Modeling Combined Immunosuppressive and Anti-inflammatory Effects of Dexamethasone and Naproxen in Rats Predicts the Steroid-Sparing Potential of Naproxen

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ABSTRACT

Dexamethasone (DEX), a widely prescribed corticosteroid, has long been the cornerstone of the treatment of inflammation and immunologic dysfunctions in rheumatoid arthritis. Corticosteroids are frequently used in combination with other antirheumatic agents such as nonsteroidal anti-inflammatory drugs (NSAIDs) and disease-modifying antirheumatic drugs to mitigate disease symptoms and minimize unwanted effects. We explored the steroid dose-sparing potential of the NSAID naproxen (NPX) with in vitro and in vivo studies. The single and joint suppressive effects of DEX and NPX on the in vitro mitogen-induced proliferation of T lymphocytes in blood and their anti-inflammatory actions on paw edema were investigated in female and male Lewis rats with collagen-induced arthritis (CIA). As expected, DEX was far more potent than NPX in these systems. Mathematical models incorporating an interaction term \( \psi \) were applied to quantitatively assess the nature and intensity of pharmacodynamic interactions between DEX and NPX. Modest synergistic effects of the two drugs were found in suppressing the mitogenic response of T lymphocytes. A pharmacokinetic/pharmacodynamic/disease progression model integrating dual drug inhibition quantitatively described the pharmacokinetics, time-course of single and joint anti-inflammatory effects (paw edema), and sex differences in CIA rats, and indicated additive effects of DEX and NPX. Further model simulations demonstrated the promising steroid-sparing potential of NPX in CIA rats, with the beneficial effects of the combination therapy more likely in males than females.

Introduction

Rheumatoid arthritis (RA) is one of the most prevalent chronic inflammatory autoimmune diseases, causing pain, swelling and stiffness of joints, as well as cartilage destruction and bone erosion. It preferentially affects synovial tissues but is also deleterious to other organ systems, producing various systemic comorbidities. With the combination of a susceptible genetic background and environmental insults, a sequence of complex immune activation cascades are involved in the disease initiation and progression of RA (Smolen et al., 2016). In particular, active immune cells present in the synovial tissues, especially T-lymphocytes, play a key role in promoting continuous autoimmune reactions by releasing various proinflammatory mediators such as cytokines and prostaglandins (PG) (Cavender et al., 1987; McInnes and Schett, 2011). Also, noticeable sex differences in the prevalence, disease course, and prognosis occur in RA, with females more affected (Jawaheer et al., 2006; van Vollenhoven, 2009). Conventional medications such as nonsteroidal anti-inflammatory drugs (NSAIDs) and corticosteroids (CS) remain first-line therapies for RA because of their rapid symptomatic effects compared with slower-acting disease-modifying antirheumatic drugs and biologic agents (Gaffo et al., 2006).

CS have long been used for treatment of RA owing to their potent anti-inflammatory and immunosuppressive properties. They not only provide rapid control and improvement of inflammatory symptoms but also exhibit disease-modifying effects (Kirwan, 1995). Dexamethasone (DEX) is one of the most frequently used therapeutic CS. As a synthetic glucocorticoid, it acts primarily through binding to the cytosolic glucocorticoid receptor and causes inhibition of transcriptional factors such as nuclear factor-kB and activator protein-1, thus suppressing the activation of genes encoding various proinflammatory mediators that aggravate inflammation and cause tissue damage (Coutinho and Chapman, 2011). The rapid actions of DEX are partly ascribed to nongenomic effects by directly regulating signal transduction pathways via membrane-associated glucocorticoid receptors and second messengers (Cato et al., 2002). Despite the significant therapeutic effects of DEX, its sustained use is limited by dose and time-dependent adverse
effects such as myopathy, growth retardation, metabolic disturbances, and osteoporosis (Saag et al., 1994). Owing to the chronic nature of RA, long-term steroid treatment remains common and necessary for alleviation of active disease activity. As a result, the undesired effects that are commonly associated with long-term and high-dose steroid therapy have a profound influence on the health-related quality of life of RA patients.

NSAIDs are widely prescribed for their potent analgesic, antipyretic, and anti-inflammatory properties. Their pharmacologic effects are mainly attributed to inhibition of cyclooxygenase (COX) and the consequent decreased synthesis of proinflammatory PG, thereby reducing pain and inflammation and improving physical function (Ricciotti and FitzGerald, 2011; Crofford, 2013). Naproxen (NPX), a nonselective COX inhibitor, has been extensively used in the symptomatic management of RA. It shows balanced inhibitory potencies on both COX isomers, constitutive COX-1 and inflammation-inducible COX-2 (Vane, 1971). NPX is better tolerated with respect to common adverse effects associated with NSAIDs (e.g., gastrointestinal bleeding and potential cardiovascular toxicity) compared with COX-selective NSAIDs (Lussier et al., 1978; Watson et al., 2002). However, like other NSAIDs, NPX is not disease modifying when used alone, thus providing the basis for its frequent use in combination therapy for RA.

Combination therapy has now become standard practice in treatment of RA. In view of the shared pharmacologic effects of DEX and NPX in inhibiting the factors promoting disease production, we hypothesize that their joint use might permit dose reduction of DEX, thereby reducing side effects without compromising efficacy. We assessed the individual and joint pharmacodynamic (PD) effects of DEX and NPX through both in vitro and in vivo studies. Their combined immunosuppressive action was first evaluated via the in vitro inhibition of concanavalin A (Con A)–stimulated whole-blood lymphocyte proliferation (WBLP). The pharmacokinetics (PK) of DEX in female and male healthy rats, the PK of NPX after coadministration of DEX, and the effects of NPX and DEX in collagen-induced arthritic (CIA) rats of both sexes were examined. We applied mathematical models incorporating a dual interaction equation to quantitatively describe the joint drug effects and the nature of drug interactions. Further model simulations were conducted to assess the potential degree of steroid dose-sparing effects of NPX.

Materials and Methods

Reagents and Chemicals
Pharmaceutic-grade dexamethasone sodium phosphate solution was obtained from Bimeda Pharmaceuticals (Dublin, Ireland). Dexamethasone (purity ≈ 98%), sodium naproxen, and Con A were purchased from Sigma-Aldrich (St Louis, MO). Dexamethasone-D5 (internal standard [IS], purity = 98%) was purchased from Toronto Research Chemicals (Toronto, ON, Canada). The RPMI 1640 medium (Invitrogen, Grand Island, NY) was used for cell culture and serial dilutions of drugs in the WBLP assay.

Animals
Male and female Lewis rats (aged 5 to 6 weeks), weighing approximately 110 g for females and 150 g for males, were purchased from Harlan (Indianapolis, IN). All rats were housed individually in the University Laboratory Animal Facility for females and 150 g for males, were purchased from Harlan (Indianapolis, IN). All rats were acclimated for 1 week before the experiments. These studies adhered to the Guide for the Care and Use of Laboratory Animals (National Research Council, 2011) and were approved by the University at Buffalo Institutional Animal Care and Use Committee.

