

総説、著書

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Ⅲ. 研究成果の刊行物・別刷

Fluorophotometric measurement of the precorneal residence time of topically applied hyaluronic acid

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ABSTRACT

Purpose: This study was performed to separately assess the aqueous flow applied with hyaluronic acid, and the behaviour of hyaluronic acid itself on the ocular surface.

Methods: Two different fluorescent dyes, fluorescein sodium dissolved in 0.1% hyaluronic acid (HA) solution and 0.1% fluorescein conjugated with hyaluronic acid (F-HA) dissolved in saline, were used. A volume of 20 µl of tested solution was applied to the eye of 10 healthy volunteers. Fluorescein sodium dissolved in saline served as a control. The fluorescent intensity of the precorneal tear film was measured at the central cornea every minute for 10 min. The turnover rate was calculated using the equation that plots fluorescent intensity against time in a semilog plot and expressed as %/min.

Results: Turnover rates of topically applied 0.1% F-HA, 0.1% HA and saline were 8.1 (SD 3.6)%/min, 21.6 (2.8)%/min, and 31.0 (3.7)%/min, respectively. The turnover rate of F-HA was significantly lower than those of HA and saline ($p = 0.00012$ and $p = 0.0000022$, respectively; Mann-Whitney test). The turnover rate of HA was significantly lower than that of saline ($p = 0.00001$; Mann-Whitney test).

Conclusion: Our results indicate that the bulk aqueous flow applied with HA and the turnover of HA itself are different. HA molecules may adhere to the ocular surface by surface-chemical and/or biochemical properties. The long retention time of HA on the ocular surface may explain the mechanism in which hyaluronic acid has been shown to enhance tear film stability for a few hours.

Dry eye is a common condition, affecting approximately 10–20% of the adult population.¹ The clinical consequences of dry eye may include symptomatic irritation, superficial punctate keratopathy, corneal erosions and possibly visual acuity problems.² Dry eye is considered to be a disorder of the tear film due to tear deficiency or excessive evaporation, which causes damage to the interpalpebral ocular surface and is associated with symptoms of ocular discomfort.³

A variety of treatment modalities have been used for the treatment of dry eye. The majority of these fall into the category of tear substitutes or replacements.⁴ Artificial tears, the most frequently used modality for the treatment of dry eye, may be effective in relieving symptoms in mild dry eye by replenishing deficient tear volume. However, in moderate and severe cases of dry eye, artificial tears alone are not enough to relieve the symptoms nor to improve superficial punctate keratopathy.⁴

Several preliminary reports in the early 1980s,^{5–7} several studies have reported that hyaluronic acid is able to improve the symptoms, signs and ocular surface damage associated with dry eye

syndrome.^{8–11} Hyaluronic acid is a glycosaminoglycan with a viscoelastic rheology. Its relatively high viscosity is believed to improve tear-film stability and to reduce washout from the ocular surface.¹² Hyaluronic acid enhances water retention on the corneal surface, and probably increases corneal wettability.¹³ In addition, hyaluronic acid promotes migration of corneal epithelial cells and accelerates the healing of corneal epithelial defects.^{14–16} Hyaluronic acid has thus become an important treatment modality for dry eye.

Conflicting results, however, have been obtained regarding the duration of the action of hyaluronic acid on the ocular surface. The residence time of topically applied hyaluronic acid assessed by the tear meniscus height and water evaporation rate from the ocular surface was less than 10 min, although this was significantly longer than that of phosphate-buffered saline.^{17–18}

Using an assessment of the tear-film breakup time, however, hyaluronic acid has been shown to enhance tear-film stability for more than a few hours.^{19–21} These observations suggest that hyaluronic acid remains on the ocular surface independent of the bulk aqueous flow.

In order to test this hypothesis, we used two different fluorescent dyes, fluorescein sodium dissolved in hyaluronic acid solution and fluorescein conjugated with hyaluronic acid dissolved in saline, in the current study. The former is a well-established dye used to assess the bulk aqueous flow, and the latter dye is used as a tracer to determine the behaviour of hyaluronic acid on the ocular surface. Although the residence time of topically applied hyaluronic acid has been investigated using ⁹⁹Tc^m as a tracer,^{22–24} we believe that this study is the first report to measure the residence time using a tracer that is associated with hyaluronic acid on the ocular surface.

SUBJECTS AND METHODS

Fluorescent dye and fluorophotometer

Fluorescein hyaluronic acid (F-HA, Mw; 800 000 Da) was purchased from Sigma-Aldrich (St. Louis, MO). F-HA, fluorescein conjugated with hyaluronic acid, was dissolved in a phosphate-buffered saline as 0.1% solution and used as a tracer of hyaluronic acid. Fluorescein sodium (0.001%; Sigma-Aldrich) and 0.1% hyaluronic acid (Mw; 800 000 Da, Sigma-Aldrich) dissolved in a phosphate-buffered saline was used as a tracer of the bulk aqueous flow in the presence of hyaluronic acid. Fluorescein sodium (0.001%) in a phosphate-buffered saline was used as a tracer of the bulk aqueous flow.

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A commercial slit-lamp fluorophotometer (Anterior Fluorometer FL-500, Kowa Co., Tokyo) was used. The illuminating light was focused as a 2-mm diameter circle on the surface of the cornea. The emitted light passed through a band-interference filter centred on 565 nm (half bandwidth 25 nm) and was directed to a photomultiplier tube with the band-interference filter centred on wavelengths 490 nm (half bandwidth 30 nm).

F-HA solution (0.5%) was diluted in a phosphate-buffered saline to produce sets of standards ranging from 0.001% to 0.5% in concentration for the calibration. A cuvette was constructed by gluing together two microscope slides and two cover glasses. The cover glasses were sandwiched by two microscope slides in order to provide space for the fluid layer to be 12–15 µm thick. A fresh one was made for each solution. Ten microlitres of calibrating fluids, containing 0.001–0.5% F-HA, was placed into a cuvette. The fluorescent intensity was measured by a slit-lamp fluorophotometer. The interaction of F-HA with the proteins was also tested using a phosphate-buffered saline containing 1% fetal bovine serum.

Measurement of residence time

Ten healthy volunteers (five male and five female) aged 27–44 years (33.8 (SD 6.8) years old, mean (SD)), who had no history of eye disease, except for refractive errors, were enrolled in this study. The principles of the World Medical Association Declaration of Helsinki were followed. The subjects received a full explanation of the procedures, and provided their informed consent for participation prior to the experiment. The protocol was approved by our institutional review board, and all subjects provided their written informed consent.

