

# IDENTIFICATION OF THE IMPURITIES OF BUDESONIDE USING SMALL PARTICLE LIQUID CHROMATOGRAPHY AND Q-TOF MASS

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Warren Potts iii; Michael Jones; Robert Plumb  
Waters Corporation, Milford, MA

## INTRODUCTION

The demands on the analytical laboratory to qualitatively and quantitatively determine active pharmaceutical ingredients and degradants continues to increase. The FDA regulations require companies to develop methods for their analysis and characterization of the APIs, as well as the impurities/degradants that could arise from the synthesis process, raw material provider, and/or storage conditions. The utilization of UV or PDA data alone for these analyses is often inadequate. Complications arise in many situations where compounds have (a) poor to no UV absorbance whether from lack of chromophores or (b) low level impurity concentrations exhibit poor UV spectral quality and peak purity becomes more difficult to identify when co-elution occurs. Mass spectral analysis becomes more essential as FDA regulations for impurity content reporting continues to decrease from 0.1% a few years ago to 0.05% today with expectations to decrease as instrumentation detection limits and techniques continue to become more sensitive. The importance of exact mass and MS fragmentation products to determine the structure of degradants/impurities provides a higher understanding to the relationship of the impurity/degradant origin. We will demonstrate the utility of a UPLC-PDA-MS system and show the significant benefits in resolution, speed, and sensitivity using the ACQUITY UPLC™ System and how this configuration will impact the identification of pharmaceuticals and their related substances.

To best illustrate this concept, we will analyze the pharmaceutical drug substance; budesonide. Budesonide is a glucocorticosteroid used for the treatment of asthma via various matrices and inhalation mechanisms. The official European Pharmacopoeia assay was used as a guidance for the redevelopment of the budesonide assay and related substances for use with UPLC-MS. The new method will be used to determine various required qualitative system performance (eg; resolution, S/N, theoretical plates, tailing, symmetry factors). We will also demonstrate the quantification benefits (eg; limits of detection, limits of quantification) of this configuration. The impurity profiles of multiple batches from three manufacturers of pharmaceutical grade budesonide will be assessed and tested for EP related impurity compliance. Exact mass MS data will be collected to determine similarities/differences between the impurity profiles of the vendor batches. This increased performance makes UPLC™/PDA/MS the ideal tool for purity profiling.

## EXPERIMENTAL

### Materials:

Budesonide: Spectrum quality products (New Brunswick, NJ); lot numbers: UI0628 (EP Rx grade); 98.0% - 102.0% and lot # RB2362 (research grade). Sigma Chemical Co. (St. Louis, MO); lot 81K1654; 99%. Molekula; batch# 52459 (Dorset, UK).

Reagents: Acetonitrile Optima; Fisher Scientific (Fairlawn, NJ); Lot#050580. Ammonium formate 97%; Sigma-Aldrich (St. Louis, MO); batch # 04507AC. Formic acid 98%-100%; Reidel-deHaen.

## INSTRUMENTATION

### UPLC Conditions

Instrument: ACQUITY UPLC  
Column: ACQUITY UPLC™ BEH C<sub>18</sub>  
Dimensions: 100 x 2.1mm, 1.7μm  
Mobile Phase: 68% 20mM Ammonium formate (pH 3.6)/32% acetonitrile  
Flow Rate: 0.60 mL/min  
Temperature: 40°C  
injection Volume: 5 μL; full loop injection mode  
Weak Wash: 68:32 (water: acetonitrile) 600μL  
Strong Wash: none  
Detection: ACQUITY PDA @ 240 nm with High sensitivity flow cell  
Software: Empower™ 2 CDS

### MS Conditions

Instrument: Waters LCT-Premier XE  
Software: Masslynx 4.1

### Tune Page Parameters:

Source: ES+  
Capillary (V): 3.2  
Sample Cone (V): 35 for reference  
20 for analyte  
Extraction Cone (V): 4.5  
Desolvation Temp (°C): 350.0  
Source Temp (°C): 150.0  
Cone Flow (L/Hr): 0.0  
Desolvation Flow (L/Hr): 800.0

### Tof Settings

Acquisition Range: 100 - 800Da  
Scan Time: 0.20s  
Interscan delay: 0.05s  
Lock mass: 100fmol leucine/enkephalin @ ~20μL/min

## UPLC-MS METHOD COMPATIBILITY

### Resulting Method

There are three published methods for the separation of budesonide and the related impurities.<sup>(1-3)</sup> Two of these methods are not "MS compatible" due to the use of phosphate buffers.<sup>(2,3)</sup> The European Pharmacopoeia requires the following system suitability specifications based on a 500μg/mL budesonide test solution and reference solutions: (a) The resolution between the R-epimer and S-epimer is not less than 1.5, (b) run time 1.5x the retention of S-epimer, (c) the symmetry factor for the R-epimer peak is less than 1.5, (d) the theoretical plates calculated for the R-epimer peak is at least 4000, (e) and after six injections of the 500μg/mL reference solution the %RSD of the sum of the peak areas of the two epimers is at most 1.0%.<sup>(2)</sup> In converting the method to UPLC we evaluated four currently available hybrid UPLC column chemistries, various pH, buffer concentrations, and temperatures. There was no significant change in separation when the pH was varied from 2.5 - 4.0, neither for pH 9.0 10mM ammonium bicarbonate nor for buffer concentrations from 10mM to 25mM. The resulting method showed the ACQUITY BEH C<sub>18</sub> 2.1 x 100mm column provided the best separation of the impurities from the budesonide API at 40°C (Figure 1). 20mM ammonium formate buffer to pH 3.2 with formic acid was chosen to keep pH consistent with the EP method.

