

Assessment of urinary oxidative stress biomarkers associated with fine particulate matter (PM2.5) exposure in Chiang Mai, Thailand (#106470)

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Assessment of urinary oxidative stress biomarkers associated with fine particulate matter (PM2.5) exposure in Chiang Mai, Thailand

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Background: Exposure to fine particulate matter (PM2.5) is known to increase oxidative stress, impacting health adversely. This study examines the relationship between PM2.5 exposure and oxidative stress biomarkers in Chiang Mai, Thailand.

Methods: A pilot prospective observational study was conducted in Samoeng District, Chiang Mai, including 25 healthy participants (age 25-60 years). Urine samples were collected during high (Feb-April 2023) and low (May-July 2023) PM2.5 seasons. PM2.5 concentrations were monitored daily from the NTAQHI website. Biomarkers analyzed included 1-hydroxypyrene (1-OHP) using HPLC, malondialdehyde (MDA) via Spectrophotometry, and 8-epi-prostaglandin F2 α (8-epi-PGF2 α) with ELISA. Statistical analysis was performed using IBM SPSS Statistics 22.0.

Results: Significant increases in urinary 1-OHP, MDA, and 8-epi-PGF2 α were observed during the high PM2.5 season compared to the low season. The mean concentration of PM2.5 was 67 $\mu\text{g}/\text{m}^3$ during high pollution and 7 $\mu\text{g}/\text{m}^3$ during low pollution. Elevated levels of these biomarkers indicate increased oxidative stress associated with higher PM2.5 exposure.

Conclusions: This study highlights a significant association between elevated PM2.5 levels and increased oxidative stress biomarkers in Chiang Mai, Thailand. The findings suggest that exposure to higher concentrations of PM2.5 contributes to oxidative stress, potentially leading to adverse health outcomes.

1 **Assessment of urinary oxidative stress biomarkers associated with fine particulate matter**
2 **(PM2.5) exposure in Chiang Mai, Thailand**

3

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25 Abstract

26 **Background:** Exposure to fine particulate matter (PM_{2.5}) is known to increase oxidative stress,
27 impacting health adversely. This study examines the relationship between PM_{2.5} exposure and
28 oxidative stress biomarkers in Chiang Mai, Thailand.

29 **Methods:** A pilot prospective observational study was conducted in Samoeng District, Chiang
30 Mai, including 25 healthy participants (age 25-60 years). Urine samples were collected during
31 high (Feb-April 2023) and low (May-July 2023) PM_{2.5} seasons. PM_{2.5} concentrations were
32 monitored daily from the NTAQHI website. Biomarkers analyzed included 1-hydroxypyrene (1-
33 OHP) using HPLC, malondialdehyde (MDA) via Spectrophotometry, and 8-epi-prostaglandin
34 F₂α (8-epi-PGF₂α) with ELISA. Statistical analysis was performed using IBM SPSS Statistics
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36 **Results:** Significant increases in urinary 1-OHP, MDA, and 8-epi-PGF₂α were observed during
37 the high PM_{2.5} season compared to the low season. The mean concentration of PM_{2.5} was 67
38 µg/m³ during high pollution and 7 µg/m³ during low pollution. Elevated levels of these
39 biomarkers indicate increased oxidative stress associated with higher PM_{2.5} exposure.

40 **Conclusions:** This study highlights a significant association between elevated PM_{2.5} levels and
41 increased oxidative stress biomarkers in Chiang Mai, Thailand. The findings suggest that
42 exposure to higher concentrations of PM_{2.5} contributes to oxidative stress, potentially leading to
43 adverse health outcomes.

44

45 Introduction

46 Air pollution, resulting from both human activities and natural sources such as vehicle and
47 industrial emissions, agricultural residue burning, biomass burning, and forest fires, has
48 significant transboundary effects, especially in regions like Southeast Asia (Thongtip, Srivichai
49 et al. 2022). Globally, ambient air pollution is a critical issue, contributing to climate change and
50 public health crises (Altuwayjiri, Taghvaei et al. 2021), and accounts for an estimated 7 million
51 premature deaths every year [WHO,2021]. In developing countries, rapid urbanization and

52 industrialization have exacerbated this problem, leading to increased exposure to pollutants like
53 PM2.5, which poses severe health risks (Sukkhum, Lim et al. 2022).

54 PM2.5, fine particulate matter or particulate matter with a diameter of less than 2.5 micrometers,
55 is particularly concerning due to its ability to penetrate deep into the respiratory system and enter
56 the bloodstream (Amnuaylojaroen and Parasin 2023). Its small size and large surface area enable
57 PM2.5 to carry various toxic substances, including polycyclic aromatic hydrocarbons (PAHs),
58 which are known to induce oxidative stress and inflammation. Epidemiological and toxicological
59 studies have linked PM2.5 exposure to a range of health issues, including cardiovascular
60 diseases, respiratory problems, and various forms of cancer (Bhatnagar 2022). Additionally,
61 oxidative stress and DNA damage are critical mechanisms through which PM2.5 exerts its
62 harmful effects, facilitated by reactive oxygen species (ROS) production (Liu, Jiang et al. 2024).

63

64 **Figure 1 Mechanism of oxidative stress induced by PM2.5 exposure, showing the imbalance**
65 **between ROS and antioxidants leading to oxidative damage.**

66

67 The mechanism of oxidative stress illustrated in Figure 1 induced by PM2.5 and PAHs is a
68 critical pathway in understanding the health impacts of air pollution. As illustrated in Figure 1,
69 the small size of PM2.5 allows it to penetrate deep into the respiratory system and enter the
70 bloodstream, where it can transport toxic substances such as PAHs. These substances can
71 generate ROS through processes like the redox cycle, leading to oxidative stress, which in turn
72 induces DNA and lipid damage. This cascade of events is linked to various adverse health
73 outcomes, including inflammation, cardiovascular diseases, and cancer (Li, Xia et al. 2008,
74 Møller and Loft 2010). Understanding this mechanism is vital for identifying potential
75 biomarkers, such as 1-hydroxypyrene (1-OHP), malondialdehyde (MDA), and 8-
76 isoprostaglandin F2 alpha (8-iso-PGF2 α), which can be used to monitor exposure and assess the
77 biological impact of PM2.5 on human health.

