Abstracts

Results She showed complete recovery after rituximab.

Conclusions Rituximab can be considered as an effective treatment modality in APLA related chorea where conventional measures have failed.

64 DYSBIOSIS AND GUT BARRIER DYSFUNCTION IN ANTIPHOSPHOLIPID SYNDROME AS REVEALED BY IGA-SEQ PROFILING

Methods Stool from 15 APS patients, 5 non-autoimmune thrombotic states, and 12 normal donors (total of 17 controls) was collected. Faecal homogenates were analysed for the gut permeability marker calprotectin and, in parallel, stained stained with PE-conjugated anti-human IgA prior to cell sorting. Faecal DNA was isolated and PCR-amplified targeting the V4 region of the 16S rRNA gene. Samples were sequenced on the Illumina MiSeq platform.

Results Faecal calprotectin and IgA-coated faecal bacterial levels were significantly higher in APS patients compared to controls (p<0.003; p<0.05). LEFSe analysis of IgA+ fractions showed that the strongest IgA-coated genus is Blautia in APS.

Conclusions These data suggest gut barrier dysfunction and aberrant IgA coating of commensals in APS. Markedly enhanced bacterial IgA coating in several APS patients supports a stronger adaptive immune response to the microbiota. Increased IgA coating of Blautia might reflect altered gut homeostasis as a Blautia species was shown to be part of proinflammatory IgA+ consortium in a recent study in IBD. To our knowledge, this study represents the first 16S rRNA profiling of IgA-coated gut commensals in patients with non-gut autoimmunity.

66 ANTIPHOSPHOLIPID SYNDROME – CLINICAL AND IMMUNOLOGICAL CORRELATIONS AND Atherosclerosis

Methods The cohort included 160 consecutive APS patients followed by the Autoimmune Diseases Unit of Hospital Clinic Barcelona from 2003 to 2013. Epidemiological, clinical, immunological and treatment features were analysed prospectively.

Results The cohort consisted of 126 (79%) women and 34 (21%) men. Mean (SD) age at diagnosis was 39 (14) years. The diagnostic causes were thrombosis (56.3%), obstetric morbidity (26.9%) and both (16.9%). 65% were primary APS, 22.5% associated with systemic lupus erythematosus (SLE), 8.8% associated with lupus-like syndrome and 3.7% associated with other diseases. Fifty-five patients were lost to follow-up (3.4% every year). In evolution, 10 primary APS patients were reclassified as SLE-associated APS and 2 patients developed catastrophic APS. Table 1 shows the frequencies of the main APS clinical manifestations at baseline and during the 10-year follow-up. At diagnosis, 95% received antithrombotic treatment: low dose antiaggregants (39.5%), oral anticoagulants (67.1%), heparin (2.6%). During the study, 72.7% of recurrences were without antithrombotic treatment and 27.3% were despite it. Eleven major bleeding episodes occurred and 2 were fatal. The global mortality rate was 6.9% and 50% in catastrophic APS. Table 2 shows the main causes of death and Figure 1 is a Kaplan-Meier survival curve.

Conclusions This study shows long-term morbidity and mortality of a large APS patient cohort and exposed the real-life experience of a referral unit.

65 ANTIPHOSPHOLIPID SYNDROME LONG TERM FOLLOW-UP: REAL LIFE EXPERIENCE OF A SINGLE CENTRE

Methods The cohort included 160 consecutive APS patients followed by the Autoimmune Diseases Unit of Hospital Clinic Barcelona from 2003 to 2013. Epidemiological, clinical, immunological and treatment features were analysed prospectively.

Results The cohort consisted of 126 (79%) women and 34 (21%) men. Mean (SD) age at diagnosis was 39 (14) years. The diagnostic causes were thrombosis (56.3%), obstetric morbidity (26.9%) and both (16.9%). 65% were primary APS, 22.5% associated with systemic lupus erythematosus (SLE), 8.8% associated with lupus-like syndrome and 3.7% associated with other diseases. Fifty-five patients were lost to follow-up (3.4% every year). In evolution, 10 primary APS patients were reclassified as SLE-associated APS and 2 patients developed catastrophic APS. Table 1 shows the frequencies of the main APS clinical manifestations at baseline and during the 10-year follow-up. At diagnosis, 95% received antithrombotic treatment: low dose antiaggregants (39.5%), oral anticoagulants (67.1%), heparin (2.6%). During the study, 72.7% of recurrences were without antithrombotic treatment and 27.3% were despite it. Eleven major bleeding episodes occurred and 2 were fatal. The global mortality rate was 6.9% and 50% in catastrophic APS. Table 2 shows the main causes of death and Figure 1 is a Kaplan-Meier survival curve.

Conclusions This study shows long-term morbidity and mortality of a large APS patient cohort and exposed the real-life experience of a referral unit.

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Background and aims The antiphospholipid syndrome (APS) is an autoimmune thrombophilic non-gut disorder with high mortality. Various pathogens have been associated with transplant antiphospholipid antibody production. We hypothesised that members of the gut microbiota in APS patients could represent a chronic trigger and exhibit heightened adaptive immune responses to the microbiota. The purpose of this study was to explore gut barrier function and faecal IgA-coated microbial composition in APS patients.

Methods Stool from 15 APS patients, 5 non-autoimmune thrombotic states, and 12 normal donors (total of 17 controls) was collected. Faecal homogenates were analysed for the gut permeability marker calprotectin and, in parallel, stained stained with PE-conjugated anti-human IgA prior to cell sorting. Faecal DNA was isolated and PCR-amplified targeting the V4 region of the 16S rRNA gene. Samples were sequenced on the Illumina MiSeq platform.

Results Faecal calprotectin and IgA-coated faecal bacterial levels were significantly higher in APS patients compared to controls (p<0.003; p<0.05). LEFSe analysis of IgA+ fractions showed that the strongest IgA-coated genus is Blautia in APS.

Conclusions These data suggest gut barrier dysfunction and aberrant IgA coating of commensals in APS. Markedly enhanced bacterial IgA coating in several APS patients supports a stronger adaptive immune response to the microbiota. Increased IgA coating of Blautia might reflect altered gut homeostasis as a Blautia species was shown to be part of proinflammatory IgA+ consortium in a recent study in IBD. To our knowledge, this study represents the first 16S rRNA profiling of IgA-coated gut commensals in patients with non-gut autoimmunity.