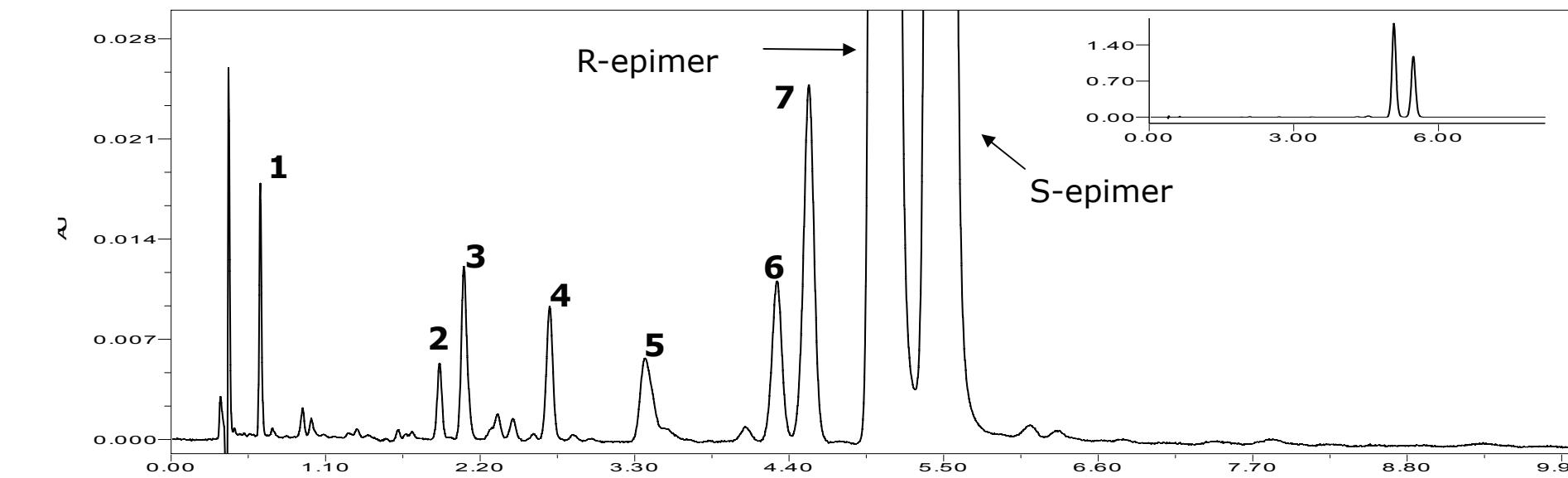


Figure 1: A 5μL full loop injection of 500μg/mL budesonide EP grade (Spectrum Quality Products) standard solution. UPLC method conditions: ACQUITY BEH C<sub>18</sub> 2.1 x 100mm; 1.7μm, 68% 20mM ammonium formate buffer to pH 3.2: 32% acetonitrile; wavelength at 240nm; temperature at 40°C; flow rate at 0.6mL/min; 11,500psi. The additional peaks 1 thru 7 are identified as impurity peaks above 0.05% area.

### Suitability Results

The %RSD of the sum of the areas of both epimers was 0.3% (n=6 injections) for the 500μg/mL budesonide EP grade standard solution. The results in the table below are for the budesonide EP 500μg/mL standard solution. The requirements of the minimum European Pharmacopoeia specifications are met.

Name	Retention Time	Resolution	Symmetry Factor	Signal/Noise	EP Plates
R- epimer	5.073	N/A	1.05	10262	17011
S- epimer	5.476	2.46	1.02	6646	17390

