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RESEARCH ARTICLE

Genotoxicity Study of Carboxymethyl Chitosan-based Hydrogel for Clinical Use

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ABSTRACT

A carboxymethyl chitosan-based hydrogel used for hemostatic and anti-adhesive on wounds was assessed by a genotoxicity test. The hydrogel was assessed for its potential to induce bacterial reverse mutation in the histidine auxotroph strains of *Salmonella typhimurium* and in the tryptophan auxotroph strain of *Escherichia coli* WP2uvrA. The study was performed with five bacterial test strains at different concentrations of hydrogel, with a negative control, and with five different types of positive controls, both in the presence and absence of a metabolic activation system. In all test strains used, there was no increase in the number of revertant colonies compared to the negative control at any concentration of test item either in the presence of or in the absence of a metabolic activation system. To evaluate additional genotoxicity of carboxymethyl chitosan hydrogel, the chromosome aberration test was performed using a cultured Chinese hamster lung cell line in the absence and presence of a metabolic activation system. The results of the bacterial reverse mutation assay indicated no mutagenic response at the concentration range tested, under the conditions of this study. The chromosome aberration test showed no significant increase of chromosome aberration in cells in metaphase arrest compared to the groups treated with vehicle control. Therefore, the carboxymethyl chitosan-based hydrogel is considered not to induce genotoxicity under the test system utilized.

## Introduction

Biocompatible hydrogels are a promising material for use as wound dressing and for hemostasis in the clinic<sup>1-3</sup>. Current development of such hydrogels is focused on adding antibacterial, antimicrobial, antioxidant, and wound debridement functions<sup>4-7</sup>. The advancement of hydrogels will result in more effective treatment of patients with acute traumatic wounds as well as chronic ulcers<sup>8-10</sup>. Before use of new chemically modified hydrogels in the clinic, extensive biological and chemical characterization is required<sup>11-13</sup>.

Various types of chitosan hydrogel have been used for wound management because of its desirable binding properties with growth factors, mechanical properties, and compatibility for combination with other gel types such as gelatin, pectin, and polyvinyl alcohol<sup>14-16</sup>. Chitosan is insoluble in water or alcohol but soluble in acid, with strong hydrogen bonding properties making the chemical structure stable<sup>17,18</sup>. Chitosan-based hydrogels are bioabsorbable, biocompatible, possess low toxicity, and can be made bioactive by binding with growth factors<sup>19,20</sup>.

A chemically modified chitosan (carboxymethyl chitosan) has been recently used for wound management and hemostasis<sup>21-23</sup>. Carboxymethyl chitosan is biocompatible, bioabsorbable, non-toxic, highly viscous, easy to synthesize and applicable for additional functions<sup>24,25</sup>. Moreover, carboxymethyl chitosan is soluble in water at neutral pH<sup>26</sup>. The similarity of its chemical structure to hyaluronic acid, makes carboxymethyl chitosan useful for visco-supplementation, anti-bacterial therapy, hemostasis, and wound closure<sup>27,28</sup>.

Carboxymethyl chitosan, a chemically modified chitosan-based hydrogel, may therefore be a promising material for wound management and hemostasis. The most recent applications of carboxymethyl chitosan derivatives showed effectiveness of antimicrobial, anticancer, antitumor, antioxidant and antifungal biological activities in various areas<sup>22</sup>. Even the clinical trial showed good clinical performance on hemostasis and anti-adhesion after endoscopic surgery<sup>29</sup>. This chemically modified chitosan is not only effective on external wound management but also promising for hemostasis and anti-adhesion during endoscopic and open surgeries.

We recently assessed carboxymethyl chitosan hydrogel for genotoxicity as an initial test of biocompatibility. Genotoxicity refers to the ability of a novel chemical compound to induce alterations at the genetic level. The *in vitro* and *in vivo* tests can be designed to evaluate compounds which induce genetic damage directly or indirectly

by various mechanisms. These tests should enable a hazard identification with respect to damage to DNA and its fixation. This study therefore addresses the genotoxicity of the chemically modified hydrogel, carboxymethyl chitosan-based hydrogel for potential clinical use in wound management and hemostasis.

## Methods

### Preparation of carboxymethyl chitosan hydrogel

Chitin (10% w/v) was mixed with 1N HCl for 1 week. After 1 week, the chitin was rinsed with water adjusted to pH 7 and dried at room temperature. The dried chitin was deacetylated by hydrolysis of the acetamide groups with NaOH (40% w/v) at 100°C for 3 hours to form chitosan. To produce carboxymethyl chitosan, chitosan was mixed with NaOH (60% w/v, 25mL), ClCH<sub>2</sub>COOH (14.5g) and CH<sub>3</sub>CH(OH)CH<sub>3</sub> (100mL) at 50°C for 7 hours (Hemostop/Surgishield, D-med, Seoul, Korea).

