

Synthetic Cathinone Impact on the Protein Profile of Intestinal Caco-2 Cells [†]

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Abstract

The global rise in New Psychoactive Substances (NPS), particularly synthetic cathinones, poses significant public health concerns due to their association with toxicity and fatalities. With oral intake a common route of consumption, understanding their effects on human intestinal epithelium is essential but remains poorly explored. This study investigates the impact of four synthetic cathinones—3-CIC (3-chloro-*N*-isopropylcathinone), 4-CIC (4-chloro-*N*-isopropylcathinone), 3-Cl-TBC (3-chloro-*tert*-butylcathinone, bupropion), and 4-Cl-TBC (4-chloro-*tert*-butylcathinone)—using Caco-2 cells, focusing on protein expression changes. Results revealed reduced protein content, with 3-CIC producing the most significant effects, including upregulation of 40–50 kDa proteins. These findings suggest pathway disruptions requiring further mechanistic investigation.

Keywords: New Psychoactive Substances; Synthetic Cathinones; drug-induced alterations; Caco-2; protein profile; SDS-PAGE

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1. Introduction

The global use of New Psychoactive Substances (NPS) has risen markedly in recent years. By the end of 2024, the European Union Drugs Agency (EUDA) had identified around 1000 NPS, including 47 newly detected compounds, 7 of which were synthetic cathinones [1]. Representing the second-largest category of NPS, synthetic cathinones exhibit psychostimulant properties comparable to those of cocaine [2], amphetamines, and MDMA [3], influencing monoamine neurotransmitter systems and causing various toxicological effects [4–6].

It has been reported in the literature, that exposure to synthetic cathinones affects multiple organs and systems. In muscle tissue, compounds such as 3-methylmethcathinone (3-MMC) and mephedrone promote myotoxicity, contributing to the muscular damage frequently associated with abuse [7]. In the liver, substances like methylenedioxypyrovalerone (MDPV) and mephedrone deplete ATP, triggering mitochondrial dysfunction,

oxidative stress, and protein changes as cells attempt to restore energy balance [8]. Similar disruptions are observed in the kidney, where cathinones disturb the regulation of autophagy and apoptosis, leading to endoplasmic reticulum stress, mitochondrial damage, and activation of pro-apoptotic pathways [9]. Extensive research in neuronal models has shown that cathinones trigger early gene expression, alter synaptic plasticity, and cause neurotoxicity influenced by metabolism [6,10]. They also disrupt protein modifications [11,12], increase oxidative stress [13], and activate neuroinflammatory responses [14,15]. Additionally, inhibition of monoamine transporters disturbs neurotransmitter balance, driving compensatory protein changes that worsen cognitive deficits and neuronal damage [16,17].

However, their effects on intestinal cells remain largely unexplored, even though the intestinal epithelium is the first physiological barrier after cathinones oral intake. This barrier not only regulates nutrient absorption but also determines the uptake and systemic distribution of xenobiotics, including drugs of abuse [18]. Alterations at the level of intestinal epithelial function may therefore alter the bioavailability of cathinones, compromise epithelial integrity, and ultimately influence the toxicological burden and systemic impact of these substances.

In this context, the present preliminary study investigates protein alterations in human intestinal epithelial Caco-2 cells following exposure to four synthetic cathinones: 3-CIC (3-chloro-*N*-isopropylcathinone), 4-CIC (4-chloro-*N*-isopropylcathinone), 3-Cl-TBC (3-chloro-*tert*-butylcathinone, bupropion), and 4-Cl-TBC (4-chloro-*tert*-butylcathinone), to characterize their impact on intestinal lining cells, assess potential consequences for bioavailability and systemic implications, and provide insights that can guide future mechanistic studies and risk assessments related to their abuse.