WBLP
The method detailed previously by Ferron and Jusko (1998) was adapted with slight modifications. Briefly, blood (about 4 ml) from male healthy rats was collected into evacuated heparinized glass tubes and diluted 1:20 (v/v) with RPMI 1640 medium supplemented with 7.5% fetal calf serum, 100 U/ml penicillin, 100 µg/ml streptomycin, 2 mM L-glutamine, 20 mM HEPES, and 0.25 mM 2-mercaptoethanol. Then 165 µl of the whole blood dilution was plated into each well of a flat bottom, 96-well plate after adding 15 µl of medium (Con A; stock concentration 133 µg/ml in medium) and 20 µl of various concentrations of NPX and DEX alone or combined (Table 1). The cultures were incubated in a 5% CO2-humidified incubator at 37°C for 72 hours, then pulsed with 1 µCi of [3H]-thymidine per well (New England Nuclear, Boston, MA) and incubated for 18 hours. The cells were harvested onto microplates (Packard Instrument, Meriden, CT), washed with 3% hydrogen peroxide, dried in an oven at 50°C for 2 hours, mildly shaken on a plate shaker for 5 minutes after the addition of liquid scintillation fluid (Microscint-O; Packard Instrument), and counted using a Top Count Microplate Scintillation Counter (Perkin Elmer, Waltham, MA).

PK and PD of DEX and NPX
The DEX dosing solution was freshly prepared by directly diluting the DEX sodium phosphate solution (equivalent to 3 mg/ml DEX) with phosphate-buffered saline (PBS). The NPX dosing solution was freshly prepared by dissolving sodium NPX into PBS (pH 8.0), and was filtered through 0.22 micron filters before use. Both drugs were administered in a volume of 1 ml/kg.

The induction of CIA in the Lewis rats was conducted following procedures described elsewhere (Earp et al., 2008b). Hind-paw swelling was used to indicate edema and was measured by digital calipers (VWR Scientific, Rochester, NY). After evaluation on day 15 for females and day 20 for males (1 day before peak disease time), the rats with a paw volume increase of at least 50% in one or two hind paws were selected and used in the subsequent PD study.

For each sex group, the CIA rats were randomly divided into four subgroups (n = 4): vehicle controls, which received only PBS intraperitoneally (i.p.); single-drug groups, which received a bolus injection of either 55 mg/kg i.p. sodium NPX PBS solution (equivalent to 50 mg/kg NPX) or 0.225 mg/kg DEX subcutaneously (s.c.); and the combined group, which received 50 mg/kg i.p. NPX immediately after s.c. injection of 0.225 mg/kg DEX. All groups received injections on day 16 for females and day 21 for males after disease induction. Paw edema was measured as the sum of the paw and ankle cross-sectional areas for each hind paw. Paw edema and body weights were monitored before disease induction (day 0) and on days 3, 7, 9, 10, 11, 13, 14, and 15 after induction for both sex groups and days 17 to 20 for CIA males. Starting from the dosing day (day 16 for females and day 21 for males), paw edema was measured before dosing and at 1, 2, 4, 6, 8, 12, 24, and 36 hours after dosing and on postdose days 3, 4, 5, 6, 7, 8, 9, and 10. The paw volumes of four female and four male healthy rats were measured through the entire study to determine the natural growth.

The PK of s.c. DEX (2.25 mg/kg) was studied in female and male healthy rats. Serial blood samples were collected from the saphenous vein at 10, 20, 30, and 50 minutes after the dose and 1, 1.5, 2, 3, 4, 6, 8, 12, and 24 hours and processed and analyzed for DEX by a liquid chromatography with tandem mass spectrometry (LC-MS/MS) method used previously by others (Samtani and Jusko, 2007) with slight modifications. Briefly, 100 µl 4% phosphoric acid was added to 100 µl of plasma and then spiked with 10 µl of IS working solution. The mixture was vortex-mixed and centrifuged at 13,000 g for 10 minutes at 10°C and then subjected to solid-phase extraction. The dried residue was reconstituted with 200 µl of acetonitrile/water (50:50, v/v) followed by vortexing and centrifugation at 13,000 g for 10 minutes at 10°C. The supernatant was transferred into vials and used for LC-MS/MS injection.

Chromatographic separations were achieved on a C8 Hydrobond AQ Column (particle size 3 µm, 150 × 2.1 mm; MAC-MOD Analytical, Chadds Ford, PA)

<table>
<thead>
<tr>
<th>Drug</th>
<th>Concentrations in WBLP Assay</th>
</tr>
</thead>
<tbody>
<tr>
<td>NPX</td>
<td>0.5, 1, 5, 10, 40, 100, 200, 500, 1000, and 2000 M</td>
</tr>
<tr>
<td>DEX</td>
<td>0.05, 0.1, 0.2, 0.5, 1, 2, 4, 10, 20, and 50 M</td>
</tr>
<tr>
<td>Combinations</td>
<td>DEX (1, 2, 5 and 10 M)+ NPX (50, 100, 150, 200, and 250 M)</td>
</tr>
</tbody>
</table>
PK Models

The NPX concentrations in combined-dosing groups were compared with those in the single-drug groups to assess the influence of single-dose DEX on the PK of NPX. Serial blood samples were collected from the saphenous vein at 15, 30, and 45 minutes and 1, 2, 4, 6, 9, 12, and 24 hours and analyzed for NPX by LC-MS/MS (Li et al., 2017a). Linearity was found over the concentration ranges of 0.2 to 100 ng/ml for DEX and 0.125 to 40 μg/ml for NPX. For both bioanalytic assays, the coefficient of variation (CV%) of intra- and interday accuracy and precision were less than 15%, and the recovery of the sample preparation method was around 90% for DEX and approached 100% for NPX.

PK Models

The plasma concentration–time profiles of NPX were described by an extended minimal physiologically based pharmacokinetic (mPBPK) model with plasma and one tissue compartment (Fig. 1). Detailed model descriptions and assumptions are provided in (Li et al., 2017b).