### Test strains and media for bacterial reverse mutation test

All test strains and media were purchased from Molecular Toxicology Inc. (MOLTOX, Boone, NC).

Base pair substitution types of *Salmonella typhimurium* such as TA100, TA1535, and *Escherichia coli* (E. Coli) WP2uvrA, and the frameshift type of *Salmonella typhimurium* such as TA98 and TA1537 were used as test strains as they are known to be very sensitive to mutagens and widely used in mutagenicity studies with bacteria<sup>30</sup>.

Each 15mL of 2.5% Oxoid nutrient broth No.2 was inoculated with test strain from the master plate and incubated in a 37°C shaking incubator at 180 rpm for 10 hours. The minimal glucose agar plates were prepared with 1.5% Difco® Bacto® agar, Vogel-Bonner medium E, and 2% glucose; 25ml dispensed per 100x15 mm petri dish. Top agar was prepared with 0.6% agar and 0.5% NaCl. This agar mix was appropriately supplemented with either 0.5mM of histidine-biotin for the *Salmonella* strains or 0.5mM tryptophan for *E. coli* strain. 0.07mL of Dimethyl sulfoxide (DMSO) was added to each 0.8mL of strain after overnight sub-culture and the mixture was stored in a deep freezer until use. Master plates of each strain were made and kept in a refrigerator to be used for each assay<sup>31</sup>.

### A metabolic activation system I

Rat liver post mitochondrial fraction (S9, Protein contents: 40.4mg/mL) with cofactor-I (S9

mix, Content of 1mL including 0.1mL S9, 8mmol MgCl<sub>2</sub>, 33umol KCl, 5mmol Glucose-6-phosphate, 4mmol NADPH, 4mmol NAHD, and 100mmol pH 7.4 Sodium phosphate) was used for a metabolic activation system (UMU genotoxicity test kit, MOLTOX). S9 fraction was prepared from male rats induced with Aroclor-1254, the xenobiotic metabolizing enzyme.

### Treatment of hydrogel extracts I

Carboxymethyl chitosan gel was incubated in saline (4g/mL) at 37° for 72 hours for testing extracts. Extracts of 0.1mL were tested with 0.5mL of S9 mix (or sodium phosphate buffer, pH 7.4), and 0.1mL of bacterial culture per test tube. The mixture was shaken for 20 minutes using a shaking incubator at 180 rpm, and then dispensed into top agar before being poured onto a minimal glucose agar plate (one tube per plate). Vehicle control groups were treated with vehicle only and each of the appropriate strain-specific positive control groups as well. The testing extracts (0.1mL) and S9 mix (0.1mL), without bacterial culture, were also plated to check sterility. After the top agar was solidified, the plates were inverted, and incubated at 37°C for about 48 hours. Then the revertant colonies were counted. All testing was performed in triplicates per concentration. The results were expressed as the mean numbers of colonies ± standard deviation from triplicate plates per dose. A positive interpretation was a reproducible increase in the number of colonies in a dose-dependent manner, in least in one test strain with or without the metabolic system, clearly exceeding 2 times the colonies in each of control<sup>32</sup>.

### Test cell line and media for chromosome aberration test

Chinese hamster lung cell line (CHL cells, ATCC, Manassas, VA) was used for the study. The culture media was Minimum Essential Medium (MEM, Gibco, Waltham, MA) supplemented with 10% fetal bovine serum (FBS, Gibco, Waltham, MA). The CHL cell was subsequently subcloned every 3-4 days<sup>33</sup>.

### A metabolic activation system and treatment of hydrogel extracts II

S9 mix was made of 0.3mL S9 (Protein contents: 31.9mg/mL), 5mmol MgCl<sub>2</sub>, 33mmol KCl, 5mmol Glucose-6-phosphate, 4mmol HEPES, and 4mmol NADH. The testing extracts were used directly at 0.5mL. 2x10<sup>4</sup> cells were plated on T-25 flasks and after 4 days, testing extracts (the same as the "hydrogel extracts I") were added in the 4.5mL culture media and incubated for 6 and 24

hours each. The cells were incubated for 6 hours, washed with 9mL PBS and cultured in the fresh culture media for 18 hours. All the cells were cultured for total of 24 hours. Saline was used as a negative control, Vehicle I, Mitomycin C (MMC, Sigma-Aldrich, St. Louis, MO), and Cyclophosphamide monohydrate (CPA, Sigma-Aldrich, St. Louis, MO) were used for each positive control I (S9-) and positive control II (S9+). All positive controls were dissolved in sterilized water and immediately used after preparation for testing. After 24-hour incubation, cells were treated with 100mL Colcemid (Thermo Fisher Scientific, Waltham, MA). After 2 hours, cells were collected and fixed onto a clean glass slide. Each slide was stained with 5% Giemsa solution for the chromosome aberrations. Two slides were prepared from each plate.

