



GENETICS IN CHOLESTATIC LIVER DISEASE: BEYOND FORTUNE TELLING

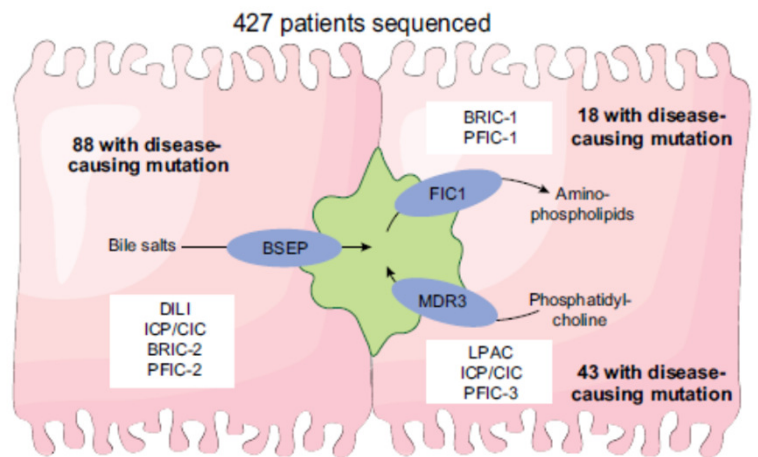
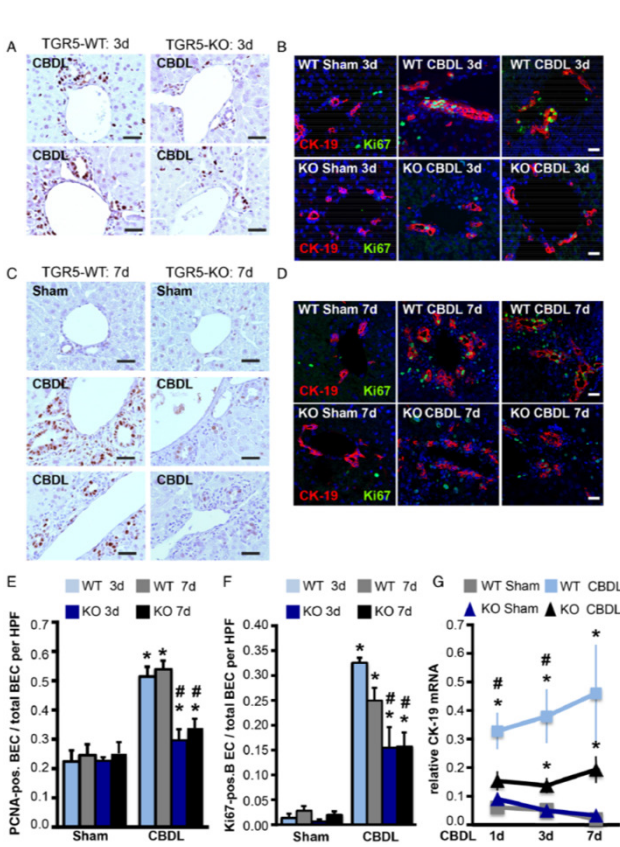
GUEST LECTURE by



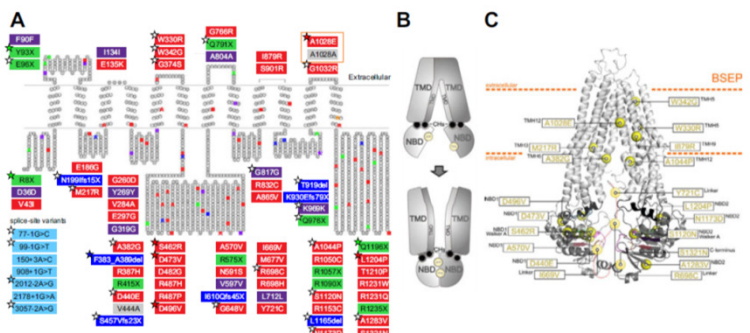
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Infectious Diseases, University Hospital
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Wednesday, 03.04.2019
11:00

MC1.G.01.005 (Seminar room 01 – Applied
Biomedicine Pathology 01, MED Campus, Neue
Stiftingtalstrasse 6, tract G, 1st floor), MUG



Sequencing of FIC1, BSEP and MDR3 in a large cohort of patients with cholestasis revealed a high number of different genetic variants.
Dröge et al. (2017) J. Hepatol. 67:1253-64



2D and 3D illustration of identified BSEP variants.
Dröge et al. (2017) J. Hepatol. 67:1253-64

Common bile duct ligation (CBDL) triggers a more pronounced cholangiocyte proliferation in TGR5-wild type (TGR-WT) as compared with TGR5-knockout (TGR5-KO) mice.

Reich et al (2016) Gut. 65:487-501