



Analysis of Nucleosides in Municipal Wastewater by Large-Volume Liquid Chromatography Tandem Mass Spectrometry

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1 **Analysis of Nucleosides in Municipal Wastewater by Large-Volume Liquid**
2 **Chromatography Tandem Mass Spectrometry**

3

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4 10 **Abstract:** Nucleosides are components of both DNA and RNA, and contain
5
6 11 either a ribose (RNA) or 2deoxyribose (DNA) sugar and a purine or pyrimidine
7
8 12 base. In addition to DNA and RNA turnover, modified nucleosides found in urine
9
10 13 have been correlated to a diminished health status associated with AIDS,
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12 14 cancers, oxidative stress and age. Nucleosides found in municipal wastewater
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14 15 influent are potentially useful markers of community health status, and as of now,
15
16 16 remain uninvestigated. A method was developed to quantify nucleosides in
17
18 17 municipal wastewater using large-volume injection, liquid chromatography, and
19
20 18 mass spectrometry. Method accuracy ranged from 92 to 139% when quantified
21
22 19 by using isotopically labeled internal standards. Precision ranged from 6.1 to
23
24 20 19% of the relative standard deviation. The method's utility was demonstrated by
25
26 21 the analysis of twenty-four hour composite wastewater influent samples that were
27
28 22 collected over a week to investigate community nucleoside excretion.
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30 23 Nucleosides originating from RNA were more abundant than DNA over the study
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32 24 period, with total loads of nucleosides ranging from 2 to 25 kg/day. Given this
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34 25 relatively high amount of nucleosides found over the study period they present an
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36 26 attractive analyte for the investigation of community health.
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28 INTRODUCTION

29 Municipal wastewater contains community scale information¹⁻³. There
30 have been numerous methods developed for the quantification of illicit drugs⁴⁻⁶,
31 personal care products^{7,8}, and pharmaceuticals^{9,10} in municipal wastewater
32 influent and effluent. Endogenous compounds such as steroids have also been
33 investigated in municipal wastewater¹¹. The concentrations of these substances
34 are converted to mass loads by the multiplication of wastewater volume in order
35 to account for dilution¹²⁻¹⁴. Community drug use, which is an important
36 indication of community health has been investigated with the use of influent
37 loads^{6,12}. Prescription pharmaceuticals also have some potential as indicators
38 of community health with the use of prescription records, dose estimation and
39 pharmacokinetic data when compared to loads. Possible prescription drug loads
40 that could be used to determine the overall health status of a community could
41 include anti-cancer, anti-viral drugs or metabolites. The usefulness of
42 prescription drug loads may be limited due to potentially sparse use throughout
43 the community.

44 Nucleosides are components of both DNA and RNA, and contain a purine
45 or pyrimidine base and either a 2deoxyribose (DNA) or a ribose (RNA) sugar.
46 Nucleosides are damaged in oxidation or alkylating reactions that produce
47 modified nucleosides which remain incorporated into DNA, repaired or ultimately
48 excreted in urine^{15,16}. Nucleoside modification is typically the result of reactions
49 that modify the base. Alkylating electrophiles commonly form adducts at N^7 , N^3 ,
50 and O^6 guanine, and at N^3 and N^1 of adenine¹⁷. Electrophilic free radicals (i.e.

