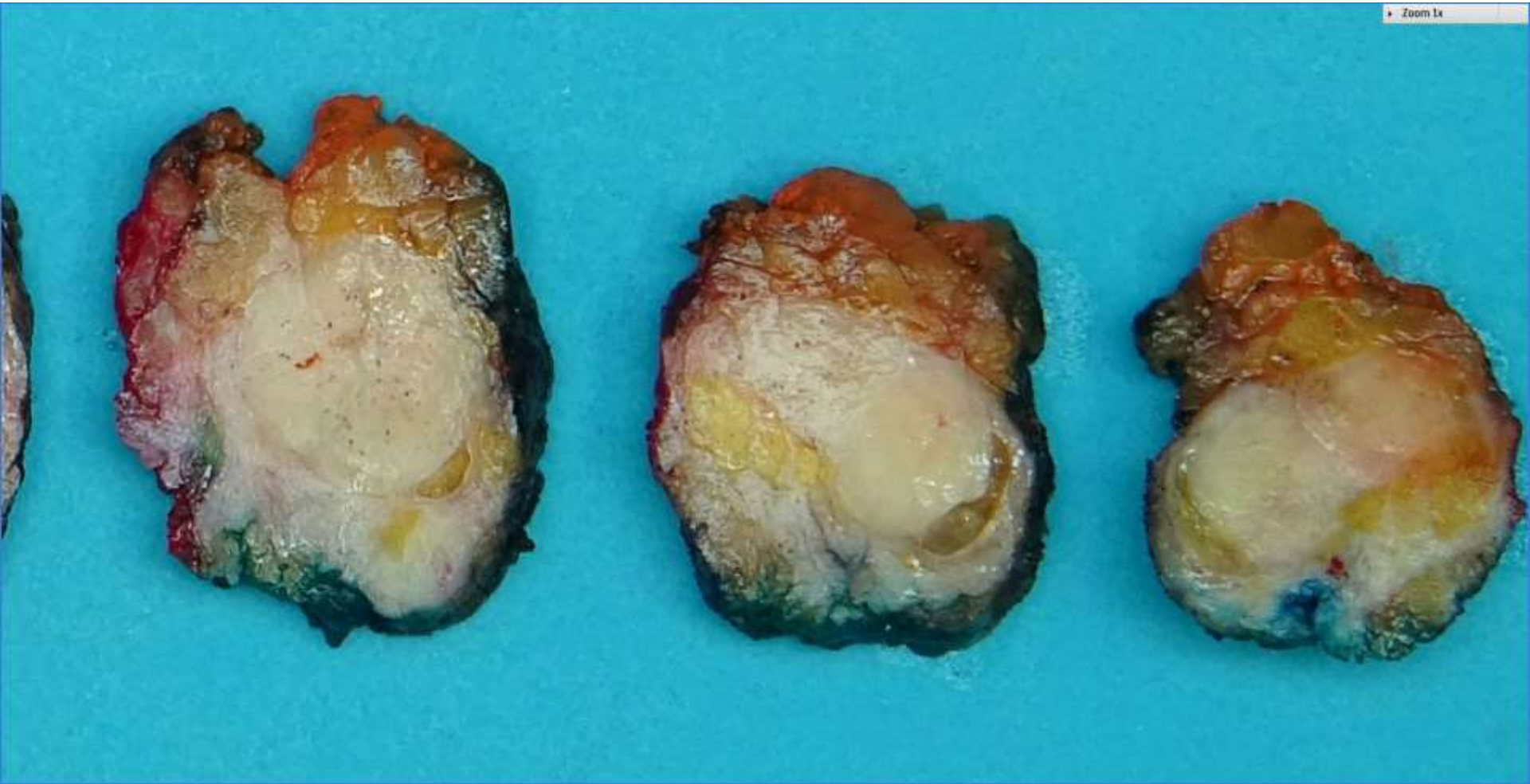


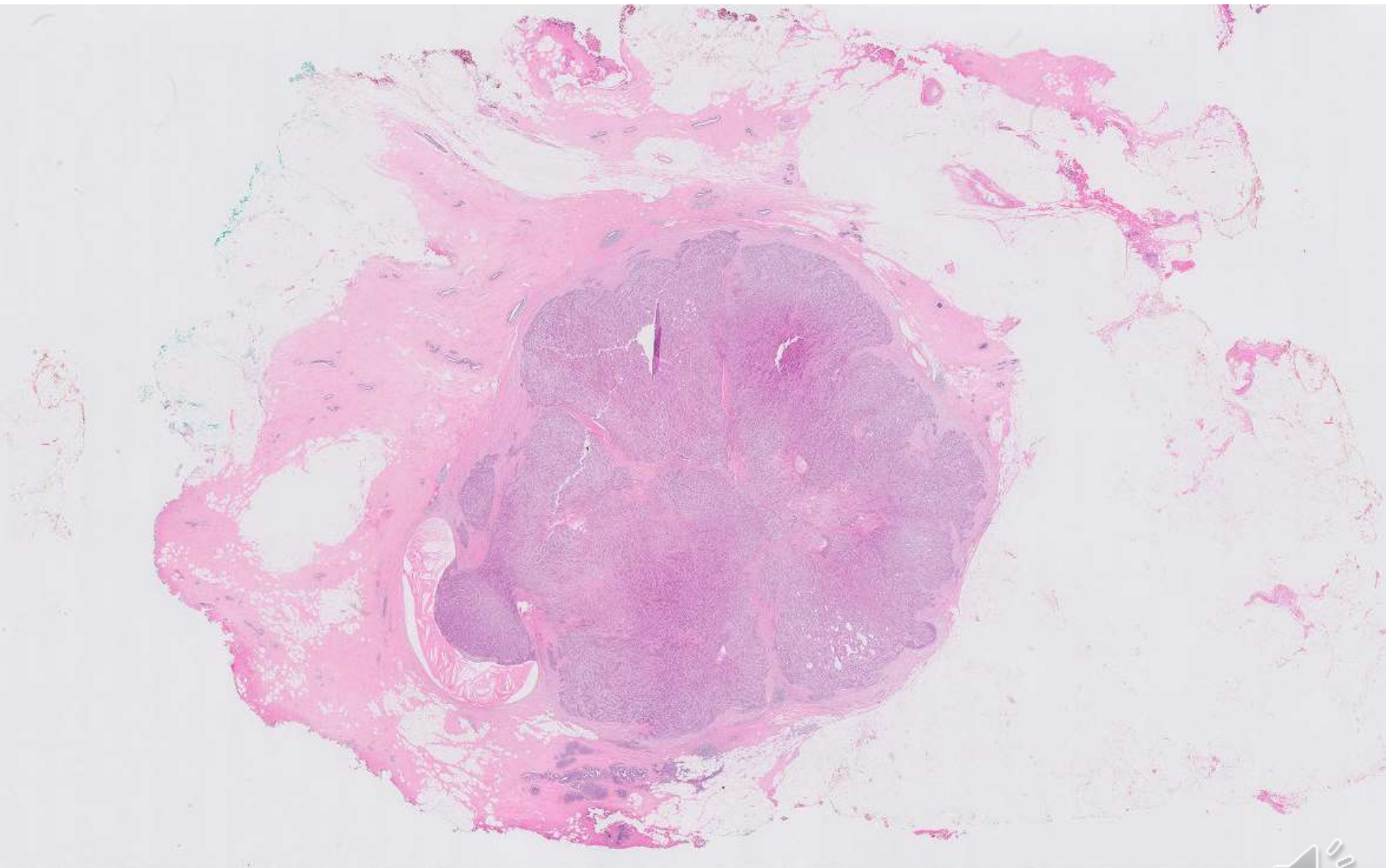
Case 16

