

COVID-19 Treatment)

# Article

# Development and Validation of a Novel HPLC Method to Analyse Metabolic Reaction Products Catalysed by the CYP3A2 Isoform: In Vitro Inhibition of CYP3A2 Enzyme Activity by Aspirin (Drugs Often Used Together in

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Abstract: Aspirin (also known as acetylsalicylic acid) is a drug intended to treat fever, pain, or inflammation. Treatment of moderate to severe cases of COVID-19 using aspirin along with dexamethasone has gained major attention globally in recent times. Thus, the purpose of this study was to use High-Performance Liquid Chromatography (HPLC) to evaluate the in vitro inhibition of CYP3A2 enzyme activity using aspirin in rat liver microsomes (RLMs). In this study, an efficient and sensitive HPLC method was developed using a reversed phase C18 column (X Bridge 4.6 mm  $\times$  150 mm, 3.5  $\mu$ m) at 243 nm using acetonitrile and water (70:30 v/v). The linearity (r<sup>2</sup> > 0.999), precision (<15%), accuracy and recovery (80-120%), limit of detection (5.60 µM and 0.06 µM), limit of quantification (16.98 µM and 0.19  $\mu$ M), and stability of the newly developed method were validated for dexamethasone and 6β-hydroxydexamethasone, respectively, following International Conference on Harmonization (ICH) guidelines. This method was applied in vitro to measure CYP3A2 activity. The results showed that aspirin competitively inhibits  $6\beta$ -hydroxylation (CYP3A2 activity) with an inhibition constant (Ki) =  $95.46 \,\mu$ M and the concentration of the inhibitor causing 50% inhibition of original enzyme activity (IC<sub>50</sub>) = 190.92  $\mu$ M. This indicated that there is a minimal risk of toxicity when dexamethasone and aspirin are co-administrated and a very low risk of toxicity and drug interaction with drugs that are a substrate for CYP3A2 in healthcare settings.

Keywords: cytochrome P450; dexamethasone;  $6\beta$ -hydroxydexamethasone; aspirin; competitive inhibitor; CYP3A activity

# 1. Introduction

Cytochrome P450 enzymes are responsible for the biotransformation of xenobiotics and the metabolism of endogenous compounds [1]. They are mainly found in the liver; however, a number of these enzymes are also expressed in the kidney, small intestine, placenta, and lungs [2]. The synthesis of these enzymes takes place endogenously, and both non-genetic and genetic factors could influence enzyme synthesis.

Many drug–drug interactions of clinical interest change the pharmacokinetic behavior of drugs due to changes in the hepatic metabolic pathway of drugs that are catalysed by CYP450 enzymes [3]. The drug interactions are caused by either CYP450 inhibition or induction. Enzyme induction or inhibition by food, drugs, and medicinal herbs, for instance, can be responsible for changes in the metabolic capacity of these enzymes [4], although inhibition is considered more important in terms of adverse clinical outcomes.

CYP450 enzyme inhibition by drugs could increase the concentration of other metabolizing drugs and could result in drug toxicity issues [5]. CYP3A is the most important family among the identified families of CYP450 enzymes involved in biotransformation



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and metabolism in humans. The CYP3A family is responsible for the metabolism of 70% of drugs as well as for indicating broad substrate specificity [6]. CYP3A is frequently associated with most drug interactions because this isoform is highly inducible and could potentially be inhibited by other drugs and herbs [6]. Of the various isoforms, CYP3A4 is the most abundant isoform present in human liver microsomes and is responsible for the metabolism of various anticancer drugs [7]. Although CYP3A4 activity exhibits 5–10-fold inter-individual variability, it still has substantial potential for clinical applications [7].

Dexamethasone is a steroidal anti-inflammatory drug that is widely used for the treatment of different conditions such as cancer, autoimmune diseases, and chronic inflammatory diseases. Dexamethasone has been shown to reduce deaths in those who have been hospitalized for severe cases of COVID-19 where patients required ventilator support [8]. The use of dexamethasone to treat multiple diseases significantly increases the risk of drug-drug interactions [9]. Tomlinson et al. (1997) stated that dexamethasone is metabolized into 6  $\alpha$  as a minor metabolite and 6 $\beta$ -hydroxymethasone as a major metabolite through CYP3A4 enzyme activity using human liver microsomes.

Previous studies have shown that ketoconazole, ellipticine, and gestodene cause the inhibition of dexamethasone 6-hydroxylation [10]. Although many studies have been conducted in human liver microsomes, rats have the closest metabolic profile to humans and show 71% sequence homology [11]. The CYP3A2 isoform in male-specific rat liver microsomes (RLMs) is responsible for the 6-hydroxylation of dexamethasone (corticosteroid) [11]. Wang et al. [12] stated that codeine administration in male rats can inhibit the metabolism of midazolam (CYP3A2 activity).