The model equations and initial conditions are

\[
\frac{dA_{a,N}}{dt} = -k_{a,N} \cdot A_{a,N} \quad A_{a,N}(0) = \text{Dose}_N \cdot F_N
\]

\[
\frac{dC_{p,N}}{dt} = \left[ k_{a,N} \cdot A_{a,N} + f_{a,D} \cdot Q_{CO} \cdot (C_{p,ISF} - C_{p,p}) \right] \cdot \frac{CL_p}{V_p} \quad C_{p,N}(0) = 0
\]

\[
\frac{dC_{ISF}}{dt} = \left[ f_{a,D} \cdot Q_{CO} \cdot (C_{p,ISF} - C_{p,p}) \right] \cdot \frac{1}{V_{ISF}} \quad C_{ISF}(0) = 0
\]

where \( A_{a,N} \) indicates the amount of NPX at the absorption site, \( k_{a,N} \) is the first-order absorption rate constant, \( C_{p,N} \) and \( C_{ISF} \) are total NPX concentrations in \( V_p \) (plasma volume) and \( V_{ISF} \) (interstitial fluid volume), \( Q_{CO} \) is cardiac plasma flow, \( f_{a,N} \) is the fraction of drug in \( Q_{ISF} \) accessing \( V_{ISF} \), \( CL_p \) is clearance of unbound NPX from plasma, \( C_{p,p} \) and \( C_{ISF} \) are the unbound NPX concentrations in plasma and ISF that could be calculated from their corresponding total concentrations using the equations, albumin concentrations, and binding parameters provided in Li et al. (2017a,b), and \( F_N \) is the bioavailability of the i.p. dose calculated to be about 0.9 from literature-reported intravenous data in rats (Lauroba et al., 1986).

The physiologic restrictions of relevant parameters are

\[ f_{a,D} \leq 1 \] and \( V_p + V_{ISF} = \text{Extracellular fluid volume (ECF)} = 206.29 \text{ ml/kg} \)

(Sah and Betts, 2012). The plasma concentration–time profiles of DEX after 2.25 mg/kg s.c. injection to female and male healthy rats were characterized by the basic mPBPK model (Cao and Jusko, 2012) shown in Fig. 1. The model equations and initial conditions describing DEX PK are:

\[
\frac{dA_{a,D}}{dt} = -k_{a,D} \cdot A_{a,D} \quad A_{a,D}(0) = \text{Dose}_D \cdot F_D
\]

\[
\frac{dC_{p,D}}{dt} = \left[ k_{a,D} \cdot A_{a,D} + f_{a,D} \cdot Q_{CO} \cdot \left( \frac{C_{p,p}}{K_p} - C_{p,D} \right) \right] \cdot \frac{CL_p \cdot C_{p,D}}{V_p} \quad C_{p,D}(0) = 0
\]

\[
\frac{dC_{t}}{dt} = \left[ f_{a,D} \cdot Q_{CO} \cdot \left( C_{p,D} - \frac{C_{t}}{K_p} \right) \right] \cdot \frac{1}{V_t} \quad C_{t}(0) = 0
\]

where \( A_{a,D} \) is the amount of DEX at the absorption site, \( k_{a,D} \) is the first-order absorption rate constant, \( C_{p,D} \) and \( C_t \) are total DEX concentrations in \( V_p \) and \( V_t \) (tissue volume), \( K_p \) is the tissue partition coefficient, \( f_{a,D} \) is the fraction of drug in \( Q_{CO} \) accessing \( V_t \), \( CL_p \) is the systemic clearance, and \( F_D \) is the bioavailability of i.m. DEX (0.86) in rats (Samtani and Jusko, 2005; Earp et al., 2008c). The physiologic restrictions of relevant parameters are:

\[ f_{a,D} \leq 1 \] and \( V_p + V_t = \text{Body weight} \)

Several modeling assumptions were made for DEX. (1) Unbound DEX concentrations equilibrate rapidly among plasma, ISF, and cell water, so the free
ISF DEX concentrations serve as the driving force for PD effects in CIA rats. (2) Protein binding of DEX is linear and occurs primarily to albumin in plasma (Cummings et al., 1990) and tissues with negligible binding to corticosteroid-binding globulin. (3) The binding of DEX to albumin in tissues occurs principally in the ISF. (4) The unbound fraction of DEX in plasma (f_{unb}) is 0.3 for both sexes.

The unbound fraction of DEX in ISF (f_{ISF}) was calculated using:

$$f_{ISF} = \frac{1}{1 + (E/P)(1 - f_{np})/f_{np}}$$

where E/P is the ratio of protein concentrations in ISF and plasma with a value of 0.9 used for arthritic rats (McNamara et al., 1983, Li et al., 2017a).

The PK/PD model structure for the single drug effects of NPX was presented in Li et al. (2017b). In this model, we assumed that the paw size increase in healthy rats due to the anti-inflammatory effects of NPX and DEX acting on paw edema production. The published model was modified to allow for the estimation of actual CPM values instead of normalized CPMs. The inhibition potency on lymphocyte proliferation of individual drug were described using

$$E = E_0 \cdot \left(1 - \frac{I_{\text{max}} \cdot C}{IC_{50} + C}\right)$$

where $E$ is the CPM in the presence of drug, $E_0$ is the baseline CPM in the absence of drug, $I_{\text{max}}$ is the maximum effect achieved by either drug, $IC_{50}$ is the drug concentration at 50% of maximum inhibition, $C$ is the total concentration of NPX or DEX in the culture medium, and $y$ is the Hill coefficient for the immunosuppressive effect. All model parameters were fitted to the CPM values for each of the four wells of each drug concentration.

The combined inhibitory effects of the two drugs on lymphocyte proliferation were characterized by a joint interaction model (Chakraborty and Jusko, 2002). The published model was modified to allow for the estimation of actual CPM values instead of normalized CPMs.

$$E = E_0 \cdot \left[1 - I_{\text{max,N}} \cdot \left(\frac{C_N}{IC_{50,N}}\right)^{\gamma_N} + I_{\text{max,D}} \cdot \left(\frac{C_D}{IC_{50,D}}\right)^{\gamma_D} + \left(I_{\text{max,N}} + I_{\text{max,D}} - I_{\text{max,N}} \cdot I_{\text{max,D}}\right) \times \left(\frac{C_N}{IC_{50,N}}\right)^{\gamma_N} \cdot \left(\frac{C_D}{IC_{50,D}}\right)^{\gamma_D} + 1\right]$$

where $\gamma_1$ is the interaction term for lymphocyte proliferation inhibition where values of 1 indicate no interaction, <1 synergism, and >1 antagonism.

**Single and Combined Drug Effects in Rat Arthritis**

Figure 1 depicts a general model scheme for the entire arthritis progression with the anti-inflammatory effects of NPX and DEX acting on paw edema production. The PK/PD model structure for the single drug effects of NPX was presented in Li et al. (2017b). In this model, we assumed that the paw size increase in healthy rats follows logistic growth. Therefore, before disease onset the rate of change of the paw volume over time in all groups is

$$\frac{d\text{Paw}}{dt} = -k_y \cdot \text{Paw} \cdot \left(1 - \frac{\text{Paw}}{\text{Paw}_{\text{ss}}}\right), \ t < t_{\text{onset}} \quad \text{Paw}(0) = \text{Paw}^0$$

where Paw is the sum of ankle and paw cross-sectional areas of a rat hindfoot; Paw_{ss} is the normal paw size at steady state. $K_e$ is the natural first-order growth rate constant of paw in healthy rats, and $t_{\text{onset}}$ is the time delay observed before disease onset.

