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Effects of stable degradation products of curcumin on cancer cell proliferation and inflammation

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1	Effects of Stable Degradation Products of Curcumin on Cancer Cell Proliferation and
2	Inflammation
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16	Title running head: Curcumin degradation
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27 Abstract

28 Curcumin is among the most promising dietary compounds for cancer prevention. However, 29 curcumin rapidly degrades in aqueous buffer at physiological pH, making it difficult to 30 understand whether the effects of curcumin are from curcumin itself or its degradation products. 31 Here we studied the anti-proliferative and anti-inflammatory effects of curcumin degradation 32 products, including its total degradation products (a mixture containing all stable degradation 33 products of curcumin) and bicyclopentadione (a dominant stable degradation compound of 34 curcumin). Curcumin potently modulated cell proliferation, progression of cell cycle, and apoptosis in MC38 colon cancer cells, and inhibited lipopolysaccharide (LPS)-induced 35 36 inflammatory responses and NF-kB signaling in RAW 264.7 macrophage cells. In contrast, neither the total degradation products of curcumin nor bicyclopentadione had such effects. For 37 38 example, after 24-h treatment in MC38 colon cancer cells, 5 µg/mL curcumin inhibited 39.2±1.8% 39 of cell proliferation, while its degradation products were inactive. Together, these results 40 suggest that the stable chemical degradation products of curcumin are not likely to play a major role in mediating the biological activities of curcumin. 41 42 43 Keywords Curcumin; degradation; bicyclopentadione; cancer; inflammation 44 45 46 47 48 49 50 51

53

Introduction

54	Curcumin is among the most promising dietary compounds for cancer prevention and treatment
55	¹ . Animal studies have shown that curcumin inhibits many types of cancers, including but not
56	limited to pancreatic cancer ² , breast cancer ³ , cervical cancer ⁴ , lung cancer ⁵ , and colorectal
57	cancer 6 . Human studies also support the anti-cancer effects of curcumin. One Phase II α human
58	clinical trial has shown that a daily intake of 4 grams of curcumin for a month caused a \sim 40%
59	reduction of aberrant crypt foci, a biomarker of colorectal cancer ⁷ . Another Phase II human
60	clinical trial has shown that a daily intake of 8 grams of curcumin displayed anti-cancer efficacy
61	in some patients with advanced pancreatic cancer ⁸ . Currently the anti-cancer effects of
62	curcumin are being evaluated in many human clinical trials.
63	
64	Until now, the molecular mechanisms of curcumin are not well understood. Recent studies
65	showed that curcumin has a poor chemical stability at physiological pH: in aqueous solution at
66	physiological pH, it has a half-life of several minutes, thus leading to the formation of various
67	degradation products ^{9, 10} . These chemical degradation products include two classes of
68	compounds: (1) alkaline hydrolysis products, where the hydroxyl ion (OH ⁻) attacks the carbonyl
69	group of curcumin, generating hydrolyzed products such as ferulic acid, vanillin, ferulaldehyde,
70	and feruloyl methane ¹¹ ; and (2) autoxidation products, where curcumin is first converted to a
71	phenolic radical, which then migrates to the conjugated heptadienedione chain and initiates a
72	chain reaction of curcumin degradation to generate cyclized compounds such as
73	bicyclopentadione (BCP) ^{9, 10, 12} . Recent studies showed that the alkaline hydrolysis products are
74	minor compounds of curcumin degradation products, and the autoxidation pathway is the
75	dominant mechanism to mediate curcumin degradation ^{9, 10, 12} . When curcumin was added into
76	mouse macrophage RAW 264.7 cells, the cellular level of BCP was increased by 1.8-fold,
77	suggesting that chemical degradation could happen in biological systems ¹³ . Because of the
78	short half-life of curcumin at physiological pH, it is difficult to understand whether the observed