COVID-19 infection has rapidly grown into a worldwide pandemic, and this has had a significant impact on human health. Dexamethasone and aspirin (Figure 1) are some of the drugs being used for the treatment of COVID-19 during the pandemic. The combined use of aspirin and dexamethasone has been shown to reduce the symptoms in moderate to severe COVID-19 infection [13]. A recent study has provided evidence in support of primary healthcare centres where they used aspirin and dexamethasone for the therapeutic management of severe COVID-19 patients [14]. Nonsteroidal anti-inflammatory drugs (NSAIDs) and aspirin are the most used therapeutic drugs worldwide [15]. Aspirin is an O-acetyl derivative of salicylic acid (acetylsalicylic acid). It is believed that it transfers this acetyl group to the amino (-NH<sub>2</sub>) and hydroxy (-OH) functionalities present in biological molecules [16]. Aspirin is also a prostaglandin synthase inhibitor that inhibits the production of prostaglandins. It has a non-selective effect on the cyclooxygenase-1 (COX-1) and cyclooxygenase-2 (COX-2) enzymes [17].



Figure 1. (a) Aspirin (inhibitor); (b) dexamethasone (substrate).

In this systematic study, a HPLC method was developed and validated to investigate the impact of aspirin on the activity of the CYP3A2 enzyme in rat liver microsomes, whereby dexamethasone was the substrate. This study promotes the safe administration of COVID-19 drugs (dexamethasone and aspirin) in clinical practice.

# 2. Results and Discussion

## 2.1. HPLC Method Development

The analytical method for the enzymatic assay was optimised and evaluated using the following chromatographic conditions: mobile phase consisting of acetonitrile and water (70% acetonitrile and 30% H<sub>2</sub>O, v/v) with an injection volume of 10 µL, a flow rate of 0.6 mL/min, column temperature set at 25 °C, and a run time of 8 min. The chosen wavelength ( $\lambda$ ) was 243 nm [10]. The selected internal standard was 4-hydroxyoctanophenone.

## 2.2. HPLC Method Validation

# 2.2.1. Linearity and Range

According to the ICH guidelines [18], the linearity was tested using different concentrations (0, 25, 50, 100, 150, and 200  $\mu$ M) of working solutions of dexamethasone and 6 $\beta$ -hydroxydexamethasone (0, 0.2, 0.4, 0.6, 0.8, and 1  $\mu$ M). Working solutions were prepared in the mobile phase (70% acetonitrile and 30% H<sub>2</sub>O, *v*/*v*) and injected into the HPLC for the construction of the calibration curve. The mean peak area obtained from the HPLC chromatograms was plotted against the concentrations of each analyte (dexamethasone and 6 $\beta$ -hydroxydexamethason) to assess the calibration graph (Figure S1). All of the obtained data were corrected for the internal standard. Table 1 represents the results of the linearity study. The r<sup>2</sup> values were within the acceptable ICH criterion (r<sup>2</sup> > 0.99), and the relative standard deviation at each concentration (%RSD < 10%) met the criteria of the ICH guidelines.

Table 1. Linearity data from the proposed analytical method.

Standards	Dexamethasone	6β-Hydroxydexamethasone
Regression equation r <sup>2</sup>	y = 0.2505x + 0.0945 0.999	y = 1.6775x + 0.0385 0.998 0.2 1M
Linear range	23–200 µM	0.2-1 µlvi

## 2.2.2. Specificity and Selectivity

Specificity/selectivity was evaluated by running the diluent blank (70% acetonitrile and 30% H<sub>2</sub>O, v/v) and internal standard solution (15  $\mu$ M) in a 1 mL HPLC vial to check that the outcomes of the analytical method were not altered by the drugs' constituents (Figure 2). Figure 3 shows the separation of the enzyme peak (NADPH (Nicotinamide Adenine Dinucleotide Phosphate Hydrogen-regenerating system) from the inhibitor (aspirin 200  $\mu$ M), metabolite (6 $\beta$ -hydroxydexamethasone), dexamethasone, and internal standard.



Figure 2. Negative control blank shows the specificity of the method.



**Figure 3.** HPLC profile of CYP3A2 assay components. (1) NADPH (Nicotinamide Adenine Dinucleotide Phosphate Hydrogen)-regenerating system, (2) aspirin, (3) 6β-hydroxydexamethasone, (4) dexamethasone, and (5) 4-hydroxyoctanophenone (IS).

### 2.2.3. Limit of Detection (LOD) and Limit of Quantification (LOQ)

The sensitivity of the analytical method was achieved by an evaluation of its LOD and LOQ. The lowest detectable concentration of the analyte is LOD. The minimum measurable concentration in the analytical method and that can be determined quantitatively with suitable precision is LOQ. The standard deviation ( $\sigma$ ) method was used to determine LOD and LOQ using the following formulae:

$$LOD = (3.3 \times \sigma/S) \tag{1}$$

$$LOQ = (10 \times \sigma/S)$$
(2)

where S is the slope of the calibration curve.

The values of LOD and LOQ for dexame thasone and  $6\beta$ -hydroxydexame thasone are presented in Table 2.