The analysis of 100 cells in mitotic and metaphase per plate was classified as follows. The frequencies of aberrations represented the frequencies of cells bearing aberrations. Evaluations were performed blindly using coded slides.

### Criteria for observation<sup>31</sup>

- Gap: The unstained region (assumed achromatic region) is located on the vertical axis of the chromatid, the width of which is about the width of the chromatid. The shape of that region should be clearly observed.
- Break: The broken part of the chromosome is not aligned with the vertical axis of the chromatid. If the part is located on the same axis, the achromatic region is two times as wide or wider than the width of the chromatid. An acentric chromosome is recorded only when it is clearly observed.
- Exchange: This is the conjugated form of two or more detached segments on one or more chromosomes, resulting from chromosome breakage. With respect to chromosome type, clearly recognizable examples such as a ring or dicentric chromosome if observed are recorded.
- Other fragmentation: There are many instances of gaps and chromosomal breaks excluding the exchange form.
- Numerical aberration: Aneuploidy is not considered and only polyploidy is documented since the CHL cell line tends to generate chromosome aberrations to some extent. The modal chromosome number of the CHL cell line is 25. Although the modal number of quadriploid is 50, polyploidy is

regarded in case the triploid has greater than 37 rods. In addition, as endoreduplication is classified as polyploid, it is important to record in case of a frequent outbreak.

**Evaluation of the test results**

For the evaluation of the frequencies of structural aberrations and of polyploidy, the following criteria, usually used for chromosomal aberration testing with CHL, were employed.

- Negative (-): less than 5%
- Equivocal ( $\pm$ ): from 5% to less than 10%
- Positive (+): 10% or more

In case there is no dose-dependent increase, or the frequency of chromosome aberrations is more than

3% in a negative control group, a re-test is performed.

**Results**

From bacterial reverse mutation test, growth of bacteria was not observed during the sterility test of the solutions for testing extracts and S9 mix. The test was performed on saline and DMSO extracts, negative and positive control both in the presence (S9+) and absence (S9-) of S9 mix.

In all test strains used, there was no increase in the number of revertant colonies compared to the negative control at any concentration of testing extracts either in the presence or in the absence of a metabolic activation system. The positive controls induced a marked increase in the numbers of revertant colonies (Table 1).

**Table 1.** Result of bacterial reverse mutation test

Test Strain	Test Sample	Dose	Colonies/plate (mean $\pm$ SD) [Factor]*	
			Without S9 mix	With S9 mix
TA98	Testing extracts	Saline Sample	26 $\pm$ 4 26 $\pm$ 6 [1.0]	26 $\pm$ 7 26 $\pm$ 4 [1.0]
TA100	Testing extracts	Saline Sample	147 $\pm$ 9 153 $\pm$ 5 [1.0]	138 $\pm$ 14 143 $\pm$ 7 [1.0]
TA1535	Testing extracts	Saline Sample	19 $\pm$ 2 16 $\pm$ 2 [0.8]	10 $\pm$ 4 10 $\pm$ 2 [1.0]
TA1537	Testing extracts	Saline Sample	12 $\pm$ 4 13 $\pm$ 4 [1.1]	17 $\pm$ 3 19 $\pm$ 4 [1.1]
WP2uvrA	Testing extracts	Saline Sample	37 $\pm$ 5 42 $\pm$ 5 [1.1]	35 $\pm$ 7 38 $\pm$ 6 [1.1]
<b>Positive controls</b>				
TA98	AF-2	0.1	331 $\pm$ 39 [12.7]	
TA100	AF-2	0.01	577 $\pm$ 32 [3.9]	
TA1535	NaN3	0.5	317 $\pm$ 24 [16.7]	
TA1537	9-AA	80.0	721 $\pm$ 20 [60.1]	
WP2uvrA	AF-2	0.01	274 $\pm$ 30 [7.4]	
TA98	2-AA	0.5		250 $\pm$ 15 [9.6]
TA100	2-AA	1.0		614 $\pm$ 11 [4.4]
TA1535	2-AA	2.0		179 $\pm$ 17 [17.9]
TA1537	2-AA	2.0		260 $\pm$ 15 [15.3]
WP2uvrA	2-AA	10.0		226 $\pm$ 20 [6.5]

\*Factor: No. colonies of treated plate / No. of colonies of negative control plate

From chromosome aberration tests, without a metabolic activation system (S9-), in the 6-hour treatment, chromosome aberration frequencies were 0.5%, 1.0%, and 45.5% at the concentration levels of negative control, test extracts, and positive control, respectively (Table 2). In the 24-hour treatment, chromosome aberration frequencies were 0.5%, 0.5%, and 39.0% at the concentration levels of negative control, test extracts, and positive control, respectively. All the results for the test

extracts were less than 5% which was evaluated as negative based on our test criteria (Table 3).