78 In Thailand, transportation, industries, and biomass burning are the main sources of emissions
79 (Sirithian and Thanatrakolsri 2022). PM2.5, PM10, ozone (O₃), and volatile organic
80 compounds (VOCs) are examples of air pollutants exceeding national ambient air quality

81 standards. Although air quality improved in 2020, levels remained above standard for over 70
82 days, primarily in northern provinces. Residents suffer respiratory issues due to haze pollution
83 during the dry season due to open burning and natural forest fires. Northern Thailand's
84 mountainous cities, surrounded by farming, generate a lot of air pollution (Amnuaylojaroen,
85 Parasin et al. 2022).

86 PAHs have been shown to **create** ROS through the redox cycle, which can induce oxidative
87 modification of DNA and lipids in vivo (Fujitani, Furuyama et al. 2023). PAHs are produced by
88 biogenesis and human activities, such as incomplete combustion of fossil fuels, industrial
89 processes, and biomass burning (Tala, Kraisitnitikul et al. 2023). Urinary 1-OHP, a metabolite of
90 pyrene, is a useful biomarker for PAH exposure (Kho, Lee et al. 2015). In several studies, 1-
91 OHP levels were utilized to biologically monitor PAH uptake from **occupational exposures**
92 (Brucker, Moro et al. 2013). The study provides an overview of 1-OHP's association with PAH
93 exposure.

94 Biomarkers like MDA and 8-iso-PGF 2α have been widely used in studies assessing oxidative
95 stress related to air pollution exposure (Zhao, Gong et al. 2018). MDA is a well-known
96 biomarker for oxidative stress, and its elevated levels have been linked to exposure to pollutants
97 like PM $_{2.5}$ and O $_3$. Studies have demonstrated increased levels of MDA in the ear and brain
98 following exposure to wood smoke, as well as in asthmatic children exposed to PM $_{2.5}$ and O $_3$
99 (Toto, Wild et al. 2022). Urinary MDA measurements have also been utilized to evaluate air
100 pollution exposure, primarily focusing on subjects with respiratory diseases, children, or the
101 elderly. Despite its potential, the application of MDA as a biomarker in epidemiological studies
102 concerning air pollution remains relatively limited.

103 On the other hand, 8-iso-PGF 2α , part of the F 2 -isoprostane family, is a nonenzymatic product of
104 arachidonic acid peroxidation and is recognized as a sensitive biomarker for oxidative status
105 (Il'yasova, Scarbrough et al. 2012). Elevated urinary levels of specific isomers, such as iPF 2α -III
106 and iPF 2α -VI, have been found in individuals with various health conditions, including
107 cardiovascular disease, Alzheimer's disease, type 2 diabetes, Down syndrome, lung disorders,
108 and heavy smokers (Zhang, Il'yasova et al. 2010). The metabolite 2,3-dinor-8-iso-prostaglandin
109 F 2α is particularly useful as a biomarker for the formation of 8-iso-PGF 2α and lipid peroxidation

110 in vivo, providing insights into the oxidative damage occurring within the body due to
111 environmental and other stressors.

112 Chiang Mai, particularly the Samoeng district, experiences significant air pollution issues,
113 primarily during the dry season due to agricultural residue burning and forest fires (Jainonthee,
114 Wang et al. 2022). The region's topography, characterized by mountains and valleys, exacerbates
115 air quality problems, trapping pollutants and leading to extended periods of poor air quality
116 (Supasri, Gheewala et al. 2023). The local population frequently experiences health issues such
117 as respiratory diseases and ocular surface diseases, associated with high levels of PM_{2.5} and
118 other pollutants. Despite these challenges, there is limited data on the health impacts of air
119 pollution in this region, particularly concerning oxidative stress biomarkers.

120 Given the significant impact of PM_{2.5} on public health and the limited data available on
121 oxidative stress biomarkers associated with air pollution in Chiang Mai, this study aims to
122 investigate the relationship between PM_{2.5} exposure and oxidative stress biomarkers in the
123 region. The primary objectives are to assess urinary levels of 1-OHP, MDA, and 8-iso-PGF_{2α},
124 evaluate their association with PM_{2.5} exposure, and explore the implications for public health in
125 the Samoeng district. This research seeks to provide valuable insights into the biological effects
126 of air pollution, contributing to developing effective strategies for mitigating its adverse health
127 impacts in northern Thailand.

128

129 **Materials and Methods**

130 **Study Area**

131 The focus area was chosen to be Chiang Mai province (Jarernwong, Gheewala et al. 2023). With
132 a total area of 20,107 km², it is the largest province in the north of Thailand. The geographical
133 location of the Chiang Mai province is shown in Figure 2. It has 25 districts, with a total
134 population of roughly 1.78 million based on the Department of Provincial Administration of
135 Thailand in 2020.

136 A unique area of Chiang Mai province, Samoeng District, was selected for sample collection
137 because of its unique environment, including higher levels of particulate matter PM_{2.5} during

138 the burning season, as well as its rural and agricultural setting, making it a perfect location to
139 study health impacts of pollution (Paesrivarotai and Tanaksaranond 2021).

140

141 **Figure 2 Map showing the study area in Chiang Mai, Thailand.**

142

143 **Study Population**

144 This is a pilot, prospective observational study including 25 healthy participants from the
145 Samoeng district of Chiang Mai province in Thailand. Participants were selected based on
146 predefined inclusion and exclusion criteria to ensure a homogeneous sample of healthy
147 individuals aged between 25-60 years. Exclusion criteria included any underlying diseases,
148 recent operations, certain chronic conditions, pregnancy, drug abuse, psychological disorders,
149 and infections.

