# USGSHydroOpt : Tools for Optical Analysis of Water

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## Contents

## 1 Introduction to USGSHydroOpt

The USGSHydroOpt package was created to streamline the process of creating optical summary variables and excitation-emission (EEMs) plots for absorbance and fluorescence data collected from water samples. Examples of optical summary variables that can be produced with this package include various absorbance

peaks, spectral slopes, fluorescence and humic index variables, and others. The motivation for producing this package is that optical summary variables can be powerful explanatory variables for biological and chemical contamination in water. Thus, the optical summary variables produced using USSHydroOpt are extremely useful for statistical modeling.

The functions in this package were designed to operate on dataframes with a standard data structures. This package is not amenable to dataframes or arrays that do not fit the prescribed formats. The example dataframes and array in this package illustrate exactly how data structures should be formatted, and the examples illustrate how the functions operate on the data structures. Depicted below is an example of an EEMs plot produced with this package. The user is encouraged to step through each piece of R code as it is introduced in this document. In many instances a sample of the function output is provided via plots and tables, but not in all cases.
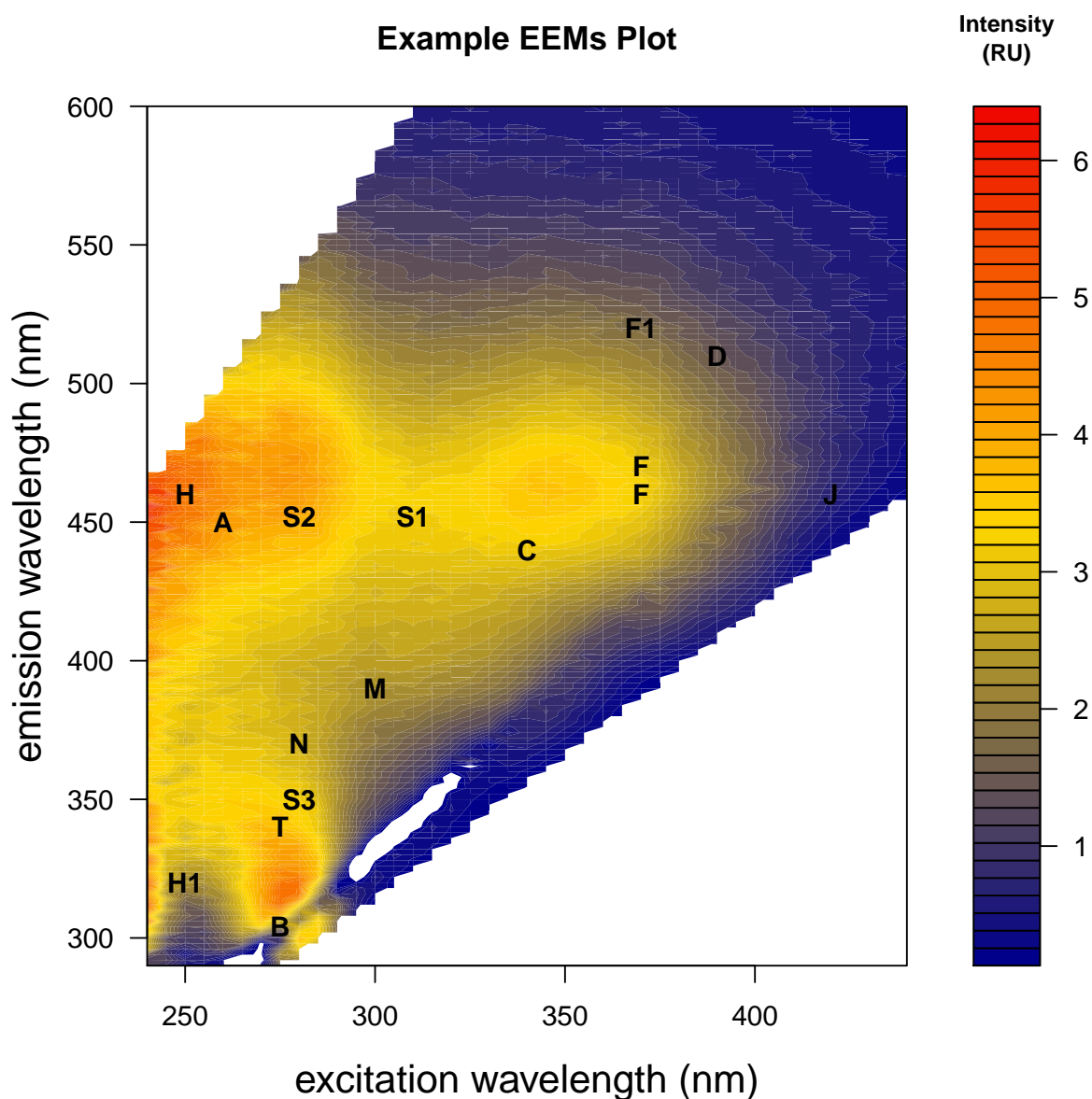
# 2 Required Data Formats

The functions contained in USGSHydroOpt operate on dataframes with defined structures. Users interested in using USGSHydroOpt should format dataframes according to the structures defined in this section.

