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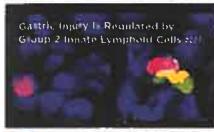
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1995 A. T. Chan and C. S. Williams

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E. Dekker, H.-M. Chiu, I. Lansdorp-Vogelaar, and On behalf of the Expert Working Group on COVID-19 of the WEO Colorectal Cancer Screening Committee

2004 Monitoring Fecal Microbiota Transplantation Practice in a Rapidly Evolving Health and Regulatory Environment

C. R. Kelly, L. A. Laine, and G. D. Wu

2007 Moving Toward Impact: An Introduction to Implementation Science for Gastroenterologists and Hepatologists

S. S. Rogal, B. J. Powell, M. Chinman, and On behalf of the Gastroenterology and Hepatology Implementation Research Group

MEETING SUMMARY

2013 Defining Endpoints and Biomarkers in Inflammatory Bowel Disease: Moving the Needle Through Clinical Trial Design

M. T. Abre, W. J. Sandborn, and the IOIBD Defining Endpoints and Biomarkers in Inflammatory Bowel Disease Writing Group



Video



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CME quiz



Editorial accompanies this article



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Cover

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- 2025 Could a Small Population of Epithelial Cells Get “Tuft” With Crohn’s Disease?**
M. J. Rosen
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A. Courtet, J. Lemale, and M. C. Carra
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- 2039 Alterations in Intestinal Microbiota of Children With Celiac Disease at the Time of Diagnosis and on a Gluten-free Diet**
K. Zafeiropoulou, B. Nichols, M. Mackinder, O. Biskou, E. Rizou, A. Karanikolou, C. Clark, E. Buchanan, T. Cardigan, H. Duncan, D. Wands, J. Russell, R. Hansen, R. K. Russell, P. McGrogan, C. A. Edwards, U. Z. Ijaz, and K. Gerasimidis

Children with celiac disease have differences in composition of intestinal microbes compared to healthy children. Some of these differences are caused by a gluten-free diet, but studies are needed to determine whether the other changes are a cause or a result of celiac disease.

- 2052 Relationship Between Combined Histologic and Endoscopic Endpoints and Efficacy of Ustekinumab Treatment in Patients With Ulcerative Colitis**
K. Li, C. Marano, H. Zhang, F. Yang, W. J. Sandborn, B. E. Sands, B. G. Feagan, D. T. Rubin, L. Peyrin-Biroulet, J. R. Friedman, and G. De Hertoghe

The drug ustekinumab induces and maintains mucosal healing in patients with UC, based on endoscopy and pathology evaluation.

Basic and Translational—Alimentary Tract

- 2065 Sex-Specific Genetic Associations for Barrett's Esophagus and Esophageal Adenocarcinoma**
 
J. Dong, C. Maj, S. Tsavachidis, Q. T. Ostrom, P. Gharakhhani, L. A. Anderson, A. H. Wu, W. Ye, L. Bernstein, O. Borisov, J. Schröder, W.-H. Chow, M. D. Gammon, G. Liu, C. Caldas, P. D. Pharoah, H. A. Risch, A. May, C. Gerges, M. Anders, M. Venerito, T. Schmidt, J. R. Izbicki, A. H. Hölscher, B. Schumacher, Y. Vashist, H. Neuhaus, T. Rösch, M. Knapp, P. Krawitz, A. Böhmer, P. G. Iyer, B. J. Reid, J. Lagergren, N. J. Shaheen, D. A. Corley, I. Gockel, R. C. Fitzgerald, Stomach and Oesophageal Cancer Study (SOCS) consortium, M. B. Cook, D. C. Whiteman, T. L. Vaughan, J. Schumacher, and A. P. Thrift

The authors identify sex-specific genetic features that are associated with BE and EA, and that may contribute to sex disparities in these diseases.

- 2077 Group 2 Innate Lymphoid Cells Coordinate Damage Response in the Stomach**
 
A. R. Meyer, A. C. Engevik, T. Madorsky, E. Belmont, M. T. Stier, A. E. Norlander, M. A. Pilkinton, W. J. McDonnell, J. A. Weis, B. Jang, S. A. Mallal, R. S. Peebles Jr, and J. R. Goldenring

We have determined that intrinsic mucosal immune cells are integral for coordinated repair of the gastric lining following severe injury.

- 2092 Increased Intestinal Permeability Is Associated With Later Development of Crohn's Disease**
 
W. Turpin, S.-H. Lee, J. A. Raygoza Garay, K. L. Madsen, J. B. Meddings, L. Bedrani, N. Power, O. Espin-Garcia, W. Xu, M. I. Smith, A. M. Griffiths, P. Moayyedi, D. Turner, E. G. Seidman, A. H. Steinhart, J. K. Marshall, K. Jacobson, D. Mack, H. Huynh, C. N. Bernstein, A. D. Paterson, The Crohn's and Colitis Canada Genetic Environmental Microbial Project Research Consortium, and K. Croitoru

Abnormal intestinal permeability appears to precede development of CD, and might be involved in pathogenesis. Strategies to restore gut barrier function might be developed to prevent CD in susceptible individuals.

- 2101 Succinate Produced by Intestinal Microbes Promotes Specification of Tuft Cells to Suppress Ileal Inflammation**
 
A. Banerjee, C. A. Herring, B. Chen, H. Kim, A. J. Simmons, A. N. Southard-Smith, M. M. Allaman, J. R. White, M. C. Macedonia, E. T. McKinley, M. A. Ramirez-Solano, E. A. Scoville, Q. Liu, K. T. Wilson, R. J. Coffey, M. K. Washington, J. A. Goettel, and K. S. Lau

See editorial on page 2025.

