

# **APPLICATIONS**

# Rapid Sample Extraction of Chloroquine, Hydroxychloroquine, and Desethylchloroquine from Serum for LC-MS/MS Analysis

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### Overview

In this fast analysis, a direct, in-plate protein precipitation and filtration is applied to increase the throughput of LC-MS/MS measurement for chloroquine, hydroxychloroquine and the metabolite desethylchloroquine from serum. The automatable sample extraction procedure produced excellent recovery values.

# **Sample Preparation**

96-Well Plate:	Impact™ Protein Precipitation Plate		
Part No.:	<u>CE0-7565</u>		
Dispense:	450 μL of Methanol into each well		
Add:	150 μL serum into each well		
Vortex:	2 minutes		
Filter:	Place an empty collection plate below the Impact protein precipitation plate and pass sample through by applying vacuum, positive pressure, or centrifuge		
Dilute:	Dilute filtrate 1:5 with aqueous 0.1 % Formic acid		
Inject:	Inject onto LC-MS/MS		

## **LC-MS/MS Conditions**

Column: Kinetex® 2.6 μm F5

**Dimension:** 50 x 2.1 mm **Part No.:** 00B-4723-AN

Mobile Phase: A: 0.1 % Formic Acid in Water

**B**: Methanol

 Gradient:
 Time (min)
 %B

 0.0
 2

 2.0
 7

 3.0
 15

 3.2
 40

 4.2
 95

 6.0
 2

Flow Rate: 0.5 mL/min
Temperature: 20 °C

Detection: MS/MS (MRM, SCIEX® 4000 QTRAP®)

Sample: 1. Desethylchloroquine

2. Hydroxchloroguine

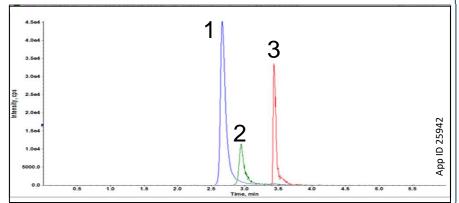
3. Chloroquine

## Recovery Value (%) and CVs (%)

	Recovery	CV (n=6)
Chloroquine (CIQ)	97.1	10.6
Hydroxychloroquine (HClQ)	95.8	16.9
Desethylchloroquine (DesClQ)	103.3	16.2

# **Advantages of the Workflow**

- 1. High-throughput
  - a. 96-well format allows for rapid, 10 minutes sample preparation of 96 samples at once
  - Analyze samples by LC-MS/MS in 6 minutes
- 2. Easily automatable sample preparation
  - a. Effectively hold the organic solvent prior to and during the protein precipitation for easy sample cleanup
  - Remove precipitated proteins while sample and Chloroquine and its metabolites pass through and into a collection plate, no transfer steps required







# PPLICATIONS

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