## 2.1 Absorbance Data

Absorbance data used by functions in USGSHydroOpt should be formatted such that each sample occupies a column, and one column contains the wavelength (nm) for which the absorbance measurement was measured (*See example Table 2.1 below*). The example absorbance dataframe included in this package is called **dfabs**. The column with the wavelengths in Table 2.1 does not need to be called "wavelengths," as it is named in the example dataframe below. Since this package was developed primarily for USGS activities, the default for naming samples is "gr" then the sample number. This convention was started by the USGS California Water Sciences Center (CA WSC) and the USGS Wisconsin Water Science Center (WI WSC) follows the same naming convention to ensure standardization.

```
     gr13307     gr13351     gr13353     gr13357 wavelengths
1  0.0001296 -0.0003505 -0.0003480 -0.0002695         750
2 -0.0002367  0.0000305 -0.0000915 -0.0001407         749
3 -0.0001582 -0.0004900 -0.0006325 -0.0001534         748
4 -0.0004642 -0.0000105 -0.0000932 -0.0000245         747
5 -0.0002551 -0.0000653  0.0000841 -0.0001615         746
6 -0.0001842 -0.0002135  0.0002082  0.0001429         745
```

## 2.2 Fluorescence Data

Fluorescence data used by functions in USGSHydroOpt should also be formatted such that each sample occupies a column, and one column contains the excitation emission wavelength pairs (nm) for which the fluorescence measurment was measured (*See example Table 2.2 below*). The example fluorescence dataframe included in this package is called **dfFluor**. The column with the excitation emission pairs in Table 2.2 does not need to be called "Wavelength.Pairs," as it is in the example dataframe below. Again since this package was developed for USGS activities, the default sample naming convention is "gr" followed by the sample number.

```
  Wavelength.Pairs gr13307 gr13308  gr13351  gr13352
1          240/290 0.09457 0.05358 -0.07268 -0.19368
2          240/292 0.07386 0.09194  0.45433  0.39722
3          240/294 0.08116 0.09930  0.49135  0.37943
4          240/296 0.12013 0.09138 -0.12248  0.09239
5          240/298 0.15264 0.15428  0.54694  0.14980
6          240/300 0.16467 0.13010  0.43319  0.33064
```

## 2.3 Spectral Slopes Data

Information on the upper and lower wavelength (nm) for which a spectral slope should be calculated needs to be stored in a dataframe if USGSHydroOpt is used. The example spectral slopes dataframe included in

this package is called **dfsags**. The dataframe should contain exactly three columns. The first column should contain the upper wavelength, the second column should contain the lower wavelength, and the third column should contain the name of the spectral slope being calculated (*See example Table 2.3 below*). The columns in Table 2.3 need to be in this exact order, although the names of the columns may be different. The data types for each column are integer, integer, and character, respectively. More spectral slopes can be added to the table than specified in the example dataframe below.

```
  wvlngth1 wvlngth2       Name
1      275      290 Sag275_295
2      290      350 Sag290_350
3      350      400 Sag350_400
4      412      676 Sag412_676
```

## 2.4   Optical Summary Data

This is the dataframe that contains many of the summary optical variables that can be produced using functions in USGSHydroOpt (*See example Table 2.4a below*). The example summary optical dataframe included in this package is called **dfsummary**. The functions in USGSHydroOpt calculate summary optical variables and add to a dataframe formatted according to Table 2.4a below. The example dataframe below is how the WI WSC stores optical summary variables. Note that this dataframe can contain other columns with metadata, for example, the sample data and time, the sample ID, or whether or not the sample went through QA/QC.

```
  GRnumber         B      T      A       J FI_2005    A254
1  gr13307  0.050997 0.1826 0.4553 0.03705   1.639 0.05228
2  gr13351 -0.245602 0.4085 3.0783 0.19110   1.456 0.43531
3  gr13353 -0.175220 0.7794 6.8624 0.56309   1.523 0.68274
4  gr13357  0.111561 0.2691 0.9451 0.06969   1.572 0.08698
5  gr13360 -0.001569 0.4593 2.8254 0.37231   1.563 0.28605
6  gr13363  0.052137 0.4892 1.9518 0.19098   1.583 0.19283
```