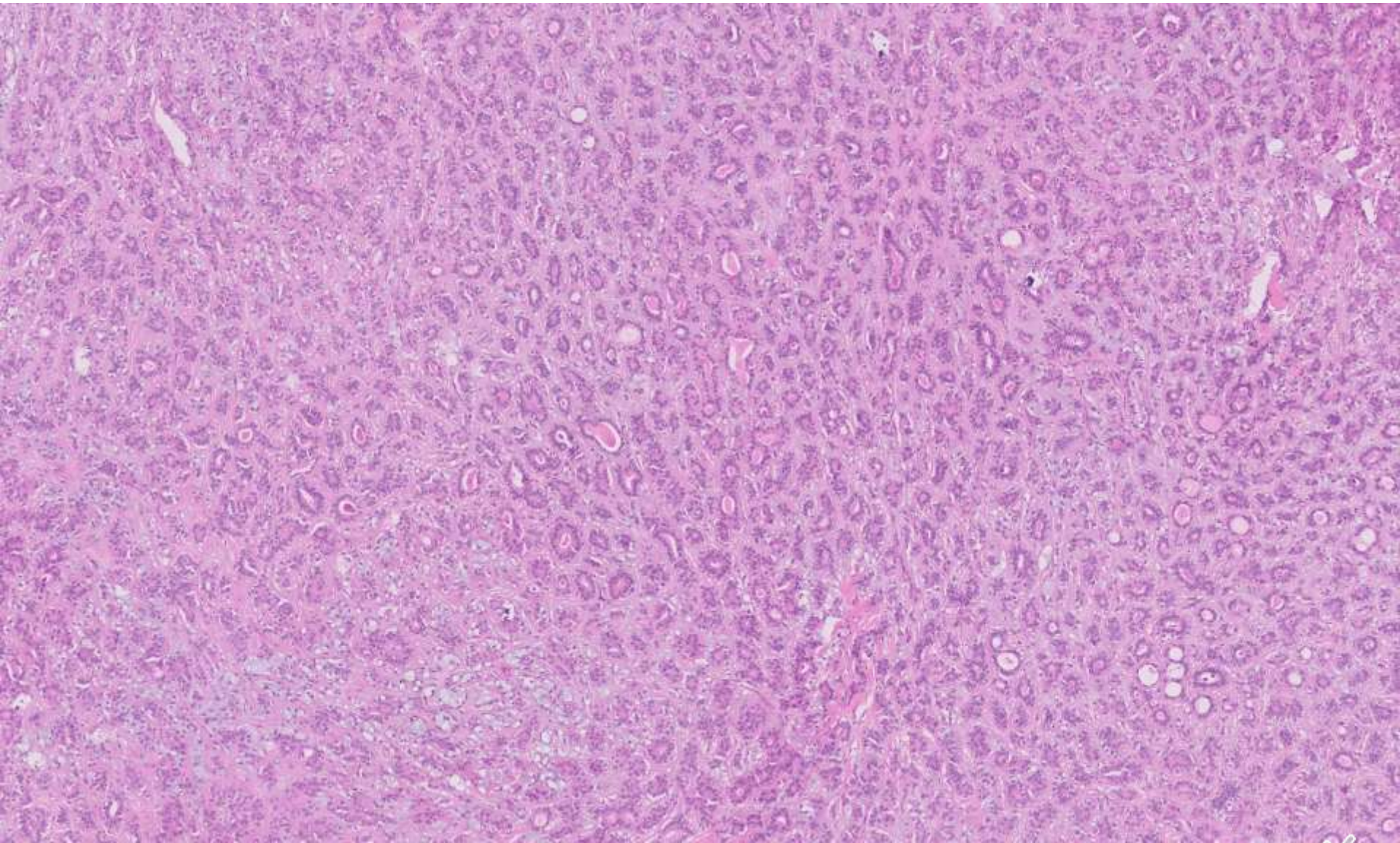
66 year old Chinese female.
Hookwire localization excision biopsy for
a right breast 1 o'clock nodule and
upper central calcifications.