In our experiments, the subjects were seated in front of the fluorophotometer. The instrument was focused on the central cornea, and the background fluorescent intensity was measured. A volume of 20 µl of tested solution was applied to the eye with an Eppendorf micropipette without making contact. The subjects were then instructed to blink several times to ensure the mixing of the dye. The fluorescent intensity of the precorneal tear film was measured at the central cornea every minute for 10 min. Repeated measurements on different days were made in some subjects to evaluate the repeatability of the test.

The turnover rate is given by the following equation, which plots fluorescent intensity against time in a semilog plot:

$$F = F_0 \exp(-kt)$$

where F is the fluorescent intensity at time (t); F_0 is the fluorescent intensity at time zero; k is the turnover rate; and t is the time in minutes.²⁵ The turnover rate was calculated using the equation and expressed as %/min. The regression fit of the log of the fluorescent intensity was recorded as the regression coefficient.

In all cases of 0.1% F-HA and 0.1% hyaluronic acid, this regression became a straight line. In cases of saline, however, this regression sometimes showed a biphasic response: an initial faster and a subsequent lower turnover rate. When the turnover rate of saline became biphasic, the subsequent lower turnover rate was used as the flow rate of saline.²⁵

All results are presented as the mean \pm 1 standard deviation (SD). Statistical significance was calculated by comparing

results using the Mann-Whitney test. A value of $p < 0.05$ was considered to indicate statistical significance.

RESULTS

Calibration of F-HA

The calibration of the fluorescent intensities against the concentrations of 0.001–0.5% F-HA is shown in fig 1. The relationship between the fluorescent intensities and the concentrations of F-HA was linear ($r^2 = 0.995$). The data generated by this method were consistent and reproducible. The fluorescent intensities of F-HA were unaffected by the presence of 1% fetal bovine serum (data not shown).

Turnover-rate measurements

A typical result of turnover-rate measurements obtained from one subject is shown in fig 2. In the presented case, the fluorescent intensities of 0.1% F-HA decayed with time at a flow rate of 7.6%/min, which was lower than those of fluorescein sodium in 0.1% hyaluronic acid (19.4%/min) and in saline (28.1%/min).

The turnover rates of topically applied F-HA, hyaluronic acid and saline obtained from 10 subjects were 8.1 (3.6)%/min, 21.6 (2.8)%/min, and 31.0 (3.7)%/min, respectively (table 1). The turnover rate of F-HA was significantly lower than those of hyaluronic acid and saline ($p = 0.00012$, and $p = 0.0000022$, respectively; Mann-Whitney test). The turnover rate of F-HA was significantly lower than that of saline ($p = 0.00001$; Mann-Whitney test).

DISCUSSION

In the current study, we used two different fluorescent dyes, fluorescein sodium dissolved in hyaluronic acid solution and F-HA solution, to separately assess the aqueous flow applied with hyaluronic acid, and the behaviour of hyaluronic acid itself on the ocular surface. Our results indicate that there are two different aspects of the duration of topically applied hyaluronic acid.

Hyaluronic acid has a high-molecular-weight, naturally occurring glycosaminoglycan. Its relatively high viscosity is believed to reduce washout from the ocular surface.¹³ The residence time of topically applied hyaluronic acid has previously been investigated by using gamma scintigraphic methods.^{22–24} Snibson and associates^{22–24} reported that hyaluronic acid had prolonged ocular residence times in comparison

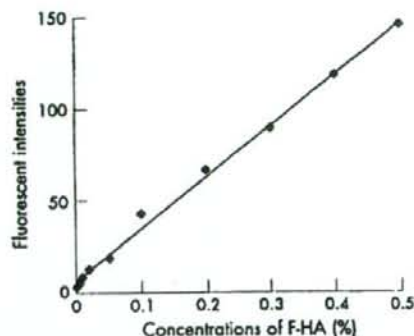


Figure 1 Calibration of the fluorescent intensities against the concentrations of fluorescein hyaluronic acid (F-HA). The relationship between the fluorescent intensities and the concentrations of F-HA was linear ($r^2 = 0.995$).

Table 1 Turnover rates of topically applied 0.1% fluorescein hyaluronic acid (F-HA), 0.1% hyaluronic acid and saline obtained from 10 subjects

Subject no.	Turnover rate (%/min)		
	F-HA	Hyaluronic acid	Saline
1	7.6	19.4	28.1
2	12.2	22.6	31.0
3	2.9	22.6	31.8
4	9.5	20.4	33.2
5	4.6	21.4	33.4
6	8.6	22.2	37.1
7	15.2	24.2	29.6
8	8.4	20.4	24.1
9	5.3	17.1	27.8
10	7.1	21.3	33.9
Mean (SD)	8.1 (3.6)	21.7 (2.8)	31.0 (3.7)

The turnover rate of F-HA was significantly lower than those of hyaluronic acid and saline ($p = 0.00012$ and $p = 0.0000022$, respectively; Mann-Whitney test).

with a buffered saline solution, a solution containing polyvinyl alcohol or hydroxypropylmethylcellulose. In the current study, the turnover rate of the 0.1% hyaluronic acid solution (21.6 (2.8)/min) was significantly lower than that of the saline (31.0 (3.7)/min). Our result is considered to be in good accordance with the previous studies using scintigraphic methods.²²⁻²⁴ This effect, however, appears to be transient, because 90% of the hyaluronic acid solution was calculated to be cleared from the ocular surface 10.7 min after instillation. This result is also consistent with the duration of topically applied hyaluronic acid assessed by the tear meniscus height and water evaporation rate from the ocular surface.^{17, 18}

The most interesting finding of the current study is that the turnover rate of F-HA (8.1 (3.6)/min) was approximately one-third of the 0.1% hyaluronic acid solution (21.6 (2.8)/min). This result indicates that the bulk aqueous flow applied with hyaluronic acid and the turnover of hyaluronic acid itself on the ocular surface are different. Besides viscosity, hyaluronic acid molecules may adhere to the ocular surface by surface-chemical and/or biochemical properties, because hyaluronic acid is known to bind with fibronectin and CD44, a cell surface adhesion molecule which has been found on corneal epithelial cells.¹⁴⁻¹⁶ Snibson and associates²⁴ made a similar consideration based on their scintigraphic results. They, however, also mentioned the limitation of their methodology and the necessity of tracers that directly associate with hyaluronic acid. We believe that F-HA is a useful tracer to determine the behaviour of topically applied hyaluronic acid. According to our

results, the times for 90% and 99% of hyaluronic acid to be cleared from the ocular surface were calculated to be 28.8 min and 57.6 min after instillation, respectively. The long retention time of hyaluronic acid in the ocular surface may explain the fact that hyaluronic acid has been shown to enhance tear-film stability for more than a few hours.¹⁹⁻²¹

Besides its biological effects on the corneal epithelial cells, hyaluronic acid appears to have two beneficial effects for the treatment of dry eye syndrome. First, it reduces the bulk aqueous flow by its viscosity and increases tear volume for a limited time, as do other viscous agents, such as chondroitin sulfate, polyvinyl alcohol and hydroxypropylmethylcellulose.²² Second, hyaluronic acid remains on the ocular surface for a longer time, in order to increase corneal wettability and to retain tear fluid on the corneal surface.¹²⁻¹³ This effect may be unique for hyaluronic acid, although it should be confirmed by further investigations.