Standards	Dexamethasone	6β-Hydroxydexamethasone
LOD	5.60 μM	0.06 µM
LOQ	16.98 μM	0.19 μΜ

Table 2. LOD and LOQ for dexamethasone and 6β-hydroxydexamethasone.

# 2.2.4. Precision

Intraday Precision of Dexamethasone

The intraday method precision was determined by analysing standard samples in triplicate at three different concentrations levels, i.e., low (40  $\mu$ M), medium, (110  $\mu$ M), and high (185  $\mu$ M), of dexamethasone. The results are summarised in Table 3. The percentage relative standard deviation (%RSD) was <5% for dexamethasone, and recovery values were found to be within the range of the ICH acceptance criterion (80–120%). The outcomes revealed that there is no large variation in the concentration of dexamethasone in the intraday analysis. The obtained values were considered satisfactory for the planned use of the method.

**Table 3.** Intraday precision and recovery for dexamethasone (n = 3).

Dexamethasone Standards	Mean (µM)	Recovery <sup>a</sup> (%)	RSD (%)
Low concentration (40 µM)	$39.28 \pm 0.90$	98.21	2.30
Medium concentration (110 µM)	$99.06\pm3.06$	90.05	3.09
High concentration (185 $\mu$ M)	$151.11\pm0.76$	81.68	0.50

Note: Recovery is the ratio of the concentration of analyte recovered to the theoretical concentration. <sup>a</sup> % recovery = (concentration of dexamethasone at 5 h/standard concentration of dexamethasone)  $\times$  100.

Interday Precision of Dexamethasone

The interday precision was determined by injecting standard sample solution for three consecutive days at three concentration levels (low (40  $\mu$ M), medium, (110  $\mu$ M), and high (185  $\mu$ M)). Each sample was run in triplicate. The results are summarised in Table 4, which shows that the recovery values are within the range of the ICH guidelines (80–120%), and the percentage relative standard deviation (%RSD) was <5% for dexamethasone. The outcomes of the experiment revealed that there is no large variation in the interday analysis.

**Table 4.** Interday precision and recovery for dexamethasone (n = 3).

Dexamethasone Standards	Mean (µM)	Recovery <sup>a</sup> (%)	RSD (%)
Low concentration (40 µM)	$45.01\pm2.09$	112.52	4.65
Medium concentration (110 μM)	$110.01\pm2.17$	100.01	1.98
High concentration (185 μM)	$178.92\pm3.13$	96.72	1.75

Note: a % recovery = (concentration of dexamethasone at 5 h/standard concentration of dexamethasone)  $\times$  100.

## Intraday Precision of 6β-Hydroxydexamethasone

The intraday precision of 6 $\beta$ -hydroxydexamethasone was assessed by measuring the low (0.3  $\mu$ M), medium, (0.5  $\mu$ M), and high (0.85  $\mu$ M) concentrations in triplicate in a batch experiment. The results are shown in Table 5 and illustrate that the recovery values (80–120%) are within the range of the ICH guidelines, and the percentage-relative standard deviation values (% RSD) are <0% for 6 $\beta$ h-hydroxydexamethasone. The experiment outcomes revealed that there is no large variation present between the concentration of the 6 $\beta$ -hydroxydexamethasone samples in the intraday precision analysis.

**Table 5.** Intraday precision and recovery for  $6\beta$ -hydroxydexamethasone (n = 3).

6β-Hydroxydexamethasone Standards	Mean (µM)	Recovery <sup>a</sup> (%)	RSD (%)
Low concentration (0.3 μM)	$0.33\pm0.01$	108.60	2.21
Medium concentration (0.5 μM)	$0.50\pm0.02$	100.01	3.64
High concentration (0.85 μM)	$1.02\pm0.09$	119.38	8.82

Note: <sup>a</sup> % recovery = (concentration of  $6\beta$ -hydroxydexamethasone at 5 h/standard concentration of  $6\beta$ -hydroxydexamethasone) × 100.

#### Interday Precision of $6\beta$ -Hydroxydexamethasone

Interday precision was evaluated by measuring the 6 $\beta$ -hydroxydexamethasone standard solution at three concentrations levels (low (0.3  $\mu$ M), medium, (0.5  $\mu$ M), and high (0.85  $\mu$ M) for three consecutive days. The results of the experiment are shown in Table 6. The %RSD (relative standard deviation) is <10% for 6 $\beta$ -hydroxydexamethasone, and recovery values are within the range of the ICH guidelines (80–120%). The experimental data revealed that no large variations are present in terms of inter-assay precision.

6β-Hydroxydexamethasone Standards	Mean (µM)	Recovery <sup>a</sup> (%)	RSD (%)
Low concentration (0.3 μM)	$0.32\pm0.02$	107.61	4.67
Medium concentration (0.5 μM)	$0.50\pm0.03$	99.28	5.90
High concentration (0.85 μM)	$0.79\pm0.03$	93.13	4.13

**Table 6.** Interday precision and recovery for  $6\beta$ -hydroxydexamethasone (n = 3).

Note: <sup>a</sup> % recovery = (concentration of  $6\beta$ -hydroxydexamethasone at 5 h/standard concentration of  $6\beta$ -hydroxydexamethasone) × 100.

