Full Length Research Paper

Effect of non-steroidal anti-inflammatory drugs and dexamethazone on the biofilm formation and expression of some adhesion-related genes of *Candida albicans* and *Staphylococcus aureus*

Rehab Mahmoud Abd El-Baky* and Sherein G. El-Gendy

1Department of Microbiology and Immunology, Faculty of Pharmacy, Minia University, Minia, Egypt.
2Department of Microbiology and Immunology, Faculty of Medicine, Assiut University, Assiut, Egypt.

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*Candida albicans* and *Staphylococcus aureus* can cause many diseases which are considered in many cases life threatening infections. Their biofilm formation on the surface of medical devices is difficult to be treated. Our study evaluates the effect of some anti-inflammatory drugs on the growth, biofilm formation and the expression of some adhesion-related genes. Antimicrobial activity of the tested drugs was determined by microbroth dilution method, their effect on biofilm formation was determined by crystal violet assay method, the mechanism of action was assessed by scanning electron microscope and ethidium bromide uptake assay. Their effects on the expression of *icaA*, *ALS1* and *HWP1* genes were determined by RT-PCR. Diclofenac Sodium had the highest antimicrobial activity followed by Meloxicam. Ethidium bromide uptake increased in the presence of diclofenac sodium in both *S. aureus* and *C. albicans* strains. All tested drugs showed significant inhibitory effect on both biofilm formation and the preformed biofilm. NSAIDs, dexamethazone and ketoconazole down-regulated *C. albicans* tested genes except for ketoprofen that up-regulated *HWP1* gene. NSAIDs and levofloxacin down-regulated the expression *icaA* gene but dexamethazone showed no effect on *icaA* gene expression. Although, dexamethazone had no antimicrobial activity, it had good anti-biofilm activity against *S. aureus* and *C. albicans*.

**Key words:** *Candida albicans*, *Staphylococcus aureus*, biofilm, *HWP1*, *ALS1*, *icaA*, Dexamethazone, NSAIDS.

INTRODUCTION

*Candida* spp. can cause opportunistic diseases especially in patients who are immune compromised, aged, receiving prolonged antibacterial and aggressive cancer chemotherapy or undergoing invasive surgical procedures and organ transplantation (Pfaller and Diekma, 2007). *C. albicans* readily forms biofilms on a wide variety of polymers used to make indwelling medical devices, such as dental materials, stents, prostheses,
implants, endotracheal tubes, pacemakers, and catheters (Douglas, 2003).

*S. aureus* is found to be common in polymicrobial biofilm infections especially with *C. albicans*. It is estimated that 27% of nosocomial *C. albicans* bloodstream infections are polymicrobial, with *Staphylococcus aureus* (Klotz and Chasin, 2007). Interestingly, the combined effect of *C. albicans* and *S. aureus* results in synergism and increased mortality in mice (Viale and Stefani, 2006).

Different studies have already described changes in gene expression levels during biofilm development (O’Gara, 2007). Adherence of microorganisms to an implanted device is a crucial step in the development of biofilm and biomaterial-entered infection (Seo et al., 2008). Adhesion of *C. albicans* to host cells depends on many adherence molecules such as the agglutinin-like sequence (ALS) family, hyphal wall protein (HWP) and cell wall glycoproteins. Also, the intercellular adhesin (ICA) operon is a virulence factor identified in staphylococci related to biofilm production. As *S. aureus* formation of biofilms requires the synthesis of polymeric N-acetylglucosamine and the enzymes responsible for its synthesis are encoded by the ica operon (Gotz, 2002).

The biofilm-associated microorganisms are refractory to both antimicrobial agents and the host immune response. The resistance of biofilm forming microorganisms to antimicrobials represents a major challenge facing most of therapeutic and prophylactic strategies. Re-evaluation of the existing drugs of known mechanisms of action may result in identifying new mechanisms and targets that can affect bacterial metabolism (Golia et al., 2011).