2. Materials and Methods

2.1. Chemicals and Biochemicals

Roswell Park Memorial Institute (RPMI) culture medium, bovine serum albumin (BSA), chloroform, Bradford reagent, Coomassie Brilliant Blue R-250, and glacial acetic acid (99.9%) were obtained from VWR. Dulbecco's Modified Eagle Medium (DMEM) and heat-inactivated Fetal Bovine Serum (FBS) were purchased from Biowest. L-glutamine (100×) was obtained from Lonza, while penicillin-streptomycin solution (50×, Pen-Strep) and Phosphate-Buffered Saline (1×, PBS) were acquired from Corning. Copper sulfate (>99%) was purchased from Sigma, methanol from Honeywell, 96% ethanol from Lab-chem, ultrapure water from Merck, and sodium hydroxide (0.66 N) from Fisher Chemical. Bolt™ MOPS SDS Running Buffer (20×) and Bolt™ LDS Sample Buffer (4×) were supplied by Novex Life Technologies.

The hydrochloride salt standards of the chlorinated cathinones used in this study were previously synthesized and characterized [19].

2.2. Protein Extraction

For protein extraction [20], five 25 cm² T-flasks of Caco-2 cells were cultured in RPMI medium supplemented with 10% FBS, 1% L-glutamine (100×), and 1% penicillin-streptomycin (50×). Cells were maintained under standard incubation conditions (37 °C, 5% CO₂, humidified atmosphere). Humidity was achieved by placing a solution of 1 g copper sulfate in 500 mL ultrapure water inside the incubator. The culture medium was refreshed every 2–3 days until cell differentiation. Cells were then exposed for 24 h to the following concentrations: 6.2 mM (3-CIC), 4.6 mM (4-CIC), 4.3 mM (3-Cl-TBC), and 2.5 mM (4-Cl-TBC). One flask was used as a control for CIC compounds and another as a control for Cl-TBC compounds. After exposure, cells were washed twice with PBS and stored at –80 °C.

Protein extraction was performed by adding 1200 μL of a solution containing ultrapure water, chloroform, and methanol (10:3:27) to promote cell lysis. Cells were detached from the surface using a cell scraper, and the suspension was sonicated on ice for 10 min.

2.3. Protein Quantification by Bradford Method

For protein quantification, 100 μL of a solution of NaOH (0.66 N) and ultrapure water (0.2 g in 10 mL) was added to 70 μL of the cell suspension to ensure complete cell lysis and protein denaturation, making them accessible for quantification. Protein concentrations were determined using a calibration curve prepared with bovine serum albumin (BSA) standards ranging from 0 to 0.5 mg/mL. For the blank, 795 μL of ultrapure water, 5 μL of ultrapure water, and 200 μL of Bradford reagent were mixed. For the calibration standards, 795 μL of ultrapure water, 5 μL of each BSA concentration, and 200 μL of Bradford reagent were used. The same procedure was applied to the samples, except that 5 μL of the cell suspension from each cathinone exposure condition replaced the BSA solution. Absorbance was measured at 595 nm.

2.4. SDS-PAGE

For SDS-PAGE electrophoresis using 4–12% Bis-Tris Plus polyacrylamide gels (1.0 mm $\times$ 12 wells), 90 μL of each previously prepared sample solution was mixed with 30 μL of Bolt™ LDS Sample Buffer (4 $\times$). Samples were boiled in a 100 $^{\circ}\text{C}$ water bath and subsequently placed on ice. One liter of Bolt™ MOPS SDS Running Buffer (20 $\times$) was prepared with ultrapure water. Gels were rinsed and assembled in a mini tank according to the manufacturer's instructions. Five microliters of Prestained Protein MW Marker and 30 μL of each sample were loaded into the wells. Electrophoresis was performed at 120 V for approximately 1 h. After the run, gels were incubated for 2 h in Coomassie Brilliant Blue R-250 staining solution, followed by overnight incubation in a destaining solution prepared with 10% absolute ethanol (99.8%), 7.5% glacial acetic acid (99%), and 82.5% ultrapure water. Gel images were captured using the ImageQuant LAS 500 system and analyzed with ImageJ software. Relative changes in protein band areas were assessed by calculating the Fold Change (FC) (Equation (1)), providing a measure to compare the relative protein expression between control and exposed cells.

$$\text{FC} = \frac{\text{Normalized band area (exposed)}}{\text{Normalized band area (control)}} \quad (1)$$

2.5. Statistical Analysis

One-Way ANOVA was applied to analyze the variance of the results. p -values lower than 0.05 ($p < 0.05$) were determined as unrevealed statistical differences.