However, also note that summary optical variable names in the dataframe must be identical to those specified in Table 2.4b below.

```
 [1] "OB1"        "OB2"        "OB3"        "S1.50"
 [5] "S2.50"      "S3.50"      "S1.25"      "S2.25"
 [9] "S3.25"      "Mrange.25"  "Mrange.50"  "B"
[13] "T"          "M"          "A"          "C"
[17] "N"          "D"          "F"          "J"
[21] "S1"         "S2"         "S3"         "H1"
[25] "H2"         "F1"         "F2"         "W"
[29] "LT1"        "LT2"        "LT3"        "LA"
[33] "HIX_2002"   "FI_2005"    "FI_2001"    "FreshI"
[37] "A254"       "A275"       "A280"       "A290"
[41] "A295"       "A350"       "A370"       "A400"
[45] "A412"       "A440"       "A488"       "A510"
[49] "A532"       "A555"       "A650"       "A676"
[53] "A715"       "Sag275_290" "Sag290_350" "Sag350_400"
[57] "Sag412_676" "Aresids"
```

## 2.5 Excitation-Emission (EEMs) Peak Data

The EEMs peak data contains three columns listing the name of a characterized EEM peak along with the corresponding wavelengths (nm) (*See example Table 2.5 below*). The example EEMs peak dataframe included in this package is called **ex_ems**. The first column, in Table 2.5, "Peak," contains the name of the characterized EEM peak. The next two columns in Table 2.5 contain the excitation and emission wavelengths (nm) at which a given peak occurs. The column names must be identical to those displayed in the example below, although the order of the columns can be different.

```
    Peak ExCA EmCA
1      B  275  304
2      T  275  340
3      M  300  390
4      A  260  450
5      C  340  440
6      N  280  370
7      D  390  510
8      F  370  460
9      J  420  460
10    S1  310  452
11    S2  280  452
12    S3  280  350
13     H  250  460
14    H1  250  320
15     F  370  470
16    F1  370  520
```

## 2.6 Optical Ratio and Signals Data

This dataframe contains one column called "ratioSignals" that contains all of the summary optical variables currently identified by the WI WSC (*See example Table 2.6 below*). An example of this dataframe included in this package is called **ratioSignals**. Note that these are the same variables as those listed in Table 2.4b. The first column must contain the various "ratioSignals" that the user desires, although the column name need not be "ratioSignals"

```
    ratioSignals keep
1            OB1   NA
2            OB2   NA
3            OB3   NA
4          S1.50   NA
5          S2.50   NA
6          S3.50   NA
7          S1.25   NA
8          S2.25   NA
9          S3.25   NA
10     Mrange.25   NA
11     Mrange.50   NA
12             B   NA
```

```
13          T   NA
14          M   NA
15          A   NA
16          C   NA
17          N   NA
18          D   NA
19          F   NA
20          J   NA
21         S1   NA
22         S2   NA
23         S3   NA
24         H1   NA
25         H2   NA
26         F1   NA
27         F2   NA
28          W   NA
29        LT1   NA
30        LT2   NA
31        LT3   NA
32         LA   NA
33   HIX_2002   NA
34    FI_2005   NA
35    FI_2001   NA
36     FreshI   NA
37       A254    1
38       A275    1
39       A280    1
40       A290    1
41       A295    1
42       A350    1
43       A370    1
44       A400    1
45       A412   NA
46       A440   NA
47       A488   NA
48       A510   NA
49       A532   NA
50       A555   NA
51       A650   NA
52       A676   NA
53       A715   NA
54  Sag275_290    1
55  Sag290_350    1
56  Sag350_400    1
57  Sag412_676    1
58     Aresids   NA
```

## 2.7 Optical Signals Data

This dataframe is similar to "ratioSignals" except it provides more metadata about peaks characterized for EEMs plots. The example signals dataframe included in this package is called **signals**. The dataframe should contain six columns (*See example Table 2.7 below*). The first column in Table 2.7, "Peak," contains the name of the characterized EEM peak. The next two columns, "Ex1" and "Ex2," contain the excitation wavelength range (nm) for a given peak. "Ex1" is the lower wavelength and "Ex2" is the upper wavelength for the excitation wavelength range (nm) for a given peak. Similarily, "Em1" and "Em2" contain the emission wavelength range (nm) for a given peak. The final column, "Source," lists the source that characterized the peak. The last column, "Source," is not required. The user must exactly replicate the column names in Table 2.7 in order for the code in USGSHydroOpt to run.