Competing interests: The authors have no proprietary interest in any materials in this manuscript.

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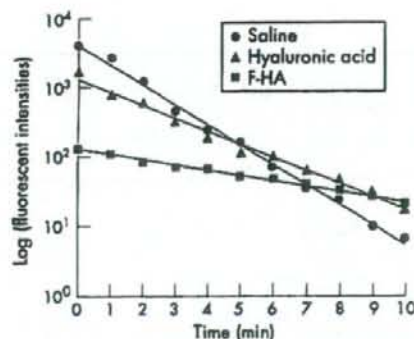


Figure 2 Typical result of the turnover rate measurements obtained from one subject. In the presented case, the fluorescent intensities of 0.1% fluorescein hyaluronic acid (F-HA) decayed with time at a flow rate of 7.6%/min, which was lower than those of fluorescein sodium in 0.1% hyaluronic acid (19.4%/min) and in saline (28.1%/min).

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In Vitro Susceptibilities of Bacterial Isolates From Conjunctival Flora to Gatifloxacin, Levofloxacin, Tosufloxacin, and Moxifloxacin

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Purpose. To evaluate and compare the in vitro susceptibilities of various fluoroquinolones (i.e., gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin) against conjunctival bacterial flora. **Methods.** Two hundred sixty-six bacterial isolates were collected from the conjunctival sacs of 251 eyes of 224 patients (118 females and 106 males) ranging in age from 6 to 91 years old, who were scheduled for intraocular surgery at National Tokyo Medical Center. The minimum inhibitory concentration (MIC) was determined by broth dilution testing. **Results.** Of 266 isolates, 258 (97.0%) strains were gram-positive bacteria and eight (3.0%) strains were gram-negative bacteria. The MIC₉₀ values of gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin against α -hemolytic streptococci were 0.39 μ g/mL, 1.56 μ g/mL, 0.39 μ g/mL, and 0.20 μ g/mL, respectively. The MIC₉₀ values of gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin against *Staphylococcus aureus* were 1.56 μ g/mL, 3.13 μ g/mL, 0.78 μ g/mL, and 0.78 μ g/mL, respectively. The MIC₉₀ values of gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin against *Staphylococcus epidermidis* were 1.56 μ g/mL, 3.13 μ g/mL, 3.13 μ g/mL, and 0.78 μ g/mL, respectively. **Conclusions.** Although the clinical usefulness and efficacy of newer fluoroquinolones remains to be defined by clinical outcomes, the current study provides data for predicting relative in vivo potency among the fluoroquinolones. **Key Words:** Bacterial flora—Conjunctiva—Drug resistance—Fluoroquinolone—Ocular infection.

Fluoroquinolones are the newest family of antibacterial agents used in the treatment of ocular infections.¹⁻⁵ In Japan, ofloxacin was the first fluoroquinolone introduced for topical ophthalmic use in 1987. Since then, six other fluoroquinolones, norfloxacin, lomefloxacin, levofloxacin, gatifloxacin, tosofloxacin, and moxifloxacin, have been approved for clinical use as eyedrops in Japan. In addition to these compounds, ciprofloxacin has been used clinically in other countries.

Double-masked, randomized clinical trials have shown that single-agent fluoroquinolone therapy using ofloxacin⁴ or cipro-

floxacin⁵ against bacterial keratitis is comparable in efficacy to combining fortified β -lactam agents and aminoglycosides. Their bactericidal activity against the most common gram-positive and gram-negative ocular pathogens is generally excellent, and their high potency has made fluoroquinolones a common choice for the treatment and prevention of ocular infections.

However, as with other antibiotic agents, continued use in a population raises the issue of emerging resistance.⁶ Since the introduction of fluoroquinolones for ophthalmic use, the reported incidence of in vitro resistance to fluoroquinolones among bacteria isolated from patients with bacterial keratitis and endophthalmitis has been steadily increasing.^{6,7} A previous study by the authors⁸ reviewed the database of bacterial flora cultured preoperatively from the conjunctival sac of 1,455 Japanese patients between 1995 and 1999. The incidence of in vitro resistance of bacterial isolates to ofloxacin increased from 13.5% in 1995 to 32.8% in 1999. Although ofloxacin was changed to levofloxacin in 2000, the incidence of resistance to levofloxacin gradually increased from 14.5% in 2000 to 20.5% in 2002.⁹

Some newer fluoroquinolones have been introduced for topical ophthalmic use: gatifloxacin, tosofloxacin, and moxifloxacin. They are sometimes categorized as third-generation fluoroquinolones (i.e., tosofloxacin) and fourth-generation fluoroquinolones (i.e., gatifloxacin and moxifloxacin).¹⁰ Although the clinical benefits of these newer fluoroquinolones have yet to be fully established, their attributes suggest a potential role for the prevention of the increasing incidence of fluoroquinolone resistance among bacterial ocular pathogens. Gatifloxacin and moxifloxacin, especially, which are called 8-methoxyfluoroquinolones, are less likely to engender resistance from single-step topoisomerase mutations. It requires a double mutation in DNA gyrase and topoisomerase IV to establish resistance to 8-methoxyfluoroquinolones.¹⁰ Other potentially beneficial features of 8-methoxyfluoroquinolones are enhanced gram-positive activity relative to older fluoroquinolones and improved drug delivery into the anterior segment of the eye.

This study compared the in vitro effectiveness of bacterial flora cultured preoperatively from the conjunctival sac of patients undergoing intraocular surgery to gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin.