With a metabolic activation system (S9+), chromosome aberration frequencies were 0.0%, 0.5%, and 54.5% at the concentration levels of negative control, test extracts, and positive control, respectively. All the results for the test extracts were less than 5% which was evaluated as negative based on our test criteria (Table 4).

**Table 2.** Result of Chromosome aberration test – without metabolic activation (S9-, 6 hours)

Treatment	Numbers of cells showing structural aberration								
	Numbers cells showing polyploid	Gap	Chromatid type		Chromosome type		Others	Total	
			Chromatid break	Chromatid exchange	Chromosome break	Chromosome exchange		-Gap	+Gap
Negative control	0	0	1	0	0	0	0	1	1
	0	1	0	0	0	0	0	0	1
	0 (0.0)	1 (0.5)	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.5)	2 (1.0)
Test sample	0	0	1	0	0	0	0	1	1
	0	1	1	0	0	0	0	1	2
	0 (0.0)	1 (0.5)	2 (1.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	2 (1.0)	3 (1.5)
Positive control (MMC)	0	3	19	24	0	0	0	43	46
	1	2	21	27	0	0	0	48	50
	1 (0.5)	5 (2.5)	40 (20.0)	51 (25.5)	0 (0.0)	0 (0.0)	0 (0.0)	91 (45.5)	96 (48.0)

MMC, mitomycin C; ( ), Average

**Table 3.** Results of Chromosome aberration test – without metabolic activation (S9-, 24 hours)

Treatment	Numbers of cells showing structural aberration								
	Numbers cells showing polyploid	Gap	Chromatid type		Chromosome type		Others	Total	
			Chromatid break	Chromatid exchange	Chromosome break	Chromosome exchange		-Gap	+Gap
Negative control	0	1	0	0	0	0	0	0	1
	0	1	1	0	0	0	0	1	2
	0 (0.0)	2 (1.0)	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.5)	3 (1.5)
Test sample	0	1	1	0	0	0	0	1	2
	0	1	0	0	0	0	0	0	1
	0 (0.0)	2 (1.0)	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.5)	3 (1.5)
Positive control (MMC)	1	4	18	20	0	0	0	38	42
	0	3	16	24	0	0	0	40	43
	1 (0.5)	7 (3.5)	34 (17.0)	44 (22.0)	0 (0.0)	0 (0.0)	0 (0.0)	78 (39.0)	85 (42.5)

MMC, mitomycin C; ( ), Average

**Table 4.** Result of Chromosome aberration test – with metabolic activation (S9+, 6 hours)

Treatment	Numbers of cells showing structural aberration								
	Numbers cells showing polyploid	Gap	Chromatid type		Chromosome type		Others	Total	
			Chromatid break	Chromatid exchange	Chromosome break	Chromosome exchange		-Gap	+Gap
Negative control	0	0	0	0	0	0	0	0	0
	0	1	0	0	0	0	0	0	1
	0 (0.0)	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.5)
Test sample	0	1	1	0	0	0	0	1	2
	1	1	0	0	0	0	0	0	1
	1 (0.5)	2 (1.0)	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.5)	3 (1.5)
Positive control (MMC)	0	3	22	31	0	0	0	53	56
	0	2	21	35	0	0	0	56	58
	0 (0)	5 (2.5)	43 (21.5)	66 (33.0)	0 (0.0)	0 (0.0)	0 (0.0)	109(54.5)	114(57.0)

MMC, mitomycin C; ( ), Average

## Discussion

Chitosan is bioabsorbable and biocompatible, and can be bound to growth factors<sup>19,20</sup>. Chitosan has bacteriostatic, antibacterial, hemostatic, wound healing, and scar reduction properties<sup>21,27,34,35</sup>. However complications have been reported because it can induce food allergy and anaphylactic shock<sup>36</sup>. On

the other hand, it has been reported that chemically modified carboxymethyl chitosan has properties similar to chitosan but is relatively safe from the side effects of chitosan, making it potentially more suitable for wound management and hemostasis for clinical use<sup>37,38</sup>. The purpose of this study was to assess the genotoxicity of carboxymethyl chitosan-

based hydrogels through bacterial reverse mutation test and chromosomal aberration test.