| | Peak | Ex1 | Ex2 | Em1 | Em2 | Source |
|---|---|---|---|---|---|---|
| 1 | OB1 | 360 | NA | 410 | 598 | Hartel Turner |
| 2 | OB2 | 360 | NA | 436 | 436 | Hartel Turner |
| 3 | OB3 | 365 | NA | 400 | 550 | Hagedorn Turner |
| 4 | S1.50 | 310 | NA | 402 | 502 | Sniffer |
| 5 | S2.50 | 280 | NA | 402 | 502 | Sniffer |
| 6 | S3.50 | 280 | NA | 310 | 390 | Sniffer |
| 7 | S1.25 | 310 | NA | 427 | 477 | Sniffer |
| 8 | S2.25 | 280 | NA | 427 | 477 | Sniffer |
| 9 | S3.25 | 280 | NA | 330 | 370 | Sniffer |
| 10 | Mrange.25 | 300 | NA | 365 | 415 | test |
| 11 | Mrange.50 | 300 | NA | 340 | 440 | test |
| 12 | B | 275 | NA | 304 | NA | CA |
| 13 | T | 275 | NA | 340 | NA | CA |
| 14 | M | 300 | NA | 390 | NA | CA |
| 15 | A | 260 | NA | 450 | NA | CA |
| 16 | C | 340 | NA | 440 | NA | CA |
| 17 | N | 280 | NA | 370 | NA | CA |
| 18 | D | 390 | NA | 510 | NA | CA |
| 19 | F | 370 | NA | 460 | NA | CA |
| 20 | J | 420 | NA | 460 | NA | CA |
| 21 | S1 | 310 | NA | 452 | NA | CA |
| 22 | S2 | 280 | NA | 452 | NA | CA |
| 23 | S3 | 280 | NA | 350 | NA | CA |
| 24 | H1 | 250 | NA | 460 | NA | Ohno2002 |
| 25 | H2 | 250 | NA | 320 | NA | Ohno2002 |
| 26 | F1 | 370 | NA | 470 | NA | Cory and McKnight, 2005 |
| 27 | F2 | 370 | NA | 520 | NA | Cory and McKnight, 2005 |
| 28 | W | 255 | 290 | 302 | 350 | |
| 29 | LT1 | 250 | NA | 340 | NA | |
| 30 | LT2 | 260 | NA | 340 | NA | |
| 31 | LT3 | 240 | NA | 340 | NA | |
| 32 | LA | 240 | NA | 440 | NA | |

## 2.8  3-Dimensional Excitation-Emission Array

A 3-D array with fluorescence data is used by many of the functions in USGSHydroOpt. The first dimension contains the excitation wavelengths (nm) as data type character at which a given fluorescence measurement was made. The second dimension contains the emission wavelengths (nm) as data type character at which a given fluorescence measurement was made. The third dimension contains the sample numbers as data type character for a given observation. The user must ensure that the third dimension of the array are sample numbers. Again, in this example the default "gr" followed by the sample number is used as a naming convention for samples. The example 3-D array with EEM data included in this package is called **a**.

To view the headers for each dimension using the following commands in R consider the example 3-D EEM array included with the USGSHydroOpt Package:

```
# this command shows the excitation wavelengths (nm)
colnames(a)

  [1] "290" "292" "294" "296" "298" "300" "302" "304" "306"
 [10] "308" "310" "312" "314" "316" "318" "320" "322" "324"
 [19] "326" "328" "330" "332" "334" "336" "338" "340" "342"
 [28] "344" "346" "348" "350" "352" "354" "356" "358" "360"
 [37] "362" "364" "366" "368" "370" "372" "374" "376" "378"
 [46] "380" "382" "384" "386" "388" "390" "392" "394" "396"
 [55] "398" "400" "402" "404" "406" "408" "410" "412" "414"
 [64] "416" "418" "420" "422" "424" "426" "428" "430" "432"
 [73] "434" "436" "438" "440" "442" "444" "446" "448" "450"
 [82] "452" "454" "456" "458" "460" "462" "464" "466" "468"
 [91] "470" "472" "474" "476" "478" "480" "482" "484" "486"
[100] "488" "490" "492" "494" "496" "498" "500" "502" "504"
[109] "506" "508" "510" "512" "514" "516" "518" "520" "522"
[118] "524" "526" "528" "530" "532" "534" "536" "538" "540"
[127] "542" "544" "546" "548" "550" "552" "554" "556" "558"
[136] "560" "562" "564" "566" "568" "570" "572" "574" "576"
[145] "578" "580" "582" "584" "586" "588" "590" "592" "594"
[154] "596" "598" "600"

# this command shows the emission wavelengths (nm)
rownames(a)

 [1] "240" "245" "250" "255" "260" "265" "270" "275" "280"
[10] "285" "290" "295" "300" "305" "310" "315" "320" "325"
[19] "330" "335" "340" "345" "350" "355" "360" "365" "370"
[28] "375" "380" "385" "390" "395" "400" "405" "410" "415"
[37] "420" "425" "430" "435" "440"

# this command shows the emission wavelengths (nm), only the
# first 20 shown for simplicity
names(a[1, 1, ])[1:20]

 [1] "gr13307" "gr13308" "gr13351" "gr13352" "gr13353"
 [6] "gr13354" "gr13357" "gr13358" "gr13360" "gr13361"
[11] "gr13362" "gr13363" "gr13364" "gr13365" "gr13374"
```

```
[16] "gr13375" "gr13433" "gr13434" "gr13435" "gr13439"
```