MATERIALS AND METHODS

Two hundred sixty-six bacterial isolates were collected from the conjunctival sacs of 251 eyes of 224 patients (118 females and 106 males) ranging in age from 6 to 91 years old, who were scheduled

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TABLE 1. Total Number of 266 Ocular Clinical Isolates

Bacteria	No. of isolates
All isolates	266 (100%)
Gram-positive species	258 (97.0%)
α -Hemolytic streptococci	43
<i>Staphylococcus aureus</i>	56
Methicillin-sensitive <i>S. aureus</i>	48
Methicillin-resistant <i>S. aureus</i>	8
<i>Staphylococcus epidermidis</i>	140
Methicillin-sensitive <i>S. epidermidis</i>	59
Methicillin-resistant <i>S. epidermidis</i>	81
<i>Enterococcus faecalis</i>	14
Others	5
Gram-negative species	8 (3.0%)
<i>Pseudomonas</i> species	6
Others	2

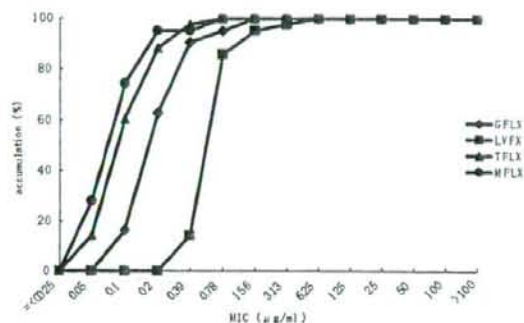


FIG. 1. The accumulation curves of the minimum inhibitory concentration (MIC) of the four fluoroquinolones against α -hemolytic streptococci ($n = 43$). Circle, moxifloxacin; triangle, tosofloxacin; diamond, gatifloxacin; square, levofloxacin.

for intraocular surgery at National Tokyo Medical Center from December 2004 to November 2005. The principles of the World Medical Association Declaration of Helsinki were followed. Each subject received a thorough explanation of the purpose of the study and all procedures involved in the study and provided written informed consent before enrollment. Approval for this investigation was granted by the Committee for the Protection of Human Subjects of National Tokyo Medical Center.

Scrapes of the inferior conjunctival fornix were taken preoperatively by using a sterile cotton swab without a topical anesthetic. The samples were immediately inoculated into the heart infusion bouillon, incubated for 24 hours at 37°C, and then inoculated into blood agar and MacConkey agar before incubating again for 24 hours at 37°C. Positive cultures were stored at -80°C until broth

dilution testing to determine the minimum inhibitory concentration (MIC).

For the broth dilution testing, frozen microdilution MIC plates with gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin were prepared according to the recommendations in the Clinical and Laboratory Standards Institute's Performance Standards for Antimicrobial Susceptibility Testing.¹¹ In this method, the concentration of antibiotic remains constant in each well, so organisms are exposed to identical concentrations for the duration of the test. Fluoroquinolone concentration ranged from 0.025 to 100 $\mu\text{g}/\text{mL}$.

The MIC of oxacillin (MIPIC) was also determined by the same method for *Staphylococcus aureus* and *Staphylococcus epidermidis*. When the MIC was greater than 4 $\mu\text{g}/\text{mL}$ for *S. aureus* and 0.5 $\mu\text{g}/\text{mL}$ for *S. epidermidis*, the strain was classified as methicillin-resistant *S. aureus* (MRSA) or methicillin-resistant *S. epidermidis* (MRSE).¹¹

RESULTS

Bacterial Isolates From Conjunctival Flora

Table 1 shows the type and frequency of the 266 ocular clinical isolates. Of the total 266 isolates, 258 (97.0%) strains were gram-positive and eight (3.0%) strains were gram-negative. *S. epidermidis* was the most common form of gram-positive bacteria with 140 strains, followed by 56 strains of *S. aureus*, 43 strains of α -hemolytic streptococci, and 14 strains of *Enterococcus faecalis*. Six of the eight strains of gram-negative bacteria were *Pseudomonas* species.

Antibacterial Activity of Fluoroquinolones

Figure 1 shows the MIC accumulation curves for the four study agents against α -hemolytic streptococci. The MIC₉₀ values of gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin were 0.39 $\mu\text{g}/\text{mL}$, 1.56 $\mu\text{g}/\text{mL}$, 0.39 $\mu\text{g}/\text{mL}$, and 0.20 $\mu\text{g}/\text{mL}$, respectively (Table 2). All strains of α -hemolytic streptococci were susceptible to all four study agents.

Figure 2 shows the MIC accumulation curves for the four study agents against *S. aureus*. Fifty-six isolates of *S. aureus* contained eight MRSA strains. The MIC₉₀ values of gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin were 1.56 $\mu\text{g}/\text{mL}$, 0.78 $\mu\text{g}/\text{mL}$, 0.78 $\mu\text{g}/\text{mL}$, and 0.78 $\mu\text{g}/\text{mL}$, respectively (Table 2). Almost all *S. aureus* strains susceptible to MIPIC were also susceptible to all four study agents, whereas MRSA strains were resistant to all four fluoroquinolones in general.

Figure 3 shows the MIC accumulation curves for the four study agents against *S. epidermidis*. One hundred forty isolates of *S. epidermidis* contained 81 MRSE strains. The accumulation curves of the MICs of all four fluoroquinolones were not linear but

TABLE 2. The Minimum Inhibitory Concentration of Four Fluoroquinolones Against Bacterial Isolates

Bacteria	MIC ₉₀ , $\mu\text{g}/\text{mL}$ (range of MIC)			
	Gatifloxacin	Levofloxacin	Tosofloxacin	Moxifloxacin
α -Hemolytic streptococci ($n = 43$)	0.39 (0.1-1.56)	1.56 (0.39-6.25)	0.39 (0.05-0.78)	0.20 (0.05-0.78)
<i>Staphylococcus aureus</i> ($n = 56$)	1.56 (0.05-50)	3.13 (0.1->100)	0.78 (\leq 0.025-25)	0.78 (\leq 0.025-50)
<i>Staphylococcus epidermidis</i> ($n = 140$)	1.56 (0.05-12.5)	3.13 (0.1-25)	3.13 (\leq 0.025-12.5)	0.78 (\leq 0.025-12.5)
<i>Enterococcus faecalis</i> ($n = 14$) ^a	0.39 (0.39-25)	1.56 (0.78-25)	0.2 (0.2-12.5)	0.2 (0.2-12.5)
<i>Pseudomonas</i> species ($n = 6$) ^a	0.78 (0.39-1.56)	0.78 (0.39-1.56)	0.2 (0.2-0.39)	1.56 (0.78-3.13)

^aFor *Enterococcus faecalis* and *Pseudomonas* species, the medians of the minimum inhibitory concentration (MIC) are listed because of insufficient sample numbers.