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3 51 superoxide anion, hydroperoxyl radical, hydrogen peroxide, and hydroxyl radical)
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6 52 attack sites of rich electron density of pyrimidine (guanine and adenine) and
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8 53 purine (cytosine, uracil and thymine) bases and form oxidative adducts ¹⁵. The
9
10 54 generation of these free radicals can be either be from endogenous and or
11
12 55 exogenous sources such as mitochondria, inflammatory cells, redox cycling
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14 56 compounds (e.g. diphenols, quinones, nitroaromatics) and metals ¹⁷. The
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16 57 modified nucleoside 8-hydroxyguanosine has been studied in urine ^{18, 19} and
17
18 58 organ tissue ²⁰ as a marker for aging. The urinary concentrations of modified and
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20 59 un-modified nucleosides have been used as markers of health status, within
21
22 60 small groups of individuals ^{21, 22}. The excretion profile of modified and
23
24 61 unmodified nucleosides have be observed to be a function of age ^{23, 24}, oxidative
25
26 62 stress ^{25, 26}, environment, cancer ^{21, 27, 28}, lifestyle ²⁹, pregnancy ³⁰ and increased
27
28 63 exposure to UV radiation ³¹. Given the variability of modifications, there are a
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30 64 number of modified nucleosides that have not been investigated due to the lack
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32 65 of analytical standards. 8-hydroxydeoxyguanosine (8OHdG) has been
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34 66 mentioned as a possible marker of community health in wastewater but never
35
36 67 fully investigated ³. While there is no “ideally” health community, an investigation
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38 68 of nucleoside loads in municipal wastewater influent could potentially be
39
40 69 compared to more traditional community (i.e. public) health data such as cancer
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42 70 occurrence, community levels of HIV, and environmental data that could impact
43
44 71 health, such as air quality or proximity to sources of pollution and or radiation.
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46 72 Thus, nucleoside loads have the potential to be complementary to these
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48 73 traditional data.
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3 74 The objective of this study is to develop a method using large-volume
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5 75 injection, liquid chromatography, mass spectrometry to quantify nucleosides and
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8 76 modified nucleosides in municipal wastewater influent. Large-volume injection
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10 77 liquid chromatography and mass spectrometry has been used for the
11
12 78 determination of contaminants and metabolites in urine ³², as well as illicit drugs
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14 79 ^{6, 33}, and steroids ¹¹, in municipal wastewater influent. Large-volume injection
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17 80 was chosen to minimize sample preparation, and to maximize sensitivity needed
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19 81 for detection and quantification. It is known that hydrophilic analytes such as
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21 82 nucleosides are difficult to separate on C8 and C18 columns, and therefore
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23 83 alternative approaches have been developed ³⁴⁻³⁶. In order to retain and
24
25 84 separate nucleosides and modified nucleosides a polar-modified column
26
27 85 employed. Nine nucleosides were chosen to a proof of concept and demonstrate
28
29 86 method applicability. Twenty-four hour composite influent samples were
30
31 87 obtained and analyzed in order to demonstrate the feasibility of the developed
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33 88 method as a step to achieve the overall goal of investigating community health
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35 89 via wastewater analyses.
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40 90 **EXPERIMENTAL**

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43 91 **Chemicals and Materials.** Adenosine (A), 2'-deoxyadenosine (2dA), guanosine
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45 92 (G), 2'-deoxyguanosine (2dG), cytidine (C), 2'-deoxycytidine (2dC), uridine (U) ,
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47 93 N²-methylguanosine (N2-MG), 7-methylguanosine (7-MG), and HPLC grade
48
49 94 ammonium acetate (>99%) were purchased from Sigma-Aldrich (St. Louis, MO,
50
51 95 USA). Internal standards of [¹³C₅] adenosine (AC5), [¹³C₅] guanosine (GC5) and
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53 96 [¹³C₅] cytidine (CC5) were purchased from Toronto Research Chemicals (North
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3 97 York, ON, CAN). LC-MS optima™ grade methanol was purchased from Fisher
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5 98 Scientific (Waltham, MA, USA). Ultra pure water was obtained via a Milli Q
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8 99 advantage 10 (EMD Millipore, Billerica, MA, USA) equipped with a Q guard T2
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11 100 purification cartridge, quantum TEX polishing cartridge and millipak express 0.22
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13 101 µm filter.

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15 102 **Liquid Chromatography.** Large volume injection liquid chromatography (LVI-
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18 103 LC) was performed using a Shimadzu liquid Chromatograph (Kyoto, JPN)
19
20 104 consisting of two LC-20AD pumps coupled with a Phenomenex (Torrance, CA,
21
22 105 USA) model DG-4400 on-line degasser, CBM-20A control unit, SIL-20AC
23
24 106 autosampler equipped with a large volume injection kit (part # 228-45405-94)
25
26
27 107 along with a climate controlled sample tray, and a CTO-20A column heater
28
29 108 equipped with a two way switching valve (part # 228-45013-94). Mobile phase
30
31
32 109 (A) consisted of 10mM ammonium acetate at pH 5.3, and mobile phase (B) was
33
34 110 100% methanol. Sample volumes of 1 mL were injected onto a 10 x 4.0 mm 5
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36 111 µm RESTEK (Bellefonte, PA, USA) Ultra aqueous C18 guard column coupled
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39 112 with a 150 x 4.6 mm 5 µm analytical Ultra aqueous C18 column.