Chitosan is an abundant polycationic biopolymer derived mainly from chitin of crustaceans by deacetylation of amino groups<sup>39</sup>. The strong hydrogen bonding makes the crystal structure very stable<sup>40</sup>. Chitosan has many useful biological properties leading to applications for wound healing and dressing, drug delivery, blood anticoagulant, antitumor agent, and artificial skin<sup>41</sup>. However, chitosan has some limitations in its use<sup>23,36,39,40,42,43</sup>. First, chitosan has low solubility in water at neutral pH and general organic solvents and is soluble only in acidic solutions with low pH. This acts as a limiting factor for the widespread use of chitosan. Second, certain crustacean sources of chitosan, such as shrimp and crab, are known allergens, which limits the use of chitosan products in patients with crustacean allergies.

Carboxymethylation makes chitosan hydrophilic and soluble at pH 6.0. Carboxymethyl chitosan is well known for its biocompatibility and relative non-toxicity, and for being bio-absorbable, highly viscous, and easily synthesized, making it applicable for the clinical use. Carboxymethyl chitosan is easy to sterilize because it has high temperature resistance<sup>44</sup>. Chitosan has been reported to have side-effects such as anaphylactic shock and food allergy. However, carboxymethyl chitosan is non-toxic and bio-absorbable, and has a molecular structure similar to that of hyaluronic acid making it suitable for joint viscosupplementation, anti-bacterial treatment, hemostasis, wound management, and scar healing<sup>27,28</sup>. Carboxymethyl chitosan has also been used for hydration and as artificial bone, artificial skin, and as a drug carrier<sup>45-51</sup>.

In this study, carboxymethyl chitosan hydrogel was evaluated for its potential to induce reverse mutation in four histidine auxotroph strains of *Salmonella typhimurium* (TA98, TA100, TA1535, TA1537) and one tryptophan auxotroph strain of *Escherichia coli* (WP2uvrA). The experiments were performed with the five test strains at concentration levels (each 4g/mL at 37°C for 72 hours) of saline and DMSO extracts, negative and positive control both in the presence (S9+) and absence (S9-) of S9 mix. We did not detect any bacterial contamination from the test item and S9 mix used in the test. We did not find any significant increase in the number of revertant colonies compared to its vehicle control at any dose in any of the strains. In addition, no antibacterial effects such as a

decrease in the number of colonies were observed in any of the strains. On the other hand, the positive controls induced a marked increase in the number of revertant colonies, as expected. These evaluations indicate that carboxymethyl chitosan hydrogel does not result in reverse mutations.

To evaluate chromosome aberration of carboxymethyl chitosan hydrogel, the test was performed using Chinese hamster lung cell line cultured in the absence (S9- Mix, 6 and 24 hours) and presence (S9+ Mix, 6 hours) of a metabolic activation system. Under the conditions of our study the carboxymethyl chitosan hydrogel did not induce statistically significant increases in the frequencies of cells with aberration in either the presence (S9+) or the absence (S9-) of a metabolic activation system (S9 Mix). Chromosome aberration frequencies were less than 5%.

Several studies have investigated the safety of carboxymethyl chitosan since Tokura et al.<sup>37</sup> first reported on the low toxicity of carboxymethyl chitosan. Yang et al.<sup>38</sup> did not detect acute in vivo toxicity in the blood system of rats after carboxymethyl chitosan was absorbed in the abdominal cavity and gradually degraded in the blood. It has also been reported that carboxymethyl chitosan is a functional biomaterial without cytotoxicity<sup>21</sup>. Fu et al.<sup>28</sup> reported that carboxymethyl chitosan implanted in the abdominal cavity of rats did not induce any significant differences in coagulation function, anticoagulant performance, fibrinolytic function, and hemorheology of rats, which supports its safety for biomedical application. Although we did not test the hydrogel in vivo, our study is consistent with these previous studies and documents the absence of significant genotoxicity.

## Conclusions

Based on the bacterial reverse mutation and chromosome aberration tests, carboxymethyl chitosan hydrogel showed no evidence of mutagenic potential at the concentration range tested in four test strains of *Salmonella typhimurium*, or in *Escherichia coli* strain WP2uvrA under the conditions of this study. Carboxymethyl chitosan hydrogel also did not have a clastogenic effect on CHL cells in vitro.

## Conflicts of Interest Statement

The authors have no conflicts of interest to declare.

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