The user should be aware of **two important caveats**: (1) There should rarely be emission wavelengths below the excitation wavelength for a given fluorescence reading. Where this occurs an NA will be found in the 3-D EEMs array. (2) Intensities at an emission wavelength that is two times the excitation wavelength will be influence by second order Rayleigh scatter.

### 2.9 Creating a 3-Dimensional Excitation-Emission Array

In Section 2.8 the format of 3-D arrays of fluorescence data that can be used with USGSHydroOpt was discussed. USGSHydroOpt can also be used to produce such arrays given the appropriate input fluorescence dataframe. Below is an example of how USGSHydroOpt is used to accomplish this task:

```
# set an arbitrary data frame (df) as dfFluor (the example
# fluorescence dataframe in USGSHydroOpt)
df <- dfFluor

# define the column in dfFluor
ExEm <- "Wavelength.Pairs"

# run the VectorizedTo3DArray function from USGSHydroOpt that
# creates a 3-D EEMs array given a vectorized fluorescence
# dataframe
aTest <- VectorizedTo3DArray(df, ExEm)
```

In the example above, dfFluor is formatted according to the fluorescence dataframe discussed in Section 2.2.

## 3 Creating Summary Optical Variables

Most of the functions included in USGSHydroOpt are for creating variables that summarize optical data. Such variables can then be used in statistical modeling efforts aimed at describing observed phenomena in aquatic ecosystems. This sections steps through the six functions that USGSHydroOpt offers for creating such summary optical variables.

### 3.1 Creating Absorbance Coefficients

The function getAbs opperates on a dataframe formatted according to Section 2.1. Specifically, it picks out absorbance coefficients for a defined set of wavelengths (nm) and adds those coefficients to an optical summary data frame formatted according to Section 2.4. Shown below is an example of how the function can be used.

```
# assign a variable dataAbs to the absorbance dataframe
# included in USGSHydroOpt
dataAbs <- dfabs

# define the column with the wavelengths in dataAbs
```

```
waveCol <- "wavelengths"

# define the wavelengths for which absorbance coefficients
# should be defined
wavs <- c(430, 530, 630, 730)

# define which columns contain the absorbance data, by
# default per WI WSC and CA WSC samples start with 'gr'
colSubsetString <- "gr"

# assign a variable dataSummary to the dfsummary dataframe
# included in USGSHydroOpt
dataSummary <- dfsummary

# assign a variable grnum to the column in dataSummary with
# sample numbers
grnum <- "GRnumber"

# use getAbs to produce absorbance coefficients
testAbs <- getAbs(dataAbs, waveCol, wavs, colSubsetString, dataSummary,
    grnum)

# note that the absorbance coefficients as defined by wavs
# have been added to dataSummary
colnames(testAbs)[69:72]

[1] "A430" "A530" "A630" "A730"
```

## 3.2 Spectral Slopes by Linear Regression

The function getExpResid computes spectral slopes by a first order decay function determined by linear regression (Helms et al. 2008).The residual at a specified wavelength (nm) is then calculated based on the spectral slope. The residual at a given wavelength and spectral slope is then added to a summary dataframe formatted according to Section 2.4. Displayed below is an example of how the function can be used. The function produces a plot with the absorbance data for each sample. On the plot the red indicates the spectral slope model, the blue is the absorbance data in rangeGap, and the black is the data in rangeReg but not in rangeGap. The dataframe dataAbs is formatted according to Section 2.1, but is shortened. If the user calls "pdf(genericName.pdf)," *See* ?pdf, then runs the function plots will be printed to the pdf in the working directory.

```
# absorbance wavelength (nm) for which residual is calculated
wavelength <- 267

# the absorbance wavelength range (nm) to be considered as a
# numeric string
rangeReg <- c(240, 340)

# the absorbance wavelength range (nm) to be considered as a
```

```r
# numeric string
rangeGap <- c(255, 300)

# assign a variable dataAbs to a shortened version of the
# absorbance dataframe included in USGSHydroOpt
dataAbs <- dfabs

# define the column with the wavelengths in dataAbs
waveCol <- "wavelengths"

# assign a variable grnum to the column in dataSummary with
# sample numbers
colSubsetString <- "gr"

# assign a variable dataSummary to the dfsummary dataframe
# included in USGSHydroOpt, column 68 or 'Aresids' is removed
# because we are computing this summary optical variable with
# this function and then adding it to dataSummary
dataSummary <- dfsummary[, -c(68)]

# assign a variable grnum to the column in dataSummary with
# sample numbers
grnum <- "GRnumber"

# use getExpResid to calculate the residual at a given
# wavelength given a spectral slope calculated per Helms et
# al. 2008.
testdfOpt <- getExpResid(wavelength, rangeReg, rangeGap, dataAbs,
    waveCol, colSubsetString, dataSummary, grnum)

# notice that the variable 'Aresids' has been added to
# dataSummary
colnames(testdfOpt)
```