3. Results

To assess the effects on intestinal Caco-2 cell protein expression, cells were exposed to four structurally related synthetic cathinones, including positional isomers: 3-CIC (3-chloro-*N*-isopropylcathinone), 4-CIC (4-chloro-*N*-isopropylcathinone), 3-Cl-TBC (3-chloro-*tert*-butylcathinone, bupropion), and 4-Cl-TBC (4-chloro-*tert*-butylcathinone). Protein content and electrophoretic profiles were subsequently evaluated.

3.1. Protein Quantification in Caco-2 Cells

Total protein concentrations were determined using the Bradford assay. A calibration curve ($R^2 = 0.9963$) was generated with bovine serum albumin (BSA) as standard as described in Section 2.3 (Figure 1).

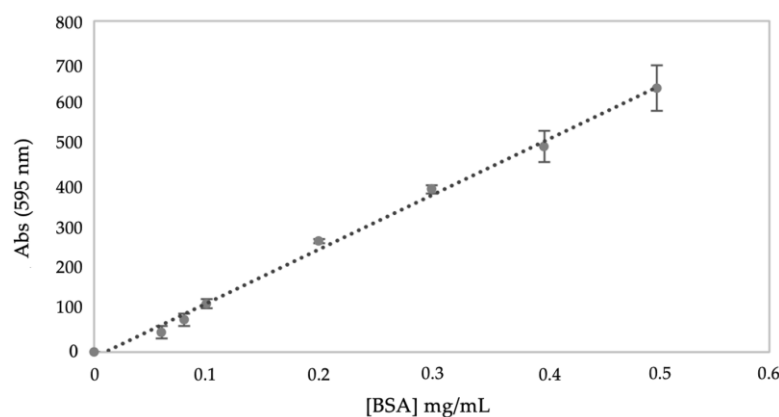


Figure 1. Calibration curve of bovine serum albumin (BSA) obtained by Bradford method. The linear equation, $\text{Abs (595 nm)} = 0.745 [\text{BSA}] (\text{mg/mL}) + 0.012$ ($R^2 = 0.998$), was applied to convert absorbance at 595 nm to protein concentration.

The calibration curve was applied to determine the content in proteins extracted from Caco-2 cells. The results shown in Figure 2 were obtained for Caco-2 cells exposed to the four cathinones under study and compared with unexposed control cells.

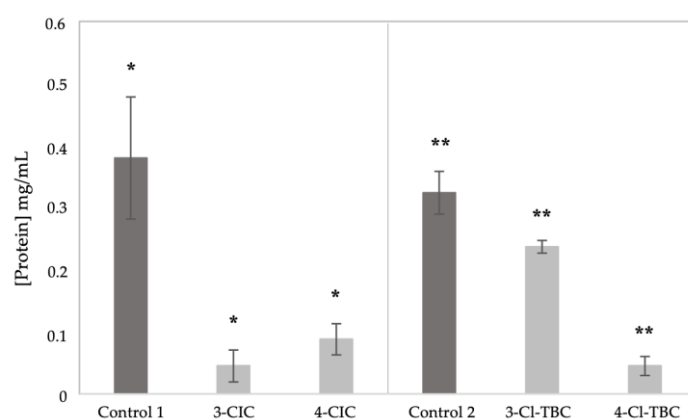


Figure 2. Protein content of Caco-2 cell extracted after 24 h exposure to 3-CIC, 4-CIC, 3-Cl-TBC and 4-Cl-TBC compared with unexposed controls. Control 1: unexposed cells for CIC assays; Control 2: unexposed cells for TBC assays. (*) and (**) Indicate statistically significant differences, compared to the controls, for $p < 0.05$.