## 2.2.5. Stability Study

Solution Stability of Substrate (Dexamethasone)

For stability testing, dexamethasone calibration curves were run for intraday and interday measurements at three concentration levels (40, 110, and 185  $\mu$ M) at ambient temperature. Stability samples were analyzed in triplicate. The results of the stability tests are shown in Table 7.

Table 7. Dexamethasone solutions stability at ambient temperature.

A malastical Damana tama	Ac	tual Concentration (µI	M)	
Analytical Parameters	Intraday	40	110	185
	0 h	35.40	88.64	176.98
Calculated Concentration	5 h	32.20	92.43	202.56
(µ111)	10 h	32.32	92.87	203.56
	0 h	88.51	80.58	95.66
% Recovery	5 h	80.50	84.03	109.49
	10 h	80.80	84.43	110.03
	0 h	111.49	119.42	104.34
% Accuracy <sup>a</sup>	5 h	119.51	115.97	90.51
·	10 h	119.20	115.57	89.97
	Interday	40	110	185
Calculated Concentration	Interday 1	32.02	94.39	181.79
(μM)	Interday 2	34.99	100.43	193.84
	Interday 3	34.12	90.60	187.22
	Interday 1	80.04	85.81	98.26
% Recovery	Interday 2	87.31	91.30	104.78
	Interday 3	85.29	82.36	101.20
	Interday 1	119.96	114.19	101.74
% Accuracy <sup>a</sup>	Interday 2	112.70	108.70	95.22
	Interday 3	114.71	117.64	98.80

Note: <sup>a</sup> % Accuracy = 100 - (calculated concentration - actual concentration/actual concentration) × 100.

The outcomes of the interday and intraday stability checks indicate that there was no variation in the concentration of dexamethasone. In the intraday stability analysis, the concentration of dexamethasone after 10 h was the same compared to the initial concentrationat at 0 h, whereas in the interday analysis, the chromatographic behaviour of dexamethasone remained the same at days 1, 2, and 3 compared to the initial concentrations. The  $r^2$  value obtained from the calibration curves of the intermediate (intraday) analysis was 0.9984

(in accordance with ICH guidelines). The calibration curve for the repeatability analysis was averaged, and the curve equation for dexamethasone was y = 0.277x + 0.6419, where the least regression square values ( $r^2$ ) were within the ICH guidelines (0.9981). Linear equations were further used to calculate the dexamethasone concentrations (low, medium, and high) and their % recovery. The results reveal that the % recovery and accuracy values remained within the range of the ICH guidelines (80–120%). Furthermore, no substantial degradation within the day (intraday) and between days (interday) was observed (Table 7), thus showing that dexamethasone was stable for up to three days at ambient temperature, which is in accordance with the stability study performed by Heda et al. (2011) [19].

## Solution Stability of 6β-Hydroxydexamethasone (Metabolite)

An evaluation of the stability of the working standard solutions of 6 $\beta$ -hydroxydexamethasone was performed at ambient temperature by intraday and interday analysis at three concentration levels (low (0.3  $\mu$ M), medium (0.5  $\mu$ M), and high (0.85  $\mu$ M)). Experiments were performed in triplicate, and the results are summarized below in Table 8.

	Ad	ctual Concentration (μΝ	(I)	
Analytical Parameters -	Intraday	0.3	0.5	0.85
	0 h	0.30	0.47	0.84
Calculated Concentration	5 h	0.29	0.44	0.77
(1111)	10 h	0.27	0.44	0.79
	0 h	98.18	94.82	99.34
% Recovery	5 h	97.22	87.58	90.45
	10 h	88.89	87.75	92.98
	0 h	101.82	105.18	100.66
% Accuracy <sup>a</sup>	5 h	102.79	112.42	109.55
	10 h	111.11	112.25	107.02
	Interday	0.3	0.5	0.85
	Interday 1	0.27	0.46	0.72
Calculated Concentration	Interday 2	0.32	0.47	0.69
(μινι)	Interday 3	0.32	0.46	0.68
	Interday 1	89.49	92.64	84.06
% Recovery	Interday 2	105.46	94.15	81.57
-	Interday 3	106.90	92.10	80.51
	Interday 1	110.51	107.36	115.94
% Accuracy <sup>a</sup>	Interday 2	94.54	105.85	118.43
-	Interday 3	93.10	107.90	119.49

Table 8. 6β-hydroxydexamethasone solutions stability at ambient temperature.

Note: a % Accuracy =  $100 - (calculated concentration - actual concentration/actual concentration) \times 100$ .

