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Rachel Bruflodt
University of Iowa

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FINGERPRINTING THE REFUGIO OIL SPILL USING TOPOGRAPHIC SIGNAL PROCESSING OF
TWO-DIMENSIONAL GAS CHROMATOGRAPHIC IMAGES

by

Rachel Bruflodt

A thesis submitted in partial fulfillment of the requirements
for graduation with Honors in the Electrical Engineering

Ananya Sen Gupta
Thesis Mentor

Spring 2018

All requirements for graduation with Honors in the
Electrical Engineering have been completed.

Xiaodong Wu
Electrical Engineering Honors Advisor

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Chromatographic Images**

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May 2017

Mentor: Ananya Sen Gupta

Abstract

Oil spill fingerprinting aims to determine whether oil samples originate from the source of the spill or from a different source. This can be done by comparing the composition of hydrocarbons in each oil sample's two-dimensional gas chromatography (GC×GC) image. In this work, we compare oil samples taken from the 2015 Refugio Oil Spill with oil samples taken elsewhere. We use an approach called Peak Topography Mapping (PTM) which aims to automatically compare oil samples using both target and non-target peaks in the GC×GC image. Results suggest that the non-target peaks analyzed in PTM can provide information beyond that of solely target-based analysis.

1. Introduction and Background

1.1. Refugio Oil Spill

On May 19th, 2015, an oil pipeline owned by Plains All American Pipeline ruptured near Refugio State Beach in Santa Barbara, CA [1]. When a spill such as this occurs, it is important to be able to determine whether found oil has come from the spill or from other sources. For this work, samples were taken from the ruptured pipeline as well as other affected areas. These samples were compared to each other and to other oil samples not from the spill.

1.2. Two-Dimensional Gas Chromatography for Oil Spills

Each oil sample has a specific hydrocarbon profile. This profile can be seen in the form of a Two-dimensional gas chromatography image (GCxGC), where the peaks of the image correspond to biomarker compounds [3,4]. By comparing their GCxGC images, we can determine if two oil samples come from the same source or different sources.

1.3. Current Methods of Comparison

There are two broad approaches for fingerprinting oil samples. One method is target-based analysis [8, 9, 10] where only the peaks corresponding to target compounds are compared. This method can sometimes fail to distinguish between different oil sources because the target peaks are very similar. Target-based analysis also ignores all non-target peaks which may also contain useful information. The other approach for fingerprinting oil spills is statistical pattern recognition [11, 12, 13]. This approach can be sensitive to shifts in retention time and can also require large sets of training data. The statistical pattern recognition approach does not distinguish between target and non-target compounds.

2. Methods

Our approach was to use both target and non-target peaks to determine the similarity of GC×GC samples. We did this using the peak topography map concepts outlined in [2, 5, 6] with some modifications.

2.1. PTM

A peak topography map (PTM) is a two-dimensional matrix representation of the peaks of a GC x GC image. The first dimension of the PTM is the first dimension retention time index of the GC x GC image, and the second dimension of the PTM contains peaks in order of their elution time along the second dimension of the GC x GC image. The PTM is normalized against the maximum peak. In this case, peaks were not considered in the PTM unless their height was at least 10% of the maximum peak height in the image. This value resulted in the most consistent number of peaks across samples expected to match. As explained in [2], if the n th column of a GC x GC image has k_n peaks, and the amplitudes and locations of the peaks are stored as $Peak_n = \{p_{1,n}, p_{2,n}, \dots, p_{k_n,n}\}$ and $Loc_n = \{m_{1,n}, m_{2,n}, \dots, m_{k_n,n}\}$, the $(l, n)^{th}$ element PTM representation is described as

$$PTM[l, n] = \begin{cases} p_{l,n} + j \times m_{l,n} & \text{if } 1 \leq l \leq k_n \\ 0 & l > k_n \end{cases}$$

2.2. Topography Partitioning

The PTMs of individual oil samples were compared using the topography partitioning described in [2]. This involves comparing peaks at the same locations in the PTM and determining the peak ratio. The peak ratio is defined as

$$\rho = \max(a, a^{-1}) \text{ where } a = \frac{p_{ref}}{p_{test}}.$$

A peak is considered “similar” if the peak ratio is below the peak threshold, and “dissimilar” if the peak ratio is above the peak threshold. For this investigation, 1.2 was used as the peak ratio threshold below

which peaks were considered to match. This number was chosen because it resulted in the largest difference between the match percentages of expected matches and expected non-matches.

2.3. Cross-PTM Score

The peak topography mapping technique was applied over 5 different coordinate boxes within each raw sample. Each box corresponds to a different group of compounds: Archeans, Monoaromatic Steranes, Hopanes, or Steranes. The Archean group is comprised of two separate coordinate boxes. The peak topography mapping technique produces a cross-PTM score, which is a percentage match value for each pair of samples. The percentage represents the combined similarity between the target and non-target peaks in the sample. As explained in [2], it is calculated as the weighted percentage of peaks that are “similar” between the pair of samples:

$$S_{\tau}(I_{test}, I_{ref}) = \frac{|\eta_{test} \in PTM_{test,aligned}(PTM_{ref}): \rho(m, n) \geq 1.2|_w}{\eta_{test} \in PTM_{test,aligned}(PTM_{ref})}$$

where I_{test} and I_{ref} are the test and reference samples, η_{test} is a node in the test PTM, and $|\cdot|_w$ is the weighted sum taken across all peaks that meet the 1.2 threshold such that larger peaks are weighted higher than smaller peaks.