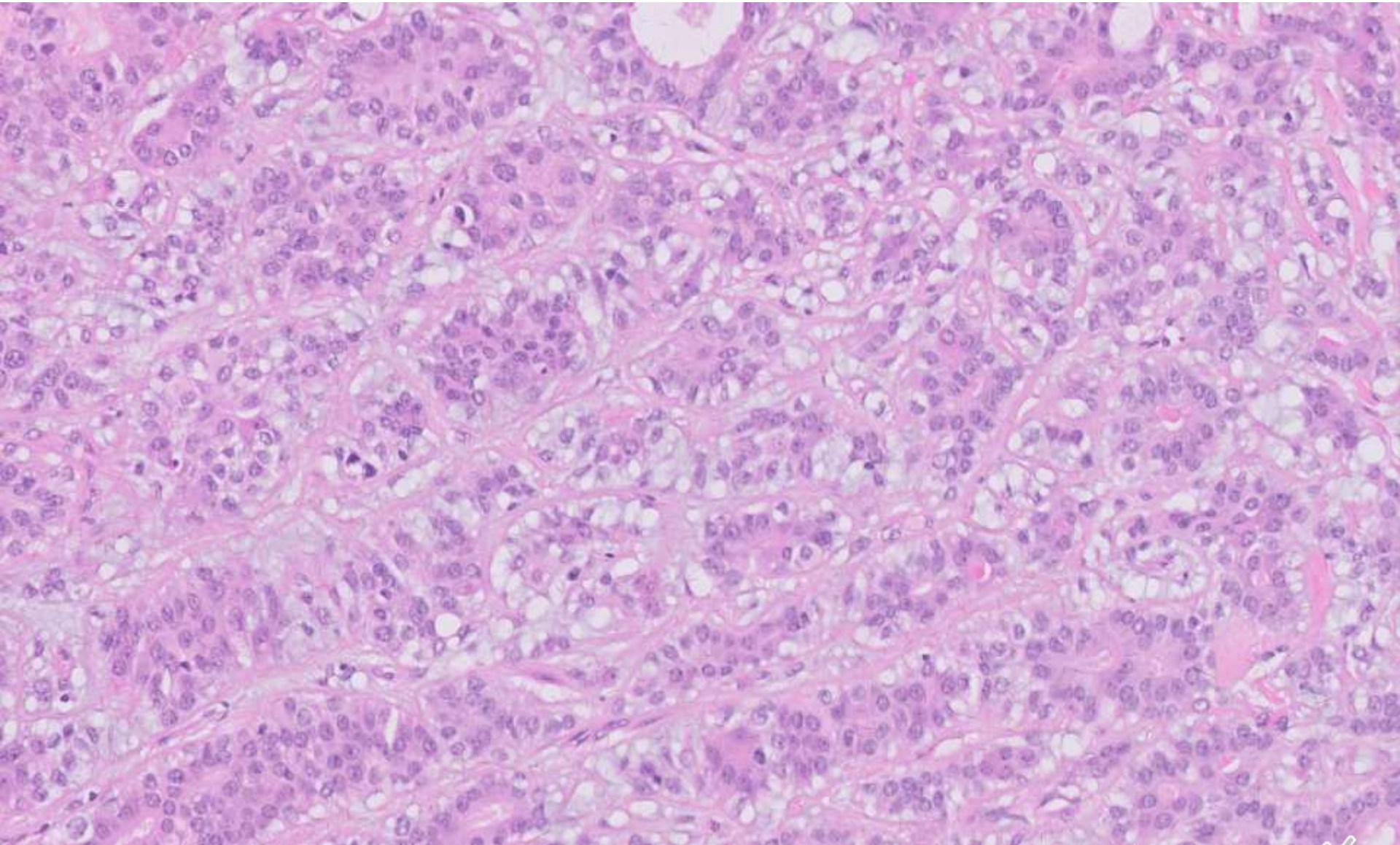
Presented by Timothy Tay

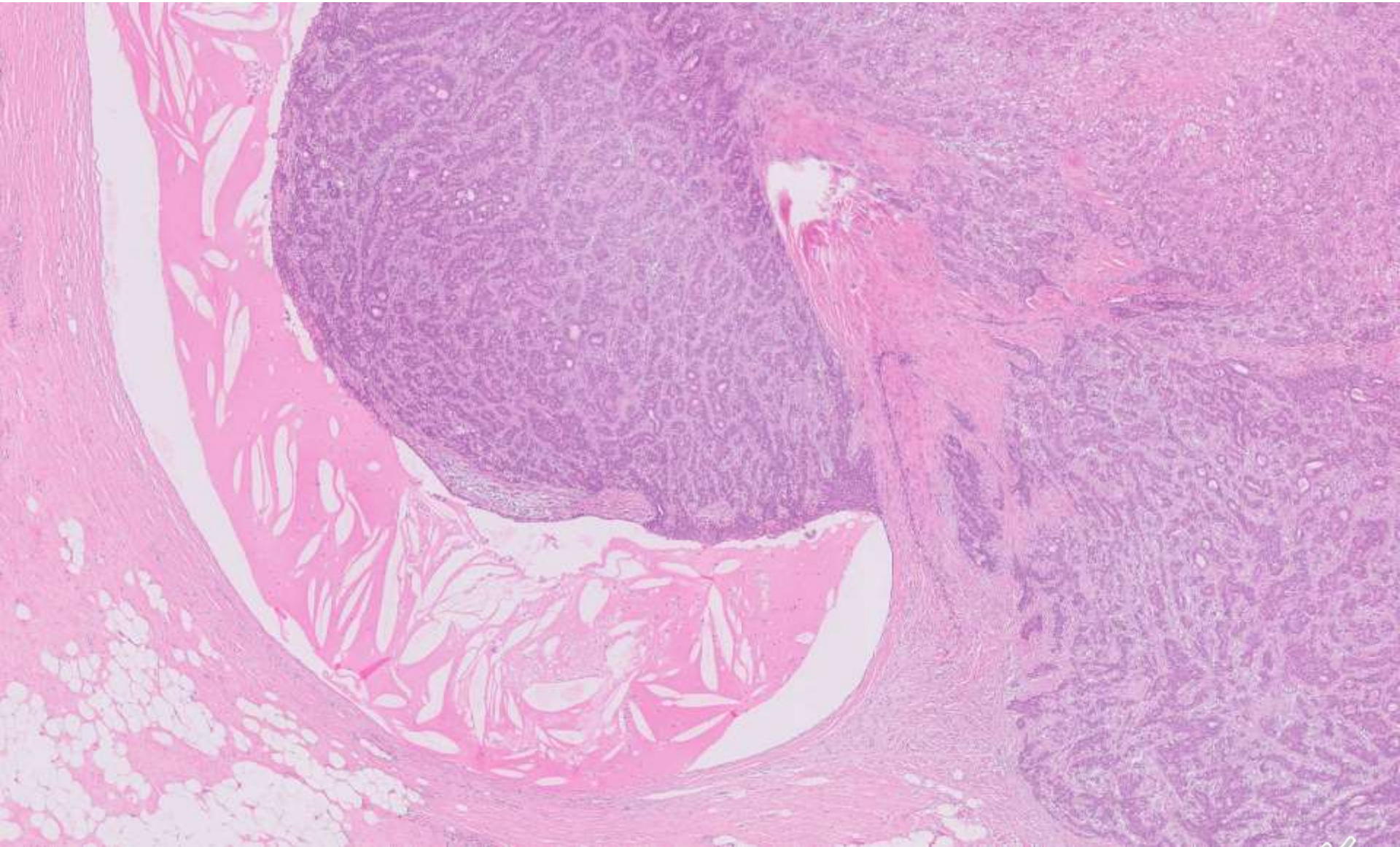


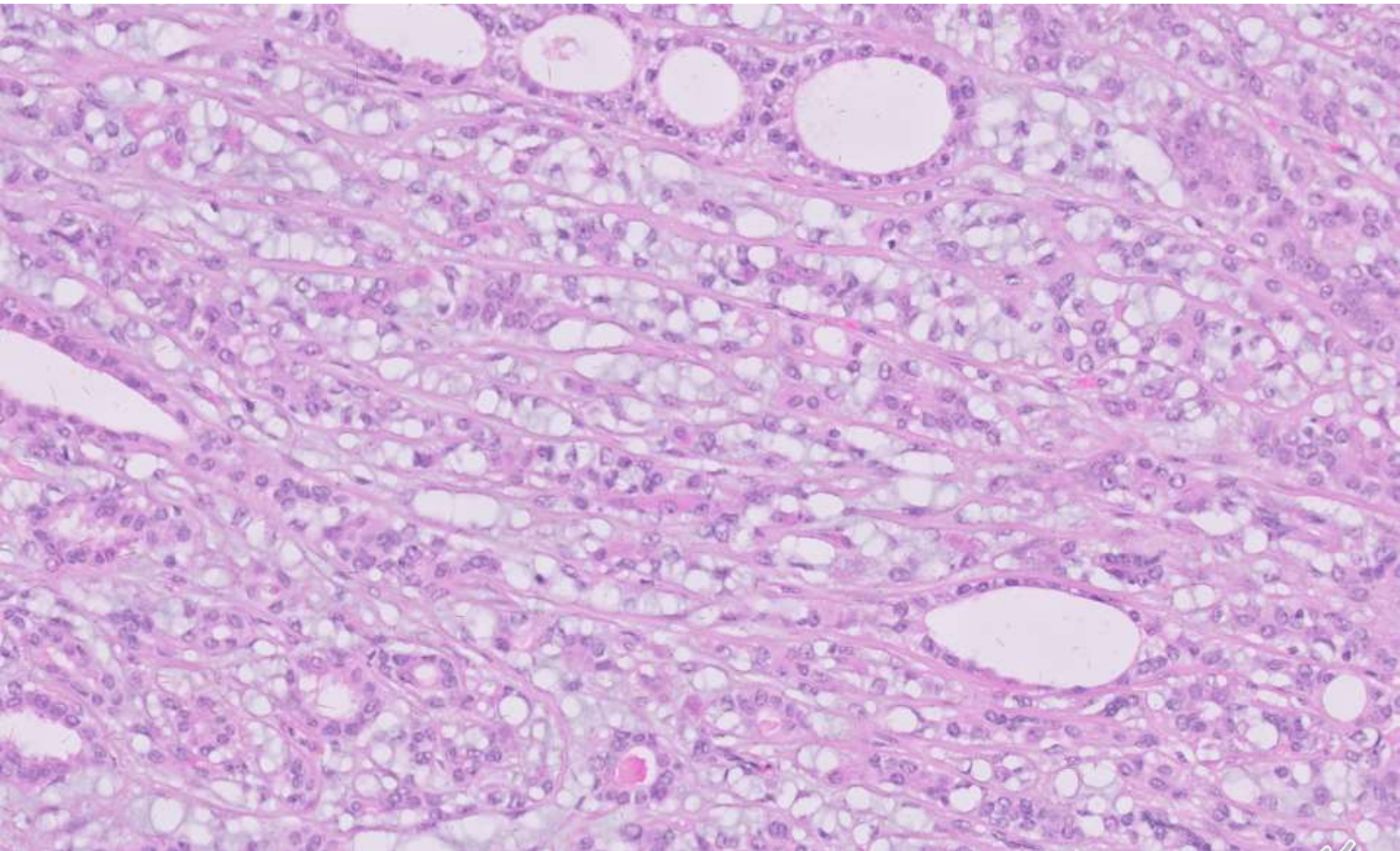












Diagnosis

- Adenomyoepithelioma



Molecular genetics

- In a study by Geyer et al. of 43 adenomyoepitheliomas of which 31 had whole exome sequencing or targeted next generation sequencing of 410 genes, *PIK3CA* hotspot mutations was found in 52% of adenomyoepitheliomas, *HRAS* mutations in 26% (p.Q61R or p.Q61K mutation), *AKT1* in 13% and *PIK3R1* mutation in 13% of tumours.
- AMEs had a lower mutation burden/fewer somatic mutations compared to invasive breast cancers.



Molecular genetics

- ER-positive and ER-negative AMEs have different mutational profiles.
- *AKT1* (p.E17K) mutation was seen only in ER-positive AMEs.
- *HRAS* and *PIK3R1* mutations were seen only in ER-negative AMEs.
- *PIK3CA* mutation was seen in both.
- *HRAS* mutations in ER-negative AME was significantly associated with necrosis and increased mitotic activity.
- ER-negative AMEs also had higher number of somatic mutations compared to ER-positive AMEs, although the difference did not reach statistical significance.







