ORIGINAL RESEARCH

Biofilm forming capacity and antibiotic susceptibility of *Staphylococcus spp*. with the icaA/icaD/bap genotype isolated from ocular surface of patients with diabetes

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Abstract

Introduction

Bacterial biofilm is an exopolysaccharide matrix that is produced by bacteria while they adhere on abiotic or biotic surfaces. The bacteria living in this matrix are more resistant to antibiotics than planctonic bacteria. The biofilm formation property of the bacteria is determined by genes; and this is related to virulence of the microorganism. In ophthalmology, biofilms form especially on abiotic surfaces such as silicon tubes, contact lenses, intraocular lenses etc.

Aim

Our aim was to investigate genotypic and phenotypic structures of biofilms that are produced by *Staphylococcus spp.*, which was obtained from the eyes of diabetic patients and determine the effect on antibiotic susceptibility.

Methods

The study group was comprised with 83 isolates from diabetic patients and 21 isolates from non-diabetic patients. Presumptive isolates were detected and confirmed by a microbial identification system VITEK II. Automated EcoRI Ribotyping was performed. Biofilm production was detected by Congo Red Agar Plate and Microtiter Plate Assay. Disc diffusion method was used for determination of antibiotic susceptibility of isolates.

Results

Out of the 83 isolates from diabetic patients, 25 were weakly (30%), 20 were moderately (24%), and 25 were strongly (30%) biofilm positive. Seven isolates of *S. aureus*, 11 isolates of *S. epidermidis*, 2 isolates of *S. warneri*, 3 isolates of *S. hominis*, and 2 isolates of *S. lugdunensis* were identified as strong biofilm producers. Out of the 83 *Staphylococcus* isolates, 37 were cefuroxime, 18 ciprofloxacin, 11 vancomycin, 12 gatifloxacin, and 18 moxifloxacin resistant. In total, 37 strains were resistant to three or more antibiotics. There was a statistically significant relation between biofilm formation and multidrug resistance (against three or more antibiotics, p<0.001). In nondiabetic patients, 15(71%) isolates were non adherent or weakly adherent, and 2(10%) were strongly adherent biofilm positive. **Conclusion**

In conclusion, bacterial conjunctival flora of patients with diabetes is likely to produce biofilm. Biofilm formation is associated with multidrug rsistance in patients with diabetes.

Keywords: Diabetes; biofilm; Staphylococcus spp.; antibiotic resistance; ocular surface

Introduction

The ocular surface has a number of defense mechanisms for the prevention of ocular infections. Protein constituents in tears (lysozyme, immunoglobulins, lactoferrin) as well as bacterial flora of the ocular surface have a major role restricting the growth of bacterial species¹. However, sources of many ocular infections, especially post-operative ones, are ocular flora members. For example, coagulase negative Staphylococci (CoNS) are by far the most isolated microorganisms from post-operative endophthalmitis²⁻⁴. Diseases that break down immunity such as diabetes mellitus (DM), acquired immunodeficiency syndrome (AIDS) and/or other systemic diseases may contribute to the occurrence of infections. In particular, DM affects cell mediated immunity and also lacrimal secretions^{5,6}. Under such circumstances, flora gain virulence, which can include biofilm production properties or antibiotic resistance. One of the main virulence factors for Staphylococci spp. predominantly found in the ocular flora is biofilm production. Biofilms, whose syntheses are

controlled by the intercellular adhesion (*ica*) genes, are non-homogeneous collections of bacteria which are bound together by an excreted matrix⁷⁻⁹. The enzyme N-acetylglucosaminyltransferase synthesizes polysaccharide intercellular adhesion (PIA), from UDP-N-acetylglucosamine in vitro that is encoded by the *i c a* locus and by the coexpression of *ica* A with *icaD* genes⁹. Biofilm formation of *Staphylococci*, which was isolated from infected eyes, has been reported previously^{10,11}. However, to our knowledge, this study is the first to show biofilm formation and genetic analysis of biofilms of *Staphylococcus* spp. that are isolated from the healthy conjunctiva of patients with diabetes. In this study we investigated genotypic and phenotypic structures of biofilm which are produced by *Staphylococcus* spp. and the effect of biofilms on antibiotic susceptibility.