## 3.3   Humification and Fluorescence Indices

Four humification and fluorescence index summary variables can be computed using the function getIndexes. The the humification index summary variable called "HIX_2002" is calculated according to Ohno (2002). The first fluorescence index summary variable "FI_2005" is computed according to Cory and McKnight (2005). The second fluorescence index summary variable, "FI_2001" is computed according to McKnight et al. (2001). The final summary variable produced by this function is the freshness index, "FreshI," is computed according to Parlanti et al. (2000). Each of these four summary variables are computed and added to the optical summary dataframe, formatted according to Section 2.4. In the example below, getIndexes is used to compute the four summary variables, which are subsequently added to an optical summary dataframe. A 3-dimensional excitation-emission dataframe with fluorescence data formatted according to Sections 2.8-2.9 is used in computing these summary variables.

```
# set a variable a as the example 3-D excitation emission
# array included with the package
a <- a

# assign a variable dataSummary to the dfsummary dataframe
# included in USGSHydroOpt
dataSummary <- dfsummary

# remove those columns with the fluorescence and humic
# indices that we are computing here
dataSummary <- dataSummary[, -c(43:46)]

# assign a variable grnum to the column in dataSummary with
# sample numbers
grnum <- "GRnumber"

# use getIndexes to compute the four humification and
# fluorescence indices
testIndexes <- getIndexes(a, dataSummary, grnum)

# note that the four indices have been added to dataSummary
colnames(testIndexes)[65:68]

[1] "HIX_2002" "FI_2005"  "FI_2001"  "FreshI"
```

### 3.4  Optical Ratios

The function getRatios uses absorbance peaks and spectral slopes in an existing summary optical data frame (e.g., dfsummary) and creates ratios between the different peaks and ratios. These ratios can be useful as predictor variables in statistical modeling efforts. In the example, below 65 different ratios are computed using an optical summary dataframe formatted per Section 2.4, and the ratioSignals example dataframe formatted per Section 2.6. The names of the calculated ratios added to the optical summary dataframe begin with "r" to signify "ratio" followed by the absorbance peak and/or spectral slope used to calculate the ratio. For example, the ratio of the absorbance peak at 254nm (A254) and the spectral slope between 350 and 400nm (Sag350_400) is called "rA254_Sag350_400."

```
# assign a variable dataSummary to the dfsummary dataframe
# included in USGSHydroOpt
dataSummary <- dfsummary

# note the number of variables in dataSummary
length(colnames(dataSummary))

[1] 68

# pick out the absorbance peaks and spectral slopes to be
# used for calculating ratios these correspond to those with
# a 1 in the 'keep' column.
sigs <- ratioSignals[which(ratioSignals[2] > 0), 1]
```

```
# assign a variable grnum to the column in dataSummary with
# sample numbers
grnum <- "GRnumber"

# use getRatios to calculate 65 different ratios of
# absorbance peaks and spectral slopes
test <- getRatios(dataSummary, sigs, grnum)

# notice that 65 ratios have been added to dataSummary
length(colnames(test))

[1] 134

# example of some ratios added
colnames(test)[69:75]

[1] "rA254_A275" "rA254_A280" "rA254_A290" "rA254_A295"
[5] "rA254_A350" "rA254_A370" "rA254_A400"
```