150 **Data Collection**

151 Data collected from participants includes demographics (age, gender, smoking, alcohol drinking,
152 underlying diseases, marital status, education, occupation, family income and financial support),
153 physical examination (height, weight, BMI, waist circumference, hip circumference, diastolic
154 BP, systolic BP, and heart rate). Urine samples were examined for biomarkers (1-OHP, MDA,
155 and 8-iso-PGF2 α) detection.

156 Urine samples were collected during high (Feb-April 2023), and low (May-July 2023) PM2.5
157 seasons, which were provided by Asst. Prof. Kanokwan Kulprachakarn, Ph.D., from the research
158 project entitled “Health Risk Assessment and Association between Metabolic and Hormonal
159 Derangements in People Exposed to In-house or Ambient PM2.5-Bond Chemicals”. The samples
160 were stored at -20 °C until further analysis.

161 The concentrations of PM2.5 was measured and recorded daily, Air pollution data on a daily
162 basis were obtained from the Northern Thailand Air Quality Index (NTAQI)

163 (<https://www2.ntaqhi.info/>). In order to conduct this study, the authors calculated the monthly
164 average.

165 **Biomarkers Analysis**

166 **Creatinine (Cr)**

167 The amount of Cr in the samples was measured using the spectrophotometric Jaffé method,
168 which relies on the reaction of Cr with picric acid in an alkaline pH solution (Campos, Guzmán
169 et al. 2011). For this purpose a creatinine assay kit (Colorimetric) (ab204537) was used which is
170 a complete kit for the quantitative determination of creatinine in urine.

171 **1-Hydroxypyrene (1-OHP)**

172 The stored urine samples were used for 1-OHP analysis with modifications by the method of K.
173 Sutan, W. Naksen, and T. Prapamontol (Sutan, Naksen et al. 2017) . Specifically, 2.5 mL of
174 urine was adjusted to pH 5.0 using 1M HCl and then transferred into a 50 mL screw cap test tube
175 containing 2.5 mL of 0.1M acetate buffer and 6.25 μ L of β -glucuronidase from *Helix pomatia*.
176 The mixture was vortexed for 10 seconds and incubated at 37°C for 2 hours. After incubation,
177 the samples were processed using a Vertipak C18 3 mL solid-phase extraction (SPE) cartridge.
178 The cartridge was pre-conditioned by rinsing it with 1 mL of methanol (three times), followed by
179 1 mL of water (three times). The samples were then loaded onto the cartridge and washed with
180 2.5 mL of water, followed by 2.5 mL of 20% methanol. Samples were loaded onto the cartridge
181 and washed with 2.5 mL of water and 2.5 mL of 20% methanol. The cartridge was allowed to
182 stand for 10 minutes before elution with 2.5 mL of pure methanol. The eluted solution was
183 filtered through a 0.2 μ m PTFE syringe filter before being evaporated to dryness under a stream
184 of nitrogen gas, then reconstituted in 100 μ L of methanol. For HPLC analysis, a 20- μ L injection
185 was performed using an HPLC system from Agilent 1260 Infinity. The mobile phase consisted
186 of 45% water (Line A) and 55% acetonitrile (Line B) at a flow rate of 0.80 mL/min.
187 Chromatographic separation was achieved using an InfinityLab Poroshell 120 EC-C18 (4.6 mm
188 x 150 mm, 5 μ m) maintained at 25 °C. The detection was carried out using a fluorescence
189 detector with an excitation wavelength of 242 nm and an emission wavelength of 388 nm, and

190 the total run time was 20 minutes. This technique was modified to efficiently process smaller
191 sample volumes while maintaining accurate and reliable quantification of 1-OHP.

192 The standard curve was created using the 1-OHP standards' concentration range of 0.00125 to
193 2.50 ng/mL. The calibration equation was obtained by linear regression, with the signal response
194 (peak area) plotted against known concentrations ($y = 1.74x + 2.24799 \times 10^{-2}$, $R^2 = 0.983$). This
195 calibration equation served as the foundation for the sample concentration calculations.
196 Specifically, the limits of detection (LOD) and limits of quantitation (LOQ) were 0.2634 and
197 0.7984 ng/ml, respectively.

198 **Thio barbituric acid-reacting substances (TBARS)**

199 TBARS were measured by the method of C. Campos et al. (Campos, Guzmán et al. 2011). In
200 summary, 140 μ L of urine was mixed in a vortex with 33 μ L of 0.01% BHT (in absolute
201 ethanol), 1 mL of 1% phosphoric acid, and 300 μ L of 42 mmol/L TBA (dissolved in water and
202 heated). Following a 45-minute incubation period in boiling water, 1.4 mL of 1-butanol was
203 added to each tube and the tubes were allowed to cool on ice. Using a UVmini-1240 Shimadzu
204 spectrophotometer (Shimadzu, Tokyo, Japan), the absorbance of the supernatant was measured at
205 535 nm after a 15-minute centrifugation ($2000 \times g$). To create the standard absorption curve,
206 MDA was dissolved in 20 mmol/L of phosphate buffer (pH 7.0).

207 **8-Epi-Prostaglandin F2 Alpha (8-epi-PGF2 α)**

208 Urinary 8-iso-PGF2 α was determined by the commercial ELISA kit (ElabScience) according to
209 the manufacturer's instruction. The ELISA kit uses the Competitive-ELISA principle, using a
210 pre-coated micro plate with 8-epi-PGF2 α . The enzyme competes with a fixed amount on the
211 solid phase supporter for specific sites. Excess conjugate and unbound sample are washed away,
212 Avidin-Horseradish Peroxidase conjugate is added, and TMB substrate solution is added. Optical
213 density (OD) is measured.

