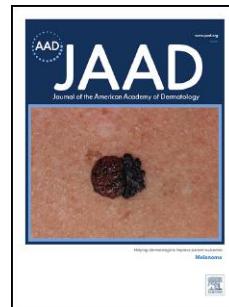


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Effect of prenatal high-dose vitamin D on childhood atopic dermatitis is modified by maternal cotinine metabolome: A secondary analysis of a randomized clinical trial

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1 **Title**

2 Effect of prenatal high-dose vitamin D on childhood atopic dermatitis is modified by maternal
 3 cotinine metabolome: A secondary analysis of a randomized clinical trial

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41 Data Protection Agency (2015-41-3696). Both parents gave written informed consent before
42 enrollment.

43 **Data sharing:** Data will be available upon request.

44 **CAPSULE SUMMARY**

45 Environmental factors are speculated to influence the risk of childhood atopic dermatitis. We found
46 a significant effect modification of the maternal tobacco exposure blood metabolome on the effect
47 of prenatal high-dose vitamin D on offspring risk of atopic dermatitis

48 The findings suggest a beneficial effect of prenatal high-dose vitamin D on atopic dermatitis until
49 school-age among mothers with a high tobacco exposure blood metabolome score suggesting a
50 possible targeted prevention strategy against smoking mothers.

51

52 **KEYWORDS**

53 Atopic dermatitis; vitamin D; primary prevention; environment; tobacco exposure; metabolomics

54

55 **ABSTRACT**

56 **Background:** Tobacco exposure has been shown to modulate the effect of vitamin D on the risk of
 57 atopic diseases. However, randomized clinical trials investigating potential effect modification
 58 between tobacco exposure and vitamin D supplementation on atopic disease risk are lacking.

59 **Objective:** We sought to investigate the potential effect modification from maternal tobacco
 60 exposure on the effect of prenatal high-dose vitamin D supplementation on risk of child atopic
 61 dermatitis (AD), asthma and allergic rhinitis (AR).

62 **Methods:** A post hoc analysis in the double-blinded COPSAC₂₀₁₀ randomized clinical trial (RCT)
 63 (NCT00856947) including 581 mother-child pairs randomized to 2800 IU/day (high-dose) vs 400
 64 IU/d (standard-dose) from pregnancy week 24. Maternal blood metabolomic profiling was
 65 performed at inclusion reflecting maternal tobacco exposure using a supervised sparse partial least
 66 square model.

67 **Results:** We found a significant effect modification from the maternal cotinine metabolome score
 68 ($p_{interaction}<0.01$) where high-dose vitamin D reduced AD until age 6 years in offspring from mothers
 69 with highest (4th quartile) cotinine metabolome score: crude; HR=0.46 (0.23-0.93), $p=0.03$, and
 70 adjusted for sex, birth season, socioeconomic circumstances, living environment, air pollution,
 71 maternal diet, vitamin D levels and fish oil intervention; HR=0.36 (0.15-0.85), $p=0.02$. Similarly,
 72 significant effect modification was demonstrated on risk of asthma ($p_{interaction}=0.03$) until age 6 years
 73 but not AR ($p_{interaction}=0.08$) at age 6 years.

74 **Conclusions:** This exploratory study of prespecified RCT outcomes demonstrates effect
 75 modifications of the maternal tobacco exposure metabolome on the primary preventive effect of
 76 prenatal high-dose vitamin D on offspring atopic disease risk suggesting a potential personalized
 77 prevention strategy targeting mothers exposed to tobacco smoking.

78

79 **INTRODUCTION**

80 Atopic dermatitis (AD) is one of the most common diseases in childhood affecting 13-20% of the
81 pediatric population with a continuous increase in prevalence^{1,2,3}. AD is a chronic inflammatory
82 skin disease characterized by itching, skin lesions and periods of exacerbations and remission and is
83 associated with other atopic diseases including asthma and allergy^{4,5}. However, there is still no cure
84 for the disease, which highlights the need for early preventive strategies⁶.