Methods

Subjects

This study is a retrospective study. Isolates in this study were

obtained from previous studies which had been stored at – 80 C. Purification was done for this study. Eighty three isolates were obtained from 50 eyes of diabetic patients (24 female, 26 male; age range: 50-88 years) with at least 10 years duration of Type II DM and without any ocular infection or allergic symptoms, and also 21 isolates of 38 eyes of non-diabetic patients (19 female, 19 male; age range: 47-89 years). The study followed the principles of the Declaration of Helsinki. Sampling technique is explained elsewhere.¹²

Bacterial identification

The swabs obtained were inoculated on to mannitol salt, nutrient and blood agar plates. The incubation temperature and duration was 37°C and 24-48 hours respectively. Growing colonies were inoculated using the same media one by one for purification. Isolates that had been obtained from plates were identified using conventional (Gram stain, catalase, oxidase, coagulase, and DNAase reaction) and molecular microbiological methods. The strains were further identified with VITEK II system (BioMerieux, Durham, NC, USA) according to the manual of manufacturer. The identification ability of these systems depends upon the number and diversity of bacteria in the databases.

DuPont Qualicon RiboPrinter® Microbial Characterization System (Oxoid, Hampshire, UK) and the standard EcoRI DNA preparation kit were used for Automated EcoRI Ribotyping according to the manufacturer's guides. The reference DuPont identification database DUP2003 was used for comparison of ribotype profiles. Each isolate was identified when there was a similarity ≥ 0.85 between the corresponding pattern and the matching pattern of the DuPont identification Library. The ribogroups were made up of the isolates automatically by the RiboPrinterTM according to the the matching similarity of the ribotype patterns. The generated Finger Printing II software was used to analyse Riboprinter® and a dendrogram was created according to Unweighted Pair Group Method using arithmetic Averages (UPGMA) and Pearson correlation coefficients (optimization 1.56%). Strains were stored in 15% glycerol and at -80° C. Working cultures were stored at 5 C and transferred periodically.

Biofilm formation

Congo red agar (CRA) method: Biofilm formation in Staphylococcus strains was detected by growth on congo red agar (CRA) plates¹³. CRA plates that were inoculated were incubated for 24 hours at 37°C. Then, they were stored at room temperature for 48 hours. Appearance of the rough black colonies indicates the slime producing strains of coagulase-negative Staphylococcus. Microtiter plate assay (MPA): A microtiter assay was used to detect biofilm production as previously described¹⁴. Staphylococcus strains were added into 10 ml of tryptic soy broth (TSB) with 0.25% glucose. Then incubated for 24 hours at 37°C.Cultures were diluted at 1:100, diluted cultures (200 µl) per well were distributed into 96 well polystyrene microtiter plates and under aerobic conditions incubation was done at 37°C for 24 hours. After incubation, they were washed with sterile phosphate buffered solution (PBS) twice, fixed in ethanol (99%) for 15 minutes, and washed again with PBS. Then 200µl of 2% crystal violet per well was used to stain the

plates for 5 minutes. Excess stain was washed out and the plates were left to dry. The bond between the dye and the adherent cells was broken with the use of 160 ml of 33% (v/v) glacial acetic acid. The optic densitometry (OD) was measured at 570 nm by using an automated microplate reader. The reading was done before glacial acetic acid addition, as in standard microtiter plate, and after glacial acetic acid addition¹⁴. Classification of the results provided by the microtiter-plate test according to Christensen et al has three categories¹⁵. However Stepeanovic et al. modified the classification¹⁴. Strains were classified as follows: OD \leq OD_c non-adherent, OD_c < OD \leq 2xOD_c weakly adherent, 2xOD_c < OD \leq 4xOD_c moderately adherent, 4xOD_c< OD strongly adherent.

All tests were performed thrice and the mean of the results was calculated.

Scanning electron microscopy analysis

Scanning electron microscopy (SEM) (GeminiSem, Zeiss, Germany) biofilm specimens were prepared by fixation, staining, drying, and conductivity coating prior to imaging under high vacuum¹⁶. Firstly, bacteria were harvested and fixed in 2.5 % glutaraldehyde in 0.1 M cacodylate buffer for 2 hours at room temperature, post fixed in 1% OsO4 (Osmium tetroxide) for 1 hour. Then bacteria were dehydrated in a series of ethyl alcohol (30, 50, 70, 90 and 100 %), with each for 15 minutes. Samples were incubated in 100 % ethanol two times for 20 minutes. CO₂ was used to dry them to a critical point. After that, according to standard procedures, they were prepared by sputtering gold film on them. Specimens were investigated with a Zeiss Ultra 50 Scanning electron microscope (SEM) operated at 5 kV accelerating tension.