After disease onset, the paw volume-time profiles in the control group with no drug are

$$\frac{d\text{Paw}}{dt} = k_y \cdot \text{Paw} \cdot \left(1 - \frac{\text{Paw}}{\text{Paw}_{\text{ss}}}\right) + k_m \cdot \text{Paw}, \ t = t_{\text{onset}}$$

$$\text{Paw}(0) = \text{Paw}^0$$

where $k_m(t)$ is a function of time and represents the production of paw edema after disease onset; $k_{\text{deg}}$ is a linear decline in $k_m$ accounting for the natural remission of arthritis; and $k_{\text{out}}$ is the first-order rate constant describing the loss of edema.

Based on the mechanisms of drug action, indirect response model I (IDR I) was applied to describe the individual inhibitory effects of NPX and DEX on paw edema production (Dayneka et al., 1993). The rate of change of paw size over time for a single drug was described as

$$\frac{d\text{Paw}}{dt} = -k_y \cdot \text{Paw} \cdot \left(1 - \frac{\text{Paw}}{\text{Paw}_{\text{ss}}}\right) + k_a \cdot \text{Paw}, \ t = t_{\text{onset}}$$

$$\text{Paw}(0) = \text{Paw}^0$$

where $\text{IC}_{50}$ is the maximum inhibition effect of either NPX or DEX on production of paw edema and $\text{IC}_{50}$ is the unbound ISF concentration ($C^*$) of drug required for 50% of maximal effect.

The combination effects of NPX and DEX were characterized using a dual drug inhibition approach (Earp et al., 2004). The paw size change over time in combined treatment groups is

$$\frac{d\text{Paw}}{dt} = -k_y \cdot \text{Paw} \cdot \left(1 - \frac{\text{Paw}}{\text{Paw}_{\text{ss}}}\right) + k_a \cdot \text{Paw}, \ t = t_{\text{onset}}$$

$$\text{Paw}(0) = \text{Paw}^0$$

where $\theta_2$ is the interaction term for inhibition of paw edema production as defined for eq. 9. The model assumes that the drugs inhibit the proinflammatory processes ($k_a$) as directly demonstrated for DEX by Earp et al. (2008b). Values of $I_{\text{max}}$ and $IC_{50}$ for each drug were generated from their individual studies, allowing $\theta_2$ to be resolved from the studies with joint NPX and DEX administration.

**Model Fitting and Data Analysis**

Naïve-pooled data from all replicates in the WBLP assay were analyzed jointly. Initial noncompartmental analysis (NCA) of the PK data of NPX and DEX was performed using Phoenix WinNonlin 6.4 software (Certara Corporation, Princeton, NJ). All PK data for NPX from our previous study (Li et al., 2017a) and this study were fitted jointly as well as DEX PK data for both sexes. The obtained PK parameter estimates were fixed in the subsequent PD analysis in CIA rats. All model fittings involved nonlinear regression using the maximum likelihood algorithm in ADAPT 5 (Biomedical Simulations Resource, Los Angeles, CA) (D’Argenio et al., 2009). All model codes are provided in the Supplemental Data. The variance model was set as

$$V_i = (\sigma_1 + \sigma_2 \cdot Y_i)^2$$
where $V_i$ represents the variance of the $i$th data point, $Y_i$ is the $i$th model prediction, and $\sigma_1$ and $\sigma_2$ are variance model parameters that were estimated together with other system parameters during model fitting. Model selection was based on the goodness-of-fit criteria which included Akaike Information Criterion (AIC), visual inspection of the fitted profiles, and CV% of the parameter estimates. Statistical analysis of PK parameters was performed using SPSS software version 22 (IBM SPSS Statistics, Chicago, IL), and $P < 0.05$ was considered statistically significant.

Results

Single and Combined Effects of DEX and NPX on WBLP. Profiles for single-drug inhibition of lymphocyte proliferation are presented in Fig. 2. Both drugs showed concentration-dependent inhibition of lymphocyte proliferation. The sigmoidal $I_{\text{max}}$ model (eq. 8) was able to capture well the concentration–response relationships of the individual agents. (See the Supplemental Data for model code for single-drug effects in WBLP.) The parameter estimates for single drugs as well as interaction parameters for both drugs and their precision are listed in Table 2.

Both NPX and DEX produced complete suppression of cell proliferation, so both $I_{\text{max}}$ values were fixed to 1.0. The IC$_{50}$ of NPX for WBLP was estimated to be 83.97 $\mu$g/ml, which is slightly lower than reported for human lymphocytes (350 $\mu$g/ml) (Panush, 1976). It is very close to the peak plasma concentration (79 $\mu$g/ml) achieved in RA patients after 500 mg, twice daily doses of NPX (van den Ouweland et al., 1987). This is in accordance with previous findings that the antiproliferative effects of NSAIDs often occur at doses/concentrations high enough to suppress inflammatory responses (Seng and Bayer, 1986). DEX was more efficacious in suppressing lymphocyte proliferation than NPX, as indicated by the 5-fold difference in their log IC$_{50}$ values. The estimated IC$_{50}$ value of DEX (1.5 ng/ml) is comparable to that using human whole blood culture (3.8 ng/ml) (Mager et al., 2003), which suggests that the antiproliferative effect and sensitivity of the mitogenic response of lymphocytes to DEX are similar across species. The Hill coefficient $\gamma$ was 3.04 for DEX, similar to that for NPX (2.92).

The fittings included baseline CPM values for complete assessment of all parameters with potential variability.

The combination effects of NPX and DEX on lymphocyte proliferation were evaluated using 20 concentration combinations. These data were fitted using the joint interaction model (eq. 9) where the interaction term $\psi$ was factored with the IC$_{50}$ of DEX because the primary immunosuppressive effects were exerted by DEX. A three-dimensional surface plot (Fig. 2C) shows the combined drug effects on lymphocyte proliferation, where the surface represents the model predictions using eq. 9 based on the estimated parameters in Table 2. (See the Supplemental Data for the model code for combined-drug effects in WBLP.) The estimated $\psi_1$ value was 0.94 (95% confidence interval [CI], 0.90–0.99), which is lower than 1. Therefore, the

Fig. 2. Inhibition of Con A–stimulated lymphocyte proliferation by DEX and NPX as single agents (A and B) and in combination (C). Symbols represent experimental data, and curves depict fitting of data using eq. 8. The interaction surface shows model predictions using eq. 9 based on the fitted parameters listed in Table 2. Red circles identify data above the surface, and green circles show data below the surface.
combination of NPX and DEX exhibits a slight synergistic effect on lymphocyte proliferation.

**PK of NPX and DEX in CIA Rats.** The PK of NPX in the combined treatment groups was compared with that in the single-drug groups from our previous study (Li et al., 2017a). The concentration–time profiles of NPX in CIA rats after giving 50 mg/kg NPX alone or together with 0.225 mg/kg DEX along with the model fittings and calculated total and unbound ISF NPX concentrations are illustrated in Fig. 3.