79 biological effects of curcumin in cell culture experiments are from curcumin itself or its 80 degradation products. With the presence of serum protein in cell culture medium, curcumin is more stable, but still could undergo degradation after long-term incubation ¹⁴. Curcumin is much 81 82 more stable in the presence of serum proteins: in cell culture medium containing 10% fetal calf serum and in human blood, the half-life of curcumin is > 8 hours ¹⁴. Our recent research showed 83 that co-addition of redox active antioxidants dramatically enhanced the chemical stability and 84 biological activity of curcumin, suggesting that curcumin, but not its degradation products, is 85 largely responsible for the observed effects ¹⁵. Until now, the biological activities of the stable 86 autoxidation products, such as BCP, are largely unknown. To this end, here we studied the 87 comparative effects of curcumin and its stable autoxidation products on cancer and 88 inflammation, in order to better understand the mechanism of curcumin. 89 90 91 92 **Materials and Methods** Chemicals 93 To synthesize curcumin, boric anhydride (0.35 g, 5 mmol) and acetylacetone (1.03 mL, 10 mmol) 94 95 in 50 mL anhydrous ethyl acetate was stirred at 50°C for 30 min. Vanillin (3.04 g, 20 mmol) and 96 tributyl borate (10.8 mL, 40 mmol) were added and the mixture was stirred at 50°C for another 30 min. n-Butylamine (0.4 mL, 5 mmol) dissolved in 15 mL anhydrous ethyl acetate was added 97 dropwise, and the reaction mixture was stirred at 80°C for 4 hours and at room temperature 98 99 overnight. To stop the reaction, 30 mL of 1N hydrochloric acid was added and stirred for 30 min. 100 The reaction mixture was extracted with ethyl acetate and dried with anhydrous magnesium 101 sulfate. Curcumin was obtained after recrystallization using methanol. The structure of synthesized curcumin was confirmed using high resolution ESI-MS and 'H NMR, as reported 102

- 103 previously 16 .
- 104

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105	Preparation and analysis of total degradation products (TDP) of curcumin
106	A 1 L solution of 25 μ M curcumin in 0.1 M phosphate buffer (2.59 g/L KH ₂ PO ₄ , 11.5 g/L
107	Na ₂ HPO ₄ , pH=7.4) was stirred at room temperature for 5 days, then extracted with ethyl acetate.
108	The organic layer, which is a mixture containing various curcumin degradation products, was
109	dried and used as total degradation products (TDP) for the biological experiments.
110	
111	TDP was dissolved in methanol and analyzed by HPLC on an Agilent 1100 HPLC system, using
112	a 4.6 x 250 mm, 5 µm Supelco Nucleosil 100-5C18 column (Sigma-Aldrich, St. Louis, MO), with
113	flow rate at 1 mL/min and detection wavelength at 254 nm. The mobile phase consisted of
114	methanol with 0.1% acetic acid (mobile phase B) and water with 0.1% acetic acid (mobile phase
115	A). The gradient was 20% solvent B increasing to 65% B in 15 min, increasing to 90% B in 3
116	min, kept at 90% B for 2 min, decreasing to 20% B in 2 min and kept at 20% B for 3 min.
117	
118	Purification of bicyclopentadione (BCP) from TDP
119	BCP, the most abundant curcumin degradation compound, was purified from TDP using a
120	Shimadzu 2020 preparative HPLC system equipped with automatic fraction collector (Shimadzu,
121	Marlborough, MA). The HPLC condition was the same as described above. The fractions
122	containing BCP were combined and extracted with ethyl acetate. The organic layer was dried
123	over anhydrous magnesium sulfate and evaporated to dryness using a rotary evaporator. The
124	purity and structure of isolated BCP was confirmed by HPLC and 'H NMR respectively, as
125	described previously ^{13, 17} .
126	
127	Cell lines

128 MC38 colon cancer cells (a kind gift from Prof. Ajit Varki at the University of California San

129 Diego ¹⁸) and RAW 264.7 murine macrophage cells were used in cellular assays. Cells were

routinely cultured in Dulbecco's Modified Eagle Medium (DMEM, Lonza, Allendale, NJ)

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ACS Paragon Plus Environment