Antibiotic sensitivity test

Antibiotic sensitivity profile of the CoNS against antibiotics, which are commonly used as eye drops, were assessed by the disc diffusion method with respect to the guidelines of Clinical Laboratory Standards Institute¹⁷. Discs contained the following antibacterial agents: gatifloxacin (5 μ g), cefuroxime (30 μ g), ceftazidime (30 μ g), vancomycin (30 μ g), gentamicin (10 μ g), amikacin (30 μ g), ciprofloxacin (5 μ g), lomefloxacin (10 μ g), moxifloxacin (5 μ g), and methicillin (5 μ g).

Detection of icaA, icaD, and bap genes encoding for biofilm using PCR

Firstly, bacterial lysates were prepared and DNA was extracted. After PCR-amplification was achieved , examination by electrophoresis on agarose gel was done as described by Arciola et al¹⁸. Genomic DNA was extracted using Fermentas Gene JETTM Genomic DNA purification kit (Thermo Fisher Scientific, Waltham, MA, USA) based on manufacturer recommendations. For detection of genes regarding biofilm formation, gene specific primers were used for PCR amplification of DNA. The primer sequences and PCR length are shown in Table 1. The reaction volume was 25 μ l that contained 10X TaqBuffer (+KCl, –MgCl₂), 25 mM MgCl₂, 2.5 mM dNTP mix, 2.5 mM forward primer, 2.5 mM reverse primer, Taq DNA polymerase (5 u/µl), nuclease- free distilled water, and template DNA. The PCR conditions for *icaA* and *icaD*

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were the following: 5 minute initial denaturation at 94°C ; 35 cycles of 94°C for 30 s, 55°C for 30 s, and 72°C for 60 s; and a five minute final extension at 72°C¹⁹. PCR conditions for the bap gene were 94°C for 5 min; 35 cycles of 94°C for 30 s, 42°C for 30 s, and 72°C for 60 s; and a five minute final extension at 72°C²⁰.

Amplification products were analyzed using 1% agarose gel electrophoresis. A further nucleotide sequence of the amplicon for bap gene was determined by sequencing. Amplicon that was purified from gel (Promega Wizard® Gel-PCR Clean Up System, Promega BioSciences, LLC. San Luis Obispo, CA, USA) was sequenced by using DNA Beckman Coulter CEQ8000 Quick Start sequencing kit (Beckman Coulter, Fullerton, CA, USA). The data obtained from this sequencing procedure was compared with NCBI (National Center for Biotechnology Information) database by BLAST programme. Our sequencing data showed 92 % similarity with Staphylococcus epidermidis biofilm-associated protein (bap) gene, complete cds (GenBank Accession Number DQ008306.1).

Statistical analysis

For statistical analysis, the Statistical Package for Social Sciences version 17 (IBM, Chicago, USA) statistical programme was used. Pearson Chi-square was performed to compare qualitative data. In order to assess the relation between the parameters, Pearson correlation analysis was performed . P<0.01 and p<0.05 were considered statistically significant.

Results

In diabetic patients, 83 conjounctival strains were isolated from 50 eyes, and 21 conjunctival strains were isolated from 38 eyes of non-diabetic subjects. The isolates were presumptively identified as *Staphylococcus* spp. According to conventional tests, isolates of diabetic patients were members of the species Staphylococcus epidermidis, Staphylococcus aureus, Staphylococcus warneri, Staphylococcus lugdunensis, Staphylococcus hominis, and Staphylococcus haemolyticus. Fig. 1 shows the RiboPrinter® Microbial Characterization System (Dupont Qualicon) results. S. epidermidis, S. aureus, S. warneri, S. lugdunensis, S. hominis, and S. haemolyticus were detected with the RiboPrinter® system. Although these genotypic identification results were almost concordant with those of VITEC II results, they were detected as S. epidermidis in 38, S. aureus in 8, S. warneri in 19, and other coagulase negative staphylococcus in the remaining 18 isolates.