214 **Ethical Considerations**

215 The study protocol was approved by Human Experimentation Committee Research Institute for
216 Health Sciences (RIHES), Chiang Mai University, Chiang Mai, Thailand on 19 January 2023
217 [Project No.03/2023]. Written informed consent was obtained from all subjects involved in the
218 study.

219 **Statistical Analysis**

220 The data are shown as mean \pm standard deviation (SD). The Shapiro–Wilk test was used to
221 determine whether the data had a normal distribution. The Mann-Whitney U-test was used for
222 nonparametric data, and the Student's t-test was utilized for parametric variables. $P < 0.05$ was
223 designated as the threshold for statistical significance. IBM SPSS Statistics 22.0 was used to
224 process the data.

225

226 **Results**

227 **Baseline Characteristics of Participants**

228 This study assessed 25 healthy participants from the Samoeng district in Chiang Mai province,
229 Thailand. The participants had a mean age of 48.1 years (SD = 14.6), with a gender distribution
230 of 60% female and 40% male. Most participants were smokers (84%), and 24% reported alcohol
231 consumption. Educational backgrounds varied, with 44% having less than a high school
232 education, 44% holding college or university degrees, and 12% having completed high school.
233 The occupational breakdown showed that 52% of participants worked in agriculture, 12% in
234 service, and 36% in other sectors. Table 1 shows the baseline characteristics of the participants in
235 terms of mean and standard deviation for **continuous** variables and frequency and percentage for
236 categorical variables.

237

238 **Table 1 Baseline characteristics of the participants (N = 25).**

239

240 **PM2.5 Concentration During High and Low Pollution Seasons**

241 The daily PM2.5 concentration data were obtained from the NTAQHI website, managed by
242 RIHES CMU, for the periods of March-April and May-July 2023. The mean concentrations of
243 PM2.5 during both high and low pollution seasons are presented in Figure 3. During the high
244 pollution season (March-April 2023), PM2.5 levels were significantly elevated, with the highest
245 concentration observed in April ($88.3 \mu\text{g}/\text{m}^3$), followed by March ($67.3 \mu\text{g}/\text{m}^3$). In contrast, the
246 low pollution season (May-July 2023) exhibited substantially lower PM2.5 concentrations, with
247 mean levels of $15.8 \mu\text{g}/\text{m}^3$ in May and $4.1 \mu\text{g}/\text{m}^3$ in July. The variation between these months
248 highlights the distinct seasonal air quality differences in the Samoeng District of Chiang Mai.

249 This variation in PM2.5 levels between the two seasons was critical for analyzing the seasonal
250 impact of air pollution on urinary oxidative stress biomarkers in participants. The average PM2.5
251 concentration for the high pollution season was calculated at $67 \mu\text{g}/\text{m}^3$, while the low pollution
252 season averaged $7 \mu\text{g}/\text{m}^3$. These differences provided a strong basis for assessing the correlation
253 between PM2.5 exposure and biomarker levels, particularly for oxidative stress markers such as
254 1-OHP, MDA, and 8-epi-PGF2 α .

255

256 **Figure 3 Mean concentrations of PM2.5 ($\mu\text{g}/\text{m}^3$) recorded during high (March-April 2023)**
257 **and low (May-July 2023) pollution seasons.** The error bars represent the standard error of the
258 mean (SEM).

259

260 **Urinary Biomarkers and PM2.5 Concentration by Season**

261 In this study, the authors observed significant differences in the urinary concentrations of
262 oxidative stress biomarkers between high and low PM2.5 seasons, which are depicted in Figure
263 4. The graph presents the median concentrations of 8-epi-PGF2 α , 1-OHP, and MDA across the
264 two seasons, highlighting the impact of seasonal PM2.5 exposure on these biomarkers.

265

266 **Figure 4 Comparison of urinary concentrations of oxidative stress biomarkers, including**
267 **A) 8-epi-PGF2 α , B) Malondialdehyde (MDA), and C) 1-hydroxypyrene (1-OHP) between**
268 **high PM2.5 and low PM2.5 seasons.** Data are presented as median values with interquartile
269 ranges (Q1, Q3) for each biomarker. The Wilcoxon signed-rank test was used to assess the
270 significance of differences between seasons, with all *p-values < 0.05 indicating statistically
271 significant increases in biomarker levels during the high PM2.5 season.

272

273 The concentration of 8-epi-PGF2 α , measured in pg/mg creatinine, was significantly higher
274 during the high PM2.5 season (median = 139.43 pg/mg, Q1 = 80.63, Q3 = 187.54) compared to
275 the low PM2.5 season (median = 54.22 pg/mg, Q1 = 26.74, Q3 = 122.76), with a p-value of
276 0.016. This indicates an elevated oxidative stress level during periods of higher air pollution.

277 The concentration of malondialdehyde (MDA), measured in μ M/mg creatinine, was also
278 significantly higher during the high PM2.5 season (median = 3.15 μ M/mg, Q1 = 2.83, Q3 =
279 4.11) compared to the low PM2.5 season (median = 2.45 μ M/mg, Q1 = 1.96, Q3 = 3.05), with a
280 p-value of 0.006. The elevated MDA levels during high PM2.5 exposure further support the
281 hypothesis that increased air pollution contributes to oxidative stress.

282 Similarly, the urinary concentration of 1-hydroxypyrene (1-OHP), expressed in mg/g creatinine,
283 showed a significant increase during the high PM2.5 season (median = 0.09 mg/g, Q1 = 0.06, Q3
284 = 0.17) compared to the low PM2.5 season (median = 0.04 mg/g, Q1 = 0.02, Q3 = 0.11), with a
285 p-value of 0.001. The increased levels of 1-OHP suggest higher internal exposure to polycyclic
286 aromatic hydrocarbons (PAHs) during periods of elevated PM2.5 levels.