Non-steroidal anti-inflammatory drugs (NSAIDs) are usually prescribed with or without antibiotics for patients had fever or pain due to an infectious condition, musculoskeletal condition (e.g. rheumatoid arthritis, gout), painful condition (e.g. metastatic bone pain, trauma, migraine headache), or the prophylaxis of ischemic heart disease. NSAIDs inhibit the cyclooxygenase enzymes COX-1 and COX-2, which are involved in the biosynthesis of mammalian prostaglandins (Pina-vaz et al., 2000).

Glucocorticoids (GC) have both immunosuppressive and anti-inflammatory actions so that they are used in the treatment of various immune-mediated inflammatory disorders, such as asthma, psoriasis, rheumatism, and multiple sclerosis (MS). GC also inhibits some transcription factors, such as nuclear factor kappa B and AP-1 (Au phan et al., 1995). As a result, the transcription of several cytokines that are involved in inflammation is decreased and the production of lipid mediators such as prostaglandins is inhibited and the expression of ICAM-1 and E-selectin on endothelial cells is reduced (Scheinman et al., 1995).

It was found that *C. albicans* can produce prostaglandins but their role in fungal biology is not yet known (Alem and Douglas, 2005). Prostaglandins can act as regulators of *C. albicans* eicosanoid pathway which have a vital role in controlling both morphogenesis and biofilm formation. Also the release of drugs that can target fungal prostaglandins pathways may combat fungal colonization and infection (Erb-Downward and Noverr, 2007). Glucocorticoids are used in many preparations in combination with antifungals and antibacterials (topical skin preparation, eye drops and ear drops). So, we thought to test the *in-vitro* activity of NSAIDs and dexamethasone on the growth, adherence, biofilm formation and the expression of some genes affect adhesion of *C. albicans* and *S. aureus* to know if the tested drugs have any additional effect on the microbial infection or not.

**MATERIALS AND METHODS**

**Microbial strains**

Two *C. albicans* and two *S. aureus* clinical strains obtained from the Department Of Microbiology And Immunology, Faculty of Pharmacy, Minia University were used in this study. *S. aureus* (ATCC 6538) and *C. albicans* ATCC 10231 were obtained from MIRCIN culture collection of the Faculty of Agriculture, Ain Shams University, Egypt.

**Drugs**

Stock solutions of Diclofenac sodium (150.4 mM), Ketoprofen (78.4 mM), Meloxicam (112 mM), Piroxicam (120 mM), Dexamethasone (4 mg/ml), Levofoxacin (122.8 mM) and Ketoconazole (74.24) (Sigma) were prepared in Dimethyl sulfoxide (DMSO).

**Determination of minimum inhibitory concentration (MIC) and minimum bactericidal concentrations (MBC) by microbroth dilution method**

According to the guidelines of Clinical and Laboratory Standards Institute (CLSI, 2008), the microbial suspensions of *S. aureus* and *C. albicans* cultures were made in sterile normal saline (0.89% NaCl wt/vol; Himedia, India) and the turbidity were adjusted to 0.5 McFarland standards (equivalent to 1.5 × 10^8 colony forming units (CFU) /ml). Two-fold serial dilutions of tested drugs were prepared in Mueller Hinton Broth (MHB; Difco Laboratories) and dispensed in 96-well microtiter plates (Tarson, India). The microbial suspensions was added to each well of the plate resulting in the final inoculum of 5 × 10^5 CFU/ml. Levofoxacin was used as standard antibacterial agent while ketoconazole was used as standard antifungal agent. DMSO was used as negative control (show no antibacterial or antifungal activity). The plates were incubated at 37°C for 18 h and were visually read for the absence or presence of turbidity. The minimum concentration of the tested agents showing no turbidity was recorded as MIC. The MBC was determined by spreading 100 μl on tryptic soy agar (Himedia, India) from the wells showing no growth. The plates were incubated at 37°C for overnight. Concentrations showed no growth on the surface of TSA was considered as MBC (CLSI, 2008).