85 The pathophysiology of AD is complex involving genetic predisposition, immune dysregulation
86 and environmental exposures such as allergens and diet^{3,7}. Maternal smoking during pregnancy and
87 around birth has been suggested as a strong risk factor for offspring risk of AD as well as for
88 childhood asthma and allergy^{8,9}. In contrast, vitamin D is hypothesized to hold a protective role
89 against AD development and disease severity as well as development of asthma/wheeze due to the
90 effects on mechanisms involved in disease progression^{10,11}. Studies have proposed an effect of
91 vitamin D on AD^{10,11,12}. Our previous findings have shown protective effects on asthma/wheeze but
92 interestingly not AD for the whole population^{13,14}. Further, a study from the Vitamin D Antenatal
93 Asthma Reduction Trial (VDAART) demonstrated interaction between maternal tobacco exposure
94 and prenatal vitamin D status on offspring atopic diseases¹⁵. We therefore hypothesize that maternal
95 tobacco exposure could modify the effect of prenatal vitamin D intervention on risk of atopic
96 diseases.

97 Because of the stigma around smoking, pregnant women are not being honest when interviewed¹⁶.
98 Therefore, we measured blood levels of cotinine as well. We previously showed that environmental
99 exposures could have effects on the blood metabolome differing from individual to individual¹⁷,
100 hence, we utilize individual blood metabolome profiles based on cotinine levels to capture systemic

101 effects of tobacco exposure. Here, we investigate whether maternal tobacco exposure and the
102 related blood metabolome profile during pregnancy could modify the effects of prenatal vitamin D
103 supplementation on risk of AD, asthma and allergic rhinitis in a double-blinded randomized clinical
104 trial (RCT).

105 **METHODS**106 The COPSAC₂₀₁₀ vitamin D RCT

107 The COPSAC₂₀₁₀ cohort has previously been described in detail including baseline characteristics
108 and successful randomization of the pregnancy double-blinded RCT^{18,19,20,21}. The vitamin D
109 intervention consisted of 2800 IU/day of vitamin D compared with 400 IU/day according to the
110 Danish recommended intake. 623 pregnant women in pregnancy week 24 were randomized. The
111 children were followed longitudinally from birth until age 6 years with 12 scheduled visits
112 including acute care visits of symptoms of asthma, allergy or eczema^{19,21}.

113 Atopic dermatitis, asthma and allergic rhinitis

114 The predefined outcomes of AD was diagnosed prospectively according to the criteria of Hanifin
115 and Rajka including typical morphology and localization of skin lesions based upon clinical
116 examination in the COPSAC clinic as previously described²². Asthma and allergic rhinitis by age 6
117 years were diagnosed in the COPSAC clinic according to previously validated algorithms¹⁹.

118 Maternal blood metabolomics

119 Maternal blood samples were collected at inclusion in pregnancy for untargeted metabolomic
120 profiling of the mothers. Profiling was performed by Metabolon (NC, USA) using HD4 liquid
121 chromatography-mass spectrometry (LC-MS)/MS platform as previously described including our
122 protocol for pre-processing of data^{23,24}. See our previous publication for more details²⁶.

123 Maternal tobacco exposure

124 Prenatal tobacco exposure was defined as maternal smoking during the first trimester of pregnancy
125 (yes/no) i.e. before the intervention and was determined by parental interviews at enrolment.

126 Cotinine levels were measured as part of the metabolomic profiling by Metabolon at pregnancy
127 week 24 and were categorized into two groups (detectable vs not detectable levels).

128 Maternal systemic low-grade inflammation

129 Using the same blood samples as for metabolomics measurements high-sensitivity C-reactive
130 protein (hs-CRP) levels were determined by a high-sensitivity electrochemiluminescence assay
131 from MesoScale²⁵.

132 Statistical analysis

133 We created a maternal metabolome score reflecting prenatal tobacco exposure based on detectable
134 blood cotinine levels (yes/no) using a supervised sparse partial least square (sPLS) model. This was
135 achieved by selecting the optimal set of metabolites through 10-fold cross-validation repeated 10
136 times based on the Area Under the Curve (AUC) as previously detailed²⁶. The final sPLS model
137 was chosen for its highest median AUC value and lowest variance across cross-validation rounds. A
138 predicted cotinine metabolome score was calculated for each individual by multiplying the original
139 data matrix by the loadings derived from the final model. Thereafter, we analyzed potential effect
140 modification from this maternal cotinine metabolome score on the effect of prenatal high-dose
141 vitamin D on the risk of AD and asthma in children until age 6 years by using Cox proportional
142 hazard regression models and on the risk of allergic rhinitis at age 6 years using logistic regression.