The NCA results for the primary PK parameters of NPX including area under the concentration–time curve (AUC), apparent clearance (CL/F), apparent volume of distribution (V/F), and terminal half-life (t_{1/2}) are listed in Table 3. No obvious visual differences are seen for the PK profiles, and also no significant differences in the PK parameters were found, suggesting that single-dose DEX did not affect the PK of NPX in CIA rats. Therefore, the PK parameter estimates of NPX obtained using the extended mPBPK model with concentration-dependent protein binding in both plasma and ISF (Fig. 1) from our previous study (Li et al., 2017b) (Table 4) were fixed and applied for the subsequent PD modeling.

The PK of DEX in female and male healthy rats receiving 2.25 mg/kg of s.c. DEX were assessed (Fig. 4) and described by the basic mPBPK model with one tissue compartment shown in Fig. 1. (See the Supplemental Data for the model code for DEX PK estimation.) The NCA results for the primary PK parameters of DEX indicate that the terminal half-life (t_{1/2}) of DEX is significantly shorter in males (Table 3), so different clearance terms (CL_p) were assigned to female and male rats in the model. The mPBPK model was able to capture all the PK data of DEX in healthy rats quite well (Fig. 4) and yielded precise parameter estimates with reasonable CV% values (Table 4).

### Table 2

<table>
<thead>
<tr>
<th>Parameter (Units)</th>
<th>Definition</th>
<th>Estimates (CV%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>I_{max}</td>
<td>Maximum inhibition of cell proliferation</td>
<td>1 (Fixed)</td>
</tr>
<tr>
<td>IC_{50} (µg/ml)</td>
<td>Total drug concentration producing 50% of maximal effect</td>
<td>83.97 (1.06)</td>
</tr>
<tr>
<td>γ</td>
<td>Hill coefficient</td>
<td>2.92 (4.06)</td>
</tr>
<tr>
<td>E₀</td>
<td>Baseline CPM in the absence of drug</td>
<td>290453 (2.12)</td>
</tr>
<tr>
<td>θ₁</td>
<td>Interaction term for inhibition of cell proliferation</td>
<td>0.94 (2.55)</td>
</tr>
</tbody>
</table>

![Fig. 3. NPX plasma concentration–time profiles in female and male arthritic rats that received 50 mg/kg i.p. doses of NPX alone (△) or jointly with 0.225 mg/kg s.c. doses of DEX (▲), and calculated ISF concentrations of total (dashed lines) and unbound (dotted lines) NPX in female and male arthritic rats. Solid curves depict model fittings jointly for all dose groups, yielding parameters listed in Table 4.](image-url)
Plasma volume ($V_p$), tissue volume ($V_t$), and cardiac plasma flow ($Q_{CO}$) are physiologic parameters applicable to rats obtained from the literature (Shah and Betts, 2012), which were fixed in the model fitting. Absorption of DEX from the s.c. injection site was rapid with a $k_a$ value of 2.09 1/h, which is slightly smaller than that obtained after i.m. dosing (5.78 1/h) (Earp et al., 2008c). The male rats showed higher clearances ($CL_p$, Males = 197.7 ml/h/kg) than females ($CL_p$, Females = 137.8 ml/h/kg), consistent with the fact that DEX exhibits sex-specific metabolite profiles in rats, where its microsomal metabolism is more rapid in males (Tomlinson et al., 1997).

The estimated tissue partition coefficient ($K_a$) is 0.63, for DEX which is close to those calculated from the steady-state distribution volume ($V_d$/Body weight = 0.7) and a literature-reported estimation method (0.5 for muscle) (Poulin and Theil, 2000). The fraction of cardiac plasma flow of DEX ($f_{DL}$) was fixed to 1, consistent with rapid diffusion of DEX into tissues with blood flow–limited distribution. Model simulations (see the Supplemental Data for the model code for DEX PK simulation) for tissue total and ISF unbound DEX concentrations in arthritic rats receiving 0.225 mg/kg of s.c. DEX are displayed in Fig. 4, with the latter used as the driving force for PD effects of DEX.

**Single and Combined Anti-inflammatory Effects of NPX and DEX.** The PK/PD/disease (DIS) progression model displayed in Fig. 1 quantitatively integrated the PK of NPX and DEX into the model for paw edema progression, where unbound ISF concentrations of NPX (Fig. 3) and DEX (Fig. 4) were used to drive their anti-inflammatory effects in arthritic rats. (See the Supplemental Data for the PK/PD/DIS progression model code.) The model was applied to jointly fit the paw volume data from all groups with model fittings illustrated in Fig. 5. The model simultaneously characterized the natural paw growth as well as progression of paw edema with or without treatments very well. All the PD/DIS parameters were estimated quite precisely with reasonable CV% values (Table 5). The disease-specific parameter estimates for male arthritic rats are comparable to those reported elsewhere (Earp et al., 2009; Lon et al., 2011).

In general, the paw edema RA progression of female and male arthritic rats in our study showed similar patterns and trends with those obtained previously (Li et al., 2017b), where females exhibited higher incidences (80% versus 60%), earlier disease onset ($t_{onset}$ 228 hours versus 307 hours), faster disease production rate (group-averaged $k_{prod}$ 2.02 versus 1.78 mm²/h), and more severe symptoms. In the absence of drug, paw size changes over time before disease onset were assumed to follow the logistic growth function (eq. 11) and increased at the same rate as in healthy animals (the estimated $k_g$ was 0.002 h⁻¹ for both sexes).

The individual anti-inflammatory effects of NPX and DEX were characterized using eq. 13. NPX was previously found to be more efficacious in male arthritic rats as demonstrated by the maximum inhibitory effect ($T_{max,N}$, 1.0 versus 0.75 for females) and concentration for 50% maximal effect (IC₅₀,N, 136 versus 221 ng/ml) (Li et al., 2017b). A pilot study assessing the effects of two single s.c. doses of DEX (0.225 and 2.25 mg/kg) on paw edema in female and male CIA rats was conducted (unpublished), and the paw edema data were described by the mPBPK/PD/DIS progression model (Fig. 1) using unbound ISF DEX concentrations as the driving force for PD. Preliminary fittings revealed values of $T_{max,D}$ that were 1.0, so it was set as 1.0 for both sexes. A lower IC₅₀,D value of 0.39 ng/ml obtained for males than females alone or joint dosing of 0.225 mg/kg s.c. DEX to arthritic rats and 2.25 mg/kg s.c. DEX to healthy rats.

