Basic hepatology

IDDF2020-ABS-0022 | THE METABOLIC SYNDROME AS A RISK FACTOR FOR NON-ALCOHOLIC FATTY LIVER DISEASE IN FILIPINO ADULTS CONSULTING IN A PHILIPPINE TERTIARY HOSPITAL: A RETROSPECTIVE COHORT **STUDY**

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Background Non-alcoholic Fatty Liver Disease (NAFLD) is a disease with increasing prevalence due to lifestyle diseases, particularly the Metabolic Syndrome. There is interest in studying its association with increased risk of progression to more serious liver disease.

Methods A retrospective approach was used to compare a total of 1,588 subjects who came for at least two wellness consults at The Medical City Center for Wellness and Aesthetics from 2013-2018.

Results On the baseline, the prevalence of the disease was 38%; and the components of the metabolic syndrome was higher among those with newly developed NAFLD on followup. NAFLD was also associated with weight gain. Multivariate analysis revealed a non-significant (p-value 0.041) adjusted odds ratio of 2.046 (1.030 - 4.063) in developing NAFLD among males with metabolic syndrome. While in women, the adjusted odds ratio in developing NAFLD was 3.886 (1.867 -8.085), with a p-value <0.001. This study estimated NAFLD prevalence in the Filipino population, with its findings consistent with previous literature in other countries.

Conclusions In conclusion, weight gain and metabolic syndrome are associated with an increased incidence of developing NAFLD, especially among adult Filipino males belonging in the older population group. NAFLD can be reversed by undergoing the proper diet and weight management.

IDDF2020-ABS-0025

THE ROLE OF GUT MICROBIOTA IN CLINICAL COMPLICATIONS AND TREATMENT RESPONSE IN ALCOHOLIC HEPATITIS - A CIRCOS®, LINEAR **DISCRIMINANT ANALYSIS EFFECT SIZE** BIOMARKER AND CONET® CO-OCCURRENCE NETWORK ANALYSIS

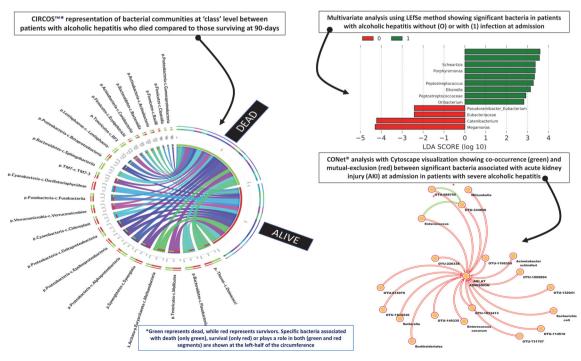
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Background In severe alcoholic hepatitis(SAH), we aimed to characterize significant bacterial communities associated with clinical events(CE) and define bacterial relationships associated with specific CE and outcomes at baseline and on corticosteroids.

Methods 16-s rRNA sequencing on stool samples(n=38) collected at admission and at last follow-up within 90-days was (N=26, 12 on corticosteroids). Visual-characterization was performed on QIIME data using CIRCOS™, linear-discriminantanalysis-effect-size (LEfSe) method was used to identify significant bacterial communities and their functional metabolites. Conet/Cytoscape® utilized to identify significant co-occurrence with respect to clinical events.

Results All were males with mean age 47.3±9.1 years, median discriminant function(DF) 64, Child-Pugh(CTP)12 and model for end-stage liver disease(MELD)25.5. At admission, 27%, 42%, and 58% had acute kidney injury (AKI), hepatic



Abstract IDDF2020-ABS-0025 Figure 1 Circos representation lefse analysis and conet result representation

Gut 2020;69(Suppl 2):A1-A95 A23 encephalopathy (HE) and infections respectively; 38.5% died at 90-days follow-up. Specific bacteria were found to be associated with HE [Bifidobacteriaceae, Coriobacteriaceae], sepsis [Veillonellaceae, Prevotellaceae], CTP>10 [Bifidobacteriaceae, Synergistaceae], MELD>25 [Dehalobacteriaceae, Turicibacteraceae] and death [Enterobacteriaceae, Peptococcaceae]. Significantly higher relative abundance(RA) of Lachnobacterium, Catenibacterium associated with HE at-admission while orallyrepresented bacteria were associated with infections at admission. Propionibacterium, Fusobacteria were associated with DF > 65 while Eubacterium, Capnocytophaga were associated with CTP. Enhydrobacter and Pediococcus were preferentially abundant post-steroid-therapy. Aerococcus was associated with post-treatment death. Prevotella was associated with survival post steroid. Upregulation of phenylpropanoid-biosynthesis (innate-immunity) in those without follow-up infections and glycerophospholipid-metabolism(cellular-integrity) in those who died were significant. Co-occurrence between Christensenella, Prevotella and mutual-exclusion between Megamonas, Citrobacter was associated with HE at admission. Mutual-exclusion between Coprococcus eutactus, Catenibacterium and Megamonas was associated with infections at admission while Enterococcus cecorum, Acinetobacter schindleri, Mitsuokella were associated with AKI at admission (figure 1).