143 We then performed stratified analyses of the effect of high-dose vitamin D dividing the cotinine
144 metabolome score into quartiles analyzing the effect on risk of AD, asthma and allergic rhinitis in
145 each group. In sensitivity analyses, we adjusted for socio-economic circumstances (household
146 income, maternal age and maternal level of education), sex, maternal 25-hydroxyvitamin D levels,
147 birth season, living environment (urban vs rural)¹⁷, air pollution (fine particulate matter <2.5 µm),
148 maternal diet based on a food frequency questionnaire at pregnancy week 24²⁷ and the embedded

149 prenatal fish-oil trial^{19,28}. Finally, we investigated the relationship between the cotinine metabolome
150 score and cotinine levels vs systemic low-grade inflammation (hs-CRP). An enrichment analysis
151 was performed for top 50 metabolites contributing to the sPLS derived cotinine-based maternal
152 metabolome model. We applied a false discovery rate (FDR) of 5% to account for multiple testing.
153 All analyses were performed with R (v4.0.3). P-values less than 0.05 were considered statistically
154 significant.

155 **RESULTS**

156 Of the 581 children, 572 (98%) had information on maternal metabolomics at pregnancy week 24
 157 including measurements of cotinine and AD during the first 6 years of life and were included in the
 158 time to first event analyses. At age 6 years, 553 (95%) children completed the clinical visit. There
 159 were no differences in baseline characteristics between the mother-child pairs in each intervention
 160 group¹⁹.

161 **Maternal tobacco exposure and risk of AD, asthma and allergic rhinitis age 0-6 years**

162 Of the 572 mothers with cotinine levels measured, 59 (10%) had a detectable cotinine level. There
 163 was a significant relationship between mothers answering “yes” to being a smoker in the first
 164 trimester (n=45, 8%) and their cotinine levels at week 24.

165 There was no association between answering “yes” to being a smoker or having a detectable
 166 cotinine level and risk of AD, asthma or allergic rhinitis age 0-6 years ($p>0.05$) and no significant
 167 interaction with the vitamin D intervention for AD ($p_{interaction}=0.15$ and $p_{interaction}=0.47$), asthma
 168 ($p_{interaction}=0.64$ and $p_{interaction}=0.86$) or allergic rhinitis ($p_{interaction}=0.99$ and $p_{interaction}=0.57$).

169 Given the trend of effect modification on vitamin D intervention on AD, we analyzed the effect of
 170 vitamin D supplementation on risk of AD among mothers having a detectable cotinine level at week
 171 24: HR; 0.70 (0.28-1.78), $p=0.46$ and mothers not having a detectable cotinine level: 1.00 (0.73-
 172 1.38), $p=0.99$.

173 **Maternal cotinine blood metabolome score**

174 The cotinine-based maternal metabolome score at week 24 showed a cross-validated median repeat
 175 AUC of 0.72 (range of 0.71-0.73) and was strongly associated with cotinine levels (**Figure 1A**).
 176 The top 50 metabolites contributing to the score with the highest loadings are displayed in **Figure**

177 **1B**, which showed highest positive loadings from xenobiotics and highest negative loadings from
 178 co-factors, vitamins and lipids including beta-cryptoxanthin and hydroxy-CMPF. The metabolite
 179 pathways are illustrated in the Sankey plot and enrichment analysis in **Figure 2** illustrating
 180 metabolite pathways, which passed the 5% FDR multiple test correction.

181 There was no direct association between the maternal cotinine metabolome score reflecting tobacco
 182 exposure and risk of AD, asthma or allergic rhinitis.