287 The Wilcoxon signed-rank test was applied to assess these differences, and all p-values were
288 found to be significant, indicating a robust association between increased PM2.5 exposure and
289 elevated urinary biomarkers of oxidative stress. These findings underscore the heightened
290 oxidative stress during periods of high air pollution, particularly in the Samoeng District, where
291 residents, predominantly farmers, are more susceptible to PM2.5 exposure due to agricultural
292 practices like stubble burning.

293 This analysis provides clear evidence of the health risks associated with seasonal variations in
294 PM2.5 levels, reinforcing the need for targeted interventions to reduce exposure, particularly in

295 vulnerable populations. The significant changes in these biomarkers reflect the biological impact
296 of air pollution and suggest potential pathways through which PM2.5 exposure may lead to
297 adverse health outcomes.

298 **Biomarkers in Relation to PM2.5 Exposure, Age, Gender, and Smoking Status**

299 To further investigate the relationship between PM2.5 exposure and oxidative stress biomarkers,
300 we employed a **Generalized Estimating Equations** (GEE) model, adjusting for potential
301 confounders such as age, gender, and smoking status as shown in Table 2. The analysis revealed
302 significant associations for some biomarkers, underscoring the potential health impacts of air
303 pollution in the study population.

304 **1-Hydroxypyrene (1-OHP)**

305 The regression analysis showed a positive association between PM2.5 levels and 1-OHP, with an
306 estimated coefficient (Exp(b)) of 1.014 (95% CI: 1.01, 1.02, $p < 0.01$). This suggests that for
307 each unit increase in PM2.5 concentration, the level of 1-OHP increases by approximately 1.4%.
308 This finding highlights the strong influence of PM2.5 on PAH metabolism, potentially leading to
309 elevated oxidative stress. Other variables, such as age (Exp(b) = 0.992, $p = 0.422$), gender
310 (Exp(b) = 1.03, $p = 0.933$), and smoking status (Exp(b) = 0.77, $p = 0.543$), did not show
311 significant associations with 1-OHP levels, indicating that PM2.5 is a more critical factor in
312 influencing this biomarker.

313 **Malondialdehyde (MDA)**

314 The association between PM2.5 and MDA was not statistically significant, with a coefficient of
315 0.998 (95% CI: 0.997, 1.00, $p = 0.051$). However, the p-value approaches significance,
316 suggesting a potential negative relationship that warrants further investigation. Interestingly, age
317 was significantly associated with MDA levels (Exp(b) = 1.01, $p = 0.007$), indicating that older
318 participants may have higher oxidative stress levels, independent of PM2.5 exposure. Gender
319 also showed a significant association (Exp(b) = 1.15, $p = 0.041$), with females exhibiting higher
320 MDA levels compared to males. Smoking status, however, did not significantly influence MDA
321 levels (Exp(b) = 1.05, $p = 0.583$).

322 **8-iso-Prostaglandin F2 α (8-iso-PGF2 α)**

323 A significant positive association was found between PM2.5 and 8-iso-PGF2 α levels (Exp(b) =
324 1.008, 95% CI: 1.00, 1.02, p = 0.045). This suggests that increased PM2.5 exposure is linked to
325 elevated levels of 8-iso-PGF2 α , a biomarker of lipid peroxidation, reinforcing the role of PM2.5
326 in promoting oxidative stress. Age and gender did not show significant associations with 8-iso-
327 PGF2 α levels (Exp(b) = 1.013, p = 0.196 and Exp(b) = 1.377, p = 0.344, respectively). However,
328 smoking status showed a marginal association (Exp(b) = 2.22, p = 0.081), indicating a trend
329 towards higher 8-iso-PGF2 α levels among smokers.

330 Overall, these findings demonstrate that PM2.5 exposure is a significant determinant of certain
331 oxidative stress biomarkers, particularly 1-OHP and 8-iso-PGF2 α . The absence of significant
332 associations with MDA may suggest different pathways or sensitivities among the biomarkers.
333 This analysis underscores the complex interactions between environmental pollutants and
334 biological responses, highlighting the importance of targeted interventions to reduce PM2.5
335 exposure and its associated health risks.

336

337 **Table 2 Multivariate regression analysis shows significant associations between PM2.5**
338 **exposure and elevated levels of urinary oxidative stress biomarkers, even after adjusting**
339 **for age, gender, and smoking status.**

340 Note: 1-OHP: 1-hydroxypyrene; MDA: Malondialdehyde; 8-iso-PGF2 α : 8-iso-prostaglandin-
341 F2 α .

342 *P-values < 0.05 indicating statistically significant association.

343

344 Discussion

345 This pilot study provides **novel** insights into the relationship between PM2.5 exposure and
346 oxidative stress biomarkers in Chiang Mai, Thailand. Our findings demonstrate that elevated
347 PM2.5 levels are significantly associated with increased concentrations of 1-OHP, MDA, and 8-
348 iso-PGF2 α , indicating a pronounced oxidative stress response. The observed increase in 1-OHP
349 and 8-iso-PGF2 α levels during periods of high PM2.5 exposure corroborates **previous studies**
350 linking air pollution to oxidative stress. PM2.5 carries various toxic substances, including PAHs,

351 which are metabolized into 1-OHP, a reliable biomarker for PAH exposure (Luo, Stepanov et al.
352 2019, Liu, Liu et al. 2023). Similarly, the significant elevation in 8-iso-PGF2 α , a marker of lipid
353 peroxidation, aligns with research highlighting the role of air pollution in inducing oxidative
354 damage and inflammation (Glencross, Ho et al. 2020, Leni, Künzi et al. 2020).

355 The correlation between PM2.5 levels and MDA, though less pronounced than for 1-OHP and 8-
356 iso-PGF2 α , still underscores the oxidative stress induced by particulate matter exposure. MDA, a
357 byproduct of lipid peroxidation, has been widely used as a biomarker for oxidative stress in
358 various studies, further validating our findings(Cui, Gong et al. 2018, Zhang, Liu et al. 2023).
359 The impact of PM2.5 on oxidative stress biomarkers, as evidenced by our study, suggests
360 potential health risks, including cardiovascular and respiratory diseases, which have been
361 extensively documented in the literature(Brook, Rajagopalan et al. 2010, Schraufnagel, Balmes
362 et al. 2019).