Conclusions Specific gut-microbiota, their interactions and metabolites are associated with complications of SAH as well as outcomes with steroid-therapy. Advanced metagenomics-based precision-medicine as add-on treatments may be a novel therapeutic area for improving disease outcomes.

IDDF2020-ABS-0151

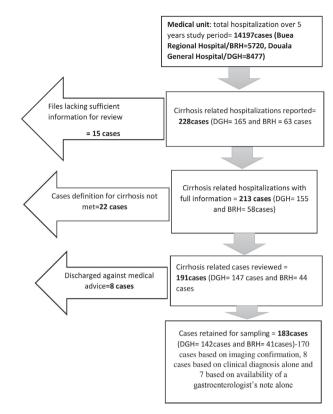
FACTORS PREDICTING IN-HOSPITAL
MORTALITY IN PATIENTS HOSPITALIZED
FOR LIVER CIRRHOSIS: A FIVE YEARS
RETROSPECTIVE REVIEW IN CAMEROON

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Background The global prevalence of liver cirrhosis has been on the rise, estimated at 4.5–9% from autopsy studies, projected at 50 million people affected in 2020 and ranked the 12th leading cause of death. However, data on studies evaluating the outcome of patients hospitalized for liver cirrhosis in SSA is scarce. We aimed to investigate the clinical and laboratory factors which predict in-hospital mortality in patients admitted for cirrhosis.

Methods This was a five years retrospective review of 183 cases hospitalized for liver cirrhosis between 1st January 2014 to 31st December 2018 in the medical units of two referral hospitals in Cameroon (figure1). Independent variables investigated included: liver disease severity; clinical complications with severities; biomarker trends (using repeated measurements during hospitalization to compute the average, peak and minimum values for each patient) for serum sodium, potassium, creatinine, C-reactive protein, Neutrophil-to-Leukocyte ratio/ NLR and Absolute Neutrophil Counts/ANC). Outcome investigated was in-hospital mortality, SPSS version 25.0 used to analyse data, logistic regression model to determine predictors of in-hospital mortality and significance set at P<0.05.



Abstract IDDF2020-ABS-0151 Figure 1 Flow chart demonstrating selection of sample population

Results Cirrhosis accounted for 16 cases per 1000 hospitalizations, with a male-to-female ratio of 1:2 and a mean age of 53 (±18) years. Most cases were hospitalized for acute decompensation (93.4%) often presenting with ascites (76.5%) and hepatic encephalopathy/HE (41.5%). In-hospital mortality was 35.6%, peaked within the first five days, with factors associated including: HE (OR, 95%CI: 3.0, 1.6–5.6 P= 0.001), high Western-Haven grade (P= 0.009), hyperkalemia (OR, 95%CI: 3.7, 2.7–4.7 P=0.011), elevated creatinine (OR,95%CI: 3.0,2.3–3.8 P=0.004) and elevated absolute neutrophil count (OR,95%CI: 2.6, 1.8–3.5 P=0.016). Following adjustments, Hyperkalaemia (OR= 10, P= 0.003) and elevated absolute neutrophil count (OR= 3.3, P= 0.047) were the independent predictors.

Conclusions In-Hospital mortality is very high in cirrhosis; it depends on a combination of factors and is predicted independently by hyperkalaemia and elevated absolute neutrophil count. Thus physicians should frequently reassess their clinical, inflammatory and metabolic status.

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SWITCHING OFF IMMUNOSUPPRESSIVE MYELOID CELLS BY TARGETING CELL CYCLE-RELATED KINASE PATHWAY: A NEW STRATEGY FOR COMBINATION IMMUNOTHERAPY

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Background The immune-checkpoint-blockade (ICB) therapy has produced promising and yet modest objective response

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