## IMPURITY PROFILE COMPARISONS

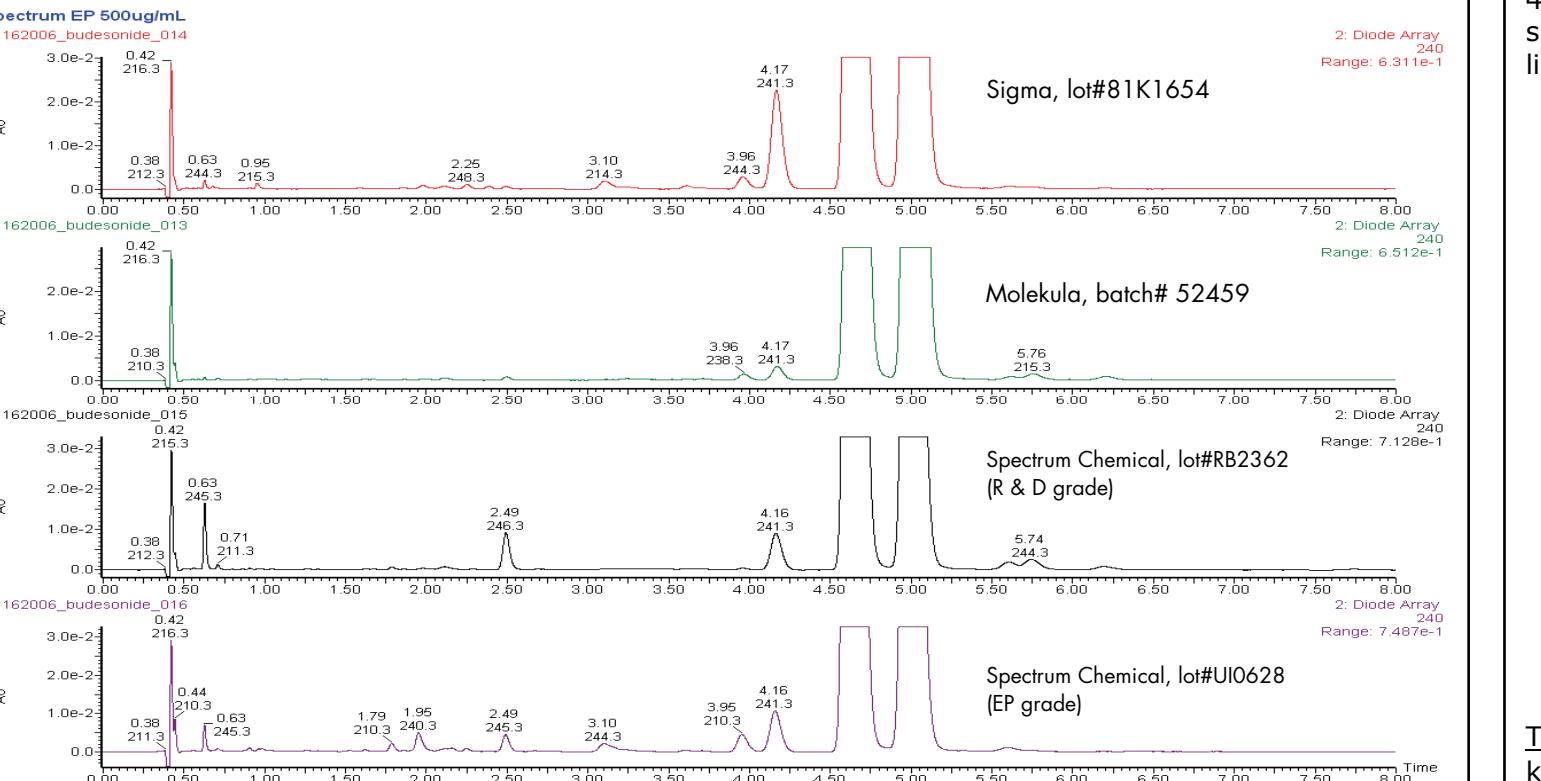
### EP Related Substances Test

The EP related substances test as described in the Budesonide EP monograph was performed on four different batch lots of budesonide which were purchased from three different vendors. Test solutions (500μg/mL) were prepared for each batch lot. Each test solution was diluted to yield two reference solutions each with concentrations of 2.5μg/mL and 7.5μg/mL with each representing 0.5% and 1.5% of the 500μg/mL solution, respectively.

European Pharmacopoeia Related Substances Test Specification	Spectrum EP grade 98% - 102%	Sigma >99% purity	Molekula 100.2%	Spectrum Research grade (no spec)
Individual Impurities (x < 2.5μg/mL. Σ of epimers areas)	Fail	Fail	Pass	Fail
Total Impurities (x < 7.5μg/mL. Σ of epimers areas)	Fail	Fail	Pass	Fail
R-epimer/S-epimer Ratio (S-epimer is 40.0% to 51% Σ of epimers areas)	59.24%/40.76%	50.49%/49.51	51.38%/48.62	58.66%/41.34
Purity	98.24%	97.99%	99.52%	98.07%

### Impurity Profile Comparisons

The impurity profiles for each lot was compared using the 500μg/mL test solution. There were varied impurities detected and at different levels for each sample when prepared fresh. It was observed after exposure to light and time, the profiles demonstrated more similarities.



## EXACT MASS

### MS Utility

Using LC/MS in purity profiling experiments aids in peak tracking during method development and facilitates a high level of confidence with known analyte identification when exact mass is employed. ACS requires < 5ppm mass accuracy for patent submission and publication. The mass accuracy of the known peaks in each of the budesonide batches are less than 3ppm (below). By coupling exact mass with tools like elemental composition, it is possible to predict molecular formulas for the unknown analytes such as the 447.2380amu.