## 3.5 EEM Peaks Computed by Average Wavelength

Various excitation-emission (EEMs) peaks can be used to identify the presence of different chemical constituents in water. An EEMs peaks occur at specific excitation and emission wavelengths (nm), or within specific excitation and emission wavelength ranges. It has become standard practice to extract these peaks from fluorescence data, and identify them on EEMs plots. When given a fluorescence dataframe formatted according to section 2.2, the function getMeanFl computes the various peaks based on a dataframe of signals formatted according to section 2.7. Many of these peaks can occur within an excitation wavelength (nm) range, and also an emission wavelength (nm) range. For peaks where this is true, the function computes the average of the excitation and/or emission wavelength (nm) range. The resultant mean excitation and/or emission wavelength (nm) is then used to compute the EEMs peak from the fluorescence data. An example of how this function can be used is illustrated below.

```
# set a variable a as the example 3-D excitation emission
# array included with the package
a <- a

# set a variable signals as the example signals dataframe
signals <- signals

# note the length of the EEMs peaks to be computed
length(rownames(signals))

[1] 32

# identify the name of the column with the EEMs Peak names
Peak <- "Peak"

# identify column with the lower excitation wavelength in the
# excitation wavelength range
```

```r
Ex1 <- "Ex1"

# identify column with the upper excitation wavelength in the
# excitation wavelength range
Ex2 <- "Ex2"

# identify column with the lower emission wavelength in the
# emission wavelength range
Em1 <- "Em1"

# identify column with the upper emission wavelength in the
# emission wavelength range
Em2 <- "Em2"

# assign a variable dataSummary to the dfsummary dataframe
# included in USGSHydroOpt
dataSummary <- dfsummary

# remove the variables in dataSummary that are going to be
# computed with testMeanFl
rm <- which(colnames(dataSummary) %in% signals$Peak)
dataSummary <- dataSummary[, -c(rm)]

# note the number of variables in dataSummary
length(colnames(dataSummary))

[1] 36

# assign a variable grnum to the column in dataSummary with
# sample numbers
grnum <- "GRnumber"

# use getMeanFl to compute the different EEMs signals and add
# them to the optical summary data frame
testMeanFl <- getMeanFl(a, signals, Peak, Ex1, Ex2, Em1, Em2,
    dataSummary, grnum)

# notice that signals have been added to dataSummary
length(colnames(testMeanFl))

[1] 68

length(rownames(signals)) + length(colnames(dataSummary))

[1] 68
```

### 3.6 Log transformation of Summary Optical Variables

Often times it is useful to log 10 tranform summary optical variables to help stabilize variance. This is especially true when using summary optical variables as predictor variables for response variables such as bacteria and viruses. The function getLog10 log10 transforms user defined optical summary variables to a log10 scale.

```
# select those variables from ratioSignals which should be
# log 10 transformmed in dataSummary
signals <- ratioSignals[which(ratioSignals[2] > 0), 1]

# note the number of variables to be log 10 transformed
length(signals)

[1] 12

# assign a variable dataSummary to the dfsummary dataframe
# included in USGSHydroOpt
dataSummary <- dfsummary

# note the number of variables in dataSummary
length(colnames(dataSummary))

[1] 68

# assign a variable grnum to the column in dataSummary with
# sample numbers
grnum <- "GRnumber"

# use getLog10 to log 10 transform signals in dataSummary
testLog10 <- getLog10(dataSummary, signals, grnum)

# note that the log 10 transformed signals have been added to
# dataSummary
length(colnames(testLog10))

[1] 80

length(signals) + length(colnames(dataSummary))

[1] 80
```