ARTICLE

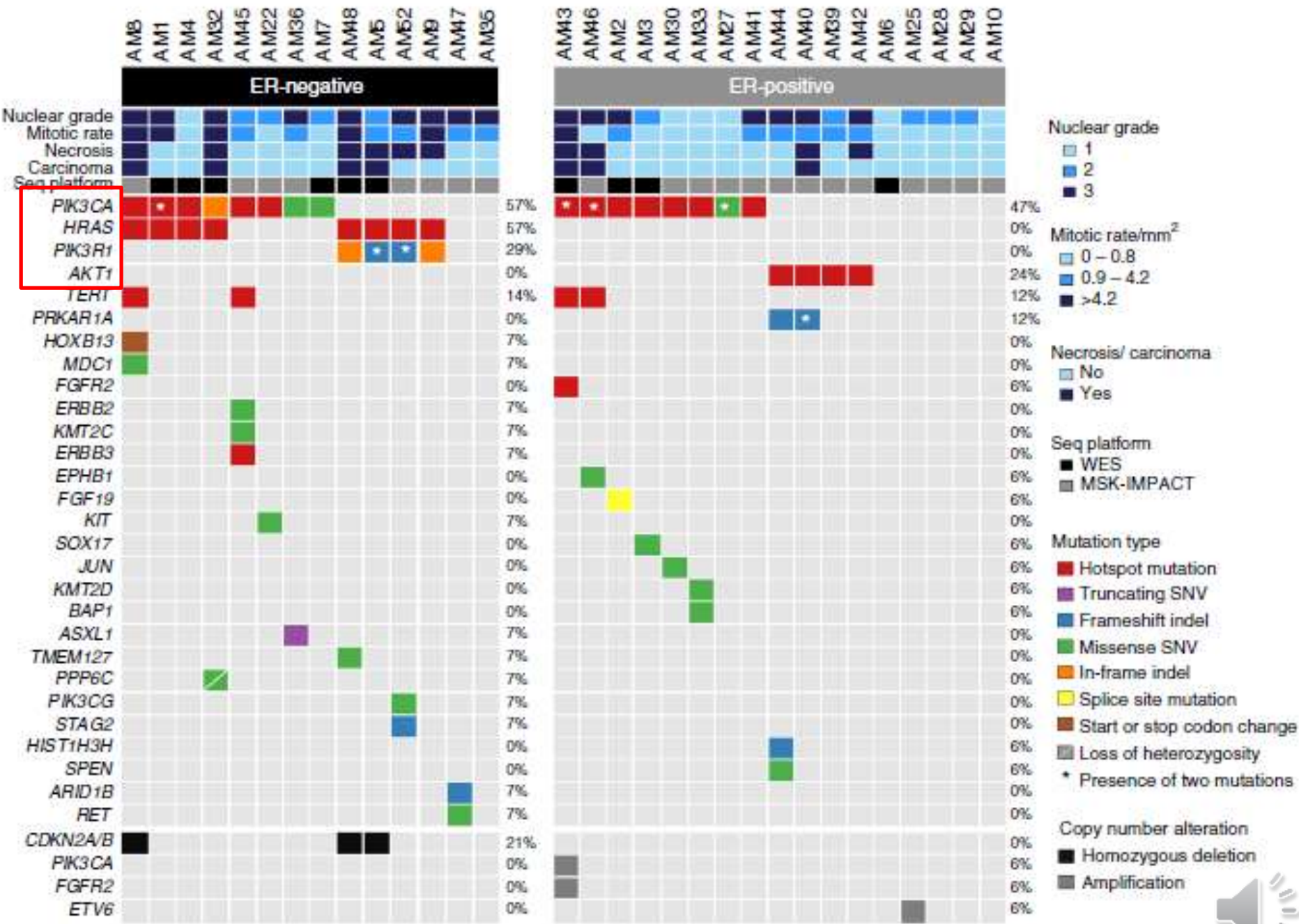
DOI: 10.1038/s41467-018-04128-5

OPEN

Recurrent hotspot mutations in *HRAS* Q61 and PI3K-AKT pathway genes as drivers of breast adenomyoepitheliomas

Felipe C. Geyer^{1,2,3}, Anqi Li^{1,4}, Anastasios D. Papanastasiou^{1,5}, Alison Smith⁶, Pier Selenica¹, Kathleen A. Burke¹, Marcia Edelweiss¹, Huei-Chi Wen¹, Salvatore Piscuoglio ^{1,7}, Anne M. Schultheis¹, Luciano G. Martelotto¹, Fresia Pareja¹, Rahul Kumar¹, Alissa Brandes¹, Dan Fan^{1,8}, Thais Basili¹, Arnaud Da Cruz Paula¹, John R. Lozada ¹, Pedro Blecua⁹, Simone Muenst⁷, Achim A. Jungbluth¹, Maria P. Foschini¹⁰, Hannah Y. Wen¹, Edi Brogi¹, Juan Palazzo¹¹, Brian P. Rubin¹², Charlotte K.Y. Ng ^{1,7,13}, Larry Norton¹⁴, Zsuzsanna Varga¹⁵, Ian O. Ellis¹⁶, Emad A. Rakha¹⁶, Sarat Chandarlapaty ⁶, Britta Weigelt¹ & Jorge S. Reis-Filho^{1,6}