### TABLE 3

Pharmacokinetic parameters (NCA) based on total plasma concentrations of NPX and DEX after 50 mg/kg i.p. NPX alone or joint dosing of 0.225 mg/kg s.c. DEX to arthritic rats and 2.25 mg/kg s.c. DEX to healthy rats

<table>
<thead>
<tr>
<th>Groups</th>
<th>AUC (μg/h/ml)</th>
<th>CL/F (ml/h/kg)</th>
<th>V/F (ml/kg)</th>
<th>$t_{1/2}$ (h)</th>
</tr>
</thead>
<tbody>
<tr>
<td>NPX (50 mg/kg)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CIA males (NPX only)</td>
<td>809.8 ± 61.0</td>
<td>61.99 ± 4.87</td>
<td>229.2 ± 48.9</td>
<td>2.55 ± 0.36</td>
</tr>
<tr>
<td>CIA males (NPX + DEX)</td>
<td>876.8 ± 77.5</td>
<td>57.39 ± 5.26</td>
<td>198.1 ± 17.6</td>
<td>2.03 ± 0.97</td>
</tr>
<tr>
<td>CIA females (NPX only)</td>
<td>936.0 ± 42.2</td>
<td>53.43 ± 2.53</td>
<td>287.1 ± 42.5</td>
<td>3.74 ± 0.64</td>
</tr>
<tr>
<td>CIA females (NPX + DEX)</td>
<td>1021.6 ± 118.7</td>
<td>49.45 ± 5.45</td>
<td>192.3 ± 35.8</td>
<td>2.70 ± 0.45</td>
</tr>
<tr>
<td>DEX (2.25 mg/kg)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Healthy males</td>
<td>10.84 ± 0.65</td>
<td>208 ± 12</td>
<td>658 ± 36</td>
<td>2.19 ± 0.12</td>
</tr>
<tr>
<td>Healthy females</td>
<td>12.89 ± 2.27</td>
<td>179 ± 35</td>
<td>890 ± 184</td>
<td>3.45 ± 0.11*</td>
</tr>
</tbody>
</table>

*AUC, area under the time–concentration curve.

<table>
<thead>
<tr>
<th>Parameter (Units)</th>
<th>Definition</th>
<th>Estimates</th>
<th>CV%</th>
</tr>
</thead>
<tbody>
<tr>
<td>$k_{DL}$ (1/h)</td>
<td>Absorption rate constant of DEX</td>
<td>2.09</td>
<td>9.33</td>
</tr>
<tr>
<td>$f_{DL}$</td>
<td>Fraction of cardiac plasma flow of DEX</td>
<td>1</td>
<td>Fixed</td>
</tr>
<tr>
<td>$CL_p$, Females (ml/h/kg)</td>
<td>Clearance of DEX in female rats</td>
<td>137.8</td>
<td>3.51</td>
</tr>
<tr>
<td>$CL_p$, Males (ml/h/kg)</td>
<td>Clearance of DEX in male rats</td>
<td>197.7</td>
<td>3.43</td>
</tr>
<tr>
<td>$K_a$</td>
<td>Tissue partition coefficient of DEX</td>
<td>0.63</td>
<td>4.35</td>
</tr>
<tr>
<td>$V_p$ (ml/kg)</td>
<td>Plasma volume</td>
<td>32.36 ³</td>
<td>Fixed</td>
</tr>
<tr>
<td>$V_t$ (ml/kg)</td>
<td>Tissue volume</td>
<td>967.6 ³</td>
<td>Fixed</td>
</tr>
<tr>
<td>$V_{ISF}$ (ml/kg)</td>
<td>ISF volume</td>
<td>173.93 ³</td>
<td>Fixed</td>
</tr>
<tr>
<td>$Q_{CO}$ (ml/h/kg)</td>
<td>Cardiac plasma flow</td>
<td>7650 ³</td>
<td>Fixed</td>
</tr>
<tr>
<td>$k_{on}$ (1/h)</td>
<td>Absorption rate constant of NPX</td>
<td>0.98 ³</td>
<td>Fixed</td>
</tr>
<tr>
<td>$f_{DL}$</td>
<td>Fraction of cardiac plasma flow of NPX</td>
<td>0.15 ³</td>
<td>Fixed</td>
</tr>
<tr>
<td>$CL_p$, NPX (ml/h/kg)</td>
<td>Unbound clearance of NPX in arthritic rats</td>
<td>1438 ³</td>
<td>Fixed</td>
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</tbody>
</table>

*Physiologic parameter values obtained from Shah and Betts (2012).

³PK parameter values of NPX obtained from Li et al. (2017b).
The strong inhibition of paw edema observed for the combined treatment groups was satisfactorily described by the mPBPK/PD/DIS progression model incorporating the joint interaction equation (eq. 14). During the model fitting, all drug-specific PD parameters ($I_{\text{max}}$ and $IC_{50}$) were fixed to the values listed in Table 5 with only the disease-specific parameters and interaction term $c_2$ estimated by simultaneously fitting all of the paw volume data from each sex group. The disease-related parameters were re-estimated to account for the between-batch variation of CIA animals. The interaction term $c_2$ was found to be 1.16 (95% CI, 0.49–1.84) for females and 0.98 (95% CI, 0.25–1.71) for males, neither of which was significantly different from 1.0, indicating that the combination of the two drugs displays simple additive effects in both sex groups.

**Simulation of Steroid-Sparing Effect of NPX in Arthritis.** The PK/PD/DIS progression model well described all the paw volume data when fitted simultaneously with fixed values of certain drug-specific parameters and yielded quite reasonable parameter estimates. Therefore, model simulations were performed based on the fitted parameters (Table 5) to compare the potential anti-inflammatory effects of different single doses of DEX and combinations with NPX on paw edema. This allowed us to evaluate the degree of potential steroid dose-sparing effects of NPX (i.e., Would giving NPX allow a substantially lower dose of DEX to be used to obtain equivalent efficacy to higher doses of DEX?). The same values of initial disease production rate constant ($k_{\text{init}}$) and paw size on day 0 ($Paw_0$) were assigned to all groups to allow more intuitive comparisons of drug effects. The model-predicted paw edema-time profiles after concurrent administration of 50 mg/kg of NPX and 0.225 mg/kg of DEX, and various single doses of DEX (0.225, 2.25, and 22.5 mg/kg) are displayed in Fig. 6. These simulations show that the combination therapy exhibits different degrees of steroid dose-sparing by NPX in female and male arthritic rats. Combined use of DEX (0.225 mg/kg) with NPX (50 mg/kg) was able to reduce paw edema to a similar extent as a 10-fold higher dose of DEX (2.25 mg/kg) for females and a 100-fold higher dose of DEX (22.5 mg/kg) for males.

**Discussion**

Corticosteroids have long been the mainstay of RA therapy owing to their potent immunosuppressive and anti-inflammatory actions. However, the clinical effectiveness of DEX and other CS is compromised by their long-term and dose-related adverse effects. We extended our previous PK/PD studies with NPX (Li et al., 2017a,b) and DEX (Earp et al., 2008a,b,c) to investigate whether better therapeutic outcomes could be achievable by a combination of DEX and NPX and to ascertain whether sex differences exist.