**Scanning electron microscopy (SEM)**

To study the effect of the tested agents on the cell morphology of *S.
aureus and C. albicans, SEM was used. After one hour contact with the MIC concentrations of the tested drugs, the cells were prefixed in 2% glutaraldehyde for one hour at 4°C. Post-fixation was done using a 2% osmium tetroxide solution. The samples were gold covered by cathodic spraying (Edwards S 150 B). Finally, the samples were examined as described with the scanning electron microscope (Stereoscan 360, Cambridge) (Benyahya et al., 1992).

Biofilm susceptibility assay

The biofilms of S. aureus strains (ATCC 6538 and two clinical strains) and C. albicans strains (ATCC 10231 and 2 clinical strains) were prepared in 96-well polystyrene microtiter plates (Tarsen, India), using a method of Wei et al. (2006) with a few modifications. The microbial suspensions were prepared from the overnight grown culture and the turbidity of the suspension was adjusted to 0.7 OD$_{610}$ (1 × 10$^6$ CFU/ml). Two fold serial dilutions for NSAIDs, dexamethasone, levofloxacin and ketoconazole were prepared in trypticase soy broth (Difco laboratories) supplemented with 0.5% glucose. Forty microliters of fresh TSB with 0.5% glucose was added to each well, followed by the addition of 60 μl of above microbial suspension. Then, different dilutions of the tested products were added and the plates were incubated for 18 h at 37°C. The plates were washed by phosphate buffer saline (PBS) and biofilms were fixed with methanol for 15 to 30 min, stained with Crystal Violet (Sigma, USA). Biofilm formation was quantified by the addition of 95% ethanol to the crystal violet stained wells and recording the absorbance at 595 nm using a microplate reader (Multiskan spectrum, Finland) (Wei et al., 2006).

The effect of the tested drugs also examined on preformed biofilms. The biofilms were prepared by inoculating the suspension of S. aureus and C. albicans strains into the wells of a polystyrene microtiter plate as mentioned above. After incubation at 37°C for 18 h, the culture supernatant from each well was decanted and planktonic cells were removed by washing the wells with PBS (pH 7.2). Two fold serial dilutions of NSAIDs, dexamethasone, levofloxacin and ketoconazole were prepared in trypticase soy broth (TSB) and 200 μl of each dilution was added to the biofilm in the wells. The plate was further incubated at 37°C for 18 h. The biofilm was fixed, stained and quantified as described above (Wei et al., 2006).

Ethidium bromide accumulation assay

The action of NSAIDs, dexamethasone, levofloxacin and ketoconazole on cell membrane permeability of S. aureus ATCC 29213 and C. albicans ATCC 10231 cells were evaluated by the method as described by Cox et al. (2000). The microbial cells were grown overnight in Muller Hinton broth (MHB), resuspended in 50-mmol/l sodium phosphate buffer. The turbidity of the suspension was adjusted to 0.7 OD$_{610}$ (1 × 10$^6$ CFU/ml) and 1 ml of it was added to flask containing 19 ml buffer and the MIC of the tested levofloxacin, ketoconazole, NSAIDs and dexamethasone at 4 mg/ml. Following 60 and 120 min incubation at room temperature, 200 μl aliquots were transferred into tubes containing 3.8 ml phosphate buffer. These tubes were stored on ice and 20 μl of staining solution, consisting of ethidium bromide (Sigma) (10 μg/ml) dissolved in milliQ water. Then, the fluorescence was measured using a Perkin Elmer 45 luminescence spectrometer connected to the FLWINLAB software (United Kingdom) (Cox et al., 2000).

Germ Tube Formation

C. albicans ATCC 10231 colonies were suspended in sterile saline and adjusted to density of 0.5 Mcfarland. Human serum was added to 1 ml of cell suspensions of C. albicans ATCC 10231. Drugs were added to the suspension at MIC concentration and saline was added to the control tube. The cell suspensions were incubated with gentle shaking at 37°C for 2 h and were examined for the presence of germ tubes by using a light microscope. Photomicrographs of colonies and invasive growth were taken with a DMRXA microscope (Leica, Germany) (Liu et al., 1994).