S. epidermidis were dominant among the isolates obtained from the conjunctiva of healthy patients with diabetes (Fig 2). From 17 eyes, 38 S. epidermidis conjunctival strains were isolated.

It was observed that 25 (30%) out of 83 isolates were weakly positive, 20 (24%) were moderately positive, and 25 (30%) were strongly positive, with the remaining 13 (16%)isolates biofilm negative with at least one of the methods used for detection of biofilm formation (Fig. 3).

According to conventional tests and the RiboPrinter® Microbial Characterization System, isolates of non-diabetic subjects were members of the species S. epidermidis, S. aureus,



Fig. 1: Riboprinter® microbial characterization system (DupontQualicon) results in diabetic patients showing ribotyping profiles of the isolates S. epidermidis, S. aureus, S. warneri, S. lugdunensis, S. hominis and S. haemolyticus



Fig. 2: Isolation rate of strains in healthy and type II diabetic cohorts

S. warneri, S. hominis, and S. saprophyticus (Fig. 2). Fifteen (71%) out of 21 isolates were non-adherent or weakly adherent, 4 (19%) were moderately adherent, and 2 (10%) were strongly adherent biofilm positives (Fig. 3).







Fig. 4: SEM microphotographs; A) A weak (S. epidermidis KA 14.5) biofilm producer strains, B) A strong (S. epidermidis KA 15.8) biofilm producer strains from diabetic patients

The rate of strong biofilm formation of *S. aureus* as the prominent pathogenic microorganism of the flora in the non-diabetic group was 20%, whereas the rate in the diabetic group was 88%. The examples of weak and strong biofilm formation by *S.epidermidis* colonies of diabetic patients are shown in Fig 4A and 4B in SEM microphotographs. Seven isolates of *S. aureus*, 11 isolates of *S. epidermidis*, 2 isolates of *S. marneri*, 3 isolates of *S. hominis*, and 2 isolates of *S. lugdunensis* from diabetics were found to be strong biofilm producers (Fig. 3).

The antibiogram study revealed that 5 isolates of *S. aureus*, 19 isolates of *S. epidermidis*, 15 isolates of *S. marneri*, and 9 isolates of *S. lugdunensis* in diabetic patients were resistant to methicillin (Fig. 5). Out of 83 *Staphylococcus* isolates, 37 were cefuroxime resistant, 18 were ciprofloxacin resistant, 9 were amikacin resistant, 11 were vancomycin resistant, 12 were gatifloxacin resistant, 16 were lomefloxacin resistant, and 18 were moxifloxacin resistant. There was no isolate resistance to gentamicin. In total, 37 strains (45%) were resistant to three or more antibiotics. Drug resistance against 6 or more antibiotics was detected in 11 (13%) isolates, against 7 or more in 3 (4%), and against 8 antibiotics in 1 (1%) (Fig. 5). The relation between biofilm formation and multidrug resistance (against three or more antibiotics)

was statistically significant (p < 0.001). In the biofilm positive coagulase negative staphylococcus strains, the resistance rate was high to methicillin (67%), cefuroxime (37%), and ceftazidime (41%), whereas the resistance rate was low against gentamicin (4%). In the biofilm negative S. epidermidis strains, the resistance rate was high to ciprofloxacin (14%) and the resistance rate was low to methicillin (3%). Biofilm producing S. epidermidis strains showed a significantly increased methicillin resistance (63%). Methicillin resistance was 50% in biofim-positive and 44% in biofilm-negative strains (Fig. 5). In non-diabetic subjects, 6 isolates were cefuroxime resistant, 8 were ceftazidime resistant, 5 were ciprofloxacin resistant, 2 were vancomycin resistant, 1 was gatifloxacin resistant, 3 were lomefloxacin resistant, and 4 were moxifloxacin resistant. There was no resistance to both gentamicin and amikacin. Only 1 isolate had resistance against 6 (5%) antibiotics. No resistance was detected against 7 antibiotics or more (Fig. 5).