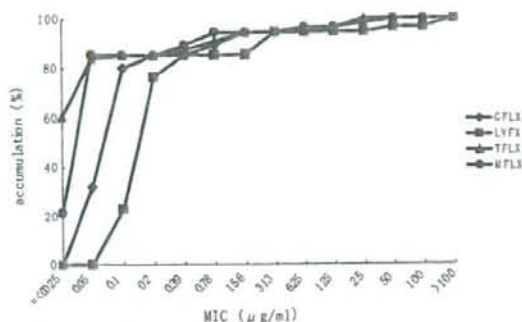


FIG. 2. The accumulation curves of the minimum inhibitory concentration (MIC) of the four fluoroquinolones against *Staphylococcus aureus* ($n = 56$). Triangle, tosofloxacin; circle, moxifloxacin; diamond, gatifloxacin; square, levofloxacin.

sigmoid and suggested the presence of drug-resistant strains. The MIC₉₀ values of gatifloxacin, levofloxacin, tosofloxacin, and moxifloxacin were 1.56 µg/mL, 3.13 µg/mL, 3.13 µg/mL, and 0.78 µg/mL, respectively (Table 2).

The MICs of fluoroquinolones and MIPIC were compared against *S. aureus* and *S. epidermidis*. The MIC of each fluoroquinolone was closely correlated with that of MIPIC for *S. aureus*, suggesting that MRSA was resistant to each fluoroquinolone. In contrast, there was no correlation between the MICs of MIPIC and the fluoroquinolones against *S. epidermidis* (Table 3).

DISCUSSION

One possible limitation of the current study is that samples originated from conjunctival cultures obtained from patients scheduled for intraocular surgery. Samples therefore should be regarded as conjunctival flora rather than ocular pathogens, because the patients did not have bacterial diseases at the time of sampling. However, in common ocular infections, such as bacterial conjunctivitis and bacterial keratitis, pathogens are frequently the normal bacterial flora residing on the ocular surface,^{12,13} even in cases of postoperative endophthalmitis, which is an infrequent but devastating form of ocular infection.¹⁴ In one study, organisms isolated from the vitreous were genetically identical to those collected from the ocular surface in 68% to 82% of patients with

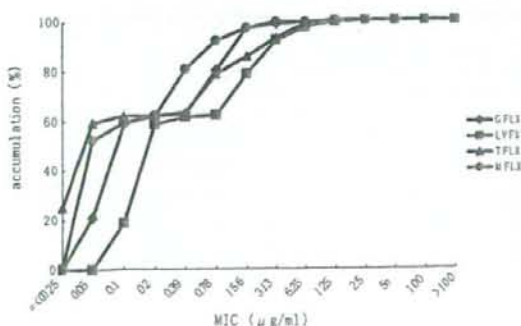


FIG. 3. The accumulation curves of the minimum inhibitory concentration (MIC) of the four fluoroquinolones against *Staphylococcus epidermidis* ($n = 140$). Triangle, tosofloxacin; circle, moxifloxacin; diamond, gatifloxacin; square, levofloxacin.

TABLE 3. Correlation Between the Minimum Inhibitory Concentration of Four Fluoroquinolones and Oxacillin Against *Staphylococcus aureus* and *Staphylococcus epidermidis*

Correlation versus oxacillin	Coefficient (P value)			
	Gatifloxacin	Levofloxacin	Tosufloxacin	Moxifloxacin
<i>Staphylococcus aureus</i> ($n = 56$)	0.85*	0.88*	0.85*	0.84*
<i>Staphylococcus epidermidis</i> ($n = 140$)	0.37	0.11	0.34	0.33

* $P < 0.0001$.

postoperative endophthalmitis,¹⁵ suggesting it is a valid approach to study in vitro susceptibility of bacteria isolated from conjunctival flora to various fluoroquinolones.

A serious concern is a drug resistance. In the current study, fluoroquinolones and MIPIC were compared with respect to their MICs against *S. aureus* and *S. epidermidis*. In *S. aureus*, the MICs of each fluoroquinolone were closely related to those of MIPIC, suggesting that MRSA was also insusceptible to most fluoroquinolones. Therefore, when ocular pathogens are identified as MRSA, the use of fluoroquinolones, even gatifloxacin and moxifloxacin, are not recommended for treatment. In contrast, there was no relationship between the MICs of MIPIC and the fluoroquinolones for *S. epidermidis*. These results indicate that some strains of *S. epidermidis* are not susceptible to fluoroquinolones, regardless of their susceptibility to MIPIC. Differences between *S. aureus* and *S. epidermidis* may be of interest when studying drug resistance mechanisms.

This study also found that some strains of *S. epidermidis* were highly resistant to levofloxacin and tosofloxacin, but not to moxifloxacin and gatifloxacin. These strains were considered to be low-level fluoroquinolone-resistant bacterial isolates. These data may favor the selection of moxifloxacin and gatifloxacin, rather than levofloxacin or tosofloxacin, for prophylactic use or the treatment of ocular infections.

In this study, newer fluoroquinolones showed more potent antibacterial activity than levofloxacin, the fluoroquinolone most commonly used in Japan. The current results support the contention that the fourth-generation fluoroquinolones have enhanced activity against gram-positive bacteria while retaining potency against most gram-negative bacteria. These newer fluoroquinolones have also improved penetration into the ocular tissue.¹⁰ Increased in vivo efficacy of these newer fluoroquinolones in some animal models of ocular infections have also been reported.¹⁶ Some caution, however, should be exercised when interpreting the data presented. The success or failure of a given therapy is not necessarily predicted by MIC₉₀ values, because the MIC₉₀ determined through in vitro assays may not directly correlate with clinical results. The actual efficacy of these newer fluoroquinolones remains to be defined by clinical outcomes. The data of the current study may be used to predict relative in vivo potency among the fluoroquinolones.

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Mutations in the quinolone resistance determining region in *Staphylococcus epidermidis* recovered from conjunctiva and their association with susceptibility to various fluoroquinolones

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ABSTRACT

Background: *Staphylococcus epidermidis* is one of the prominent pathogens in ocular infection. The prevalence of mutations in the quinolone resistance determining region (QRDR) area in *S epidermidis* isolated from the ocular surface and its association with fluoroquinolone resistance has not been fully elucidated.

Methods: Mutations in the QRDR of *gyrA*, *gyrB*, *parC*, and *parE* genes of 138 isolates of *S epidermidis* recovered from the human conjunctival flora were analysed. The minimal inhibitory concentrations (MICs) of four fluoroquinolones (levofloxacin, gatifloxacin, moxifloxacin and tosofloxacin) against these isolates were also determined using agar dilution methods.

