Estimation of trace vapor concentration-pathlength in plumes for remote sensing applications from hyperspectral images

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Objective and Outline

- Objective
 - Show that end members extracted from the off-plume can be used as a basis for the quantification of plume analytes
 - Allows non-negativity constraints to be employed
 - Show how mismatch between the plume temperature and estimator spectra affect the quantification error
- Remote sensing problem
- IR-SAGE (Synthetic Scene Generator)
- Quantification Algorithm
- End-Member Extraction
- Results
- Conclusions



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Remote Sensing Problem

- Remote sensing for chemical analytes can be split into 3 tasks:
 - Detection (where is the plume?),
 - Classification (what is in the plume?), and
 - Quantification (how much is in the plume?).
- Interest is in quantification
 - quantification is limited to concentration-pathlength since the actual pathlength is usually unknown





Problem with Measured Data

- Hyperspectral images are rarely well-characterized
- There are a large number of factors influencing the measured radiance
 - plume location, contents, quantities, temperature,
 - atmosphere down- and up-welling radiance, and transmittance,
 - ground emissivity and temperature ...
- are *unknown*
- so how to judge performance of algorithms?



Synthetic Data

- IR-SAGE: InfraRed Systems Analysis in General Environments
 - flexible synthetic scene generator (in MATLAB)
 - · combines physical models and measured spectra
 - everything is known
 - can be used to test detection, classification, and quantification algorithms



Radiance Model



How to Quantify?

- Lot's of parameters, so ...
 - split the problem into parts, and assume some parameters are known
 - (future work)
 - assume plume location is known (detection)
 - assume plume analytes are known (classification)
- Assume that on-plume pixel "looks like" an off-plume pixel if there were no plume
 - on-plume "background" lies in subspace spanning off-plume pixels



Rearrange Radiance Model for τ_p

as $\tau_p \to 1$ (no plume), $L_{on}(\upsilon) \to L_{off}(\upsilon) \equiv L_{bkg}(\upsilon)$

$$\tau_p = e^{-\mathbf{c}\mathbf{S}^T} = \frac{L_{on}(\upsilon) - L_u(\upsilon) - \tau_a^{up}B(\upsilon,T_p)}{L_{bkg}(\upsilon) - L_u(\upsilon) - \tau_a^{up}B(\upsilon,T_p)}$$



Assumptions

- Assumed we know: $\overline{L}_{u,off}$, $\overline{\tau}_{a,off}^{up}$, T_p
- Don't require knowledge of: T_g and ε_g
- Taylor Series, substitute known parameters

$$\mathbf{cS}^{T}(\upsilon) = \frac{L_{on}}{d(\upsilon)} - \frac{L_{bkg}}{d(\upsilon)} \qquad \qquad d(\upsilon) = \overline{L}_{u} + \overline{\tau}_{a}^{up} B(T_{p}) - \overline{L}_{bkg}$$

$$\mathbf{c}\tilde{\mathbf{S}}^{T}\left(\boldsymbol{\upsilon}\right) = L_{on} - L_{bkg}$$



Quantification: ELS

• Extended mixture model (extended least squares) which can be solved for [c t] using least squares

$$\mathbf{l}_{on} = \begin{bmatrix} \mathbf{c} & \mathbf{t} \end{bmatrix} \begin{bmatrix} \tilde{\mathbf{S}} & \mathbf{P}_{bkg} \end{bmatrix}^{T}$$

• End-member extraction for background basis

$$\mathbf{l}_{on} = \begin{bmatrix} \mathbf{c} & \mathbf{c}_{bkg} \end{bmatrix} \begin{bmatrix} \tilde{\mathbf{S}} & \mathbf{S}_{bkg} \end{bmatrix}^T$$

• can apply non-negativity constraints



End-Member Extraction

- Initial solution: extremes of mean centered 1-norm
- ALS with non-negativity constraints
- Sequential for "smaller" factors
 - extracted 14 to 53 factors
- Provides a basis on which non-negativity constraints can be used in quantification

$$\mathbf{L}_{off} = \mathbf{c}_{bkg} \mathbf{S}_{bkg}^T + \mathbf{E}$$



End-Member Extraction





Quantification: GLSI

• ELSI predictions were good for narrow-featured spectra but not as good for broad-featured spectra

$$\mathbf{c}\tilde{\mathbf{S}}_{i}^{T}(\upsilon) = \left[L_{on}(\upsilon) - \overline{L}_{off}(\upsilon)\right] - \left[L_{off}(\upsilon) - \overline{L}_{off}(\upsilon)\right]$$
$$\mathbf{W} = \left[\mathbf{L}_{off} - \overline{\mathbf{L}}_{off}\right]$$
$$\hat{\mathbf{c}} = \left(\mathbf{I}_{on} - \overline{\mathbf{L}}_{off}\right) \mathbf{W}^{-1} \begin{bmatrix}\tilde{\mathbf{S}} & \mathbf{S}_{bkg}\end{bmatrix} \left(\begin{bmatrix}\tilde{\mathbf{S}} & \mathbf{S}_{bkg}\end{bmatrix}^{T} \mathbf{W}^{-1} \begin{bmatrix}\tilde{\mathbf{S}} & \mathbf{S}_{bkg}\end{bmatrix}\right)^{-1}$$