Molecular genetics

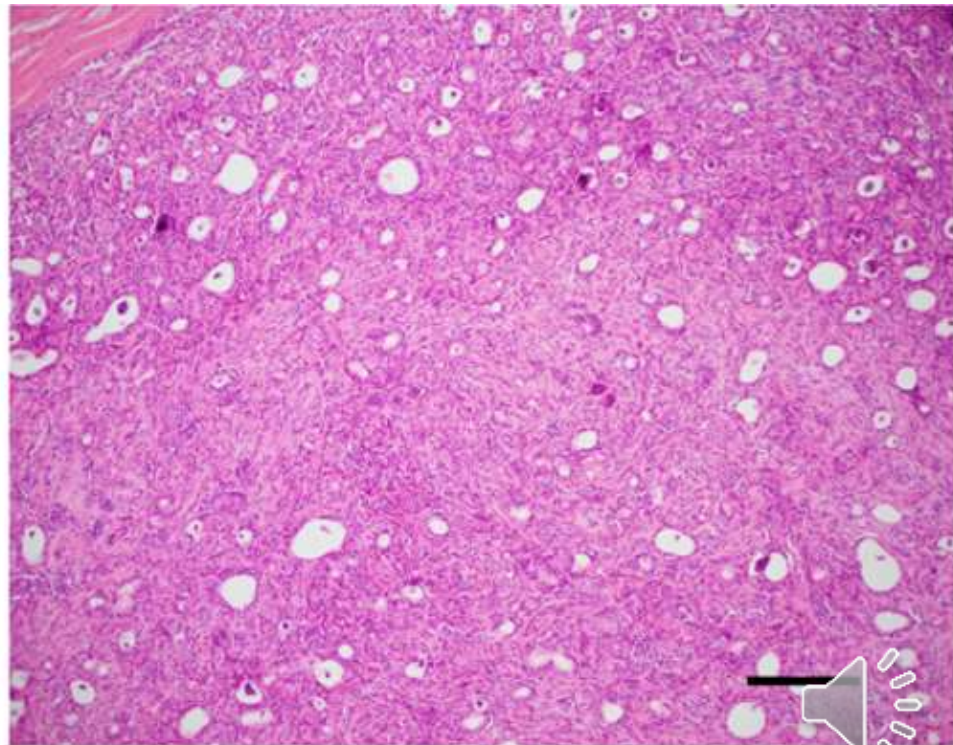
- Interestingly, papillomas without atypia which overlaps morphologically with ER-positive AME with papillary architecture have also been found to harbour *PIK3CA* and *AKT1* mutation in a separate study.
- *HRAS* p.Q61 mutations which occur in ER-negative tumours have also been reported in morphologically similar epithelial-myoepithelial carcinomas of the salivary gland.
- In a subsequent study by the same group, an AME that was wildtype for *PIK3CA*, *AKT1* and *HRAS* mutations was found to harbour a *HMGA2-WIF1* fusion, previously described in pleomorphic adenomas of the salivary gland.



ARTICLE OPEN

Assessment of *HMGA2* and *PLAG1* rearrangements in breast adenomyoepitheliomas

Fresia Pareja¹, Felipe C. Geyer¹, David N. Brown¹, Ana P. Martins Sebastião¹, Rodrigo Gularte-Mérida¹, Anqi Li¹, Marcia Edelweiss¹, Arnaud Da Cruz Paula¹, Pier Selenica¹, Hannah Y. Wen¹, Achim A. Jungbluth¹, Zsuzsanna Varga², Juan Palazzo³, Brian P. Rubin⁴, Ian O. Ellis⁵, Edi Brogi¹, Emad A. Rakha⁵, Britta Weigelt¹ and Jorge S. Reis-Filho¹



Prognosis

- Majority of AMEs are benign and excision with clear margins are curative.
- Tumours with extensive intraductal growth or satellite nodules may recur
- AMEs with benign histology have also been reported to metastasize to distant sites in very rare instances.





Breast
Pathology
Course 2020 @21

Thank You



Division of Pathology
Singapore General Hospital


SingHealth DukeNUS
ACADEMIC MEDICAL CENTRE
PATHOLOGY