183 **Maternal cotinine metabolome score and effect of prenatal high-dose vitamin D on risk of AD,
 184 asthma and allergic rhinitis age 0-6 years**

185 We investigated potential effect modification of the maternal cotinine metabolome score reflecting
 186 tobacco exposure on the effect of prenatal high-dose vitamin D supplementation on risk of AD age
 187 0-6 years and found a significant interaction between these ($p_{interaction}<0.01$). A similar interaction
 188 was observed after adjustment for all covariates described²⁸. Then, we investigated the effect of
 189 high-dose vitamin D on risk of AD age 0-6 years (age at onset) stratified by quartiles of the
 190 maternal cotinine metabolome score and found a significant beneficial effect of the supplement
 191 among children whose mothers had the highest (4th quartile) score (n=143): HR; 0.46 (0.23-0.93),
 192 $p=0.03$ (**Figure 3**). The results were even stronger after covariate adjustments showing a larger
 193 reduced risk in the 4th quartile: 0.36 (0.15-0.85), $p=0.02$. This finding was further supported by an
 194 effect of vitamin D supplementation on risk of having an AD diagnosis (yes/no) during the first 6
 195 years of life in the same group of children, i.e., born to mothers with the highest metabolome score:
 196 OR; 0.43 (0.19-0.95), $p=0.04$ (**Figure 4**).

197 Thereafter, we similarly investigated effect modification of the metabolome score on the effect of
 198 prenatal high-dose vitamin D on asthma risk age 0-6 years using Cox regression ($p_{interaction}=0.03$)

199 and allergic rhinitis at age 6 years using logistic regression ($p_{interaction}=0.08$). We found no
200 significant effects on asthma or allergic rhinitis in the stratified analyses ($p>0.05$).

201 **Maternal cotinine metabolome score and systemic low-grade inflammation**

202 Finally, we investigated the relationship between the maternal metabolome score based on cotinine
203 levels vs hs-CRP levels (pg/mL) at the same timepoint at week 24 in pregnancy and found a
204 significant positive association: beta-estimate; 0.49 (0.32-0.66), $p=1.8e-10$. In contrast, there was no
205 association between maternal cotinine levels (detectable vs not detectable) and hs-CRP levels at
206 week 24; 0.04 (-0.25-0.32), $p=0.79$.

207 **DISCUSSION**

208 In this study, we demonstrated significant effect modification by the maternal blood cotinine
 209 metabolome profile on the effect of prenatal high-dose vitamin D vs placebo on time to AD and
 210 asthma diagnosis during the offsprings' first 6 years of life. We found a beneficial effect of the
 211 supplement in children of mothers who had the highest blood metabolome score reflecting tobacco
 212 exposure. Finally, we found a significant effect modification from the maternal cotinine
 213 metabolome score on risk of asthma.

214 **Strengths and limitations**

215 This study is strengthened by the RCT study design with successful randomization procedure¹⁹
 216 mitigating the risk of residual confounding²⁹. Further, the COPSAC₂₀₁₀ cohort is well known for its
 217 high follow-up rate, and we here report an unparalleled 95% clinical follow-up at age 6 years with
 218 12 scheduled clinical visits from birth throughout early childhood, which allows for thorough
 219 registration of symptoms and close monitoring of AD, asthma and allergic rhinitis. COPSAC
 220 physicians were responsible for the clinical diagnostic procedure at the COPSAC research unit
 221 ensuring diagnostic accuracy and consistency. Finally, the population is representing the Danish
 222 background population by including an unselected group of pregnant women allowing for
 223 generalization of our findings.

224 The main limitation is the relatively few mothers reporting being actively smoking or exposed to
 225 smoking with a detectable cotinine level in the blood, which reduced the statistical power for the
 226 effect modification analyses. However, we used the information from cotinine blood measurements
 227 to create a cotinine-based metabolome score reflecting tobacco exposure for each pregnant women,
 228 which was used for the effect modification analysis and which we hypothesized was a more precise
 229 measure of how tobacco exposure affected the individual mothers.

230 **Interpretation**

231 Here, we found a protective effect of vitamin D supplementation among a specific group of children
232 whose mothers metabolome reflected a high tobacco exposure during pregnancy compared with the
233 null finding we and others have previously reported^{19,14,30}. The reported effect modification and
234 significant effect of the high-dose vitamin D intervention on AD in the children born to mothers
235 whose metabolome were most similar to cotinine reflecting maternal tobacco exposure, suggests a
236 possible targeted intervention strategy that should be confirmed in trials among a high-risk group of
237 smoking mothers, since we were limited by a low prevalence of 8% of mothers smoking compared
238 to other populations reporting between 14-40% dependent on socio-economic group¹⁶. This has
239 previously been done with prenatal vitamin C supplementation among a population of pregnant
240 smokers (n=251) showing reduction in asthma symptoms with supplementation³³ and a similar
241 sample size may be required to demonstrate effects of prenatal vitamin D supplementation as well,
242 which is more than 5 times larger than our sample size of pregnant smokers. However, we
243 demonstrated effect modification from the cotinine metabolome score reflecting environmental
244 tobacco exposure in the whole metabolome and interestingly, the results were unchanged after
245 adjustment for several socioeconomic and environmental exposures.