Tuft cell subpopulations in the intestinal tract respond differentially to external stimuli. Therapeutic expansion of tuft cells in models of small intestinal inflammation ameliorates disease.

- 2116 Single-Cell Transcriptional Analyses Identify Lineage-Specific Epithelial Responses to Inflammation and Metaplastic Development in the Gastric Corpus**
 
K. A. Bockerstett, S. A. Lewis, C. N. Noto, E. L. Ford, J. B. Saenz, N. M. Jackson, T.-H. Ahn, J. C. Mills, and R. J. DiPaolo

This study identified cells and proteins that contribute to development of cancer in the inflamed stomach.

- 2130 S100A8 and S100A9 Are Important for Postnatal Development of Gut Microbiota and Immune System in Mice and Infants**
 
M. Willers, T. Ulas, L. Völlger, T. Vogl, A. S. Heinemann, S. Pirr, J. Pagel, B. Fehlhaber, O. Halle, J. Schöning, S. Schreek, U. Löber, M. Essex, P. Hombach, S. Graspeuntner, M. Basic, A. Bleich, K. Cloppenborg-Schmidt, S. Künzel, D. Jonigk, J. Rupp, G. Hansen, R. Förster, J. F. Baines, C. Härtel, J. L. Schultze, S. K. Forslund, J. Roth, and D. Viemann

The authors found proteins that modulate the immune system, that are present in higher levels of full-term infants than preterm infants, and that are important for development of a healthy intestinal microbiota and immune system.

2146 The Long Noncoding RNA CCAT2 Induces Chromosomal Instability Through BOP1-AURKB Signaling

B. Chen, M. P. Dragomir, L. Fabris, R. Bayraktar, E. Knutsen, X. Liu, C. Tang, Y. Li, T. Shimura, T. C. Ivkovic, M. Cruz De los Santos, S. Anfossi, M. Shimizu, M. Y. Shah, H. Ling, P. Shen, A. S. Multani, B. Pardini, J. K. Burks, H. Katayama, L. C. Reineke, L. Huo, M. Syed, S. Song, M. Ferracin, E. Oki, B. Fromm, C. Ivan, K. Bhuvaneshwar, Y. Gusev, K. Mimori, D. Menter, S. Sen, T. Matsuyama, H. Uetake, C. Vasilescu, S. Kopetz, J. Parker-Thornburg, A. Taguchi, S. M. Hanash, L. Girnita, O. Slaby, A. Goel, G. Varani, M. Gagea, C. Li, J. A. Ajani, and G. A. Calin

The authors identified a gene product that causes chromosomes to become unstable in colon cells, promoting development of colorectal cancer.

2163 In Colorectal Cancer Cells With Mutant KRAS, SLC25A22-Mediated Glutaminolysis Reduces DNA Demethylation to Increase WNT Signaling, Stemness, and Drug Resistance

C. C. Wong, J. Xu, X. Bian, J.-L. Wu, W. Kang, Y. Qian, W. Li, H. Chen, H. Gou, D. Liu, S. T. Y. Luk, Q. Zhou, F. Ji, L.-S. Chan, S. Shirasawa, J. JY. Sung, and J. Yu

The authors identified an altered metabolic pathway in colorectal cancer cells that increases their proliferation and survival. Strategies to block this pathway might be developed for treatment of colorectal and other cancers.

2181 Clostridioides difficile Toxin A Remodels Membranes and Mediates DNA Entry Into Cells to Activate Toll-Like Receptor 9 Signaling

X. Chen, X. Yang, J. de Anda, J. Huang, D. Li, H. Xu, K. S. Shields, M. Džunková, J. Hansen, I. J. Patel, E. U. Yee, D. T. Golenbock, M. A. Grant, G. C. L. Wong, and C. P. Kelly

This study identified a mechanism by which a bacterial toxin activates an inflammatory response in the intestine.

2193 Linking Strain Engraftment in Fecal Microbiota Transplantation With Maintenance of Remission in Crohn's Disease

L. Kong, J. Lloyd-Price, T. Vatanen, P. Seksik, L. Beaugerie, T. Simon, H. Vlamakis, H. Sokol, and R. J. Xavier

This manuscript represents the first study to use metagenomics to investigate the effects of fecal microbiota transplant (FMT) in Crohn's disease (CD). This approach enables far greater resolution than previously available, revealing strain- and species-level differences induced by FMT.

Basic and Translational—Liver**2203 Cooperation Between Distinct Cancer Driver Genes Underlies Intertumor Heterogeneity in Hepatocellular Carcinoma**

P. Molina-Sánchez, M. Ruiz de Galarreta, M. A. Yao, K. E. Lindblad, E. Bresnahan, E. Bitterman, T. C. Martin, T. Rubenstein, K. Nie, J. Golas, S. Choudhary, M. Bárcena-Varela, A. Elmas, V. Miguela, Y. Ding, Z. Kan, L. T. Grinspan, K.-L. Huang, R. E. Parsons, D. J. Shields, R. A. Rollins, and A. Lujambio

This novel collection of murine hepatocellular carcinoma models and corresponding cell lines establishes the role of cancer driver genes in promoting inter-tumor heterogeneity and enables mechanistic and translational studies.

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J. M. Kolb, S. Han, F. I. Scott, C. C. Murphy, P. Hosokawa, and S. Wani, on behalf of the Early Onset Esophageal Adenocarcinoma Study Group
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P. E. Young, M. Tadros, and S. Mago

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