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41 113 **Mass Spectrometry.** Detection was performed using an AB SCIEX
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44 114 (Framingham, MA, USA) model 3200 Q TRAP equipped with a Turbo V™ ion
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46 115 source with an electrospray ionization probe operated in positive ion mode.
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48 116 Mass spectrometric parameters (Table 1) were controlled (along with the LC)
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51 117 using Analyst version 1.5.1. Ion source temperature was set to 375°C, ionization
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53 118 potential set to 5 kV, nebulization gas was set to 20 arbitrary units, desolvation
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3 119 gas was set to 45 arbitrary units, curtain gas was set to 25 arbitrary units and
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5 120 CAD gas set to high.
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8 121 **Wastewater flow.** Flow data was recorded using a Foxboro 9300 series flow
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10 122 meter (Houston, TX, USA), which is operated and maintained by the municipal
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12 123 wastewater treatment plant. All flow data was recorded by the minute for the
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14 124 duration of the study. Precipitation data was collected via daily climate reports
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16 125 published by the National Oceanic and Atmospheric Administration for Lawrence,
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18 126 Kansas (<http://www.nws.noaa.gov/climate/index.php?wfo=top>).
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22 127 **Sample Collection.** Composite (twenty-four hour) wastewater samples were
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24 128 collected using an in-line sampling device installed within the municipal
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26 129 wastewater treatment plant after influent screening. Sample collection was
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28 130 started on the morning of September 30, 2014 (Monday) and ended on October
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30 131 5, 2014 (Saturday). Samples were collected in a volume-dependent manner,
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32 132 with one sub-sample being collected after every 6.9×10^4 L of influent. 50 mL
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34 133 aliquots of composite samples were collected at 4°C in HDPE centrifuge tubes,
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36 134 transported directly to the lab, and immediately frozen at -20°C until analysis.
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39 135 **Standard and Sample Preparation.** Standard calibration and quality control
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41 136 solutions were prepared for analysis at concentrations ranging from 3×10^3 to
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43 137 2×10^5 ng/L using 10 mM ammonium acetate spiked with the internal standard
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45 138 mix to a final concentration of 5×10^3 ng/L. All samples were prepared by thawing
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47 139 to room temperature and then centrifugation at 4×10^3 g for 30 min using a VWR
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49 140 clinical 100 centrifuge (Radnor, PA, USA) in 15 mL tubes. 1425 μ L of sample
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51 141 was then transferred into a 1.5 mL autosampler vial, spiked with 75 μ L's of
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3 142 internal standard mix (AC5, GC5, CC5) to a final concentration of 5×10^3 ng/L
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6 143 and then vortexed.

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8 144 **Quantification and Identification.** All analytes were infused into the mass
9
10 145 spectrometer at a concentration ~ 1 mg/L made up in 10 mM ammonium acetate
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12 146 and methanol 1:1 (v/v) using an analytical syringe and integrated syringe pump
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15 147 to optimize mass spectrometric parameters. MRM transitions that were the most
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17 148 abundant were chosen for quantification (Table 1). Analyte responses were
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19 149 normalized to internal standard responses and all linear analyte calibration
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22 150 regression lines had a coefficient of determination (R^2) of >0.98 .

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24 151 **Quality control.** Blanks were made up of 10mM ammonium acetate, and quality
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27 152 control samples were calibration standard solutions. Blanks and quality control
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29 153 samples were analyzed following eight wastewater samples. Quality control
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31 154 samples made up 25% of the total sample sequence, which was completed in
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33
34 155 one day. Rejection criteria of the quality control samples were $\pm 15\%$ of the
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36 156 initial calculated concentrations at the beginning of the run sequence.

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38 157 **Standard Addition, Accuracy, Precision.** The use of a solvent-based
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40 158 calibration curve was determined appropriate for quantification of wastewater
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42 159 samples by using standard addition. Internal standards were deemed
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44 160 appropriate for quantification based on if there was a statistically significant
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46 161 difference (95%CI) between the solvent-based calibration derived value of the
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48 162 initial concentration in the sample and quantification using a calibration curve
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50 163 within the matrix using a 95% CI of the slope. The initial concentration of the
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52 164 sample was determined in the matrix by extrapolating the linear regression to the
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3 165 x intercept by using the additional concentrations. Wastewater samples were
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5 166 initially analyzed along with five different concentrations to extrapolate to the x
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8 167 intercept by calibration in the wastewater matrix. The accuracy of the method
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10 168 was determined by analyzing an over-spiked sample 8 times (Table 2).