246 By using mechanistic data layers, it allows for a deeper understand of the interplay between prenatal
247 tobacco exposure and high-dose vitamin D intervention. The effect modification was not observed
248 for maternal reported tobacco smoking or blood cotinine levels underscoring the utility of blood
249 metabolomics profiling to capture individual adverse effects of environmental tobacco exposure.
250 We have previously used this approach with success to understand the effect of living environment
251 on common infection risk¹⁷ and the role of infant gut virome on preschool asthma³⁴. The added
252 strength of utilizing this metabolome score compared with smoking habits from interviews or a
253 single cotinine measure alone is that we are hereby able to characterize the whole metabolomic

254 profile reflecting the individual adverse effects of environmental tobacco exposure, which can then
255 be applied on all individuals in the study population. This is even more important when being
256 reduced in sample size due to a low number of smoking mothers as our initial analyses indicated an
257 effect of vitamin D among mothers with a detectable cotinine level although not statistically
258 significant. Interestingly, our cotinine metabolome score was also highly associated with systemic
259 low-grade inflammation, which was not the case when analyzing cotinine as a single metabolite
260 measure, which came as a surprise given that a relationship between smoking and increased
261 inflammation previously has been reported^{35,36}. It is likely that other environmental factors, which
262 we were not able to identify with our data, are also reflected in the cotinine metabolome score as
263 this is driven by several metabolites from different pathways as demonstrated in our enrichment
264 analysis. However, it does not seem to reflect maternal diet, socioeconomic status, living
265 environment or air pollution.

266 The mechanisms behind the reported effect modification in this trial is unclear and can only be
267 speculated. It may be related to the immunomodulatory, anti-inflammatory and antioxidant effects
268 of vitamin D³⁷ since tobacco exposure is known for increasing inflammation and produce oxidative
269 stress³⁸. In support of this theory, the metabolite that contributed to the model with the highest
270 negative loading was the antioxidant beta-cryptoxanthin suggesting an inverse relationship between
271 this antioxidant and tobacco exposure. In addition, we reported a positive association between the
272 maternal cotinine metabolome score and low-grade inflammation, but we did not find levels of hs-
273 CRP to associate with AD, asthma or allergic rhinitis risk in the offspring. Others have
274 demonstrated that tobacco exposure decreases the conversion of inactive to active vitamin D in
275 human epithelial cells suggesting a reduction in activity of the vitamin from smoking³⁹.
276 In conclusion, this study of prespecified atopic outcomes of the high-dose vitamin D RCT in
277 COPSAC₂₀₁₀ demonstrates effect modification of a maternal tobacco exposure blood metabolome

278 profile on the effect of prenatal vitamin D supplementation on AD risk that may hold potential for a
279 personalized prevention strategy.

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395 **FIGURE LEGENDS**

396 **FIGURE 1**

397 Sparse partial least squares (sPLS) model predicting cotinine exposure during pregnancy from week 24 metabolomic profiles. A) Overview
398 of top 50 metabolites with corresponding loadings. B) The cross-validated prediction with AUC and p-value, which were strongly
399 associated with detectable cotinine levels.

400 **FIGURE 2**

401 Sankey plot and enrichment analysis of top 50 metabolites contributing to the sPLS derived cotinine-based maternal metabolome model.
402 fdr=false discovery rate.

403 **FIGURE 3**

404 Effect of high-dose vitamin D vs placebo on risk of atopic dermatitis (AD) until age 6 years in a time to first event (age at onset) analysis.
405 Estimates, 95% CI and p-values from a Cox proportional hazard regression model.

406 **FIGURE 4**

407 Effect of high-dose vitamin D vs placebo on risk of atopic dermatitis (AD) between age 0-6 years (yes/no) in each quartile of the maternal
408 cotinine metabolomics score “fingerprint”. Estimates from a logistic regression model.