131	supplemented with 10% Fetal Bovine Serum (FBS, Corning Inc., Corning, NY) in a 37 °C
132	incubator under an atmosphere with 5% CO ₂ .
133	
134	Cell proliferation assay
135	MC38 cells (6000 cells/well) in DMEM medium were plated into 96-well plates and allowed to
136	attach overnight. Cells were treated with 10-40 μ g/mL test compound or DMSO vehicle (0.1%)
137	for 24, 48, and 72 h. Cell proliferation was assessed by 3-(4,5-dimethylthiazol-2-yl)-2,5-
138	diphenyltetrazolium bromide (MTT) assay.
139	
140	Flow cytometry analysis of cell cycle and apoptosis
141	MC38 cells (200,000 cells/well) in DMEM medium were plated in 6-well plates and allowed to
142	attach overnight. Cells were treated with 5 μ g/mL test compounds or DMSO vehicle (0.1%) for
143	24 h. For cell cycle analysis, cells were collected, washed by ice-cold PBS, and fixed in 70%
144	ethanol at -20°C overnight. Cells were then centrifuged, washed by PBS, and stained with
145	propidium iodide solution (10 μ g/mL propidium iodide, 100 μ g/mL RNAse) for 30 min at room
146	temperature. For apoptosis analysis, cells were collected, washed by ice-cold DMEM fortified
147	with 2% FBS medium, centrifuged, and stained with propidium iodide and Annexin V solution
148	(10 μ g/mL propidium iodide, 0.5 μ g/mL Annexin V) for 10 min at room temperature. After
149	staining, cells were analyzed using the BD LSRFortessa™ cell analyzer (BD Biosciences) and
150	data were processed using the FlowJo software.
151	
152	Griess assay of LPS-induced nitric oxide (NO) production in RAW 264.7 cells
153	RAW 264.7 cells in 96-well plates were treated with 2.3-9.2 $\mu\text{g}/\text{mL}$ test compounds and 0.5
154	μ g/mL LPS for 24 h, then NO concentration in the cell culture medium was analyzed by a Griess
155	assay as previously described ¹⁹ , and cell proliferation was measured by MTT assay.

157 Immunoblotting

158 Immunoblotting was conducted using standard procedures. Briefly, after cellular treatments in RAW 264.7 cells, the medium was decanted and cells were washed with cold PBS buffer then 159 160 lysed. Nuclear extracts were collected using a nuclear extraction kit (Cayman Chemicals, Ann 161 Arbor, MI) according to the manufacturer's instructions. Cell lysates were resolved using SDS/PAGE and transferred onto nitrocellulose membranes. Membranes were blocked with 5% 162 163 BSA (Sigma Aldrich, St. Louis, MO) for 1 h at room temperature then probed with antibodies against iNOS, COX-2, IκBα, phospho-IKK, NF-κB p65 (Cell Signaling Technology, Danvers, 164 MA), Iamin A/C (Santa Cruz), and β -actin (Sigma-Aldrich). Membranes were then probed with 165 LI-COR IRDye[®] 800CW Goat anti-Rabbit and IRDye[®] 680RD Goat anti-Mouse secondary 166 antibodies. The Odyssey imaging system (LI-COR, Lincoln, NE) was used to detect the 167 168 antibodies. ImageJ software was used for western blot quantification.

169

Data Analysis 170

All cell culture experiments have been performed in at least three independent experiments. 171

The results are expressed as means \pm standard deviation (SD), where each of the mean is 172

derived from an independent experiment. All statistical calculations were performed using 173

174 ANOVA followed by Tukey's honest significance difference test with the assistance of R

statistical software. If a significant difference among the interaction between treatment and dose 175

was found, the interaction was separated by either treatment or dose and analyzed again with 176

ANOVA followed by Tukey's. Statistical significance was considered significant at p < 0.05. 177

178

- **Results** 180
- 181 Preparation of curcumin, total degradation products of curcumin, and BCP

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We chemically synthesized curcumin, since most commercial samples of curcumin contain other curcuminoids such as demethoxycurcumin and bisdemethoxycurcumin. The purity of the synthesized curcumin was >99% as assessed by HPLC and NMR. To study the biological activities of curcumin degradation products, we prepared the total degradation products of curcumin (TDP), which was a mixture containing all stable degradation products of curcumin after incubating curcumin in phosphate buffer, and BCP, which is was the most abundant degradation compound of curcumin and was purified from TDP by HPLC (**Figure 1**).