The results of the 6 $\beta$ -hydroxydexamethasone stability test showed that there were no obvious changes perceived in the chromatographic behavior and elution profile of the metabolite. In the intraday stability analysis, the chromatographic behaviour of 6 $\beta$ hydroxydexamethasone after 10 h remained the same compared to the initial concentration at 0 h whereas, in the interday analysis, the concentration of 6 $\beta$ -hydroxy dexamethasone stayed the same at days 1, 2, and 3, compared to the initial concentrations. The calibration curve was plotted for intermediate analysis, and the straight-line equation was y = 2.4559x + 0.0767. The averaged calibration curve of the metabolites was constructed using three days of calibration data, and the averaged straight-line equation was (y = 2.0023x + 0.0364). All of the the % recovery and % accuracy values were in the range specified by the ICH guidelines (80–120%), demonstrating the fact that the metabolite solution was stable. Therefore, the results indicate that 6 $\beta$ -hydroxydexamethasone solution was stable at ambient temperature during intraday and interday analysis, which is in accordance with the literature [7]. It is evident that metabolite sample solutions need to be kept and run on HPLC for longer than overnight for the incubation process to see the inhibition.

## 2.2.6. Robustness of the Method

The robustness of the method (50  $\mu$ M aspirin, 0.2  $\mu$ M 6 $\beta$ -hydroxydexamethasone, 25  $\mu$ M dexamethasone and 15  $\mu$ M internal standard) was tested by changing the following parameters: increasing the wavelength by 5 nm, increasing the temperature by 5 °C, and increasing the flowrate. Therefore, replicate injections (n = 3) of standard multianalyte solution were performed. The observations were made based on peak areas and changes in the retention time. Table 9 summarises the effects of the wavelength, temperature, and flow rate variation on the peak area and retention time of the compounds.

**Table 9.** Evaluation of robustness parameters: (A) Change in temperature; (B) change in wavelength; (C) Change in flow rate.

Analytics of Interact	Average t <sub>R</sub>	Average Peak Area	Resolution	
Analytes of Interest —	Normal Conditions (0.6 mL/min, 243 nm and 25 $^\circ$ C)			
Aspirin	2.04	16,162.67		
6β-Hydroxydexamethasone	2.64	127,567.67	All compounds were well	
Dexamethasone	3.08	58,676.00	resolution was achieved.	
4-Hydroxyoctanophenone	6.67	31,991.33		
		A: Temperature (30 °C)		
Aspirin	2.04	12,351.00	All compounds were	
6β-Hydroxydexamethasone	2.64	120,462.67	separated well, with a faster	
Dexamethasone	3.07	55,890.67	elution pattern as the	
4-Hydroxyoctanophenone	6.51	29,137.67	- temperature increased.	
		B: Wavelength (248 nm)		
Aspirin	2.08	10,039.33	All four compounds were	
6β-Hydroxydexamethasone	2.67	75,995.00	separated, but there was a	
Dexamethasone	3.08	49,674.00	decrease in intensity of	
4-Hydroxyoctanophenone	6.61	53,310.00	— metabolite peak.	
		C: Flow rate (0.8 mL/min)		
Aspirin	1.42	6458.00	Peaks were separated with a	
6β-Hydroxydexamethasone	1.99	97,253.00	0.8 mL/min flow rate. All	
Dexamethasone	2.33	38,095.00	compounds showed a faster	
4-Hydroxyoctanophenone	5.02	22,253.33	— and narrow elution pattern.	

The outcomes of the robustness test showed that the developed method could optimally perform when small changes are made to parameters such as the wavelength, temperature, and flow rate. Good dexamethasone,  $6\beta$ -hydroxydexamethasone, and internal standard separation were achieved for all of the temperature, wavelength, and flow rate variations.

## 2.3. Optimisation of Incubation Time for Incubation System In Vitro

The incubation time was optimised (Figure 4). The  $6\beta$ -hydroxydexamethasone formation rate from dexamethasone (50  $\mu$ M) by cytochrome P3A2 was linear and took place over 40 min. Thus, the optimal incubation time for CYP3A2 was 40 min.



Figure 4. Effects of incubation time on 6β-hydroxydexamethasone production.

# 2.4. Inhibitory Effects of Aspirin on CYP3A2 Activity in Rat Liver Microsomes

To see the effects of aspirin on rat CYP activities, different concentrations of dexamethasone (10, 20, 30, 40, and 50  $\mu$ M) in the presence of 0, 50, 100, and 200  $\mu$ M aspirin were investigated. Microsomal proteins (0.5 mg/mL) were incubated for 40 min at 37 °C with dexamethasone, 3.0 mM magnesium chloride (MgCl<sub>2</sub>), 1.0 mM Ethylenediaminetetraacetic acid (EDTA), 1.0 mM Nicotinamide Adenine Dinucleotide Phosphate Hydrogen (NADPH), and 0.067 M phosphate buffer (pH 7.4). Aspirin competitively inhibited the production of 6 $\beta$ -hydroxydexamethasone, as presented in Figure 5 and Table 10, whereas the effect of aspirin on 6 $\beta$ -hydroxydexamethasone production when using 10–50  $\mu$ M substrate is shown in Figure 6. Aspirin (0–200  $\mu$ M), even at lower than therapeutically relevant concentrations (150–300  $\mu$ g/mL), causes a 50% inhibition of CYP3A2 enzyme activity (IC<sub>50</sub>), as presented in Table 10.