Fig. 5: Antimicrobial resistance to antibiotics of biofilm-positive *Staphyloccocus spp.* strains in healthy and type II diabetic cohorts

All isolated strains from diabetic patients were positive for *icaA* and *icaD*. The biofilm-producing and non-producing *Staphylococcus* spp. had both genes, giving a 188-bp band for the *icaA* gene and a 198-bp band for the *icaD* gene (Fig. 6 A and B). The molecular weight marker kit was used to determine the size of standard bands, then the image analyzer system designated the expected lengths to the bands that were obtained by amplification of the extracted DNA.

DNA extracted from 8 *S. aureus*, 18 *S. warneri*, 5 *S. hominis*, 12 *S. lugdunensis*, 1 *S. haemolyticus*, and 39 *S. epidermidis* isolates were tested for *icaA*, *icaD* detection , and the bap gene by PCR using gene-specific primers (Fig 6 C). All isolates had *icaA* and *icaD* positivity, and one of the tested isolates was found to be bap positive. Sequencing was done for the bap gene. Sequencing of this amplicon showed 92% similarity with *S. epidermidis* biofilm-associated protein (bap) gene, complete cds (GenBank Accession Number DQ008306.1).

All of the isolates of non-diabetic subjects were *icaD* and *icaA* positive. No bap positivity was detected.



Fig. 6: PCR amplified product of (A) icaA gene. M; pBR322 DNA/ AluI Marker (Fermentas) (B) icaD gene. M; pBR322 DNA/AluI Marker (Fermentas) and (C) bap gene. M; DirectLoad 1 kb DNA Ladder (Sigma-Aldrich).

Discussion

Coagulase negative staphylococci are types of flora found in the eyelids, meibomian glands, and especially on the ocular surface. It has been shown that these bacteria enter the eye during and after intraocular surgery, which may cause endophthalmitis, a devastating complication of operations^{4,21-23}. However, this complication is rare because of the equilibrium between host defense, bacterial virulence, and antibiotic resistance. Recent studies show that one of the major virulence factors for bacteria, especially for staphylococci, is biofilm production. Microorganisms, which produce biofilms are also more prone to develop resistance to antibiotics^{7,24,25}.

The rate of strong biofilm formation in diabetic patients was 30%, whereas the rate is 10% in non-diabetic subjects. On the one hand, we found that, in diabetics, the rate of strong biofilm formation of *S. aureus* as the prominent pathogenic microorganism was 88% whereas it was 20% in the control group. This study may suggest that bacteria from the ocular surface in patients with diabetes are likely to produce biofilm. Moreover, non-biofilm producer strains are more susceptible to antibiotics than the biofilm producers.

In previous studies, the rates for biofilm production of *S. epidermidis* were 34-74% in infected eyes and 18-46% in healthy conjunctivas according to CRA methods^{10,11,26,27}. In this study, the biofilm production rate of all *Staphylococcus* and all CoNS are 77% and 74%, respectively, according to CRA methods and 79% in diabetics. Our rate is higher than the rate found in previous studies in which healthy eyes were evaluated and similar to those with ocular infections, though our samples were taken from healthy eyes. This may be attributed to the presence of diabetes in the cohort, probably facilitating the rate of biofilm production.

Studies concerning biofilm production from healthy eyes are limited in literature. Suzuki et al. reported that their biofilm production rate for *S. epidermidis* was 46% and they noticed that all black colonies in the CRA were *icaA* positive²⁶. Verdayes et al. also reported a lower rate (18%) for biofilm production; however their rates for *icaA* positivity for biofilm positive isolates were 88% and *icaD* positivity were $100\%^{27}$. In this study, both *icaA* and *icaD* positivity rates of biofilm plus strains are 100%. In addition to the ica genes, we investigated isolates for the bap gene, which we detected in one isolate. Also in non-diabetics, the rates for *icaA* and *icaD* positivity in all isolates is 100% but no positivity for bap is detected. Due to the small sample size of control subjects in this study, it was not feasible to compare the results of diabetic patients with the control subjects statistically. Therefore, we only presented the findings of the healthy control subjects.