As shown in Figure 2, exposure to all four cathinones resulted in a general reduction of total protein content compared with their respective controls. Among the compounds tested, 3-CIC, 4-CIC and 4-Cl-TBC caused the most pronounced decrease, suggesting a stronger impact on cellular protein homeostasis. These findings indicate that for Cl-TBC positional isomers exert distinct effects on protein levels in Caco-2 cells.

3.2. Electrophoretic Profiling and Quantitative Analysis of Caco-2 Effects on Proteins

To further investigate which proteins were affected by cathinones exposure, protein extracts from Caco-2 cells were analyzed by SDS-PAGE to examine changes in band patterns and relative abundance (Section 2.4).

The protein profiles obtained by SDS-PAGE of the extracted proteins from Caco-2 cells exposed to 3-Cl-TBC and 4-Cl-TBC and unexposed control (Control 2) and, from Caco-2 cells and cells exposed to 3-CIC and 4-CIC and unexposed control (Control 1) are shown in Figure 3.

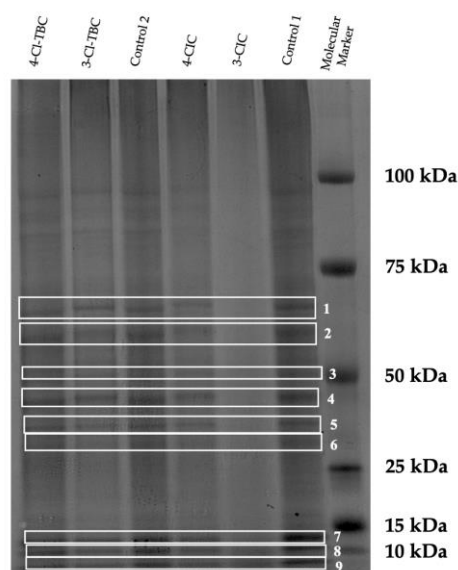


Figure 3. SDS-PAGE 4–12% (Bis-Tris), electrophoretograms of proteins extracted from Caco-2 cells exposed or unexposed to cathinones for 24 h. Control 1: unexposed cells from CIC assay day; Control 2: unexposed cells from TBC assay day. Molecular weight marker is shown on the left.

The electrophoretograms revealed alterations in protein profiles in cathinones-exposed cells compared with their respective controls, with structural isomers producing distinct effects on specific bands, especially for the bands identified in Figure 3 from 1 to 8. In order to evaluate those differences, band areas were quantified by densitometry. Band areas were normalized and expressed as Fold Change (FC) defined as the ratio of the normalized areas of the exposed cells to the normalized area of the corresponding control cells. The average molecular weight (MW) was also determined for the bands 1 to 8 (Table 1).

Table 1. Fold Change (FC) and average molecular weight (MW) values for the bands 1–8 identified in Figure 3 Caco-2 protein profiles after 24 h exposure to 3-CIC, 4-CIC, 3-Cl-TBC and 4-Cl-TBC.

Bands	Fold Change				MW (kDa)
	3-CIC	4-CIC	3-Cl-TBC	4-Cl-TBC	
1	0.80	1.01	1.41	1.11	58.88
2	14.38	1.18	1.33	1.17	50.12
3	20.07	1.21	2.25	0.97	41.69
4		0.88	1.02	1.25	38.90
5		1.47	1.16	1.12	33.88
6		1.18	0.78	0.88	28.84
7		0.76	0.79	0.79	14.45
8		0.60	0.76	0.94	13.80
9		1.23	0.63	0.68	12.59

As shown in Table 1, cathinone exposure induced both up- and down-regulation of specific protein bands compared with unexposed controls. 3-CIC produced the most striking effects, with fold changes exceeding 14- and 20-fold for proteins in the 50 and 41 kDa range (bands 2 and 3, respectively) and 3-Cl-TBC a 2-fold increased also for the latter. 4-CIC and the TBC derivatives induced more moderate increases or decreases in the selected bands, suggesting a milder compound-specific modulation of protein expression.