RT-PCR analysis of C. albicans adhesion-related genes

Total RNA was extracted from C. albicans biofilms using QIAGEN RNasy Mini kit (cat no.74524, Germany) according to the guidelines of manufacturer. RNA concentrations and RNA purity were determined using a spectrophotometer (GeneQuant 1300, Germany). An equal amount of RNA was subjected to cDNA synthesis using the cDNA Reverse Transcriptase reagent kit (Applied Biosystems, cat no. 1311190, USA). Real-time PCR primers were designed for the target genes ALS1, HWP1 using Real Time Equipment (Strategene MXP 3000, Germany). The β-actin gene (ACT1) was used as an endogenous reference gene. The sequences of the primers are shown in Table 1. QuantiTect®SYBR® Green PCR Kit (Applied Biosystems, cat no. 204141, USA) was used. All PCR reaction mixtures contained: 10 μl QuantiTect®SYBR® Green PCR Master Mix (2X), 2 μl first strand cDNA, 0.5 μl each primer, 0.4 μl ROX Reference Dye (50X) and RNase-H2O to the final volume of 20 μl. The program for amplification was 95°C for 15min as an initial denaturation step to

<table>
<thead>
<tr>
<th>Primer</th>
<th>Sequence (5'-3')</th>
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<tbody>
<tr>
<td>icaA</td>
<td>TGGATGTGGTTCCAGAAACATTTGGGAG</td>
</tr>
<tr>
<td>ALS1</td>
<td>CCAACTGAAACACCA</td>
</tr>
<tr>
<td>HWP1</td>
<td>CGTTTCCCAATTTCACGTGT</td>
</tr>
<tr>
<td>β-actin (ACT1)</td>
<td>CACGCGTTGCTTCCAAAGA</td>
</tr>
</tbody>
</table>

Table 1. Primers used for RT-PCR experiments.
activate the HotStartTaq® DNA polymerase, followed by 40 cycles of PCR consisting of 94°C for 15 s, 60°C for 30 s and 72°C for 30 s. Negative controls (water as template) were included in each run. After amplification, a melting curve was analyzed to confirm the specificity of the primers. Expression of each investigated gene was normalized to the housekeeping ACT1 gene and analyzed using comparative Ct method (ΔΔCt). Expression of ALS1 and HWP1, genes from cells grown under drug treatment was indicated. Each experimental condition was performed in duplicate and each experiment was repeated twice (Ding et al., 2014).

RT-PCR analysis of S. aureus adhesion-related gene

TaqMan®Fast Universal quantitative Real-time PCR for icaA gene. Taqman primers and probe are summarized in Table 1. Probe was labeled with the reporter dye 6-carboxyfluorescein (6'-FAM) at the 5' end and with the quencher dye 6-carboxy-tetramethylrodamine (TAMRA) at the 3'end. TaqMan®Fast Universal PCR Kit (Applied Biosystems, cat no.4351891, USA) was used. Thermal cycling conditions were as follows: 10 min at 95°C followed by 45 repeats of 15 s at 95°C, and 1 min at 60°C. Data collection was performed during each annealing phase. Each experimental condition was performed in duplicate and each experiment was repeated twice (Paniagua-Contreras et al., 2012).

Statistical analysis

Data were described as mean ± SD. All statistical analyses were performed by statistical analysis computer software package SPSS 17.0 (SPSS Inc., IL, USA). Student’s t-test or one-way ANOVA were used to compare the biofilm formation, planktonic growth, and the gene expression of C. albicans and S. aureus strains in the presence or absence of drugs. Results with a p-value less than 0.05 were considered statistically significant.

