High Epidermal Growth Factor Receptor Mutation Rates in Peruvian Patients With Non-Small-Cell Lung Cancer: Is It a Matter of Asian Ancestry?

In a recent article, Lopez-Chavez et al 1 reported a high mutational rate of epidermal growth factor receptor (EGFR) in Peruvian patients (37%) that is higher than in other Latin American countries such as Mexico, Bolivia, Venezuela, and in a mixture of Latinos in the United States. High mutational rates of the EGFR gene in Peruvian patients were reported previously in independent cohorts. Mas et al 2 reported a frequency of 39.3% (n = 122), and Arrieta et al 3 reported a frequency of EGFR mutations in Peruvian patients is higher than other reports, these rates could be explained by environmental factors.

However, ancestry could also play an important role in explaining this fact. We would like to point out two events that could lead to a gene flow explaining the high prevalence of *EGFR* mutations in Peruvian patients. Population of the Americas in the late Pleistocene epoch by migrants from Asia through the Bering land bridge shaped the genetic pool of Native Americans. The second event occurred after slavery was abolished in Peru and a massive wave of Chinese workers reached the Peruvian coast (approximately 100,000 between 1849 and 1880), with a Peruvian population estimated at 2 million in 1850.^{4,5}

Joseph A. Pinto Luis A. Mas Henry L. Gomez

Joseph A. Pinto, Luis A. Mas, and Henry L. Gomez Oncosalud-AUNA, San Borja: Luis A. Mas and Henry L. Gomez, Instituto Nacional de Enfermedades Neoplasicas, Lima, Peru Corresponding author: Henry L. Gomez, Department of Medical Oncology, Oncosalud-AUNA, Av. Guardia Civil 571, San Borja, Peru; e-mail: hgomezmoreno@ gmail.com.

AUTHOR CONTRIBUTIONS

Manuscript writing: All authors
Final approval of manuscript: All authors

AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

The following represents disclosure information provided by authors of this manuscript. All relationships are considered compensated. Relationships are self-held unless noted. I = Immediate Family Member, Inst = My Institution. Relationships may not relate to the subject matter of this manuscript.

Although there are not many projects that are evaluating Asian ancestry markers in Latin American countries, data for ancestry admixture proportions for Mexico, Colombia, and Peru (0.012, 0.021, and 0.035, respectively) suggest a correlation between ancestry proportion and rate of *EGFR* mutations (Fig 1). ^{1,6-8}

On the other hand, the *Helicobacter pylori* bacterium accompanied humans in the migration waves. These bacteria are not only a chronic pathogen in humans, but also coevolve with their hosts and have been used to trace human migration routes. Work by Devi et al of with Peruvian strains of *H. pylori* found considerable homology with Asian strains. Another interesting fact is the high prevalence of human T-cell lymphotropic virus, ranging from 7% to 25%, in several Peruvian cities. This pattern is typical of some Asian countries such as Japan.

High rates of *EGFR* mutations in Peruvian patients with non–small-cell lung cancer could be a signature of Asian ancestry in the Peruvian population.

DOI: https://doi.org/10.1200/JG0.2016.008201 Published online on jgo.org on January 18, 2017.

For more information about ASCO's conflict of interest policy, please refer to www.asco.org/rwc or ascopubs.org/jco/site/ifc.

Joseph A. Pinto

No relationship to disclose

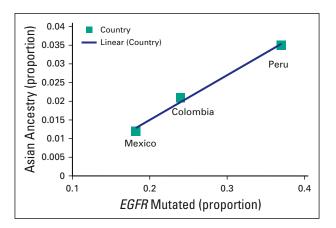
Luis A. Mas

No relationship to disclose

Henry L. Gomez

No relationship to disclose

Fig 1. Graph showing a correlation between the proportion of Asian ancestry in Latin American countries and rates of *EGFR* mutations in non–small-cell lung cancer. *EGFR*, epidermal growth factor receptor.



REFERENCES

- 1. Lopez-Chavez A, Thomas A, Evbuomwan MO, et al: *EGFR* mutations in Latinos from the United States and Latin America. J Glob Oncol 2:259-267, 2016
- 2. Mas L, Gómez de la Torre JC, Barletta C: Estado mutacional de los exones 19 y 21 de EGFR en adenocarcinoma de pulmón: Estudio en 122 pacientes peruanos y revisión de la evidencia de eficacia del inhibidor tirosina kinasa erlotinib. Carcinos 2:52-61, 2011
- 3. Arrieta O, Cardona AF, Martín C, et al: Updated frequency of EGFR and KRAS mutations in nonsmall-cell lung cancer in Latin America: The Latin-American Consortium for the Investigation of Lung Cancer (CLICaP). J Thorac Oncol 10:838-843, 2015
- 4. La Torre Silva R: La inmigración China en el Perú (1850-1890). Boletín de la Sociedad Peruana de Medicina Interna 5: 1992 http://sisbib.unmsm.edu.pe/BvRevistas/spmi/v05n3/Inmigraci%C3%B3n.htm
- 5. Gootenberg P: Población y etnicidad en el Perú republicano (siglo XIX): Algunas revisiones. Lima. 1995 http://repositorio.iep.org.pe/handle/IEP/318
- 6. Silva-Zolezzi I, Hidalgo-Miranda A, Estrada-Gil J, et al: Analysis of genomic diversity in Mexican Mestizo populations to develop genomic medicine in Mexico. Proc Natl Acad Sci USA 106: 8611-8616, 2009
- 7. Sandoval JR, Salazar-Granara A, Acosta O, et al: Tracing the genomic ancestry of Peruvians reveals a major legacy of pre-Columbian ancestors. J Hum Genet 58:627-634, 2013
- 8. Rishishwar L, Conley AB, Wigington CH, et al: Ancestry, admixture and fitness in Colombian genomes. Sci Rep 5:12376, 2015
- 9. Falush D, Wirth T, Linz B, et al: Traces of human migrations in Helicobacter pylori populations. Science 299:1582-1585, 2003
- 10. Devi SM, Ahmed I, Khan AA, et al: Genomes of Helicobacter pylori from native Peruvians suggest admixture of ancestral and modern lineages and reveal a Western type cag-pathogenicity island. BMC Genomics 7:191, 2006