363 Our results also reveal a significant association between age and MDA levels, with older
364 individuals exhibiting higher levels of oxidative stress. This finding is consistent with previous
365 studies showing age-related susceptibility to oxidative damage due to the cumulative effects of
366 environmental exposures over time (Weary 2023) . Gender differences were observed, with
367 females showing higher MDA levels, a finding that resonates with research suggesting that
368 hormonal differences, particularly the presence of estrogen, may influence oxidative stress
369 responses(Viña, Borrás et al. 2005, Berry 2022).

370 Interestingly, no significant association was found between smoking status and the oxidative
371 stress biomarkers studied, although a trend was noted with 8-iso-PGF2 α . This may be attributed
372 to the small sample size and the possibility that oxidative stress induced by PM2.5 exposure
373 could overshadow the effects of smoking. The literature reports mixed results, with some studies
374 showing a synergistic effect of smoking and air pollution on oxidative stress, while others have
375 not (Peretz, Kaufman et al. 2008, Makra, Puskás et al. 2015).

376 Overall, our findings are consistent with a growing body of evidence linking air pollution,
377 particularly PM2.5, to oxidative stress and the development of non-communicable diseases
378 (NCDs). Oxidative stress is a key mechanism through which PM2.5 exposure contributes to the
379 pathogenesis of cardiovascular and respiratory diseases, as well as cancer (Wang, Ma et al.
380 2019). The specific biomarkers analyzed in our study—1-OHP, MDA, and 8-iso-PGF2 α —are

381 widely recognized as indicators of oxidative stress, providing a robust framework for assessing
382 the biological impact of air pollution (Zhao, Xu et al. 2023).

383 The findings of this study have important public health implications, particularly in regions like
384 Chiang Mai, where seasonal air pollution is a significant concern. The association between
385 PM_{2.5} exposure and oxidative stress biomarkers suggests that residents in highly polluted areas
386 are at an increased risk of developing oxidative stress-related diseases. This underscores the need
387 for targeted interventions to reduce air pollution exposure, particularly during the burning
388 season. Additionally, the use of oxidative stress biomarkers in epidemiological studies could
389 provide valuable insights into the long-term health effects of air pollution and inform public
390 health policies aimed at mitigating these risks.

391 The strengths of this study lie in its focused, area-specific approach, targeting the Samoeng
392 District, a region previously unexplored in this context. This district, primarily inhabited by
393 farmers, presents a unique population with heightened exposure to PM_{2.5}, particularly due to
394 stubble burning, making the findings highly relevant to local environmental health concerns. The
395 study's design, with repeated measurements during different PM_{2.5} seasons, allows for the
396 observation of seasonal variations in oxidative stress biomarkers, providing a comprehensive
397 view of the impact of air pollution on health. Additionally, the use of multiple biomarkers—8-
398 epi-PGF₂α, MDA, and 1-OHP—strengthens the evidence of oxidative stress due to PM_{2.5}
399 exposure, and the significant correlations observed add to the robustness of the study's findings.

400 However, this study has some limitations. One of the primary limitations is the small sample
401 size, which limits the generalizability of the findings. As a pilot study, the results should be
402 interpreted with caution, and larger studies are needed to validate these findings. Furthermore,
403 the study did not account for other potential confounding factors, such as diet, physical activity,
404 and socioeconomic status, which could influence oxidative stress levels. The cross-sectional
405 design of the study also limits the ability to draw causal inferences between PM_{2.5} exposure and
406 oxidative stress biomarkers.

407 Future research should focus on expanding the sample size and incorporating longitudinal data to
408 better understand the temporal relationship between PM_{2.5} exposure and oxidative stress.

409 Additionally, exploring the role of other potential confounders, such as genetic susceptibility and
410 lifestyle factors, could provide a more comprehensive understanding of the factors influencing

411 oxidative stress. Investigating the combined effects of air pollution and other environmental
412 exposures, such as indoor air pollution and chemical contaminants, could also shed light on the
413 cumulative impact of multiple stressors on oxidative stress and health outcomes.

414

415 **Conclusion**

416 In conclusion, this study highlights the significant impact of PM_{2.5} exposure on oxidative stress
417 biomarkers, with higher levels observed during the high PM_{2.5} season in the Samoeng District.
418 The strong associations found between PM_{2.5} and biomarkers such as 8-epi-PGF₂ α , 1-OHP, and
419 MDA reinforce the health risks posed by air pollution, particularly in rural areas prone to stubble
420 burning. These findings underscore the urgent need for targeted strategies to mitigate air
421 pollution and protect public health in vulnerable communities.

422

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429

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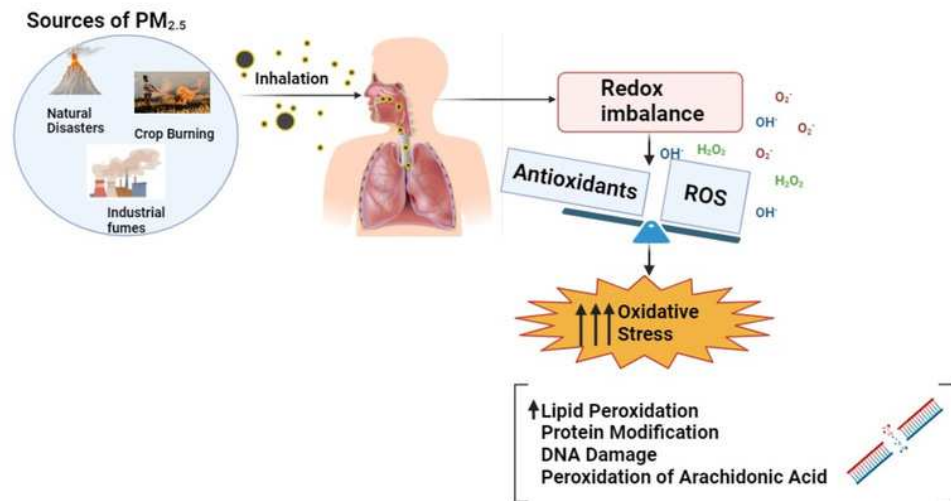
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548

Figure 1

Mechanism of oxidative stress induced by PM2.5 exposure, showing the imbalance between ROS and antioxidants leading to oxidative damage.



