



**DEPARTMENT OF GENETICS
&
HUMAN GENETICS INSTITUTE OF NEW JERSEY**

RESEARCH SEMINAR

**“Challenges to globalizing genetic
predictions of disease risks”**



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Human genomes have been shaped by natural selection, population bottlenecks, and archaic admixture. These evolutionary processes contribute to hereditary differences in disease risk across populations. Here, I explore how disease risks have changed over time and address some of the challenges of applying precision medicine approaches to diverse global populations.

Noon, Monday, December 6, 2021

[Zoom Registration Link](#) | Meeting ID: 986 4197 9928 | Passcode: Genetics

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