First, the single and combined immunosuppressive actions of DEX and NPX were evaluated by the in vitro WBLP assay, which measures the mitogenic response of T lymphocytes to drugs. This assay is convenient to assess the nature and intensity of drug-drug interactions as it mimics the natural milieu and requires small blood volumes. The

(0.62 ng/ml) (Table 5) suggested that DEX exhibited modestly stronger potency in males, consistent with the effects of NPX.

**Fig. 4.** Total plasma concentration–time profiles of DEX in female and male healthy rats that received 2.25 mg/kg of s.c. doses of DEX (○), and model-simulated total tissue (dashed lines) and unbound ISF concentrations (dotted lines) of DEX in female and male arthritic rats that received 0.225 mg/kg of s.c. doses of DEX. Solid curves depict model fittings jointly for all healthy animal groups, yielding parameters listed in Table 4.
relative immunosuppressive potency of several therapeutic CS were studied previously using this assay (Mager et al., 2003). DEX inhibits T-lymphocyte proliferation by suppressing nuclear factor-κB, causing reduction of interleukin-2 (IL-2) gene expression, which is responsible for the proliferation and differentiation of T lymphocytes (Toribio et al., 1989).

Insufficient attention has been paid to the immunologic actions of NSAIDs although considerable evidence has shown their capability of suppressing lymphocyte activation (Kelly et al., 1979; Ali et al., 1984; Seng et al., 1990; Cicala et al., 2000). The commonly acknowledged mechanism of action of NSAIDs is to block the synthesis of PG via COX inhibition. However, PG has been reported to inhibit the

![Fig. 5. Disease progression of paw edema in female (upper panels) and male (lower panels) arthritic rats after no drug ( ), 50 mg/kg of NPX ( ), 0.225 mg/kg of DEX ( ), and the combination of NPX 50 mg/kg and DEX 0.225 mg/kg ( ); time courses of paw volumes in male and female healthy rats ( ). Lines are model fittings of all data jointly, yielding parameters listed in Table 5.](image)

### TABLE 5

<table>
<thead>
<tr>
<th>Parameters (Units)</th>
<th>Definition</th>
<th>Estimates (CV%)</th>
<th>Females</th>
<th>Males</th>
</tr>
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<tr>
<td>Disease-specific parameters</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$t_{onset}$ (h)</td>
<td>Time of disease onset</td>
<td>258 (0.44)</td>
<td>307 (0.60)</td>
<td></td>
</tr>
<tr>
<td>$k_{on}$ (1/h)</td>
<td>Loss of edema rate constant</td>
<td>0.015 (6.07)</td>
<td>0.012 (5.37)</td>
<td></td>
</tr>
<tr>
<td>$k_{deg}$ (1/h)</td>
<td>Loss of production rate constant</td>
<td>0.001 (5.98)</td>
<td>0.001 (8.18)</td>
<td></td>
</tr>
<tr>
<td>$k_{g}$ (1/h)</td>
<td>Natural growth rate constant</td>
<td>0.002 (36.0)</td>
<td>0.002 (29.5)</td>
<td></td>
</tr>
<tr>
<td>$P_{awg}$ (mm$^2$)</td>
<td>Paw size at steady-state</td>
<td>72.16 (6.64)</td>
<td>103.1 (6.60)</td>
<td></td>
</tr>
<tr>
<td>$k_{0,D}$ (mm$^2$/h)</td>
<td>Disease production rate constant at $t_{onset}$ for control group</td>
<td>2.18 (3.94)</td>
<td>1.72 (4.26)</td>
<td></td>
</tr>
<tr>
<td>$P_{aw0,D}$ (mm$^2$)</td>
<td>Paw size on day 0 for control group</td>
<td>58.8 (1.43)</td>
<td>73.4 (1.58)</td>
<td></td>
</tr>
<tr>
<td>$k_{in0,D}$ (mm$^2$/h)</td>
<td>Disease production rate constant at $t_{onset}$ for DEX treatment group</td>
<td>1.93 (4.11)</td>
<td>1.82 (4.27)</td>
<td></td>
</tr>
<tr>
<td>$P_{aw0,C}$ (mm$^2$)</td>
<td>Paw size on day 0 for DEX treatment group</td>
<td>58.9 (1.26)</td>
<td>73.6 (1.42)</td>
<td></td>
</tr>
<tr>
<td>$k_{in0,C}$ (mm$^2$/h)</td>
<td>Disease production rate constant at $t_{onset}$ for NPX treatment group</td>
<td>2.06 (4.19)</td>
<td>1.79 (4.39)</td>
<td></td>
</tr>
<tr>
<td>$P_{aw0,N}$ (mm$^2$)</td>
<td>Paw size on day 0 for NPX treatment group</td>
<td>60.0 (1.29)</td>
<td>70.2 (1.55)</td>
<td></td>
</tr>
<tr>
<td>$k_{in0,N}$ (mm$^2$/h)</td>
<td>Disease production rate constant at $t_{onset}$ for combined treatment group</td>
<td>2.00 (4.33)</td>
<td>1.80 (4.50)</td>
<td></td>
</tr>
<tr>
<td>$P_{aw0,B}$ (mm$^2$)</td>
<td>Paw size on day 0 for combined treatment group</td>
<td>57.7 (1.28)</td>
<td>74.8 (1.52)</td>
<td></td>
</tr>
<tr>
<td>Drug-specific parameters</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$I_{max,N}$ (%)</td>
<td>Maximum inhibition of NPX on production of paw edema</td>
<td>0.75 (Fixed)</td>
<td>1 (Fixed)</td>
<td></td>
</tr>
<tr>
<td>$IC_{50,N}$ (ng/ml)</td>
<td>Unbound ISF concentration of NPX at 50% maximum inhibition</td>
<td>221 (Fixed)</td>
<td>136 (Fixed)</td>
<td></td>
</tr>
<tr>
<td>$I_{max,D}$ (%)</td>
<td>Maximum inhibition of DEX on production of paw edema</td>
<td>1 (Fixed)</td>
<td>1 (Fixed)</td>
<td></td>
</tr>
<tr>
<td>$IC_{50,D}$ (ng/ml)</td>
<td>Unbound ISF concentration of DEX at 50% maximum inhibition</td>
<td>0.62 (Fixed)</td>
<td>0.39 (Fixed)</td>
<td></td>
</tr>
<tr>
<td>Drug interaction parameter</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$\phi_2$</td>
<td>Interaction term for inhibition of paw edema production</td>
<td>1.16 (28.82)</td>
<td>0.98 (37.28)</td>
<td></td>
</tr>
</tbody>
</table>
mitogen-induced proliferation of T cells (Goodwin et al., 1977). Therefore, the antiproliferative effect of NSAIDs may involve COX-independent mechanisms. These drugs have been shown to inhibit DNA synthesis and cause cell growth arrest in the G1 phase (Bayer and Beaven, 1979). The nature and intensity of interactions between DEX and NPX on lymphocyte proliferation were evaluated. The slightly synergistic effect of NPX on the antiproliferative action of DEX ($\phi_1 < 1$) is promising in that combination therapy would add to and possibly potentiate the immunosuppressive activity of DEX.

