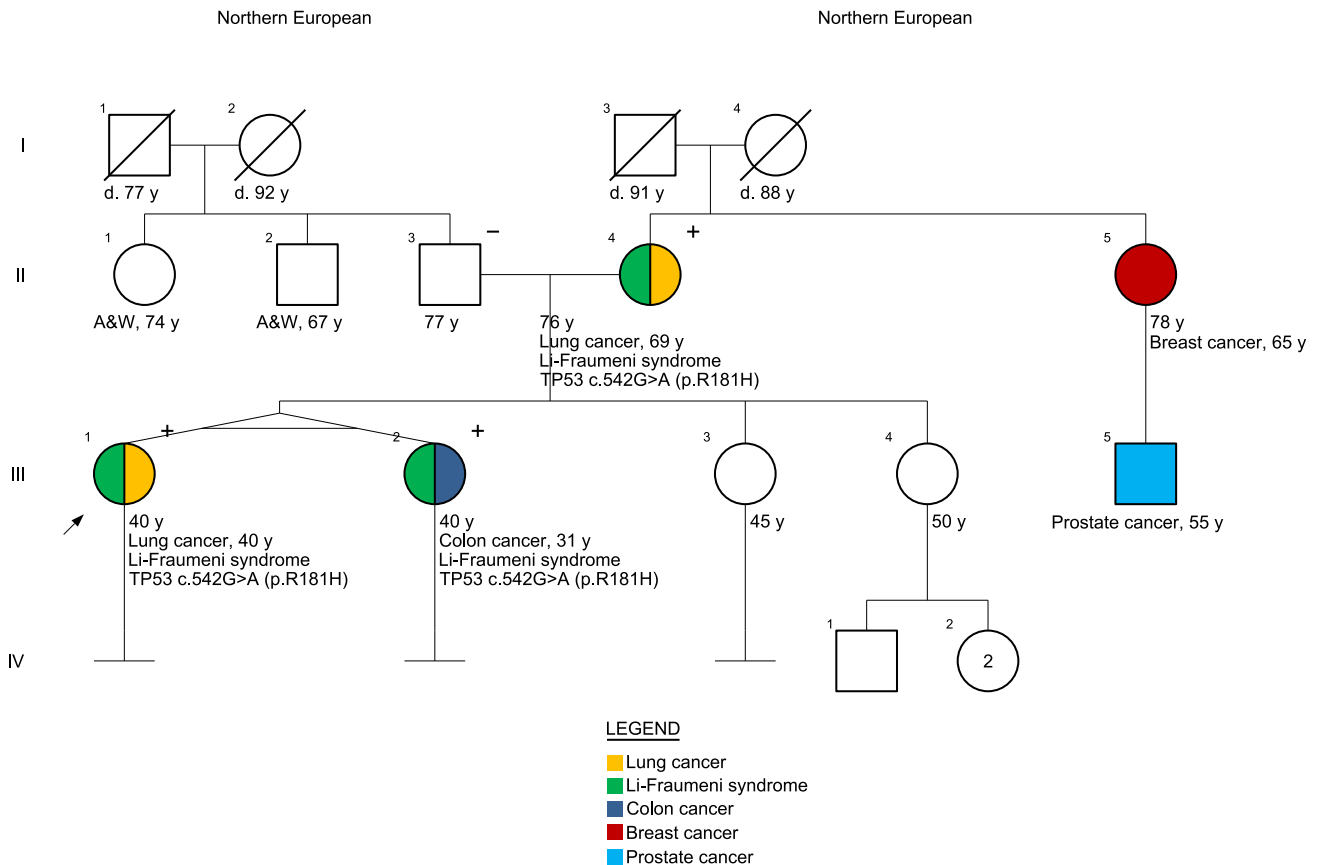


Figure 2. Family pedigree. A&W, alive and well; d, died.