## 4  Plotting Excitation-Emission (EEMs) Spectra

One of the most useful visualizations of fluorescence data is creating a contour plot including the intensities of all excitation-emission wavelength (nm) pairs for a given sample. The function plotEEMs2 creates contour plots of EEMs spectra and also labels important EEM peaks. Displayed below is an example of how plotEEMs2 can be used to plot EEMs spectra. The 3-D Excitation-Emission array discussed in Sections 2.8-2.8 is used to produce the EEMs plot.

```r
# choose a sample of interest from the 3-D EEMs array
GRnum <- c("gr14500")

# create a matrix from the 3-D EEMs array with only data for
# sample GRnum
mat <- a[, , GRnum]

# define the excitation wavelengths (nm) as Ex data type
# numeric
Ex <- as.numeric(names(a[, 1, 1]))

# define the emission wavelengths (nm) as Em data type
# numeric
Em <- as.numeric(names(a[1, , 1]))

# define the number of numeric color levels for contour plot
nlevels <- 50

# set a variable Peaks as the example dataframe ex_ems
Peaks <- ex_ems

# define the column in Peaks which contains the name of EEMs
# peaks
peakCol <- "Peak"

# define the column in Peaks which contains the excitation
# wavelength for a given Peak
peakEx <- "ExCA"

# define the column in Peaks whcih contains the emission
# wavelength for a given Peak
peakEm <- "EmCA"

# assign the main plot title
mainTitle <- paste("EEM spectra for", GRnum)

# use plotEEMs2 to create a contour plot of EEM spectra
exampleEEMs2 <- plotEEMs2(mat = mat, Ex = Ex, Em = Em, nlevels = nlevels,
    Peaks = Peaks, peakCol = peakCol, peakEx = peakEx, peakEm = peakEm,
    mainTitle = mainTitle)
```
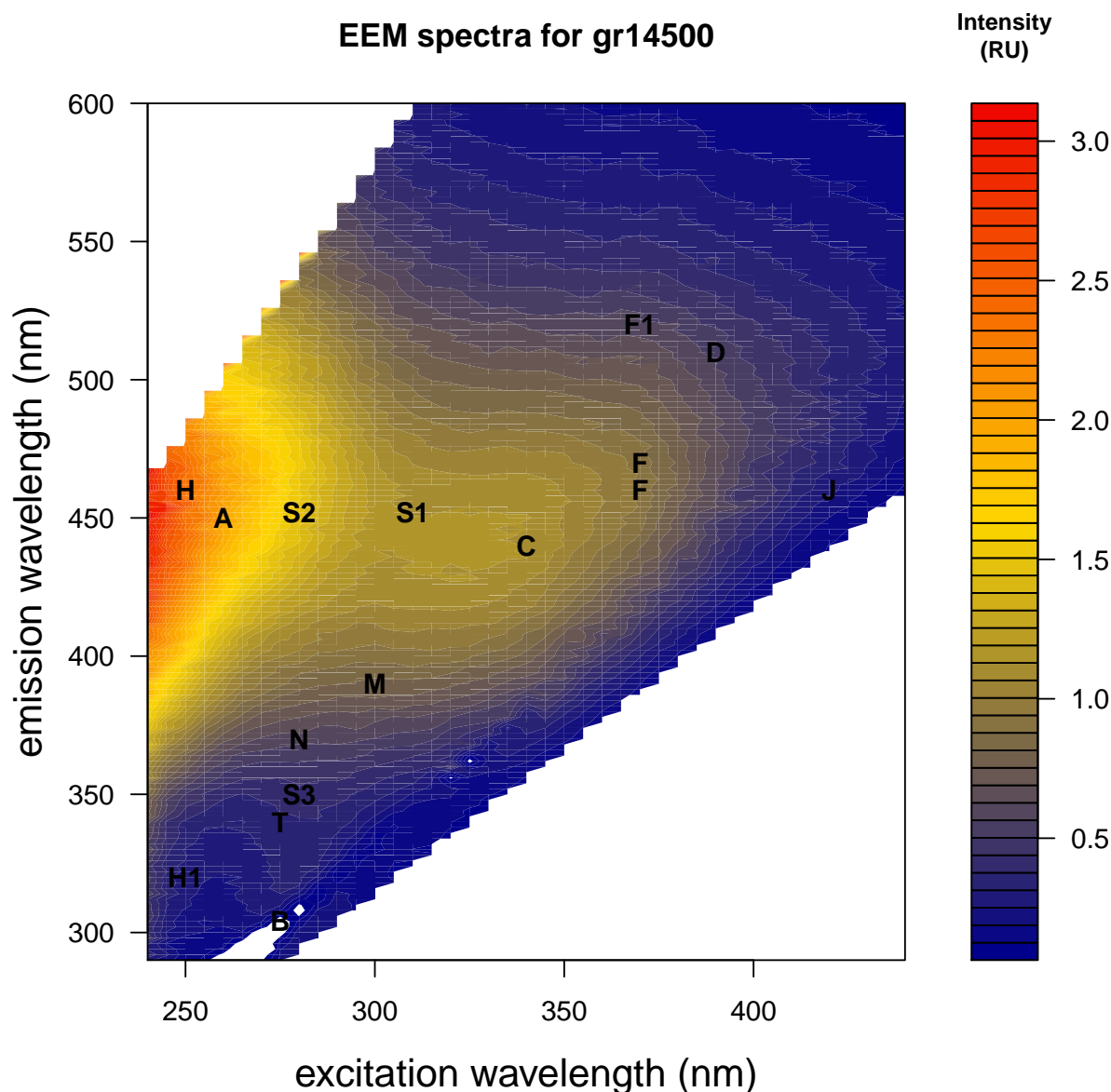
EEM spectra for gr14500

# 5 An Example Workflow Leveraging USGSHydroOpt

# 6 Conclusion

USGSHydroOpt can produce optical summary variables for absorbance and fluorescence spectra. Such predictor summary optical variables are useful for explaining biological and chemical contamination in water. This package also contains a useful function for visualizing fluorescence spectra by EEMs contour plots. A downfall of the package is that it was designed for data structures that must follow standardized formats, and consequently it is not ammenable to data structures that deviate from the prescribed format.

# 7 References

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