Results: The MIC₉₀ values of levofloxacin, gatifloxacin, moxifloxacin and tosofloxacin were 3.13, 1.56, 0.78 and 3.13 µg/ml, respectively. The MIC values of all fluoroquinolones showed a bimodal distribution (susceptible strain and less susceptible strain). Mutations with amino acid substitution in the QRDR were present in 70 (50.7%) isolates. 19 different combinations of mutations were detected: 3 isolates (2.2%) had four mutations, 8 (5.8%) had three mutations, 43 (31.2%) had double mutations and 16 (11.6%) had single mutations. Isolates with mutations in the QRDR of both *gyrA* and *parC* ($n = 53$) were less susceptible to fluoroquinolones.

Conclusions: The present findings show that approximately half the *S epidermidis* isolates from the normal human conjunctiva have mutation(s) in the QRDR. The presence of mutations in both *gyrA* and *parC* is strongly associated with reduced susceptibility to fluoroquinolones.

Staphylococcus epidermidis is one of the most prominent causes of conjunctivitis, keratitis and endophthalmitis.¹⁻⁴ Although the relative frequency of different organisms as causative agents in keratitis varies during different periods and in different geographical regions, *S epidermidis* is among the most frequently encountered organisms in clinical studies conducted in the USA, Germany and Japan.²⁻⁴ It is the most common bacterial isolate in most large studies of acute postoperative endophthalmitis.⁵⁻⁸

The fluoroquinolones are the newest family of antibacterial agents used in the treatment of ocular infections.⁹⁻¹¹ In Japan, ofloxacin was the first fluoroquinolone introduced for topical ophthalmic use in 1987. Since then, six other fluoroquinolones—norfloxacin, lomefloxacin, levofloxacin (LVFX), gatifloxacin (GFLX),

tosofloxacin (TFLX) and moxifloxacin (MFLX)—have been approved for clinical use as eye drops in Japan. In addition to these compounds, ciprofloxacin has been used clinically in other countries. Their bactericidal activity against the most frequently observed Gram-positive and Gram-negative ocular pathogens is generally excellent and their high potency has made them a common choice for the treatment and prevention of ocular infections.

However, as with other antibiotic agents, continued use in a population raises the issue of emerging resistance.¹²⁻¹⁴ Since the introduction of fluoroquinolones for ophthalmic use, the reported incidence of in vitro resistance to fluoroquinolones in bacteria isolated from cases with bacterial keratitis and endophthalmitis has been steadily increasing. A previous study reviewed the database of bacterial flora cultured from the conjunctival sac of 1455 Japanese patients scheduled for intraocular surgeries between 1995 and 2002.¹⁴ The incidence of in vitro resistance of bacterial isolates to ofloxacin increased from 13.5% in 1995 to 32.8% in 1999. Moreover, when ofloxacin was replaced by LVFX in 2000, the incidence of resistance to LVFX gradually increased from 14.5% in 2000 to 20.5% in 2002.

The primary targets of fluoroquinolones are two essential enzymes of bacterial cells, DNA gyrase and topoisomerase IV.¹⁵⁻¹⁷ In *S epidermidis*, DNA gyrase is composed of the *GyrA* and *GyrB* subunits encoded by the *gyrA* and *gyrB* genes, respectively. Topoisomerase IV is composed of *ParC* and *ParE* subunits encoded by *parC* and *parE* genes, respectively. In most bacterial species, mutations occur in the highly conserved quinolone resistance-determining regions (QRDR) of the genes that encode DNA gyrase and topoisomerase IV. In *Staphylococcus aureus*, several studies have shown that a combination of mutations in both genes can cause high-level resistance even to the newer fluoroquinolones.¹⁸⁻²¹ However, the prevalence of mutations in the QRDR in *S epidermidis* isolated from the ocular surface and its association with fluoroquinolone resistance have not been fully investigated.¹⁵⁻¹⁷ The present study analysed mutations in the QRDR of *gyrA*, *gyrB*, *parC* and *parE* genes of 138 isolates of *S epidermidis* recovered from conjunctival flora. The susceptibility of these isolates to LVFX, GFLX, MFLX and TFLX was also determined.



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Table 1 Primers used in the study. Nucleotide positions are indicated according to GenBank sequence number NC 002976 (*S. epidermidis* RP62A)

Target gene	Primer sequence (5' to 3')	Product size (bp)	Position
<i>gyrA</i>	ATGCGTGAATCATCTTAGACTATGC	284	2 609 699–2 609 724
	GAGCCAAAGTTACCTTGACC		2 609 441–2 609 460
<i>gyrB</i>	CAGCATTAGACGTTCAAG	251	2 610 508–2 610 528
	CCAATACCCGTACCAATGC		2 610 278–2 610 297
<i>parC</i>	TCGCAATGATTCAAGTGGG	197	939 185–939 204
	ATCGTTATCGATACTACATT		939 361–939 381
<i>parE</i>	AAGCTCAACAAGCACGCGAGGCTG	324	938 196–938 219
	TTAAAGTCAGTACCAACCAGCAC		938 493–938 520

METHODS

Bacterial isolates and susceptibility testing

One hundred and thirty-eight isolates of *S. epidermidis* were collected from the conjunctival sac of 138 eyes of 129 patients who were scheduled for intraocular surgery at the National Tokyo Medical Center between November 2004 and June 2005. The mean (SD) age of the patients was 70.7 (14.9) years (range 6–91 years). The patients had not received either ophthalmic or systemic antibiotics prior to bacterial sampling.

Scrapes of the inferior conjunctival fornix were taken in the absence of topical anaesthetic using a sterile cotton swab. The samples were immediately inoculated into Mueller-Hinton (MH) agar and incubated at 35°C in air for 16–20 h for the selection of staphylococci. The MicroScan WalkAway-96 (Baxter Japan, Tokyo) with MicroScan Rapid Pos Combo Panel (Baxter) was used for the identification of *S. epidermidis*. Positive cultures were stored at –80°C until the agar dilution testing to determine the minimum inhibitory concentration (MIC).

MICs for LVFX, GLFX, MFLX and TFLX were determined by the agar dilution method in accordance with the recommendations of the Japanese Society of Chemotherapy.²³ The bacterial suspensions in saline were inoculated on MH agar plates supplemented with defined concentrations of drugs. The plates

were incubated at 35°C under aerobic conditions and MICs were determined after 20–24 h of incubation. Drug concentrations ranged from 0.025 µg/ml to 100 µg/ml in twofold increments except for TFLX (0.025 µg/ml to 25 µg/ml) because of its limited solubility.

