

EPIGENETIC PROGRAMMING OF ASTHMA BY IN UTERO EXPOSURE TO p,p'-DDE

Eva Morales, Center for Research in Environmental Epidemiology (CREAL), Barcelona, Catalonia, Spain

Nadia Vilahur, Center for Research in Environmental Epidemiology (CREAL), Barcelona, Catalonia, Spain

Mariona Bustamante, Center for Research in Environmental Epidemiology (CREAL), Barcelona, Catalonia, Spain

Georgia Escaramis, Center for Genomic Regulation (CRG), Barcelona, Catalonia, Spain

Magda Montfort, Center for Genomic Regulation (CRG), Barcelona, Catalonia, Spain

Maties Torrent, Area de Salut de Menorca, IB-SALUT, Menorca, Spain

Xavier Estivill, Center for Genomic Regulation (CRG), Barcelona, Catalonia, Spain

Jordi Sunyer, Center for Research in Environmental Epidemiology (CREAL), Barcelona, Catalonia, Spain

Background and Aims: Environmental epigenetic is emerging in asthma. Early-life exposure to dichlorodiphenyldichloroethylene (p,p'-DDE) impacts the immune system development and the risk of asthma by unknown mechanisms. We investigated if prenatal exposure to p,p'-DDE induces epigenetic reprogramming involving aberrant DNA methylation of specific genes mechanistically related to asthma.

Methods: Data came from 404 children of a population-based birth cohort of the INMA Project. p,p'-DDE was measured in cord blood. Wheezing phenotype at age 6 years was defined according to parental report of wheezing episodes on each interviewer-led annual questionnaire. The coding variant Ile105Val from *GSTP1* was genotyped. In a subset (n=128), gene-specific methylation was screened along 1505 genomic CpG loci using Illumina GoldenGate panel in DNA from peripheral blood cells collected at age 4 years. Subsequently, a pyrosequencing assay was used to validate the results from the Illumina screening.

Results: Higher levels of p,p'-DDE were associated with higher risk of persistent wheeze among *GSTP1* Val carriers (OR for an interquartile range increase in p,p'-DDE=1.77; 95% CI 1.17-2.68), but not among the Ile homozygotes (OR=1.05; 95% CI 0.80-1.39; p for interaction=0.069). Higher levels of p,p'-DDE were associated with a decrease in methylation of a CpG site in arachidonate 12-lipoxygenase (*ALOX-12*) (beta(sd) for an interquartile range increase in p,p'-DDE =-1.7(0.6), p=0.060). The decrease in methylation was accounted for *GSTP1* Val carriers compared with the Ile homozygotes (beta(sd)=-4.1(0.8) vs. -0.8(0.8); p for interaction=0.007). Higher levels of methylation in *ALOX-12* at age 4 years were associated with a lower probability of being classified as persistent wheezing at age 6 years (OR=0.94; 95% CI 0.90-0.98).

Conclusions: Epigenetic changes in *ALOX-12*, a gene involved in the immune response of the airway, could mediate the lifelong effect of in utero exposure to p,p'-DDE on the risk of asthma in childhood.