189

Curcumin degradation products have weaker anti-proliferative effects than curcumin 190 We compared the effects of curcumin, TDP, and BCP on cancer cell proliferation. MC38 colon 191 cancer cells are among the most widely used cells to study colon carcinogenesis ¹⁸. Curcumin 192 193 potently inhibited MC38 colon cancer cell proliferation in a dose- and time-dependent manner: 194 at a dose of 10 µg/mL, curcumin inhibited ~60% of MC38 proliferation after 24 h treatment, and inhibited >95% of MC38 proliferation after 48-72 h treatment (Figure 2). This is consistent with 195 previous studies which showed potent anti-proliferative effects of curcumin in various cancer 196 197 cells. In contrast to the potent action of curcumin, the degradation products of curcumin, 198 including TDP and BCP, had dramatically reduced anti-proliferative effects. For example, TDP 199 at a dose of 10 µg/mL had no inhibitory effect on MC38 proliferation after 24 h treatment, and 200 only inhibited ~20% of cell proliferation after 48-72 h treatment. BCP was completely inactive to inhibit cancer cell proliferation in the tested dose range (10-40 µg/mL) after 24-72 h treatment 201 (Figure 2). Together, these studies showed that curcumin, but not its stable autoxidation 202 products, inhibited cancer cell proliferation. 203

204

Curcumin degradation products have weaker effects on cell cycle progression and
 apoptosis than curcumin

207 We further compared the effects of curcumin, TDP, and BCP on two critical processes involved 208 in regulation of cell proliferation: the progression of cell cycle and apoptosis. After 24 h 209 treatment, curcumin significantly induced G2 cell cycle arrest and apoptosis in MC38 colon 210 cancer cells; in contrast, neither TDP nor BCP had such effects (Figure 3). These results further 211 support that curcumin, but not its degradation products, played a major role in modulating 212 cancer cell proliferation and associated cellular responses. 213 214 Curcumin degradation products have weaker anti-inflammatory effects than curcumin We compared the effects of curcumin, TDP, and BCP on LPS-induced inflammatory responses 215 in RAW 264.7 macrophage cells, which are widely used to study inflammatory responses. 216 Consistent with previous studies, curcumin inhibited LPS-induced NO production in a dose-217 218 dependent manner. In contrast, TDP had a much weaker inhibitory effect and BCP had no effect 219 (Figure 4A). Consistent with the NO result, curcumin dose-dependently inhibited LPS-induced 220 expression of iNOS, which is the major enzyme involved in production of NO; while TDP or BCP 221 had no effect (Figure 4B). We also found that curcumin, but not its degradation products, 222 inhibited LPS-induced expression of inflammatory protein COX-2 (Figure 4C). Together, these 223 results showed that curcumin, but not its degradation products, have anti-inflammatory effects. 224 225 Curcumin degradation products have weaker inhibitory effects on NF-kB signaling than 226 curcumin We compared the effects of curcumin, TDP, and BCP on LPS-induced NF-kB signaling in RAW 227 264.7 macrophage cells. Consistent with previous studies, curcumin inhibited LPS-induced IKK 228 229 phosphorylation, IkBa degradation, and nuclear translocation of p65, demonstrating that 230 curcumin inhibited LPS-induced activation of NF-kB signaling. In contrast, neither TDP nor BCP 231 had such inhibitory effect (Figure 5). Together, these results further suggest that curcumin, but not its degradation products, have anti-inflammatory effects. 232

233 Discussion

Due to the rapid degradation of curcumin at physiological pH, recent research suggests that the

resulted degradation products, including the reactive and unstable degradation products, could

contribute to the observed biological activities of curcumin ^{20, 21}. Here in this study, we compared

the anti-proliferative and anti-inflammatory effects of curcumin with its stable degradation

products. Our central finding is that compared with curcumin, the stable degradation products of

239 curcumin such as TDP and BCP have dramatically reduced biological effects in vitro,

suggesting that these stable degradation products are not likely to play a major role in mediating

the anti-proliferative and anti-inflammatory effects of curcumin.