RESULTS AND DISCUSSION

The attachment of microbial cells to surfaces and accumulation of these cells in forming multilayered cell clusters are key steps in any infection (Ji et al., 1995). In this regard, adhesion formation is considered as one of the major virulence factors in C. albicans and S. aureus. C. albicans is the fourth leading cause of bloodstream infections and the third most commonly isolated organism from intravascular catheters and is associated with the highest incidence of mortality (Crump and Colligan, 2000). C. albicans and S. aureus readily form biofilms on a wide variety of polymers used to make indwelling medical devices (Kojic and Darouiche, 2004). There is some evidence to suggest that a large proportion of device-related C. albicans infections involve biofilms (Dominic et al., 2007). In addition, severe catheter-related S. aureus infections have been reported in many studies to be important causes of morbidity, mortality, and a source of concern in the primary and emergency care context over the past decade (Parker and Doebbeling, 2012).

In recent years, due to the increased resistance of many bacteria to the commonly used antimicrobial agents, attention has shifted to drugs belonging to different pharmacological classes for possible antimicrobial activity. Many studies showed that NSAIDs have antibacterial activity, antifungal activity and decrease adherence and biofilm formation by bacteria and fungi (Ashraf et al., 2015; Mohsen et al., 2015). Our results showed that Meloxicam and Diclofenac sodium had the highest activity against the tested strains. Diclofenac sodium exhibited higher anti-candidal activity than that showed by Ketoconazole while Levofloxacin had the highest antibacterial activity against S. aureus followed by Meloxicam and Diclofenac sodium. Dexamethasone showed no activity against the tested strains (Table 2). Also, we tested the synergistic activity among NSAIDs, dexamethasone and the tested standard antimicrobials; it was found that the tested NSAIDs increased the activity of the tested antimicrobials but dexamethasone showed no increase in their activity (data not shown). The combined effect of antifungal or antibacterial agents and COX inhibitors in Candida infections was reported in many studies (Pina-vaz et al., 2000; Abdelmegeed and Shaaban, 2013).

The increase in the uptake of ethidium bromide is an evidence to the effect of the tested agents on the membrane integrity in comparison to the unexposed cells
It was found that Meloxicam and Levofoxacin showed no increase in the uptake of ethidium bromide by *S. aureus*. For *C. albicans*, Meloxicam showed no effect while Ketoconazole showed the highest disruptive effect on the membrane integrity followed by Diclofenac sodium. As observed, Diclofenac sod had the highest effect on the membrane integrity of both *S. aureus* and *C. albicans* which may explain its antimicrobial effect on both organisms (Figures 1 and 2). Also, the antifungal activity of Diclofenac sodium against *C. albicans* may be due to that their log P (4.75) which are close to the log P of Ketoconazole (4.35). Niazi et al. (2010) explained that the antifungal activity of two Schiff bases may be due to their logP (3.77 and 3.74) which are close to Ketoconazole (4.35). Many studies agreed with our results, as they reported that the antimicrobial activity of NSAIDs may be due to their inhibition of bacterial DNA synthesis or impairment of membrane activity (Niazi et al., 2010; Dutta et al., 2004).

By testing the effect of NSAIDs, Dexamethasone, Levofoxacin and Ketoconazole on the ability of the tested strains to form biofilm, it was found that meloxicam showed the highest ability to inhibit biofilm formation by both *S. aureus* and *C. albicans* strains followed by Diclofenac sodium, Dexamethasone, Levofoxacin (in case of *S. aureus*) or Ketoconazole (in case of *C. albicans*), Ketoprofen and Piroxicam (Figure 3). On the other hand, Diclofenac sodium exhibited the highest disruptive effect on the preformed biofilm formed by the tested strains followed by Meloxicam, Dexamethasone, Levofoxacin (in case of *S. aureus*) or Ketoconazole (in case of *C. albicans*), Ketoprofen and Piroxicam (Figure 4).