• GLS, after the last ELSI iteration, resulted in better estimates for broad-featured



Algorithm Testing

- Image 128x128 x 536 (800 to 1335 cm⁻¹)
 - 1, 2, 4, 8, 128 different background materials \mathcal{E}_{g}
 - temperature variability, T_p - T_g (12 to 32 K)
 - plume 16x128 pixels
 - plume spectra at 298 K, estimator at 298 K: matched
 - plume spectra at T_p , estimator at 298 K: unmatched
 - 5 analytes in the plume, 9 to 26 ppm \cdot m
 - atmospheric variability
 - 0.112 cm⁻¹ res, 0.06 cm⁻¹ spacing, convoluted to 1 cm⁻¹



Unconstrained vs. Constrained







Matched vs. Unmatched

• temperature mismatch \Rightarrow higher estimation error





Unmatched - Matched

• Mismatch generally results in higher error



Model Bias

- Estimates of **c** tend to have a low bias due to model assumptions
 - Taylor series approx.
- Temperature mismatch can result in a high bias that partially offsets the model bias





Conclusions

- End member extraction can be used to obtain a basis on which non-negativity constraints can be applied
- Temperature mismatch between the plume temperature and the spectrum used for estimation generally results in higher estimation error



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Extra Slides



Summary of Assumptions

- plume location is known*
- plume analytes are known*
- for off-plume \overline{L}_{u} ,** $\overline{\tau}_{a}^{up}$,* and \overline{L}_{bkg}^{*} are known
- T_p is known^{*}
- L_{bkg} lies in subspace spanned by off-plume \mathbf{P}_{bkg}

* same assumption as present state-of-the-art

** present state of the art assumes T_g is known



Scenarios

side-looking



down-looking



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Noise Model



Quantification



ples for down-Ig scenarios be used for side-, upne: know where the me is (detection), and

know what is in the me (classification)



Not all terms are known...

- Approximate $e^{-\mathbf{cS}^T} \approx 1 \mathbf{cS}^T$
- isolate \mathbf{cS}^{T}
- and assume $L_u(v) \approx \overline{L}_{u,off}(v)$ $\tau_a^{up}(v) \approx \overline{\tau}_{a,off}^{up}(v)$ T_p known
- yielding

$$\mathbf{cS}^{T}(\upsilon) = \frac{L_{on} - L_{bkg}}{\overline{\tau}_{a.off}^{up} B(T_{p}) + \overline{L}_{u.off} - L_{bkg}}$$



Algorithm Testing

- Image 128x128 x 536 (800 to 1335 cm⁻¹)
 - 1, 2, 4, 8, 128 different background materials \mathcal{E}_{g}
 - T_g (284 to 300 K), T_p - T_g (12 to 32 K)
- Plume 16x128 (9 to 26 ppm·m)
 - 1 to 4 analytes in the plume
- US 76 Standard Atms
 - 101 different FASCODE realizations
 - 1 % T and 3 % C variation w/in each layer
 - H_20 , CO_2 , O_3 , N_2 , CO, CH_4 , O_2 , and 25 others
- All spectra
 - 0.112 cm⁻¹ res, 0.06 cm⁻¹ spacing
 - convoluted to 1 cm⁻¹



"Mathematical Construct"





Results: 4 Dissimilar Analytes





Results: GLSI / SOA

- Compare GLSI to State-of-the-Art (SOA)
- For 4-analyte plume of dissimilar analytes



Results: Num. of Backgrounds

- Compare GLSI vs number of backgrounds
- For 4-analyte plume of dissimilar analytes

RMSEP (ppm·m)	Number of Backgrounds					
Analyte	<u>128</u>	8	4	2	1	
NH3	1.0	0.85	0.85	0.93	0.94	
NBUTANOL	16	4.7	3.4	3.9	3.9	
CH4	39	32	32	36	38	
F113	1.0	0.65	0.63	0.67	0.66	

Estimation vs. NAS

• Net Analyte Signal accounts for spectral modification and number of basis vectors in \mathbf{P}_{bkg}