242

Chemical degradation of curcumin in aqueous buffer occurs through multiple mechanisms. The 243 244 first identified mechanism is the hydrolysis pathway: a hydroxyl ion in the aqueous buffer attacks 245 the carbonyl group of curcumin and generates break-down products such as vanillin, ferulic acid and feruloyl methane ¹⁴. Substantial studies have investigated the biological activities of 246 hydrolysis-derived degradation products of curcumin, and found that these compounds such as 247 248 vanillin and ferulic acid have dramatically reduced biological activities compared with curcumin. For example, in terms of cell proliferation in peripheral-blood mononuclear cells, the IC₅₀ value 249 of vanillin was ~260 times higher than that curcumin 22 . In addition, recent studies showed that 250 the hydrolysis-derived compounds are minor degradation products of curcumin^{13, 17, 21}. 251 Therefore, it is unlikely that these hydrolysis-derived degradation products contribute to the 252 biological activities of curcumin. 253 254 255 Besides the hydrolysis pathway, recent research showed that curcumin degradation is mainly 256 mediated by a radical-dependent autoxidation pathway, leading to formation of cyclized compounds such as BCP^{13, 17, 21}. The roles of the autoxidation pathway in biological activities of 257

curcumin remain largely unknown, and here we studied the biological effects of stable

259 autoxidation-derived degradation products of curcumin. In our experiment, we treated cells with 260 curcumin in cell culture medium containing 10% FBS, at this condition curcumin is relatively stable (half-life > 8 hours, based on previous study 14). Therefore, the biological effects of 261 262 curcumin observed in our study, in particular the NF-κB assays in which the cells were treated 263 curcumin for 1-2 hours, were at least in part mediated by intact curcumin. Consistent with previous studies for the anti-cancer and anti-inflammatory effects of curcumin²³, our results 264 showed that curcumin inhibited cell proliferation, and modulated cell cycle progression and 265 266 apoptosis in MC38 colon cancer cells, and inhibited LPS-induced inflammatory responses (NO, 267 iNOS, COX-2, and NF-kB signaling) in RAW 264.7 cells. In contrast, its stable autoxidation products such as TDP (mixed degradation products of curcumin, in which BCP is the most 268 abundant compound) and purified BCP have dramatically reduced effects in these cellular 269 270 assays. In particular, BCP, the most abundant compound in TDP, was completely inactive in 271 these assays. TDP still has some effects to suppress cell proliferation and production of NO. this effect is at least in part mediated by the remaining curcumin. Previous studies have 272 characterized the chemical components of curcumin degradation products in detail ^{13, 17, 21}, 273 274 though we need to address that due to the difference in terms of preparation method, the 275 degradation products prepared by us could have some unidentified compounds. The different effects of curcumin and BCP are likely due to the presence and number of α , β -unsaturated 276 carbonyl groups in their chemical structures. Curcumin has two α , β -unsaturated carbonyl 277 groups, which are thiol-reactive and have been suggested to be the pharmacophore of curcumin; 278 in contrast, BCP has no reactive α , β -unsaturated carbonyl group, which could in part explain 279 why this compound has dramatically reduced anti-proliferative and anti-inflammatory effects 280 compared with curcumin. Due to the lack of potent effects of TDP and BCP, it is not likely that 281 282 these compounds play a major role in mediating the activities of curcumin. We need to point out 283 that here we studied the anti-proliferative and anti-inflammatory effects of stable degradation products of curcumin. It is likely that during the radical-dependent autoxidation process of 284

curcumin, some highly reactive intermediates could be formed and could play a role in the
biological activities of curcumin ²¹. In fact, a study showed that curcumin alone or BCP could not
increase topoisomerase II-mediated DNA cleavage. When combining curcumin and potassium
ferricyanide, an oxidizing agent, cleavage was increased, suggesting that curcumin's biological
effects could be due to reactive and unstable intermediates ²⁴. Further studies are needed to
better characterize the molecular mechanisms of curcumin.