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12 169 **Limits of detection, quantification and ion suppression.** Limits of detection
13
14 170 (LODs) were determined by spiking concentrations of internal standard into a
15
16 171 wastewater sample and analyzing four times to determine the lowest
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18 172 concentration that yielded a S/N (peak to peak) ≥ 3 . LOQs were defined as the
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20 173 concentration of the lowest standard with a S/N ≥ 10 . The percent of ion
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22 174 suppression was determined subtracting the ratio of the average area counts of
23
24 175 internal standard peaks (n=8) in 10 mM ammonium acetate, by the average area
25
26 176 counts of internal standard in wastewater (n=9) and multiplying by 100 (Equation
27
28 177 1). The method detection limit (MDL), and method quantification limit (MQL) was
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30 178 calculated by multiplying the LOD (ng/L) or LOQ (ng/L) by 100 and dividing the
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32 179 product by the percent of the sample recovery (100%) multiplied by the
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34 180 concentration factor of one.

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36 181 **Analyte Stability During Collection and Storage.** Analyte stability during
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38 182 sample collection was determined by collecting 500 mL of fresh wastewater
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40 183 influent and storing an aliquot at 4°C over 24 h, and another aliquot at 24°C
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42 184 (room temperature) for three hours. Samples at 4°C were collected initially and at
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44 185 12, and 24 h and directly frozen. Samples held at 24°C were taken initially and
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46 186 after three hours and then directly frozen. Samples were then analyzed in
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48 187 quadruplicate during one analytical sequence.
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3 188 **RESULTS AND DISCUSSION**
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5 189 **Large volume injection liquid chromatography.** The Liquid chromatograph
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8 190 flow path configuration was optimized to minimize gradient delay and to speed up
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10 191 column equilibration, effectively decreasing total analysis time (Figure 1). A two-
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12 192 way switching valve was installed inside the column heater and between the
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14 193 solvent mixer and autosampler in order to bypass the autosampler after the
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16 194 sample volume was injected onto the column. The design of the autosampler is
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18 195 such that the sample is at the head of the injection needle and directly injected
19
20 196 onto the column. After sample injection the initial mobile phase composition of
21
22 197 0% B flowed at 0.4 mL/min through the autosampler and column to waste. At 5
23
24 198 min after sample injection, mobile phase was diverted away from the
25
26 199 autosampler and directly through the column using the switching valve located
27
28 200 inside of the column heater. When using a 1 mL injection volume, a switch in the
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30 201 flow path away from the autosampler was done 5 min after injection so that the
31
32 202 injection loop containing the sample was flushed with two volumes (2 mL) to
33
34 203 insure no carryover. The %B mobile phase was then linearly increased to 80%
35
36 204 starting at 5 and ending at 12 min. Given the time when the front of the sample
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38 205 reaches the column (2.5 min) and then completely passed through the column
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40 206 (6.25 min using a flow rate of 0.4 mL/min, ~1.5 mL void volume and 1 mL sample
41
42 207 volume) the valve leading to the mass spectrometer ion source was set to divert
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44 208 flow after injection until 10.3 min in order to keep salts and unretained
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46 209 compounds from fouling the instrument. At 10.3 min the mass spectrometer
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48 210 divert valve diverted mobile phase from inject (waste) to load (into ESI probe).
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3 211 The mobile phase was then held at 80% B until 13.5 min and then linearly
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5 212 ramped to 0% B at 13.75 min. The mass spectrometer divert valve switched
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7 213 from load back to inject at 15 min. The flow rate was then increased to 0.6
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9 214 ml/min at 15.10 min and held until 19.50 min when the switching valve diverted
10
11 215 flow back to the autosampler, which was done in order to speed up column
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13 216 equilibration. The earliest analyte, cytidine (Figure 2), has an elution time of
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15 217 approximately 1 min after the divert valve switch, which is 6 min after sample
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17 218 injection.