To further examine the PD interactions between DEX and NPX, the joint anti-inflammatory effects of DEX and NPX were investigated in vivo using CIA rats, a well-established animal model that mirrors human RA (Holmdahl et al., 2001). Similar to humans, the CIA rat model exhibits consistent sex differences in the disease progression of arthritis in that females exhibit higher incidences, earlier disease onset, and worse symptoms. We used this animal model previously to assess the PK/PD of both NPX (Li et al., 2017b) and DEX (Earp et al., 2008a). NPX exhibited moderate but significant inhibitory effects on paw edema production in both sexes with slightly stronger efficacy in male rats, and DEX showed potent anti-inflammatory effects in male arthritic rats, with 60% reduction of paw edema compared with untreated controls. In particular, DEX-dependent transrepression of production of the cytokines tumor necrosis factor-$\alpha$, IL-1$\beta$, and IL-6 plays a central role in the alleviation of paw edema and bone erosion in arthritic rats (Earp et al., 2008a).

Both female and male CIA rats were studied so that the influence of sex in the anti-inflammatory actions of DEX with NPX could be compared. The proposed PK/PD/DIS progression model (Fig. 1) characterized well the anti-inflammatory effects of DEX and NPX both as single agents and in combination. The PK models include an extended mPBPK model with nonlinear binding for NPX adapted from our previous study (Li et al., 2017b) and a basic mPBPK model for DEX. The individual drug effects described by the inhibition model was linked to the calculated unbound ISF concentrations of both NPX and DEX. This was done for several reasons. As shown previously elsewhere (Li et al., 2017b), the unbound NPX concentration–time profiles in ISF closely resembled those in synovial fluid, which better correlates with the clinical efficacy of NPX as the synovium is the proposed site of action in RA (Jalava et al., 1977; Netter et al., 1989; Bertin et al., 1994; Day et al., 1995). DEX must permeate cells to bind to receptors and/or alter transcriptional factors, thereby exerting its effects. Rapid equilibration of unbound DEX concentrations between plasma, ISF, and cell water was thus assumed. According to their corresponding mechanisms of action, the joint effect of DEX and NPX was exerted on the production process of paw edema in a joint and noncompetitive manner and was therefore modeled using the dual-drug inhibition approach (Chakraborty and Jusko, 2002; Earp et al., 2004).

The hepatic metabolism of NPX is mainly mediated by cytochrome P450 (CYP) enzymes 1A2 and 2C9 (Miners et al., 1996), which were reported to be induced by DEX in vitro (Gerbal-Chaloin et al., 2002; Vrzal et al., 2009). The NCA results for the primary PK parameters of NPX with or without joint dosing of DEX (Table 3) indicates that the single-dose of DEX had no significant influence on the PK of NPX in either female or male CIA rats.

Previous studies have shown that the PK of DEX in CIA male rats was not altered by RA (Earp et al., 2008c), and DEX concentrations in healthy male rats after s.c. and i.m. dosing have been shown to be identical (Earp et al., 2008a). The same situation was assumed here for female rats, so DEX PK in healthy rats was used to model the PK in arthritic rats. In addition, the PK parameters after i.m. dosing of DEX in female rats were found to be in good agreement with those in males, with 86% bioavailability for both sexes (Samtani and Jusko, 2005).

We also assumed that the PK of DEX was not altered by NPX. Therefore, the same PK parameter values of DEX were applied for both single and combined drug groups. Male arthritic rats exhibited better responses to NPX, which can be partially explained by the higher unbound ISF concentrations of NPX (Li et al., 2017b). A similar sex-specific response was observed for DEX, consistent with previous findings that the anti-inflammatory actions of CS are more effective in males because they have more glucocorticoid-responsive genes (Duma et al., 2010). The IC$_{50}$ values of NPX in both sexes are close to the peak unbound NPX concentrations in synovial fluid after a single dose of 500 mg NPX to arthritic patients (0.13 $\mu$g/ml) (Day et al., 1995), further confirming the feasibility of using the unbound NPX concentrations in ISF/synovial fluid to mimic the biophase concentrations.

In line with the in vitro results, DEX was more efficacious in reducing paw edema production in both sexes than NPX. For combination treatments in CIA rats, the interaction term $\phi_2$ incorporated in the inhibition of paw edema production (eq. 14) is similar to the $\phi_1$ in eq. 9, which reflects the magnitude by which the IC$_{50}$ of DEX is altered.
by NPX. The estimated $\psi_3$ values signified additive effects of combination therapy in both sexes. Nevertheless, model simulations of the anti-inflammatory effects of various dosing regimens demonstrated promising steroid-sparing potential of NPX (Fig. 6), especially for male arthritic rats. The occurrence of additivity is sufficient to produce steroid-sparing effects. The $\psi$ parameter is an empirical term that used to signify the nature of pharmacologic interactions (synergistic, additive, or antagonistic) when detailed mechanisms are unclear. Further studies with the measurements of the molecular factors that drive chronic arthritis and that are relevant to the effects of DEX and NPX, such as the cytokines tumor necrosis factor-$\alpha$, IL-1$\beta$, IL-6, and prostaglandin $E_2$, are necessary to better understand the sex-specific anti-inflammatory actions and the underlying mechanisms of interaction.

Overall, the drug-specific parameters ($I_{\text{max}}$ and $IC_{50}$) of both drugs obtained in vivo were comparable to the in vitro values except that the maximum inhibition of NPX on paw edema in female arthritic rats was less than 1.0 ($I_{\text{max}} = 0.75$). This difference is quite reasonable because the whole blood used in the in vitro study was from healthy male rats and, more importantly, the drug effects in vivo incur many additional complicating factors. In our study, the PD models were directly fitted to the actual measured CPM and pooled paw volume data without normalization by the control data, thus avoiding the impact of the variability from control groups (Woo et al., 2009).

In conclusion, the single and joint effects of DEX and NPX on the in vitro mitogen-stimulated proliferation of T lymphocytes and in assessing paw edema in female and male CIA rats were well described by mathematical models incorporating dual interaction equations for inhibition. With an interaction term $\psi$, the proposed models also indicated the slight extent to which the potency ($IC_{50}$) of DEX is altered by NPX. Our study has thus mainly demonstrated additive immunosuppressive and anti-inflammatory effects of the drug combination. With the promising steroid-sparing potential of NPX, the combined use of DEX and NPX should permit dose reduction of steroids. The models of the anti-inflammatory effects of various dosing regimens demonstrate additive immunosuppressive effects in rats with collagen-induced arthritis. It J Pharmacol Exp Ther 326:546–554.


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