Figure 5. Representative Lineweaver–Burk plot for CYP3A2 enzyme inhibition on dexamethasone metabolism into  $6\beta$ -hydroxydexamethasone with 0, 50, 100, and 200  $\mu$ M aspirin in rat liver microsomes. Average data are taken from triplicate measurements.

(Inhibition Parameters)					
Aspirin Concentration	Κ <sub>m</sub> (μΜ)	$V_{max}$ ( $\mu M^{-1} \cdot min^{-1}$ )	$Cl_{int}$ ( $\mu M^{-2} \cdot min^{-1}$ )	á	%Inhibition
0 μM Aspirin	$21.23\pm0.51$	$0.0127 \pm 1.53  imes 10^{-4}$	$0.0006 \pm 1.10  imes 10^{-5}$	-	-
50 µM Aspirin	$23.83\pm0.31$	$0.0123 \pm 1.15  imes 10^{-4}$	$0.0005 \pm 4.00  imes 10^{-6}$	$1.03\pm0.01$	$12.44 \pm 1.20$
100 µM Aspirin	$26.13\pm0.70$	$0.0127 \pm 7.94  imes 10^{-4}$	$0.0005 \pm 5.00  imes 10^{-5}$	$1.02\pm0.06$	$23.29\pm3.30$
200 µM Aspirin	$32.57\pm0.35$	$0.0123 \pm 1.73  imes 10^{-4}$	$0.0004 \pm 1.00  imes 10^{-5}$	$1.04\pm0.01$	$53.64 \pm 1.76$

**Table 10.** Pharmacokinetic parameters of inhibitory effects of aspirin on enzyme metabolism. The mean result is taken  $\pm$  SD (n = 3). Note: p < 0.001.



**Figure 6.** Effect of aspirin on CYP3A2 activity in rat liver microsomes. Control activity was taken as 100%. Average data are taken from triplicate measurements.

The inhibition of hepatic cytochrome P450 activity is one of the most significant mechanisms of a drug interaction. Severe adverse events have been associated with drug interactions caused during coadministration [20]. Thus, regulatory authorities need to conduct interaction studies both in vitro and in vivo during drug development.

Aspirin and dexamethasone have been used for the treatment of COVID-19 during the pandemic (2020), and the associated use of aspirin and dexamethasone causes a reduction in COVID-19 mortality. This research is the first to investigate the effects of aspirin on the metabolism of dexamethasone (CYP3A2 enzyme activity).

This study has demonstrated that dexamethasone is metabolised in rat liver microsomes to  $6\beta$ -hydroxydexamethasone. The in vitro findings indicated that aspirin at doses between 0–200  $\mu$ M acts as a competitive inhibitor and may weakly inhibit cytochrome P4503A2 enzyme activity. The V<sub>max</sub> (V<sub>max</sub> is the maximum rate of a reaction catalysed by an enzyme) for the inhibition experiments at three aspirin concentration levels (50, 100, and 200  $\mu$ M) remained the same as it did for negative control (without Inhibitor). Whereas higher K<sub>m</sub> (the concentration of substrate requiring the half-maximal activity of the enzyme) values were obtained in the presence of the inhibitor, as shown in Table 10. Alpha prime ( $\alpha'$ ) indicates the effect of an inhibitor on an enzyme's affinity for its substrate, and similarly, the effect of the substrate on the enzyme's affinity for the inhibitor) [21]. The CL<sub>int</sub> (the intrinsic ability of hepatic CYP450 enzymes to metabolise the drug) of drugs is often predicted based on in vitro data obtained from kinetic analysis (Michaelis–Menten). It was calculated as a ratio of in vitro kinetic constants  $V_{max}$  and  $K_m$  [22], as presented in Table 10.

The results show that aspirin is a competitive inhibitor that binds to the active site of the CYP3A2 enzyme and decreases the activity of the CYP3A2 isoform, which is responsible for the 6 $\beta$ -hydroxylation of dexamethasone in male rats, with the K<sub>i</sub> (the binding affinity between the enzyme and inhibitor) = 95.46  $\pm$  4.25  $\mu$ M and IC<sub>50</sub> (inhibitor concentration required to inhibit 50% of the enzyme activity) = 190.92  $\pm$  8.50  $\mu$ M. A compound with an IC<sub>50</sub> value below 1  $\mu$ M is considered to be a strong inhibitor, and it is considered to be a weak inhibitor if the IC<sub>50</sub> value is more than 50  $\mu$ M [23]. Therefore, aspirin has a weak inhibitory effect on CYP3A2 isoform activity.

These in vitro findings would be useful for future in vivo studies in the healthcare sector. The simple metabolic profile of dexamethasone shows that this steroid could be useful as an in vivo probe for CYP3A4 [10]. Further in vitro and in vivo clinical studies on the potential risks associated with the interactions of dexamethasone and aspirin in humans are required.