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22 219 **Method validation.** The accuracy which was determined by calculating the
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24 220 percent difference between quantification using a solvent based calibration curve
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26 221 and calibrating in the matrix by standard addition were 100 +/- 20% except for
27
28 222 2dC (100 +/- 39%) (Table 2). Precision was also determined using the same set
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30 223 of samples and ranged from 6.05 %RSD in U to 18.59 %RSD in 7-MG (Table 2).
31
32 224 Calibration standards were linear from 100 to 2×10^5 ng/L. LOD was observed to
33
34 225 be 50 ng/L for AC5 and CC5, while GC5 was 100 ng/L. The LOQ was
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36 226 determined to be 100 ng/L for AC5 and CC5, and 1×10^3 ng/L for GC5. The
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38 227 LOQ of for analysis was 3×10^3 ng/L, which was the concentration of the lowest
39
40 228 standard. Linear ranges for all calibration curves used to analyze samples were
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42 229 from 3×10^3 to 2×10^5 ng/L for all analytes. The MDL and MQL are 5% higher than
43
44 230 the LOD and LOQ respectively due to the total recovery of the sample (100%)
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46 231 and a concentration factor of 0.95 (1425 μ L of sample with the addition of 75 μ L
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48 232 of IS). The percent of ion suppression was calculated to be 31.97 +/- 12.67
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50 233 (95%CI, Equation 1).
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3 234 Samples collected and stored at 4°C generally displayed a slight increase
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5 235 in concentration. 2dG concentrations were significantly (95%CI) higher within the
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8 236 24 h sample, while A displayed significantly (95%CI) lower concentrations for the
9
10 237 12 h and 24 h samples. The samples stored at 22°C resulted in a significant
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12 238 (95%CI) increase in 2dG between initial sample and after 3 h, while over the
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14
15 239 same time a decrease in A was observed.
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17 240 **Method demonstration.** Twenty-four hour composite wastewater samples were
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19 241 collected starting on the morning of September 30, 2014 (Monday) and ending
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21 242 on the morning of October 5, 2014 (Saturday). The calculated concentration
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23 243 (ng/L) of every nucleoside was above the LOD and LOQ except for N²MG and 7-
24
25 244 MG which were below the LOD (Figure 3). The volume of wastewater entering
26
27 245 the treatment plant was multiplied by the concentrations of all analytes to
28
29 246 calculate loads (mg) (Figure 4). The loads of nucleosides containing ribose,
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31 247 were significantly higher than nucleosides containing 2deoxyribose on every day
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33 248 sampled (Figure 4). The total uncertainty of calculated loads were calculated
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35 249 using the sum of the analytical uncertainty which is analyte dependent, the
36
37 250 uncertainty associated with the flow meter (0.25%) and estimated sampling
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39 251 uncertainty of 5% (Equation 2). This would indicate that there are more
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41 252 nucleosides originating from RNA than DNA, which makes sense given that the
42
43 253 turnover of DNA is slower than RNA²⁸. Wastewater influent flow displayed
44
45 254 diurnal variation on four out of the six days (Figure 4), with the other two days
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47 255 (October 2 to 3) appearing to have been influenced by rain events (Figure 4).
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49 256 During October 2 and October 3 there seems to be a significant amount of
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3 257 infiltration of stormwater into the system given that this is not a combined
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5 258 (stormwater and wastewater) sewer system. The loads of nucleosides are
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8 259 lowest on October 1 when there was no precipitation (Wednesday) and increase
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10 260 until sampling is complete on October 4 (Saturday).

11
12 261 In order to estimate the average mass of A, U, C and N²-MG (which are
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14 262 ribose containing nucleosides) excreted in a day, published concentrations of
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16 263 nucleosides in urine ²¹ were multiplied by average estimated daily volume of
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18 264 urine (1.1 L) ³⁷, then multiplied by a population estimate for Lawrence ³⁸, Kansas
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20 265 USA (93,742). This estimate ranged from 0.2 to 45, with a mean of 8 kg/day for
21
22 266 the sum of A, U, C and N²-MG ^{21, 39}. These values are similar to the loads of A,
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24 267 U, C found during the entirety of this study, which range from 1.2 to 16, with an
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26 268 average of 8.4 kg (Table 3). The highest loads (25 kg, Table 3) were calculated
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28 269 on October 4 (Saturday) and could potentially be an indication of commuters
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30 270 staying within the city on their workday off, thus the total amount of nucleosides
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32 271 may represent possible population markers. Although the usefulness of
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34 272 nucleosides (especially nucleosides 2deoxyribose nucleosides) as population
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36 273 markers are plausible, an investigation into their full potential is beyond the
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38 274 scope of this study.

