

Data Sharing Statement

Mitchell. Germline Variants Incidentally Detected via Tumor-Only Genomic Profiling of Patients With Mesothelioma. *JAMA Netw Open*. Published August 09, 2023.

doi:10.1001/jamanetworkopen.2023.27351

Data

Data available: Yes

Data types: Other (please specify)

Additional Information: Other data will be made available

How to access data: Germline sequence variants - for upload to databases such as Clinvar as needed

When available: With publication

Supporting Documents

Document types: None

Additional Information

Who can access the data: Anyone requesting data from Clinvar team

Types of analyses: Public database - Clinvar

Mechanisms of data availability: With Investigator support