Our study showed that Dexamethasone affect the membrane permeability of the *C. albicans* while showed slight effect of *S. aureus* permeability in comparison to controls and its effect on biofilm formation and preformed biofilm are close to that shown by Ketoconazole and Levofoxacin. SEM was used to verify the effect of the tested drugs on the biofilm and the cell morphology of the tested strains. SEM graphs showed that the tested anti-inflammatory drugs caused morphological deformities (*S. aureus* and *C. albicans*) and pore formation on the surface of *C. albicans*. Also, the graphs showed that Dexamethasone caused swelling and pore formation to *C. albicans* cells (Figure 5). But Dexamethasone did not affect the cell morphology of *S. aureus*. The effect of NSAIDs on *C. albicans* and *S. aureus* cell morphology were also detected (Ashraf et al., 2015; Mohsen et al., 2015).

The ability of germ tube formation by *C. albicans* in the presence of tested drugs was determined. Diclofenac, Ketoconazole, Meloxicam (at MIC) and Dexamethasone (1 mg/ml) were found to inhibit germ tube formation but Ketoprofen and Piroxicam decreased the amount of germ tubes (Figure 6). Abdelmegeed and Shaaban (2013) reported that cultures treated with some NSAIDs exhibited a dose-dependent inhibition of germ tube
formation following 3 h incubation at 37°C. They reported significant reduction in germ tube formation under treatment with 1 and 10 mM anti-inflammatory drugs for 3 h especially with Ketorolac, Piroxicam and Diclofenac. Brooks and Day (1991) reported the antigerm tube effect of Diclofenac sodium suggesting that the antigerm tube effect observed in vitro might also be relevant in vivo (Brooks and Day, 1991).

Several studies demonstrated that local steroid treatment, even without antibiotics, can substantially reduce *S. aureus* colonization and the therapeutic effect of early combined therapy was superior to that in the steroid-only group; however, at the end of the treatment period, there was no significant difference between the 2 treatment groups in clinical scoring (Gong et al., 2006). According to our results, Dexamethazone had no antibacterial activity, slight effect on permeability of *S. aureus* cell membrane but had significant inhibitory effect on biofilm formation. So, we suggest that its effect may be due to low log P (partition coefficient) (1.83) which increase the wettability of tissue culture plate (TCP) surface and decrease the hydrophobic interaction between *S. aureus* surface and the plastic surface of TCP. Also, its low log P decreased its permeability through the hydrophobic membrane of *S. aureus* (Wang et al., 2012). Its effect on *C. albicans* may be due to its effect on prostaglandin production. The role of prostaglandin EX (PGEx) in *C. albicans* in biofilm development and fungal pathogenesis has been demonstrated (Erb-Downward and Noverr, 2007). Moreover it has been investigated by Alem and Douglas (2005) that prostaglandin synthesis by both planktonic and biofilm cells was sensitive to the cyclooxygenase inhibitors aspirin, Diclofenac, and Etodolac. Also, they found that NSAIDs that showed more activity on COX2 (Meloxicam, Diclofenac sodium) had the highest ability to decrease germ tube and biofilm formation (Alem and Douglas, 2005). This explain the variation in activity between the tested drugs as meloxicam and Diclofenac sodium which showed higher effects on germ tube and biofilm formation while Piroxicam showed the lowest effect as it has a COX 1: COX 2 activity ratio of 250:1 (Frölich, 1997). Dexamethazone cause the repression of COX2 and prostaglandin E2 release in human pulmonary A549 cells by transcriptional and post transcriptional mechanisms. So, we suggested that the inhibitory effect of Dexamethazone may be due to its effect on COX2 and PGE2 that play an important role in hyphae formation and host cell damage (Mishra et al., 2014).