Taken together, these findings demonstrate that cathinones not only reduce overall protein content in Caco-2 cells but also alter the expression of specific proteins in a manner

dependent on their structural features. Among the tested compounds, 3-CIC consistently caused the strongest alterations, while its positional isomer, 4-CIC, and the TBC derivatives elicited weaker or variable responses. These results provide a first insight into the ability of synthetic cathinones to differentially modulate intestinal epithelial protein profiles, potentially impacting intestinal barrier function and bioavailability.

4. Discussion

Overall, exposure of Caco-2 cells to synthetic cathinones resulted in a reduction of total protein content, indicating that these compounds broadly affect protein homeostasis in intestinal lining cells. This decrease may be associated with cathinone-induced cytotoxicity through inhibition of protein synthesis pathways [21,22]. This general effect was accompanied by changes in specific proteins, with some proteins showing decreased abundance while others were upregulated.

Among the cathinones tested, 3-CIC exposure was seen to produce the strongest alterations in the protein profile of Caco-2 cells, with increased expression of proteins in the 50 and 41 kDa range. Although their identity remains to be determined, such upregulation may indicate activation of cellular defense pathways, possibly linked to stress or detoxification responses [23]. As several cathinones have been associated with oxidative stress and mitochondrial dysfunction, the observed increases could correspond to protective mechanisms against reactive oxygen species [24]. Previous studies have also reported 3-CIC as a potent synthetic cathinone in terms of cytotoxicity and functional effects in neuronal models [19]. The present findings extend these observations to the intestinal epithelium, suggesting that 3-CIC may exert significant effects beyond neuronal systems.

The findings also suggest that even closely related isomers can differentially affect protein expression in intestinal epithelial cells. Although 3-CIC and 4-CIC share the same molecular formula and only differ in the position of the chlorine substituent, their effects on the Caco-2 protein profile were clearly distinct, with 3-CIC producing stronger and more consistent alterations. A similar pattern was observed for the *tert*-butylcathinone derivatives, where 3-Cl-TBC and 4-Cl-TBC induced more moderate and variable changes compared with the isopropylcathinone analogues. These results indicate that subtle structural differences among cathinones can lead to divergent cellular responses, a phenomenon that may be critical for understanding their toxicological properties.

Given that many synthetic cathinones are designed by making small structural modifications to circumvent regulation, these observations highlight the importance of studying positional isomers in greater detail. Clarifying how such differences in structure translate into differences in protein expression and cellular impact will be essential for anticipating the biological activity of new derivatives as they continue to emerge on the illicit market.

5. Conclusions

Protein quantification in Caco-2 cells exposed to the cathinones revealed an overall reduction in protein content compared to unexposed cells, particularly following exposure to 3-CIC and 4-CIC. Regarding the effects of cathinones on cellular protein expression, the most pronounced changes were observed by exposure to 3-CIC, which induced an increase in the expression of certain proteins with molecular weights between 40 and 50 kDa.

Further studies will be needed to identify the proteins involved and to validate the observed changes with complementary approaches. Despite being only preliminary, the present results provide novel evidence that synthetic cathinones alter protein expression in intestinal epithelial cells. Importantly, the observation that positional isomers can

produce distinct effects underlines the need to clarify how subtle structural differences shape cellular responses. These findings highlight the value of extending this line of investigation to better understand the potential impact of cathinone exposure at the intestinal barrier, which may influence bioavailability and, ultimately, the systemic toxicity and health risks associated with their abuse.

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