291

292 Our current finding is consistent with our recent study which showed that when curcumin degradation is suppressed, the anti-proliferative effects of curcumin are significantly enhanced 293 ¹⁵. Indeed, our recent study showed that the co-addition of a various redox active antioxidants, 294 295 dramatically enhanced curcumin stability in aqueous buffer. We further showed that when MC38 296 cells are treated with curcumin in basal cell culture medium (without FBS), curcumin rapidly 297 degraded and had a weak anti-proliferative effect while co-addition of these redox active antioxidants, which enhanced chemical stability of curcumin, significantly increased the anti-298 proliferative effect of curcumin¹⁵. Together, these results suggest that curcumin, but not its 299 300 degradation products, has potent anti-proliferative effects. Co-addition of antioxidants increases 301 levels of curcumin and reduces levels of its degradation products in cellular systems, leading to 302 enhanced activities. It remains to test whether co-addition of antioxidants could be used as a practical strategy to enhance the effects of curcumin in cell culture and animal models. 303

304

In conclusion, here our results showed that the stable degradation products of curcumin,

including TDP (mixed degradation products of curcumin, in which BCP is the most abundant

307 compound) and purified BCP have dramatically reduced anti-proliferative and anti-inflammatory

308 effects compared with curcumin, suggesting that these compounds are not likely to contribute to

the effects of curcumin. A limitation of the study is that we only performed our study in cultured

310 cells with short treatment time. It is important to validate the results in animal models, in

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311	narticular with	long_term	treatment of	curcumin to	mimic	human co	nsumntion c	of curcumin
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- 312 Previous study has shown that long-term treatment of curcumin has potent biological effect on
- 313 inflammation, cancer, and immune responses ²⁵⁻²⁷. Further studies are needed to understand
- how curcumin exerts various beneficial effects despite its poor chemical and metabolic stability.
- 315

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325 Figure Legends

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327 Figure 1. Preparation and analysis of curcumin, total degradation products of curcumin (TDP),

and bicyclopentadione (BCP). (*A*) Scheme to prepare curcumin, TDP and BCP, and (*B*) HPLC

analysis of curcumin, TDP, and BCP.

330

Figure 2. Curcumin degradation products have weaker anti-proliferative effects than curcumin.

332 MC38 cells were treated with curcumin, TDP, BCP, or DMSO vehicle (0.1%) in complete DMEM

medium for 24-72 h and cell proliferation was assessed by MTT assay. The results are

expressed as % of cell viability of compound-treated cells to vehicle-treated cells, expressed as

mean \pm SD from at least three independent experiments. * P < 0.05.

336

Figure 3. Curcumin degradation products have weaker effects on cell cycle progression and apoptosis than curcumin. MC38 cells were treated with 5 μ g/mL of curcumin, TDP, BCP, or DMSO vehicle (0.1%) in complete DMEM medium for 24 h, then cell cycle progression (*A*) and apoptosis (*B*) were analyzed. *Left*: representative images of flow cytometry analysis. *Right*: quantification of cell population. The results are mean ± SD from at least three independent experiments, * P < 0.05.

343

Figure 4. Curcumin degradation products have weaker anti-inflammatory effects than curcumin. RAW 264.7 cells were treated with curcumin, TDP, BCP, or DMSO vehicle (0.1%) and 0.5 μ g/mL LPS in complete DMEM medium for 24 h. (*A*) NO in cell culture medium, the results are expressed as % of NO production of compound-treated cells to vehicle-treated cells. (*B-C*) expressions of iNOS and COX-2, *Top panel*: representative western blot, *Bottom panel*: quantitative analysis of protein expression, the results are mean ± SD from at least three independent experiments, * P < 0.05.

351	
352	Figure 5. Curcumin degradation products have weaker inhibitory effects on NF-kB signaling
353	than curcumin. RAW 264.7 cells were pre-treated with curcumin, TDP, BCP, or DMSO vehicle
354	(0.1%) for 1.5 h, then stimulated with 0.5-2 $\mu g/mL$ LPS in complete DMEM medium for 20-30
355	min. (A) representative western blots of phospho-IKK, IKB α , and nuclear p65, (B) quantitative
356	analysis of protein expression, the results are mean \pm SD from at least three independent
357	experiments, * P < 0.05.
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