Due to the observed effect of the tested drugs on biofilm formation and the preformed biofilms, we tried to determine the potential molecular mechanism behind the ability of the tested agents to prevent biofilm formation by both *C. albicans* and *S. aureus*. We have provided evidence that there were variable effects of the tested agents on *S. aureus* biofilm formation.
Figure 3. Effect of the tested agents on biofilm formation of both *S. aureus* (A, B & C) and *C. albicans* (D, E & F) strains. The results are expressed as average optical density readings for crystal violet assays compared to growth control. The biofilm of *S. aureus* and *C. albicans* were reduced with all tested agents (*P* < 0.01) while Piroxicam showed low effect on *C. albicans* (*P* < 0.05) compared with those of control. Values are mean (±SD) from four independent determinations (Student’s t test).
Figure 4. Effect of the tested agents on preformed biofilm of both *S. aureus* (A, B & C) and *C. albicans* (D, E & F) strains. The results are expressed as average optical density readings for crystal violet assays compared to growth control. The biofilm of *S. aureus* and *C. albicans* were reduced with all tested agents ($P < 0.01$) while Piroxicam showed low effect on *C. albicans* ($P < 0.05$) compared with those of control. Values are mean (±SD) from four independent determinations (Student’s t test).
drugs on the tested genes in comparison to control (C. albicans in the presence of serum). All tested drugs down-regulated the expression of the adhesion-related gene ALS1 (1.0643 to 2 fold). Dexamethasone was found to down-regulate the expression of ALS1 by 2 fold while Ketoconazole by 1.197 fold. On other hand, it was found...
that Ketoprofen up-regulated the expression of HWP1 gene by 3.2 fold. Meloxican down-regulated HWP1 expression by 4.11 fold followed by Ketoconazole (3.89 fold) and Piroxicam (3.81) (Figure 7). The morphological switch from yeast-to-hypha is one of the most important biological features that enable C. albicans to colonize, invade, and survive in the host tissues during infection. C. albicans, the yeast-to-hypha transition is triggered by various environmental causes, such as serum, N-acetylglucosamine, neutral pH, high temperature, starvation, CO2, and adherence (Biswas et al., 2007). In responses to a wide variety of stresses, including nutritional and environmental stresses, C. albicans initiate morphological changes which are controlled by a complex network of parallel pathways (e.g. MAPK and cAMP-PKA pathways) (Garcia-Rodriguez et al., 2005). Thus, any interference with the expression of genes involved in the MAPK cascades and cAMP-PKA pathway can block filamentation in C. albicans. The ALS family gene is related to the growth and morphological change of C. albicans. ALS family especially, ALS1 and ALS3 are important in the yeast-to-hypha transformation of C. albicans (Hoyer, 2001). Another well-characterized C. albicans hypha specific gene is the glycosylphosphatidylinositol (GPI)-linked cell surface protein HWP1. Functional analyses showed that this protein is required for the adherence of fungal cells to epithelial cells as well as for normal biofilm and hypha formation repression effect of Diclofenac sodium on the regulation of genes controlling C. albicans morphogenesis was also reported (Loza et al., 2004; Zhou et al., 2012).

S. aureus biofilm formation requires the production of polymeric N-acetylglucosamine that is controlled by icaABCD operon. Meloxicam was found to down-regulate the expression of icaA gene by 17.46 fold followed by levofloxacin (15.62 fold) and diclofenac sod. (7.042 fold) but dexamethasone showed no significant effect on its expression (Table 3).

Conclusion

NSAIDs have antibacterial and antifungal activity, can affect membrane permeability, germ tube formation and the expression of some genes controlling adhesion. Dexamethazone showed significant effect on biofilm formation of both S. aureus and C. albicans which may
Figure 7. Effect of NSAIDs, dexamethasone and levofloxacin on the expression of icaA gene in S. aureus (A) and the effect of NSAIDs, dexamethasone and ketoconazole on the expression of ALS1 (C) and HWP1 (D) genes in C. albicans. ACT1 housekeeping gene (B).
be a result of its Log P and the inhibition of PGEs release and affected gene expression of ALS1 and HWP1 of C. albicans. These effects of the tested drugs in addition to its anti-inflammatory properties may result in good response to the antimicrobial therapy in-vivo.

**Conflict of Interests**

The authors have not declared any conflict of interests.

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