## 3. Materials and Methods

# 3.1. Chemicals

The compound  $6\beta$ -Hydroxydexamethasone was purchased from Cayman Chemical Company (Ann Arbor, MI, USA). Dexamethasone from Tokyo Chemical Industry CO.LTD (Tokyo, Japan), and 4'-hydroxyoctanophenone with a purity greater than 99% were obtained from Alfa Aesar (A Johnson Matthey Company, London, UK). Potassium phosphate monobasic, potassium phosphate dibasic, glucose-6-phosphate (G-6-P), glucose-6-phosphate dehydrogenase (G-6-PDH), phosphoric acid (85% *w/w*), NADP<sup>+</sup> (Nicotinamide Adenine Dinucleotide Phosphate), EDTA (Ethylenediaminetetraacetic acid), and magnesium chloride (MgCl<sub>2</sub>) were purchased from Merck. Diethyl ether was purchased from Fischer Scientific (Bishop Meadow Road, Loughborough, UK), and ethyl acetate was purchased from VWR chemicals (France). HPLC-grade acetonitrile and water were obtained from Merck, Co. (Old Brickyard, Gillingham, UK).

The pooled liver microsomes from male rats (Sprague Dawley) were stored at -80 °C and were purchased from Merck (Old Brickyard Road, Gillingham, UK).

#### 3.2. Instruments

A 570 pH Meter from JANEWAY Limited (Beacon Road, Stone, Staffordshire, ST15 0SA, UK) was purchased. A high-performance liquid chromatographic system (LC-2010A HT Shimadzu Corporation, Kyoto, Japan) was used for the analysis and was equipped with a degasser, a UV detector, a low-pressure quaternary pump, a LC column oven, and an autosampler. A WATERS (Waters Corporation, 34 Maple St., Milford, MA, USA) C18 column (15 mm  $\times$  4.6 mm, 3.5 µm particle size) was used for chromatographic separation. The chromatographic data were processed using Shimadzu HPLC 1 LabSolutions (software processing system). The shaking incubator used for the incubation of the tubes was from Eppendorf Ltd. (Stevenage, UK).

## 3.3. Cytochrome P450 Assay

#### 3.3.1. Dexamethasone 6β-Hydroxylation Assay for CYP3A2

Assay development and validation were carried out using a LC-2010A HT Module HPLC system (Shimadzu, Toyko, Japan). Target components (4-hydroxyoctanophenone used as an internal standard, aspirin as the inhibitor, dexamethasone as the CYP3A2 substrate, and  $6\beta$ -hydroxydexamethasone as the CYP3A2 metabolite) were separated on a C18 column.Good separation of the compounds of interest was achieved using the optimised acetonitrile/water (70%/30%, v/v) mobile phase. The retention time ( $t_R$ ) of the four compounds is shown in Table 11. The chromatographic separation was performed at 25 °C and at a flow rate of 0.6 mL/min. A 10 µL solution volume was injected for HPLC analysis, and all the components were detected at the 243 nm wavelength. Table 11 provides

information concerning efficiency (N), where N =  $5.54 \times (t_R/W_{\frac{1}{2}})$  and  $W_{\frac{1}{2}}$  = width at half peak height; plate height (H), where H(cm) = column length/N; asymmetry factor (AsF), where AsF = B/A, and A is the distance from the leading edge of the peak to the midpoint of the peak measured at 10% of peak height, and B is the distance from the midpoint of the peak to the trailing edge of the peak measured at 10% of the peak height; and resolution (R<sub>s</sub>), where Rs =  $2\Delta t_R/0.5(W_1 + W_2)$ , and W is the width at the peak base. As can be seen from Table 11, these chromatographic parameters show that the methodology has been suitabialy optimised, with total run time under 8 min.

	Aspirin	6β-Hydroxydex- amethasone	Dexame- thasone	Internal Standard
Retention Time (t <sub>R</sub> )	2.09	2.65	3.08	6.66
Efficiency (N)	1078	970	3363	6137
Plate Height (H)	$1.39 imes10^{-2}$	$1.55  imes 10^{-2}$	$4.46 imes10^{-3}$	$2.44 imes10^{-3}$
Resolution (R <sub>s</sub> )	1	.58 1.3	33 1	1.01
Asymmetry Factor (AsF)	0.97	1.03	1.07	1.07

Table 11. Column efficiency, plate height, asymmetry, and resolution of compounds of interest.

#### 3.3.2. Microsomal Incubations Procedure

Microsomal protein (0.5 mg/mL) was incubated at 37 °C, with a serial range of dexamethasone (10, 20, 30, 40, and 50  $\mu$ M), magnesium chloride (3.0 mM), NADPH (1.0 mM), Glucose-6-Phosphate (5 mM), (Glucose-6-Phosphate Dehydrogenase (1.7 units/mL), Ethylenediaminetetraacetic acid (1.0 mM EDTA), and 0.067 M potassium phosphate buffer (pH 7.4) in a final volume of 500  $\mu$ L. A serial range of aspirin (0, 50, 100, and 200  $\mu$ M, dissolved in mobile phase) was added to the incubation mixture in triplicate. Incubations were for 40 min and were initiated by the addition of NADP<sup>+</sup> (Nicotinamide Adenine Dinucleotide Phosphate) to the mixture after the pre-incubation of all of the components for 5 min in a water bath (T = 37 °C).