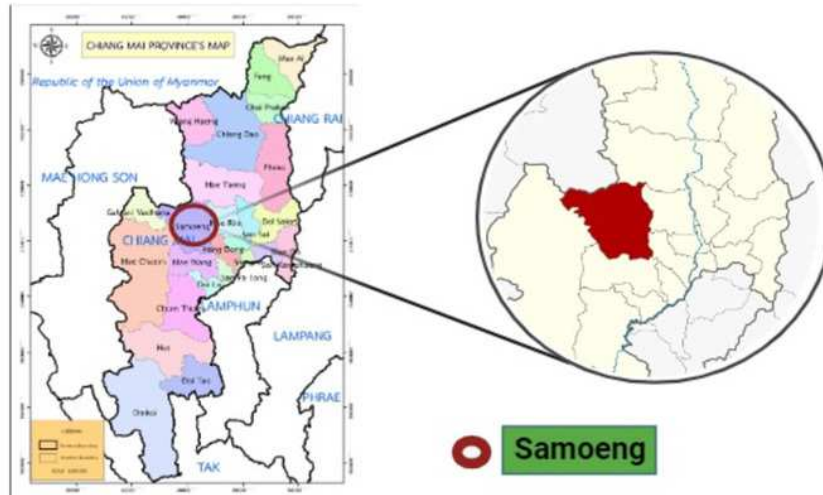
1

2 **Figure 1 Mechanism of oxidative stress induced by PM_{2.5} exposure, showing the imbalance**
3 **between ROS and antioxidants leading to oxidative damage.**

4

Figure 2

Map showing the study area in Chiang Mai, Thailand.



1

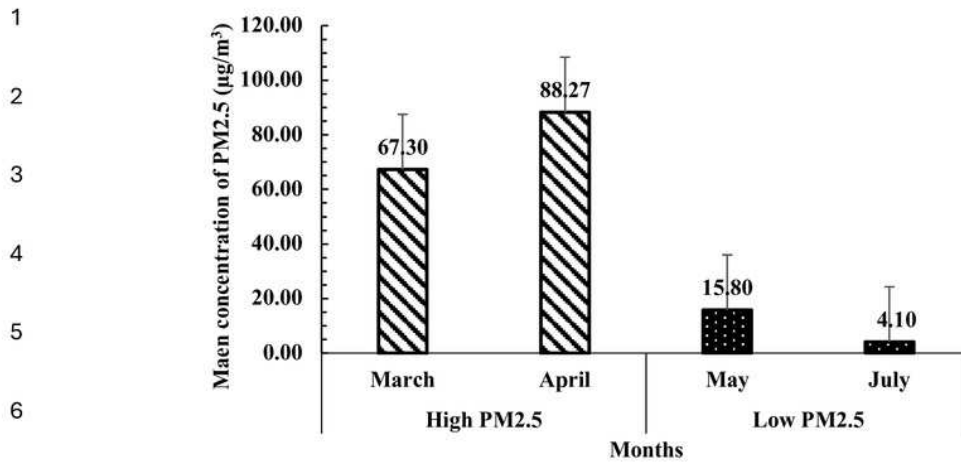
2 **Figure 2 Map showing the study area in Chiang Mai, Thailand.**

1

Figure 3

Mean concentrations of PM_{2.5} ($\mu\text{g}/\text{m}^3$) recorded during high (March-April 2023) and low (May-July 2023) pollution seasons.

The error bars represent the standard error of the mean (SEM).

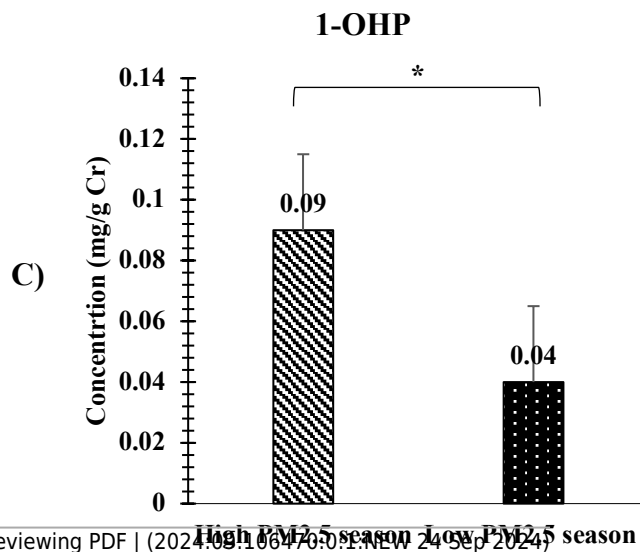
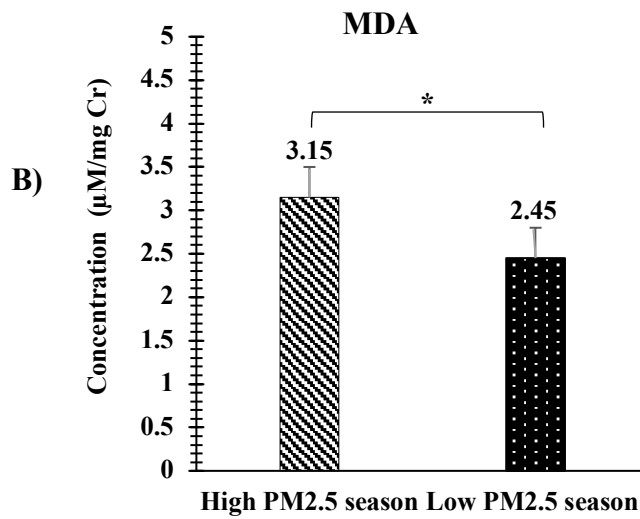
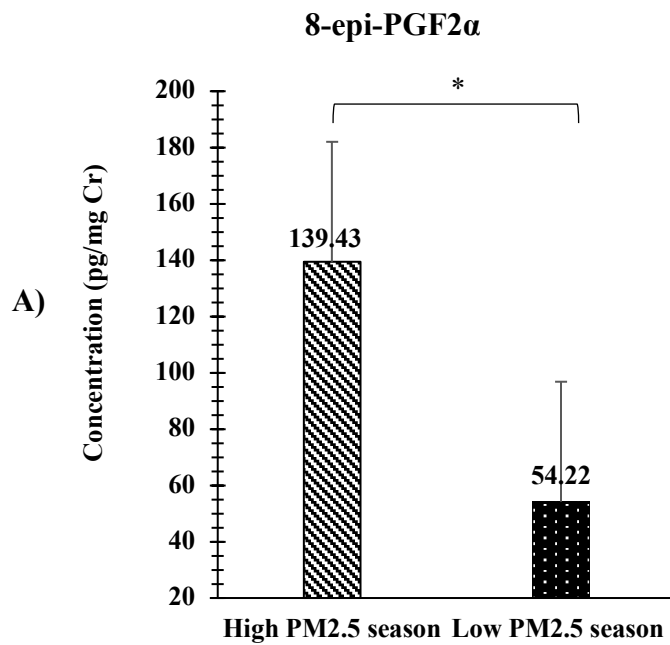