3. Results

Expanding on the results in [7], we provide results based on 15 GC×GC samples, some of which were taken from the area of the Refugio spill.

3.1. Matching Samples

Table 1 provides the percentage match of each sample to RS029 in each compound region. RS029 is a sample taken from the source of the Refugio spill. The rows

highlighted green indicate the samples which were expected to match RS029 and the rows highlighted orange indicate the samples which were expected to not match RS029. Table 2 provides the number of peaks found in each sample for each compound region and uses the same color coding scheme. Table 3 provides descriptions for the samples expected to match RS029.

Table 1 Percent matches to RS029

Sample	Archean Index (243,440) to (260,540) match	Archean Index (212,610) to (225,645) match	Monoaromatic Steranes (80,25) to (130, 183) match	Hopanes (75,185) to (140,585) match	Steranes (140,90) to (180,346) match
Sample RS029	100%	100%	100%	100%	100%
Sample RS004	99.9971%	100%	81.3603%	89.8363%	87.3532%
Sample RS005	100%	95.5480%	66.7552%	64.4538%	83.7939%
Sample RS007	96.0916%	92.0283%	14.5531%	26.7380%	30.2062%
Sample RS009	94.1948%	95.8331%	25.2361%	27.2325%	33.0696%
Sample RS028	95.6403%	96.3337%	2.1340%	25.9239%	31.1829%
Sample RS036	85.1580%	91.8798%	25.1399%	26.7581%	31.9805%
Sample RS023	17.7005%	41.0260%	6.4315%	3.2441%	23.8260%
Sample RS070	85.7867%	45.1885%	10.4183%	0.0146%	24.2280%
Sample RS071	14.5794%	71.8709%	5.5889%	3.1727%	19.7485%
Sample Seep_Stringer	59.3328%	78.5981%	99.9916%	99.9860%	99.7506%
Sample Seep_Surface	5.2606%	46.8358%	68.7344%	83.9838%	6.7912%
Sample SRM1582_A	5.9012%	15.5048%	9.6640%	15.8478%	27.2685%
Sample SRM1582_B	13.7645%	9.8375%	7.4759%	18.6454%	26.1028%
Sample SRM1582_C	5.8993%	15.6132%	7.7073%	18.2553%	25.3634%

Table 2 Number of peaks

Sample	Archean Index (243,440) to (260,540) number of peaks	Archean Index (212,610) to (225,645) number of peaks	Monoaromatic Steranes (80,25) to (130, 183) number of peaks	Hopanes (75,185) to (140,585) number of peaks	Steranes (140,90) to (180,346) number of peaks
Sample RS029	28	34	68	25	256
Sample RS004	27	34	81	28	224
Sample RS005	27	36	84	29	214
Sample RS007	26	35	94	37	229
Sample RS009	26	36	93	35	229
Sample RS028	26	36	78	38	266
Sample RS036	26	37	83	37	215
Sample RS023	41	36	262	40	368
Sample RS070	81	32	175	47	281
Sample RS071	41	35	261	44	317
Sample Seep_Stringer	37	36	72	27	221
Sample Seep_Surface	16	16	82	30	543
Sample SRM1582_A	37	34	309	48	354
Sample SRM1582_B	37	34	309	45	351
Sample SRM1582_C	38	34	310	46	347