DNA amplification and sequencing of QRDR

The isolates were suspended in tryptic soy broth and cultured overnight. Genomic DNA was extracted using the Wizard SV 96 genomic DNA purification system (Promega KK, Japan). One µl of the genomic DNA solution was applied in 20 µl of amplification mixture (5 pM each primer, 1.6 µl dNTP mixture, 2 µl Ex Taq buffer and 0.1 µl LA Taq (Takara Bio Inc, Japan)). Polymerase chain reaction (PCR) amplification was performed with the primers as shown in table 1. PCR primers were selected from the published sequences of *S. epidermidis* RP62A. Each reaction was amplified with the following temperature profiles: 30 cycles at 94°C for 30 s, 55°C for 30 s and 72°C for 1 min. The amplified DNA products were separated and identified by 2% agarose gel electrophoresis.

PCR products were purified using ExoSAP according to the manufacturer's instructions (GE Healthcare Bio-Sciences KK, Japan). PCR-amplified DNA was sequenced by the dye

Table 2 Mutations in the quinolone resistance determining regions of *gyrA*, *parC* and *parE* in 70 strains of *Staphylococcus epidermidis*

Mutation type	No of isolates	Mutation			
		<i>gyrA</i>	<i>gyrB</i>	<i>parC</i>	<i>parE</i>
1	28	Ser84Phe	–	Ser80Tyr	–
2	1	Ser84Phe	–	Ser80Tyr + Asp84Val + Ala85Ser	–
3	4	Ser84Phe	–	Ser80Phe	–
4	4	Ser84Phe	–	Ser80Phe + Asp84Tyr	–
5	1	Ser84Phe	–	Ser80Phe + Asp84Asn	–
6	2	Ser84Phe	–	Asp84Tyr	–
7	5	Ser84Tyr	–	Ser80Phe	–
8	3	Ser84Tyr	–	Ser80Phe	Asp434Asn
9	1	Ser84Tyr + Glu88Lys	–	Ser80Phe + Asp84Ala	–
10	2	Ser84Tyr	–	Ser80Tyr	–
11	1	Ser84Tyr	–	Ser80Ile	–
12	1	Ser84Ile	–	Ser80Phe	Asn404Ser + Asp434Asn
13	1	–	–	Ser80Tyr	Asn404Ser
14	1	–	–	Ser80Phe	–
15	1	–	–	Asp69Asn	–
16	1	–	–	Ser81Pro	–
17	1	–	–	Asp84Gly	–
18	11	–	–	–	Asn404Ser
19	1	–	–	–	Asp434Asn

Table 3 Susceptibility of strains of *S. epidermidis* to four fluoroquinolones

	No of isolates with the following MIC ($\mu\text{g/ml}$)										MIC ₅₀	MIC ₉₀	
	0.025	0.05	0.1	0.2	0.36	0.76	1.56	3.13	6.25	12.5			25
All isolates (n = 138)													
LVFX			26	55	4		23	19	7	3	1	0.2	3.13
GFLX		29	52	4	1	24	24	2	1	1		0.1	1.56
MFLX	1	72	9	3	26	16	7	3		1		0.05	0.78
TFLX	35	47	3		2	21	10	10	8	2		0.05	3.13
Wild type (n = 68)													
LVFX			24	44								Mode	0.2
GFLX		26	42										0.1
MFLX	1	60	7										0.05
TFLX	30	38											0.05
Mutations in <i>parC</i> and/or <i>parE</i> (n = 17)													
LVFX			1	11	4			1				Mode	0.2
GFLX		2	10	4			1						0.1
MFLX		11	2	3		1							0.05
TFLX	4	9	3			1							0.05
Mutations in both <i>gyrA</i> and <i>ParC</i> (n = 53)													
LVFX			1				23	18	7	3	1	Mode	1.56
GFLX		1			1	24	23	2	1	1			0.78
MFLX		1			26	15	7	3		1			0.39
TFLX	1				2	20	10	10	8	2			0.78

GFLX, gatifloxacin; LVFX, levofloxacin; MFLX, moxifloxacin; TFLX, tosufloxacin.

terminator method in both the forward and reverse directions. Using Phred/Phrap/Polyphred software, the quality score of each base was calculated. Sample sequences were compared with a reference sequence and mutations were detected. The strain *S. epidermidis* ATCC 35984 (RP62A) was used as a reference.

RESULTS

The mutations identified in the QRDR of the *gyrA*, *gyrB*, *parC* and *parE* genes are summarised in table 2. Nineteen different combinations of mutations were identified in 70 isolates, whereas no mutations were detected in 68 isolates. Three isolates (mutation profile type 2, 9 and 12) had four amino acid substitutions, 8 isolates (mutation profile type 4, 5 and 8) had three amino acid substitutions, 43 isolates (mutation profile type 1, 3, 6, 7, 10, 11 and 13) had double amino acid substitutions and 16 isolates (mutation profile type 14–19) had single amino acid substitutions.

In the *gyrA* gene, a single-point mutation was found in 53 isolates at codon 84. Double-point mutations in the *gyrA* gene were identified in 1 isolate at codons 84 and 88 (mutation profile type 9). No mutations were found in the QRDR area of the *gyrB* gene. In the *parC* gene, single-point mutations were found in 51 isolates at codons 69, 80, 81, 84 or 85. Double-point mutations were identified in 6 isolates at codons 80 and 84 (mutation profile type 4, 5 and 9). Triple-point mutations were identified in 1 isolate at codons 80, 84 and 85 (mutation profile type 2). In the *parE* gene, single-point mutations were found in 16 isolates at codon 404 or 434. Double-point mutations were identified in 1 isolate at codons 404 and 434.

