Background and aims Bronchopulmonary dysplasia (BPD) is an important morbidity in premature infants with a multifactorial etiology. In recent years, both genetic and epigenetic mechanisms were suggested to play an important role in BPD development. NO (Nitric Oxide) which is produced along with L-Citrulline by the oxidation of L-Arginine and catalysed by three different isoforms of NOS (NOSynthase), is a short-lived free radical involved in diverse physiological and pathological processes. It consists of 3 types such as neuronal NOS, endothelial NOS and inducible NOS. All the NOS genes are expressed in airway epithelial cells and they are important for physiological functions in the airways. The aim of this study was to investigate possible association between eNOS gene polymorphism and development of BPD in preterm infants.

Methods One hundred and twenty two blood samples DNA isolation was carried out using the PureLink™ Genomic DNA Mini Kit and the concentration of the DNA samples was measured by nanophotometerImplent P 300. For the SNP analysis of eNOS exons37999833 optimised primers were used. Real Time Polymerase Chain Reaction (QRT-PCR) was carried out in a CFX96 thermocycler. Chi-square $\chi^2$ test, Fisher's exact test, the odds ratio and confidence intervals were calculated for the comparisons of allelic and genotype frequencies.

Results Comparison of the allele frequency distribution revealed the presence of G allele as a highly significant risk factor for development of BPD ($p = 0.000^*$; OR 4.07, 95% CI 2.066–8.009) compared to the T allele. The distribution of the T allele in eNOS was found to be similarly distributed amongst BPD (51.9%) and healthy control groups (48.1%). This study demonstrated that the frequency of the GG genotype (25.37%) of the eNOSgene was higher in babies with BPD rather than TT (53.6%) and TG (59.4)genotype, when these genotypes were compared with the healthy control groups. No healthy infants were seen to carry the GG genotype ($p = 0.000^*$; OR 1.89, 95% CI 1.514–2.148). The TT genotype ($p = 0.019^*$; OR 0.39, 95% CI 0.180–0.870) also displayed a susceptibility for developing BPD. Heterozygous TG genotype ($p = 0.631$; OR 0.63, 95% CI 0.527–2.873) was not associated with the development of BPD.

Conclusion To our best of knowledge, noninvestigation of the eNOS gene polymorphism has previously been documented in BPD. Thefindings of this study demonstrated that the GG genotype in eNOS gene was highly significant for BPD.

Background Positive anti-tTG levels have been associated with reduced weight and bone mineral density (BMD) in clinical celiac disease (CD). However, effects in subclinical CD are not known.

Objective To assess associations between anti-tTG levels, growth and BMD in children with subclinical CD.

Methods In a population-based prospective cohort study, serum samples were analysed for anti-tTG levels at 6 years of age. (n = 4,442) Children were categorised into 2 groups: negative (< 7 U/ml, n = 4,249), or positive anti-tTG (>7 U/ml, n = 57). Positive tTG levels were further categorised into 2 categories based on the > 10 times upper limit of normal (ULN) levels (70 U/ml). Height, weight and BMI z-scores were obtained using Dutch reference growth charts. BMD was measured by Dual-energy X-ray absorptiometry (DEXA). Multivariable linear regression and linear mixed models were performed.

Results Delayed growth in weight SDS/year (-0.05; 95% CI -0.09,-0.01) and BMI SDS/year (-0.10; 95% CI -0.18,-0.01) from 6 months until 6 years was observed in children with positive anti-tTG levels. Height growth tended to be delayed over time (-0.02 SDS/year; 95% CI -0.06, 0.02). A lower height (-0.29; 95% CI -0.55,-0.04), weight (-0.38; 95% CI -0.64,-0.12), BMI (-0.26 95% CI -0.49,-0.03) and BMD (-0.26; 95% CI -0.45,-0.08) at 6 years of age was found in children with positive anti-tTG levels.

Conclusion Positive anti-tTG levels in children without gastrointestinal symptoms at 6 years of age were associated with reduced BMD and delayed growth trajectories until 6 years. This suggests that subclinical CD has consequences for BMD and normal growth.

Background Necrotizing enterocolitis (NEC) may further increase the susceptibility to neurological damage, possibly via gut-derived inflammatory signals. To investigate this, we test if NEC severity and intestinal permeability in formula-fed preterm pigs is associated with histopathology, microglial activation, and increased proinflammatory cytokine levels in the hippocampus.

Methods Forty-four preterm piglets were fed increasing doses of formula and euthanized on day five. Macroscopic NEC lesions were scored in five regions of the gut (stomach, proximal, middle, and distal small intestine, colon). Intestinal permeability was assessed by urinary lactulose-mannitol-ratio. Hippocampal IL-1β and IL-8 levels were determined by ELISA. Histopathology, neurodegeneration, and microglia were investigated by analyses of hematoxylin-cosin, Fluoro-jade B (FJB), and Iba-1 stained coronal sections.

Results Proximal, middle, and distal small intestinal NEC score, and intestinal permeability correlated positively with IL-8 levels (all p < 0.05) but not with IL-1β. In preterm piglets with severe NEC lesions, numerous shrunken, hyperchromatic neurons were observed. Neurodegeneration was confirmed by positive FJB staining. Iba-1 positive cells with a morphology resembling...