We noticed a correlation between biofilm formation capacity and antibiotic resistance. We found there was no vancomycin resistance in non-biofilm producer strains; however, vancomycin resistance was found in 12 of the biofilm producer strains. For fluoroquinolones, which are frequently used as eye drops, resistance was also similar. In diabetics, the number of resistant strains in non-biofilm producers for ciprofloxacin was 5, lomefloxacin was 2, moxifloxacin was 1, and gatifloxacin was 0. These numbers for isolates in biofilm producer strains were 12, 15, 15, and 12, respectively. In non-diabetic subjects, similar to the nonbiofilm producer strains of diabetic patients, the number of resistant strains in biofilm producer strains for ciprofloxacin was 5, for lomefloxacin it was 3, for moxifloxacin it was 4, and for gatifloxacin it was 1.

We found that biofilm producing strains are all susceptible to gentamicin. Multidrug resistance was found to be related to biofilm formation. Antibiotic susceptibility rates of biofilm producing strains from patients with diabetes have not been previously published. Catalanotti et al. reported that biofilm producing microorganisms, which were isolated from the conjunctiva of soft contact lens users, had higher antibiotic resistance than non-biofilm producers¹⁰. They noticed that all S. epidermidis were susceptible to gentamicin, netilmicin, ofloxacin, neomycin, and kanamycin¹⁰. Alabiad et al. showed the fluoroquinolone resistance of conjunctival cultures to be 33% from patients who had undergone intravitreal injection²⁸. Blanco et al. reported that staphylococci isolates from chronic conjunctivitis were highly susceptible to vancomycin and moxifloxacin²⁹. Methicillin sensitive S. epidermidis had low susceptibility to ciprofloxacin and erythromycin. Methicillin resistant S. epidermidis was only susceptible to moxifloxacin and vancomycin²⁹. However, these studies did not mention the biofilm states of the isolates and whether these isolates were obtained from patients with diabetes or not. The relationship between biofilm formation and multidrug resistance has been reported previously^{26,31,32}.

We have isolated non-biofilm producing strains which were *icaA* and *icaD* positive. Similar findings were reported from Hou et al previously, and they noted that the bacterial strains required genetic capability for biofim production; it is not implied that the biofilms will certainly form¹¹.

Diabetes Mellitus (DM) is a systemic disease which damages cell-mediated immunity. Because of diabetes, retinopathy may develop. DM may also cause cataract. Besides these diseases, DM may cause some changes on the ocular surface and tear film⁶. Thus, microbial flora of the ocular surface varies. There are studies on the ocular surface flora of diabetic patients, which have shown that culture positivity and CoNS rates are higher in patients with diabetes; and the flora of the conjunctiva in diabetic patients is different from non-diabetic patients³³⁻³⁵. However, to the best of our knowledge, biofilm capacity of the ocular surface bacteria of patients with diabetes has not been reported before. In order to reduce or inhibit biofilm production in diabetic patients, specific strategies have not been defined yet. During biofilm formation, the process of

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bacterial attachment may be affected by species of bacteria, the surface condition, environmental factors (presence of glucose, etc.), growth medium, and essential gene products³⁶. Antibiotics inhibit biofilm formation *in vitro* when added into media at the same time as the microorganism, however the effect of antibiotics is limited on preformed biofilm³⁷. On the one hand, any agent that can change the viscoelastic property of the biofilm, chelating agents (EDTA), and polyamines (norspermidine) is helpful in inhibiting biofilm formation^{38,39}. Therefore, besides an adequate metabolic control of glucose in diabetic patients, any measure targeting one of these factors may be helpful in the future for controlling or inhibiting biofilm production.

In this study, we report the staphylococcal characteristics of the ocular flora. We found CoNS (90%) as a dominant group of the staphylococcal flora, and the main member of this group is *S. epidermidis* (45%). There are few studies about biofilm producing microorganisms that are isolated from patients' bodies with diabetes except the eyes. Podbielska et al. reported that 59% of *S. aureus* and 75% of *S. epidermidis* from feet of diabetic patients had produced biofilms.⁴⁰ However, they did not mention the *ica* status of the isolates.

Conclusion

Bacterial conjunctival flora of patients with diabetes may be more likely to produce biofilms. Biofilm formation is associated with multidrug resistance. Further studies with larger control groups may improve our understanding of biofilm formation and properties in diabetic patients. Patients with a long duration of diabetes should be evaluated carefully before planned surgery. It should be kept in mind that antibiotic resistance patterns may be vital for the success of surgery.

Conflicts of Interest

All contributing authors declare no conflicts of interest.

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