Discussion

Li-Fraumeni syndrome is an autosomal-dominant condition characterized by germline mutations of the *TP53* tumor suppressor gene resulting in heightened cancer risk. Characteristic malignancies include sarcoma, leukemia, breast cancer, primary brain tumors, and adrenal cortical tumors. Lung cancer, in particular *EGFR*-mutant NSCLC, and colorectal cancers have also been reported.^{1,2} The present case supports a single prior report describing multiple lung cancers harboring different *EGFR* mutations within a single Li-Fraumeni family.²

The occurrence of two distinct somatic *EGFR* mutations within this genetic context further establishes the association of *TP53* germline mutations with this genomic alteration.

Most *EGFR*-mutant NSCLC cases reported within Li-Fraumeni cohorts have occurred after earlier cancers, most often breast cancer.^{1,2} Such patterns have historically led to a two-hit hypothesis, with a germline mutation providing the first hit and additional environmental exposure (such as ultraviolet radiation exposure, radiation therapy, or chemotherapy administered for a prior cancer) providing the second hit. Nevertheless, in the

present case, the NSCLC was the first (and to date only) cancer diagnosis for both the patient and her mother. Among Li-Fraumeni-associated breast cancer cases, HER2-positive cases seem over-represented, providing evidence that HER family protein kinases across tumor types may be particularly susceptible to *TP53*-related alterations.⁵

Conclusion

History of lung cancer in first-degree relatives seems to confer increased lung cancer risk, independent of age, sex, and smoking.⁶ Attention to familial lung cancer patterns may help identify individuals at heightened risk for lung cancer who do not otherwise meet age- or smoking-related eligibility for screening. It may also contribute to further understanding of the development of *EGFR* mutations and other genomic alterations in this disease.

CRediT Authorship Contribution Statement

David E. Gerber, Mitchell S. von Itzstein, Shelby Edmondson: Conceptualization.

Shelby Edmondson, Mitchell S. von Itzstein, Brian Reys, Melissa Mayer, Jeffrey Gagan, David E. Gerber: Investigation.

David E. Gerber, Mitchell S. von Itzstein, Melissa Mayer: Project administration.

Shelby Edmondson, David E. Gerber: Roles/Writing—original draft.

Shelby Edmondson, Mitchell S. von Itzstein, Brian Reys, Melissa Mayer, Jeffrey Gagan, David E. Gerber: Writing—review and editing.

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Supplementary Data

Note: To access the supplementary material accompanying this article, visit the online version of the *JTO Clinical and Research Reports* at www.jtocrr.org and at <https://doi.org/10.1016/j.jtocrr.2022.100368>.

References

1. Michalarea V, Calcasola M, Cane P, Tobal K, Izatt L, Spicer J. EGFR-mutated lung cancer in Li-Fraumeni syndrome. *Lung Cancer*. 2014;85:485-487.
2. Mezquita L, Jove M, Nadal E, et al. High prevalence of somatic oncogenic driver alterations in patients with NSCLC and Li-Fraumeni syndrome. *J Thorac Oncol*. 2020;15:1232-1239.
3. Valdez JM, Nichols KE, Kesserwan C. Li-Fraumeni syndrome: a paradigm for the understanding of hereditary cancer predisposition. *Br J Haematol*. 2017;176:539-552.
4. Kratz CP, Achatz MI, Brugieres L, et al. Cancer screening recommendations for individuals with Li-Fraumeni syndrome. *Clin Cancer Res*. 2017;23:e38-e45.
5. Gallardo-Alvarado LN, Tusie-Luna MT, Tussie-Luna MI, et al. Prevalence of germline mutations in the TP53 gene in patients with early-onset breast cancer in the Mexican population. *BMC Cancer*. 2019;19:118.
6. Jonsson S, Thorsteinsdottir U, Gudbjartsson DF, et al. Familial risk of lung carcinoma in the Icelandic population. *JAMA*. 2004;292:2977-2983.