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The CORE-IBD Collaborators: C. Ma, J. Hanzel, R. Panaccione, W. J. Sandborn, G. R. D'Haens, V. Ahuja, R. Atreya, C. N. Bernstein, P. Bossuyt, B. Bressler, R. V. Bryant, B. Cohen, J.-F. Colombel, S. Danese, A. Dignass, M. C. Dubinsky, P. R. Fleshner, R. B. Gearry, S. B. Hanauer, A. Hart, P. G. Kotze, T. Kucharzik, P. L. Lakatos, R. W. Leong, F. Magro, J. Panés, L. Peyrin-Biroulet, Z. Ran, M. Regueiro, S. Singh, A. Spinelli, A. H. Steinhart, S. P. Travis, C. J. van der Woude, B. Yacyshyn, T. Yamamoto, M. Allez, W. A. Bemelman, A. L. Lightner, E. Louis, D. T. Rubin, E. J. Scherl, C. A. Siegel, M. S. Silverberg, S. Vermeire, C. E. Parker, S. C. McFarlane, L. Guizzetti, M. I. Smith, N. Vande Casteele, B. G. Feagan, and V. Jairath

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A.-M. Globig, L. S. Mayer, M. Heeg, G. Andrieux, M. Ku, P. Otto-Mora, A. V. Hipp, K. Zoldan, A. Pattekar, N. Rana, C. Schell, M. Boerries, M. Hofmann, C. Neumann-Haefelin, A. Kuellmer, A. Schmidt, T. Boettler, V. Tomov, R. Thimme, P. Hasselblatt, and B. Bengsch

This study identifies the exhaustion status of a population of CD8⁺ immune cells that express the regulating molecule CD39 as linked to the clinical course of Crohn's disease.

Functional GI Disease**982****Efficacy of Fecal Microbiota Transplantation for Patients With Irritable Bowel Syndrome at 3 Years After Transplantation**

M. El-Salhy, R. Winkel, C. Casen, T. Hausken, O. H. Gilja, and J. G. Hatlebakk

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This study applied a protocol for fecal microbiota transplantation to irritable bowel syndrome with high response rates and sustained effects. Ten possible were identified, who were correlated to fecal microbiota transplantation effects.

995**Risk Factors for Abdominal Pain-Related Disorders of Gut-Brain Interaction in Adults and Children: A Systematic Review**

J. K. Zia, A. Lenhart, P.-L. Yang, M. M. Heitkemper, J. Baker, L. Keefer, M. Saps, C. Cuff, G. Hungria, E. J. Videlock, and L. Chang

This comprehensive systematic review assessed multiple risk and protective factors for abdominal pain disorders of gut-brain interaction and symptom persistence in both adults and children.

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1024 Integrated Analysis of Colorectal Cancer Reveals Cross-Cohort Gut Microbial Signatures and Associated Serum Metabolites

 *R. Gao, C. Wu, Y. Zhu, C. Kong, Y. Zhu, Y. Gao, X. Zhang, R. Yang, H. Zhong, X. Xiong, C. Chen, Q. Xu, and H. Qin*

Cross-cohort gut microbiome signatures and their substantial serum metabolic links for CRC are identified and a composite microbial–metabolic predictive model is established, which may assist its diagnosis and therapeutics.

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 *N. Nagata, S. Nishijima, T. Miyoshi-Akiyama, Y. Kojima, M. Kimura, R. Aoki, M. Ohsugi, K. Ueki, K. Miki, E. Iwata, K. Hayakawa, N. Ohmagari, S. Oka, M. Mizokami, T. Itoi, T. Kawai, N. Uemura, and M. Hattori*

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 *M. Krishnamurthy, D. O. Kechelle, T. Broda, X. Zhang, J. R. Enriquez, H. A. McCauley, J. G. Sanchez, K. McCracken, J. Palermo, A. Bernieh, M. H. Collins, I. H. Thomas, H. C. Neef, A. Heider, A. Dauber, and J. M. Wells*

We used pluripotent stem cell-derived stomach and intestinal organoids to identify new pathologies and improve care for patients with PDX1 mutations.

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 *R. Bhatia, C. M. Thompson, E. J. Clement, K. Ganguly, J. L. Cox, S. Rauth, J. A. Siddiqui, S. S. Mashiana, M. Jain, T. A. Wyatt, H. S. Mashiana, S. Singh, N. T. Woods, K. K. Kharbanda, S. K. Batra, and S. Kumar*

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