Ice-cold-grade acetonitrile containing 15  $\mu$ M 4-hydroxyoctanophenone (as an internal standard) was added to terminate the reaction. Dexamethasone and metabolite were extracted with ethyl acetate (3 mL) and then with diethyl ether (3 mL). The polar extracts were evaporated to dryness, and the residues were dissolved in the mobile phase (70% acetonitrile and 30% water, v/v) and made up to 1000  $\mu$ L in volume. An amount of 10  $\mu$ L of each sample was injected into the HPLC instrument for analysis.

#### 3.4. Preparation of Standard Substrate and Metabolite Solutions

For the cytochrome P3A2 enzyme assay, 4-hydroxyoctanophenone was used as an internal standard. An amount of 0.0010 g of the powder was dissolved in acetonitrile in a 10 mL volumetric flask. The final stock (15  $\mu$ M) was prepared by adding 165  $\mu$ L of the stock in a volumetric flask containing 49 mL and 835  $\mu$ L of mobile phase (70% methanol + 30% water, *v*/*v*).

Aspirin (0.0018 g) (C = 1000  $\mu$ M) was weighed accurately and dissolved in a 10 mL volumetric flask in methanol. Serial dilutions of aspirin (200, 100, and 50  $\mu$ M) were performed. An amount of 0.0039 g of dexamethasone (C = 1000  $\mu$ M) was weighed and added to a 10 mL volumetric flask containing methanol. Different concentrations of dexamethasone (50, 40, 30, 20, and 10  $\mu$ M) from the stock solution were prepared in the mobile phase (70% methanol + 30% water, v/v). The metabolite (6 $\beta$ -hydroxydexamethasone) stock solution of 2  $\mu$ M was prepared in the mobile phase (70% methanol + 30% water, v/v), and serial dilutions (0.2, 0.4, 0.6, 0.8, and 1  $\mu$ M) were prepared from the stock.

## 3.5. Optimisation of Incubation Time In Vitro

The dexamethasone (substrate) and protein conentrations were fixed in the incubation system, and samples were incubated for 10, 20, 30, 40,50, 60, 70, and 100 min. The samples were prepared and incubated as described in Section 3.3. The concentrations of the produced metabolite  $6\beta$ -hydroxydexamethasone were calculated from the standard calibration curve. The optimal incubation time was determined by the linear relationship between the time and metabolite concentration.

#### 3.6. Data Analysis

Data were analysed using Microsoft Excel 2010 software for validation and the kinetic parameters. All results are shown as mean  $\pm$  S.D. The least-square regression analysis was performed to calculate the concentration of the metabolite produced by the CYP reaction. Secondary Lineweaver–Burk plots and Michaelis–Menten plots were plotted to find kinetic parameters such as K<sub>i</sub>, V<sub>max</sub>, K<sub>m</sub>, Cl<sub>int</sub>, and  $\alpha$ . Inhibition was assumed to be competitive based on the obtained data and based on a visual inspection of the Lineweaver–Burk plot. The inhibitor concentration that would cause a 50% reduction in enzyme activity (IC<sub>50</sub>) was calculated by plotting the percentage of the remaining control enzyme activity versus the inhibitor concentration.  $K_m$  values were used to calculate the percentage of inhibition.

#### 4. Conclusions

In conclusion, a HPLC method was developed, and analytical parameters including linearity, precision, % recovery, linear regression, LOD, and LOQ were derived for dexamethasone and  $6\beta$ -hydroxydexamethasone. All of the analytical parameters were validated in accordance with the ICH guidelines. In vitro incubation assays using rat liver microsomes were adopted to determine the effects of aspirin on dexamethasone metabolism (CYP3A2 activity), as aspirin and dexamethasone have been used together for COVID-19 treatment. Our findings revealed that aspirin acts as a competitive inhibitor and that it has a weak inhibitory effect on dexamethasone metabolism in rat liver microsomes. The outcomes of the study further suggest the safe use of aspirin and dexamethasone in clinical practice. Nevertheless, further in vivo inhibition studies are required to consider this interaction and its implications more completely for patient care.

**Supplementary Materials:** The following supporting information can be downloaded, Figure S1: Typical HPLC chromatogram of CYP3A2 assay components obtained with 0.8 mL/min flow rate. The peaks marked are: (1) aspirin, (2)  $6\beta$ -hydroxydexamethasone, (3) dexamethasone, (4) internal standard, respectively.

**Author Contributions:** All authors conceived and designed the experiments. A.H. performed the experiments. J.B. and D.P.N. made significant contributions to the data analysis and interpretation. A.H. wrote the first draft of the manuscript. The submitted version of the manuscript was revised and approved by all authors. All authors have read and agreed to the published version of the manuscript.

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Sample Availability: Samples of the compounds are available from the authors.

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