

Stanford



Erna Forgo

- Affiliate, Dean's Office Operations - Dean Other
- Fellow in Pathology
- Resident in Pathology

Bio

BIO

Dr. Erna Forgó is a Gastrointestinal & Hepatobiliary Pathology Fellow. She completed her Anatomic and Clinical Pathology Residency Training at Stanford University School of Medicine. Her clinical interests include Gastrointestinal & Hepatobiliary Pathology and Gynecologic Pathology.

BOARDS, ADVISORY COMMITTEES, PROFESSIONAL ORGANIZATIONS

- Member, United States and Canadian Academy of Pathology (USCAP) (2016 - present)
- Member, American Society of Clinical Pathology (ASCP) (2013 - present)
- Member, American Society of Clinical Oncology (ASCO) (2014 - present)

Publications

PUBLICATIONS

- **DNA Methylation Profiling of Uterine Sarcomas**
Forgo, E., Lang, A., Natu, V., Longacre, T., Bennett, J., Quick, C., Parra-Herran, C., Nucci, M., Kolin, D., Howitt, B.
NATURE PUBLISHING GROUP.2020: 1056–57
- **INSM1 Expression in Clear Cell Sarcoma of Kidney and other BCOR-Rearranged Tumors**
Wang, H., Forgo, E., Clay, M., Al-Ibraheemi, A., Charville, G.
NATURE PUBLISHING GROUP.2020: 835–36
- **5-hmC Immunohistochemistry is a Reliable Marker for Predicting Leiomyomas with Fumarate Hydratase Mutations**
Sharma, A., Forgo, E., Howitt, B., Rabban, J., Garg, K., Bennett, J.
NATURE PUBLISHING GROUP.2020: 1134–35
- **A human lung tumor microenvironment interactome identifies clinically relevant cell-type cross-talk.** *Genome biology*
Gentles, A. J., Hui, A. B., Feng, W., Azizi, A., Nair, R. V., Bouchard, G., Knowles, D. A., Yu, A., Jeong, Y., Bejnood, A., Forgo, E., Varma, S., Xu, et al
2020; 21 (1): 107
- **Detection of Circulating Tumor DNA in Patients With Uterine Leiomyomas** *JCO PRECISION ONCOLOGY*
Przybyl, J., Spans, L., Lum, D. A., Zhu, S., Vennam, S., Forgo, E., Varma, S., Ganjoo, K., Hastie, T., Bowen, R., Debiec-Rychter, M., van de Rijn, M.
2019; 3
- **Morphologic, Immunophenotypic and Molecular Features of Hypermutation in Colorectal Carcinomas with Mutations in DNA Polymerase # (POLE).** *Histopathology*
Forgó, E., Gomez, A. J., Steiner, D., Zehnder, J., Longacre, T. A.
2019
- **Plurihormonal pituitary neuroendocrine tumor with Pit1 and SF-1 coexpression: A novel entity**
Jonathan, L., Forgo, E., Katznelson, L., Harsh, G., Vogel, H.

OXFORD UNIV PRESS INC.2018: 523

- **Combination Approach for Detecting Different Types of Alterations in Circulating Tumor DNA in Leiomyosarcoma** *CLINICAL CANCER RESEARCH*
Przybyl, J., Chabon, J. J., Spans, L., Ganjoo, K. N., Vennam, S., Newman, A. M., Forgo, E., Varma, S., Zhu, S., Debiec-Rychter, M., Alizadeh, A. A., Diehn, M., van de Rijn, et al
2018; 24 (11): 2688–99
- **Practical Approach to the Use of Helicobacter Immunohistochemistry Based on a Single-Institution Retrospective Quality Assurance Review**
Forgo, E., Longacre, T., Martin, B.
NATURE PUBLISHING GROUP.2018: 774–75
- **Clinicopathologic Features of Mismatch Repair Protein-Deficient Non-Colorectal Gastrointestinal, Esophageal and Pancreatic Adenocarcinomas**
Forgo, E., Charville, G., Longacre, T.
NATURE PUBLISHING GROUP.2018: 261
- **PAX7 Expression in Rhabdomyosarcoma, Related Soft Tissue Tumors, and Small Round Blue Cell Neoplasms.** *American journal of surgical pathology*
Charville, G. W., Varma, S., Forgó, E., Dumont, S. N., Zambrano, E., Trent, J. C., Lazar, A. J., van de Rijn, M.
2016; 40 (10): 1305-1315
- **Transcriptome sequencing analysis of four psammomatous meningiomas**
Samghabadi, P., Forgo, E., West, R., Vogel, H.
LIPPINCOTT WILLIAMS & WILKINS.2016: 575
- **Correlation of Protein Expression with Chromosomal Copy Number Alterations in the Progression of Early Breast Neoplasia to DCIS**
Forgo, E., Varma, S., West, R. B.
NATURE PUBLISHING GROUP.2016: 41A
- **CDX2 as a Prognostic Biomarker in Stage II and Stage III Colon Cancer** *NEW ENGLAND JOURNAL OF MEDICINE*
Dalerba, P., Sahoo, D., Paik, S., Guo, X., Yothers, G., Song, N., Wilcox-Fogel, N., Forgo, E., Rajendran, P. S., Miranda, S. P., Hisamori, S., Hutchison, J., Kalisky, et al
2016; 374 (3): 211-222
- **Molecular subtyping of leiomyosarcoma with 3' end RNA sequencing.** *Genomics data*
Guo, X., Forgó, E., van de Rijn, M.
2015; 5: 366-367
- **Chromosomal copy number alterations for associations of ductal carcinoma in situ with invasive breast cancer** *BREAST CANCER RESEARCH*
Afghahi, A., Forgo, E., Mitani, A. A., Desai, M., Varma, S., Seto, T., Rigdon, J., Jensen, K. C., Troxell, M. L., Gomez, S. L., Das, A. K., Beck, A. H., Kurian, et al
2015; 17
- **Clinically Relevant Molecular Subtypes in Leiomyosarcoma.** *Clinical cancer research*
Guo, X., Jo, V. Y., Mills, A. M., Zhu, S. X., Lee, C., Espinosa, I., Nucci, M. R., Varma, S., Forgó, E., Hastie, T., Anderson, S., Ganjoo, K., Beck, et al
2015; 21 (15): 3501-3511
- **Correlation of Protein Expression With Chromosomal Copy Number Alterations for Risk Classification of Ductal Carcinoma In Situ**
Forgo, E., Varma, S., West, R.
NATURE PUBLISHING GROUP.2015: 45A
- **Chromosomal copy number alterations (CNAs) for risk assessment of ductal carcinoma in situ (DCIS)**
Afghahi, A., Forgo, E., Mitani, A., Desai, M., Varma, S., Seto, T., Jensen, K. C., Gomez, S., Das, A. K., Beck, A. H., Kurian, A. W., West, R. B.
AMER SOC CLINICAL ONCOLOGY.2014