8 **Figure 3 Mean concentrations of PM2.5 ($\mu\text{g}/\text{m}^3$) recorded during high (March-April 2023)**
9 **and low (May-July 2023) pollution seasons. The error bars represent the standard error of the**
10 **mean (SEM).**

Figure 4(on next page)

Comparison of urinary concentrations of oxidative stress biomarkers, including A) 8-epi-PGF2 α , B) Malondialdehyde (MDA), and C) 1-hydroxypyrene (1-OHP) between high PM2.5 and low PM2.5 seasons.

Data are presented as median values with interquartile ranges (Q1, Q3) for each biomarker. The Wilcoxon signed-rank test was used to assess the significance of differences between seasons, with all *p-values < 0.05 indicating statistically significant increases in biomarker levels during the high PM2.5 season.



26 **Figure 4 Comparison of urinary concentrations of oxidative stress biomarkers, including A)**
27 **8-epi-PGF2 α , B) Malondialdehyde (MDA), and C) 1-hydroxypyrene (1-OHP) between high**
28 **PM2.5 and low PM2.5 seasons.** Data are presented as median values with interquartile ranges
29 (Q1, Q3) for each biomarker. The Wilcoxon signed-rank test was used to assess the significance
30 of differences between seasons, with all *p-values < 0.05 indicating statistically significant
31 increases in biomarker levels during the high PM2.5 season.

Table 1 (on next page)

Baseline characteristics of the participants (N = 25).

1 **Table 1 Baseline characteristics of the participants (N = 25).**

Characteristic	N (%)
Age (years)	
Mean \pm SD	48.1 \pm 14.6
Gender	
Male	10 (40.0%)
Female	15 (60.0%)
Smoking	
Yes	21 (84.0%)
No	4 (16.0%)
Alcohol Consumption	
Yes	6 (24.0%)
No	19 (76.0%)
Education Level	
Less than high school	11 (44.0%)
High school	3 (12.0%)
College/University	11 (44.0%)
Occupation	
Agriculture	13 (52.0%)
Service	3 (12.0%)
Others	9 (36.0%)

2

Table 2 (on next page)

Multivariate regression analysis shows significant associations between PM2.5 exposure and elevated levels of urinary oxidative stress biomarkers, even after adjusting for age, gender, and smoking status.

1-OHP: 1-hydroxypyrene; MDA: Malondialdehyde; 8-iso-PGF2 α : 8-iso-prostaglandin-F2 α .

*P-values < 0.05 indicating statistically significant association.

1 **Table 2 Multivariate regression analysis shows significant associations between PM2.5**
 2 **exposure and elevated levels of urinary oxidative stress biomarkers, even after adjusting**
 3 **for age, gender, and smoking status.**

Dependent Variable	Independent Variables	Coefficient (Exp(b))	Standard Error	z-value	p-Value	95% CI
1-OHP	PM2.5	1.014	0.003	4.25	<0.01*	1.01,1.02
	Age	0.992	0.01	-0.80	0.422	0.97,1.01
	Gender (Female)	1.03	0.33	0.08	0.933	0.55,1.91
	Smoking Status (Smoker)	0.77	0.33	-0.61	0.543	0.33,1.80
MDA	PM2.5	0.998	0.0008	-1.95	0.051	0.997,1.00
	Age	1.01	0.002	2.70	0.007	1.00,1.01
	Gender (Female)	1.15	0.079	2.04	0.041	1.01,1.32
	Smoking Status (Smoker vs Non-Smoker)	1.05	0.10	0.55	0.583	0.87,1.27
8-iso-PGF2 α	PM2.5	1.008	0.004	2.00	0.045*	1.00,1.02
	Age	1.013	0.014	1.29	0.196	0.99,1.03
	Gender (Female)	1.377	0.465	0.95	0.344	0.71,2.67
	Smoking Status (Smoker)	2.22	1.017	1.75	0.081	0.91,5.45

4 1-OHP: 1-hydroxypyrene; MDA: Malondialdehyde; 8-iso-PGF2 α : 8-iso-prostaglandin-F2 α .

5 *P-values < 0.05 indicating statistically significant association.

6

7