The MICs of the four tested fluoroquinolones against *S. epidermidis* are shown in table 3. All four fluoroquinolones had a

bimodal distribution in all isolates (n = 138). Isolates with no mutations in the QRDR (wild type; n = 68) were susceptible to fluoroquinolones. The modes (the number that appears the most) were 0.2 $\mu\text{g/ml}$ for LVFX, 0.1 $\mu\text{g/ml}$ for GFLX, 0.05 $\mu\text{g/ml}$ for MFLX, and 0.05 $\mu\text{g/ml}$ for TFLX. Isolates with mutations restricted in the QRDR of *parC* and/or *parE* (n = 17) showed similar susceptibilities to fluoroquinolones as wild type strains except for one strain with mutation profile type 18. The modes were 0.2 $\mu\text{g/ml}$ for LVFX, 0.1 $\mu\text{g/ml}$ for GFLX, 0.05 $\mu\text{g/ml}$ for MFLX and 0.05 $\mu\text{g/ml}$ for TFLX. Isolates with mutations in the QRDR of both *gyrA* and *parC* (n = 53) were less susceptible to fluoroquinolones. The modes were 1.56 $\mu\text{g/ml}$ for LVFX, 0.78 $\mu\text{g/ml}$ for GFLX, 0.39 $\mu\text{g/ml}$ for MFLX and 0.78 $\mu\text{g/ml}$ for TFLX. Of these 53 isolates, 51 had amino acid substitutions at GyrA84 and ParC80. One isolate (mutation profile type 9) with two amino acid substitutions both in GyrA and ParC had the highest MICs (25 $\mu\text{g/ml}$ for LVFX, 12.5 $\mu\text{g/ml}$ for GFLX, MFLX and TFLX, respectively).

DISCUSSION

The primary targets of fluoroquinolones are two essential enzymes of bacterial cells, DNA gyrase and topoisomerase IV.^{18–20} In most bacterial species the mutations in the genes that lead to fluoroquinolone resistance are limited to a few point mutations at restricted positions of the genes called QRDR. The present study revealed that approximately half (50.7%) of *S. epidermidis* isolates in the human conjunctival flora have mutation(s) in the QRDR area of *gyrA*, *gyrB*, *parC* and *parE* genes.

Fluoroquinolone resistance has been studied intensively in *S. aureus*.^{18–21} The genes encoding topoisomerase IV in *S. aureus* are called *griA* and *griB*, which are analogous to *parC* and *parE* in *S. epidermidis*, respectively. Fluoroquinolone resistance in *S. aureus*

is generally associated with two single-point mutations in *gyrA* at codon 84, and in *griA* at codon 80 or 84. *S. aureus* isolates with higher levels of resistance are associated with the second mutation in *griA* at codon 80 or 84, depending on the position of the first mutation. When the second mutation in *gyrA* occurs at codon 85 or 88, in addition to the first mutation at codon 84, the strain shows the highest fluoroquinolone resistance even to newer fluoroquinolones.²¹

The present QRDR sequencing results indicate that the major mechanism of fluoroquinolone resistance in *S. epidermidis* is analogous to that of *S. aureus*. Isolates with mutations restricted to the QRDR of *parC* and/or *parE* ($n = 17$) in this study were similarly susceptible to fluoroquinolones as wild type strains. However, the presence of two mutations ($n = 53$) in both *gyrA* gene (located at codon 84) and *parC* gene (located at codon 80) have been found to be associated with the development of fluoroquinolone resistance.^{19, 16}

In this study only one isolate (mutation profile type 9), which was highly resistant to all four fluoroquinolones tested, had two amino acid substitutions both in GyrA and ParC. Previous studies have shown that isolates of *S. epidermidis* and *S. aureus* with two amino acid substitutions both in GyrA and ParC (GrlA in *S. aureus*) have the highest fluoroquinolone resistance. The isolates with this mutation type are reported to be relatively rare in *S. epidermidis*^{15, 16} and to account for less than 10% in *S. aureus*.^{18, 20} However, a high prevalence (50%) of two amino acid substitutions in both GyrA and GrlA has recently been reported.²¹ The empirical use of newer fluoroquinolones without a proper clinical indication may produce additional resistant strains of *S. epidermidis*, as has already occurred with *S. aureus*.

One possible limitation of the present study was that the patients were scheduled for intraocular surgery. Bacterial isolates therefore represent conjunctival flora rather than ocular pathogens. However, in common ocular infections such as bacterial conjunctivitis and bacterial keratitis, pathogens are frequently the normal bacterial flora that reside on the ocular surface.²⁻⁶ This is true even in cases of postoperative endophthalmitis, in which *S. epidermidis* is the most common bacterial isolate from vitreous aspirates.^{7, 8} Organisms isolated from the vitreous were genetically identical to those collected from the ocular surface in 68–82% of patients with postoperative endophthalmitis,⁷ suggesting that the study of in vitro susceptibility to various fluoroquinolones is valid.

Drug resistance is a serious concern in treating ocular infections. The current study showed that approximately half the *S. epidermidis* isolates from the conjunctival flora have mutation(s) in the QRDR. Both *gyrA* gene and *parC* gene are associated with the development of fluoroquinolone resistance.

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Ethics approval: The principles of the World Medical Association Declaration of Helsinki were followed. Each subject received a thorough explanation of the purpose of the study and all procedures involved in the study, and provided written informed consent prior to enrolment. Approval for this investigation was granted by the Committee for the Protection of Human Subjects at National Tokyo Medical Center.

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**Conjunctival Fixation Sutures for Refractory Superior Limbic
Keratoconjunctivitis**

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Running head: Fixation Sutures for Superior Limbic Keratoconjunctivitis

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Case Report

A 75-year-old Japanese woman was referred to us for persistent irritation in September 2006. She reported a history of foreign body sensation in the left eye for 4 years. Slit lamp examination revealed hypertrophy, hyperemia, and pronounced fluorescein staining of the superior bulbar conjunctiva in the left eye (Figure 1-A, B). Schirmer's test results were 2 mm in the right eye and 1 mm in the left eye. The diagnosis of superior limbic keratoconjunctivitis (SLK) with aqueous deficiency dry eye was made based on these findings. Initially, treatment was attempted with 0.1% fluorometholone and 0.1% sodium hyaluronate eyedrops without success. Punctal plugs inserted to the upper and the lower puncta provided little symptomatic relief.

We applied fixation sutures to the superior bulbar conjunctiva in December 2006 to attempt to treat this refractory condition. First, the patient received a topical and subconjunctival injection of lidocaine 2%. A traction suture was made by placing a 6-0 silk suture at the limbus at the 12 o'clock position, so as to rotate the eye downward (Figure 2). Stretching the redundant superior bulbar conjunctiva with a spatula, anchoring sutures were then placed using 10-0 nylon sutures at a location 10-12 mm from the limbus to fixate the conjunctiva with the sclera. Two stitches were placed nasally from the superior rectus muscle, and three stitches were placed temporally. Postoperatively, the patient received 0.5% levofloxacin and 0.1% fluorometholone eyedrops four times daily for 2 weeks. Two weeks after the procedure, the patient reported complete resolution of symptoms. Slit-lamp examination revealed a marked reduction in conjunctival hyperemia and fluorescence staining of the affected area (Figure 1-C,D). The patient's findings remained stable through over one year of follow-up.