39
40 275 Nucleosides are components of all living organisms and some non-living
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42 276 organisms (i.e. viruses), thus the total loads of nucleosides in wastewater could
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44 277 be influenced by unused food ⁴⁰ disposal, and biofilm ⁴¹ (i.e. bacteria). Direct
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46 278 disposal is a possible interference for the interpretation of loads of illicit drugs,
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48 279 pharmaceuticals, as well as other potential analytes that are not directly excreted
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3 280 as a metabolite originating from human use. Nucleosides in this study were
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5 281 observed to be relatively stable in wastewater at 4°C and room temperature
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8 282 without the influence of biofilm. Given these factors future investigations related
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10 283 to comparing the loads of nucleosides between communities may have to take
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12 284 nucleoside transformation and or addition via biofilm into consideration.

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14
15 285 **Conclusions.** The goal of this study was to develop an analytical method to
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17 286 quantify nucleosides in municipal wastewater. The method developed in this
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19 287 study was applied to analyze 24 h composite samples, which were used to verify
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21 288 the presence and to quantify nucleosides in wastewater and to investigate their
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23 289 stability. This method offers an insight into the total amounts of selected
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25 290 nucleosides found in municipal wastewater influent. Thus, the method developed
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27 291 here is a complimentary tool for the future investigation of community health. For
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29 292 further analyses and interpretation it may be beneficial to employ non-targeted
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31 293 detection to obtain a profile of excreted nucleosides which could be used to
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33 294 detect nucleosides for which standards would be expensive or impossible to
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35 295 obtain. Ultimately analytical methods such as the one developed in this study
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37 296 need to provide data of any proposed health marker in wastewater, which will
38
39 297 lead to investigations and correlations of these markers to known health
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41 298 stressors and human health endpoints.

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43
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 306 represent the official views of the National Institutes of Health.

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308

309 **Table 1.** Mass spectrometric parameters for MRM scans (collision cell exit

310 potential of 2V for all analytes) Transitions used for quantification are labeled with

311 an *.

Identifier	Precursor ion (m/z)	Product ion (m/z)	Dwell time (msec)	Declustering potential (V)	Collision energy (V)
A-1*	268.3	119.2	125	35	50
A-2	268.3	136.1	125	35	30
2dA-1*	252.2	136.1	125	35	35
2dA-2	252.2	119.2	125	35	75
AC5-1*	273.5	136.1	125	35	27
AC5-2	273.5	119.2	125	35	61
G-1*	284.0	135.2	125	75	50
G-2	284.0	152.3	125	35	30
2dG-1*	268.4	152.3	125	35	20
2dG-2	268.4	135.2	125	35	60
GC5-1*	289.4	152.5	125	35	35
GC5-2	289.4	135.5	125	35	45
C-1*	244.2	112.0	125	15	35
C-2	244.2	95.2	125	15	65
2dC-1*	228.5	95.2	125	15	65
2dC-2	228.5	112.3	125	15	35
CC5-1*	248.9	112.0	125	15	35
CC5-2	248.9	95.2	125	15	65
U-1*	245.1	113.2	125	15	35
U-2	245.1	70.4	125	20	50
N ² -MG-1*	298.3	149.2	125	75	50
N ² -MG-2	298.3	110.4	125	75	50
7-MG-1*	299.4	149.2	125	75	50
7-MG-2	299.4	124.3	125	75	50

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315 **Table 2.** Accuracy as determined as the percent difference between solvent
 316 based calibration curve quantification and standard addition, and precision
 317 determined from multiple analysis of one wastewater sample.

Identifier	Accuracy (mean % difference n=8)	Precision (%RSD n=8)
A	108	6.7
2dA	99	8.1
G	92	15
2dG	116	10
C	103	6.1
2dC	139	11
U	98	6.1
N2-MG	102	12
7-MG	115	19

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320 **Table 3.** Daily loads of all nucleosides collected during study period.