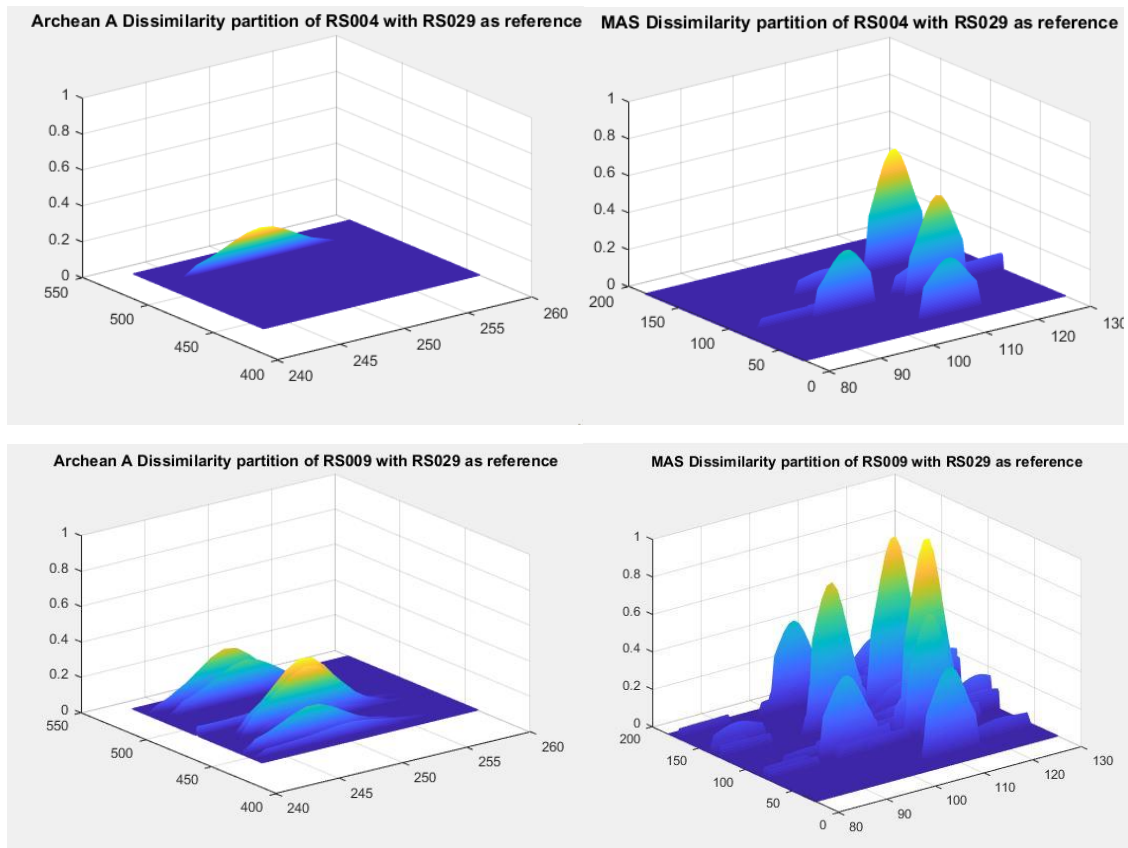
Table 3 Sample descriptions

Sample	Description
RS004	Minimal weathering; collected day 0
RS005	Minimal weathering; collected day 0
RS007	Some on-water weathering; collected day 1
RS009	Some on-water weathering; collected day 1
RS028	Some weathering; collected day 1
RS036	Some on-water weathering; collected day 2

3.2. Differences in Match Percentages Across Compound Regions

Table 1 shows that RS029 matches with itself 100% in each compound region as expected. It also shows that the regions which most clearly differentiate between expected matches and non-matches are the two Archean regions. The match percentages for the samples expected to match RS029 are much lower in the Monoaromatic Sterane, Hopane, and Sterane regions. Samples collected after day 0 show the most dramatic drop in match percentages from the Archean regions to the other regions. The two samples collected on day 0 maintain relatively high match percentages in the non-Archean regions. Figure 1 shows the dissimilarity partitions for the samples RS004 and RS009 in the first Archean region and the Monoaromatic Sterane (MAS) region. That is, it shows the peaks in those regions of those two samples that did not match corresponding peaks in RS029.

Figure 1 Dissimilarity partitions



The figure shows that the major nonmatching peaks in the Archean region are fewer and generally smaller than the major nonmatching peaks in the non-Archean Monoaromatic Sterane region. Similar results are seen across all samples in all Archean and non-Archean regions. This accounts for the strong match percentages between RS029 and expected matches to RS029 in the Archean regions. This also accounts for the less strong match percentages in non-Archean regions.

4. Discussion

4.1. SRM1582 Discrepancies

The three samples SRM1582_A, SRM1582_B, and SRM1882_C represent the same sample which was processed three different times to produce three different GC×GC images. The discrepancies in match percentages for these three samples in Table 1 are explained by small differences in peak values in the raw images. The three images have enough small differences that a few peaks in one SRM image may match corresponding peaks in the RS029 image, while the same peaks in the other SRM samples do not. This is consistent with the fact that the largest match percentage discrepancies occur in the compound regions with the fewest number of peaks, meaning that an occasional inconsistently matched peak has a larger effect on the overall match percentage. When the peak topography technique is applied to compare the SRM samples to each other, they all match each other with a match percentage of 100%.

4.2. Effects of Weathering

Figure 1 may show some potential effects of weathering. The sample with minimal weathering collected on day 0 (RS004) has a few, mostly small peaks in its dissimilarity partition for the MAS region. The sample collected on day 1 with some on-water weathering (RS009) on the other hand, has many small peaks in its dissimilarity partition with some larger ones. Many small peaks are characteristic of the dissimilarity partitions for weathered samples in the Monoaromatic Sterane, Hopane, and Sterane compound regions. Dissimilarity partitions for weathered samples in the Archean regions contain less peaks, and have smaller peaks overall. Dissimilarity partitions in the Archean regions therefore show less of these effects possibly influenced by weathering. More varied samples are needed to further understand the effect of weathering in each compound region.

5. Conclusions

The cross-PTM scores for the Archean compound region were able to differentiate oil samples that had come from the Refugio spill from oil samples that had not. The distinction was less clear when comparing cross-PTM scores for the Hopane, Sterane, and Monoaromatic Sterane regions.

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