Date	A	2dA	G	2dG	C	2dC	U	Sum (kg)
29-Sep	4.1	0.37	5.7	0.69	2.4	0.23	2.8	16
30-Sep	4.1	0.37	5.5	0.65	2.1	0.34	2.3	15
1-Oct	0.60	0.08	0.54	0.09	0.22	0	0.44	2
2-Oct	2.6	0.27	2	0.36	1.3	0.15	1.7	8
3-Oct	4.7	0.30	4.7	0.46	2.2	0.18	3.0	15
4-Oct	9.7	0.40	7.4	0.79	3	0.36	3.5	25

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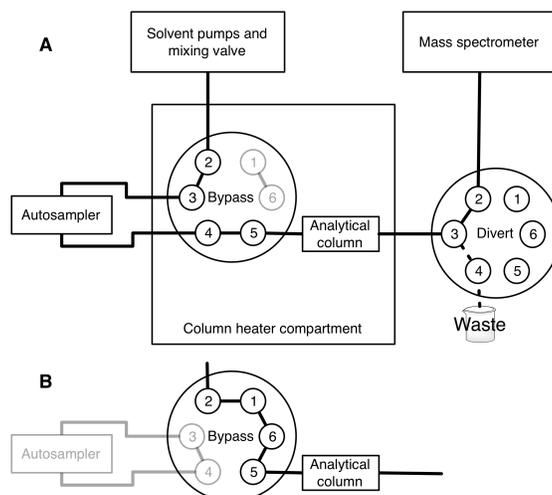
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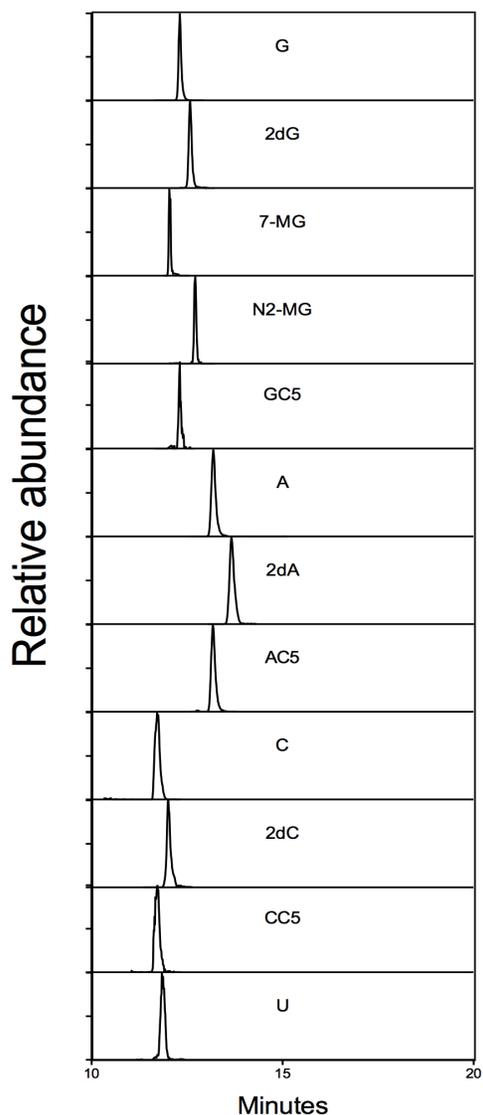
Figure 1. Schematic diagram showing the flow path while the sample is being loaded onto the column from 0-5 min (A), and when the flow path is routed around the autosampler 5-19.5 min (B). Faded areas of the “bypass” valve indicate ports of the valve not being used at each respective time. The “divert” valve was set to send flow into the mass spectrometer ionization source from 10.3-15 min.

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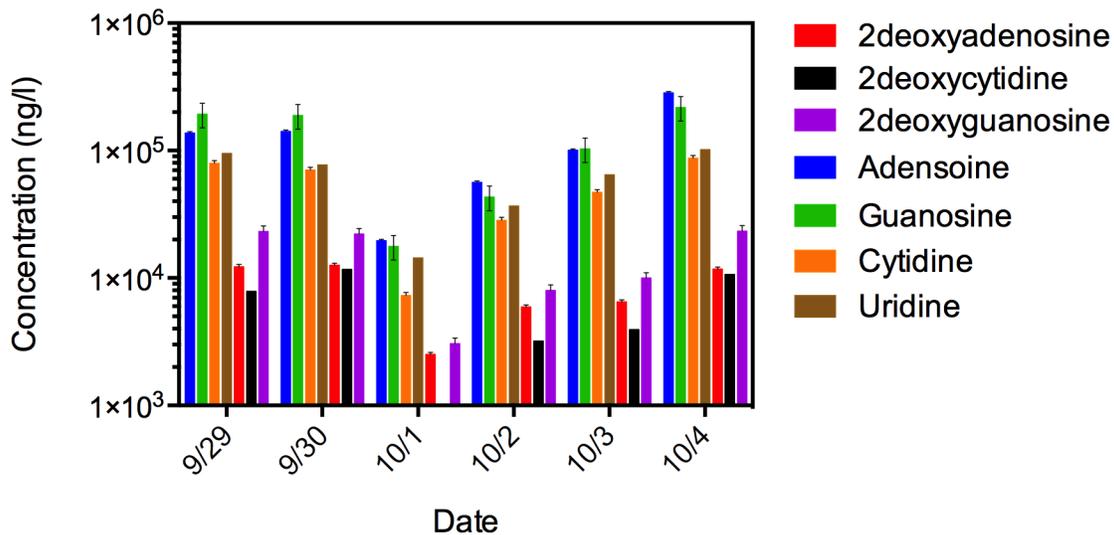


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340 **Figure 2.** Example of typical wastewater sample chromatogram showing relative
341 abundance (0-100%) of guanosine (G), 2deoxyguanosine (2dG),
342 7methylguanosine (7-MG), N2methylguanosine (N²-MG), guanosine C5 (GC5),
343 adenosine (A), 2deoxyadenosine (2dA), adenosine C5 (AC5), cytidine (C),
344 2deoxycytidine (2dC), cytidine C5 (CC5), uridine (U).

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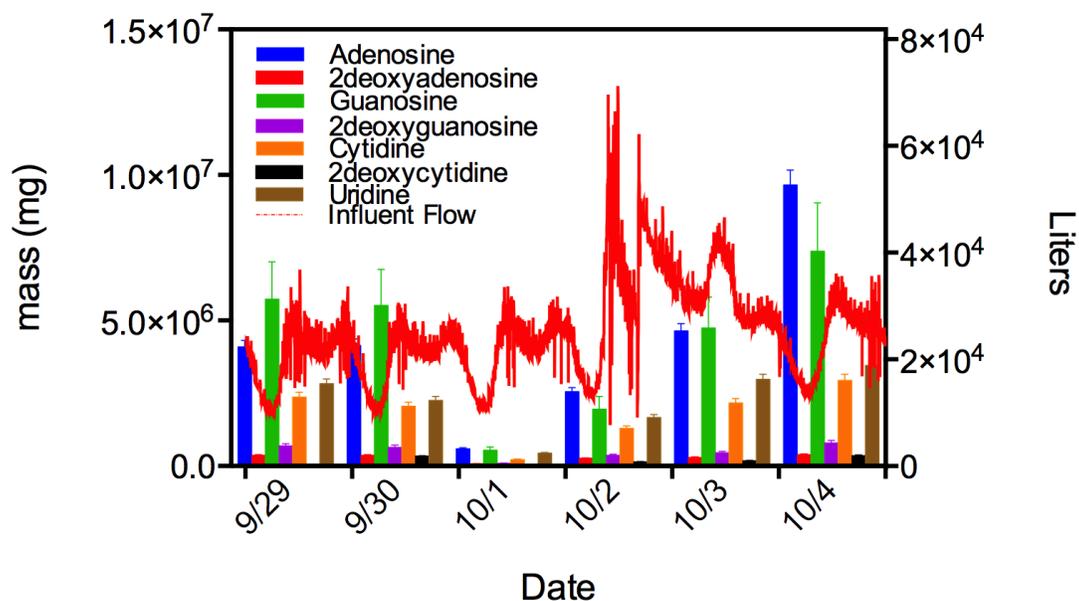
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Figure 3. Bar graph showing the concentration of nucleosides in wastewater influent samples from September 29 to October 4, 2014

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358 **Figure 4.** Loads (mg) of nucleosides and influent flow (L) from September 29
359 through October 4, 2014, error bars represent total uncertainty which is the sum
360 of analytical, flow meter, and sampling uncertainty (%RSD). Total daily rainfall
361 was 0.25, 22, 43, and 1 mm starting on September 30th and ending on October
362 3rd.

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365 **Equation 1.** Calculation of ion suppression (%) in wastewater compared to 10

366 mM ammonium acetate buffer.

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$$\text{Ion suppression \%} = 1 - \frac{\text{IS peak area in wastewater}}{\text{IS peak area in buffer}} \times 100$$

368

369 **Equation 2.** Calculation of total uncertainty from analysis, sampling, and flow. ¹²

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$$\text{Total uncertainty} = \sqrt{(\text{RSD analytical})^2 + (\text{RSD sampling})^2 + (\text{RSD flow})^2}$$

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