Article

Studying Vegetation Salinity: From the Field View to a Satellite-Based Perspective

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Academic Editors: Lenio Soares Galvao and Prasad S. Thenkabail

Received: 6 December 2016; Accepted: 19 January 2017; Published: 1 February 2017

Abstract: Salinization of irrigated lands in the semi-arid Jezreel Valley, Northern Israel results in soil-structure deterioration and crop damage. We formulated a generic rule for estimating salinity of different vegetation types by studying the relationship between Cl/Na and different spectral slopes in the visible–near infrared–shortwave infrared (VIS–NIR–SWIR) spectral range using both field measurements and satellite imagery (Sentinel-2). For the field study, the slope-based model was integrated with conventional partial least squares (PLS) analyses. Differences in 14 spectral ranges, indicating changes in salinity levels, were identified across the VIS–NIR–SWIR region (350–2500 nm). Next, two different models were run using PLS regression: (i) using spectral slope data across these ranges; and (ii) using preprocessed spectral reflectance. The best model for predicting Cl content was based on continuum removal reflectance ($R^2 = 0.84$). Satisfactory correlations were obtained using the slope-based PLS model ($R^2 = 0.77$ for Cl and $R^2 = 0.63$ for Na). Thus, salinity contents in fresh plants could be estimated, despite masking of some spectral regions by water absorbance. Finally, we estimated the most sensitive spectral channels for monitoring vegetation salinity from a satellite perspective. We evaluated the recently available Sentinel-2 imagery’s ability to distinguish variability in vegetation salinity levels. The best estimate of a Sentinel-2-based vegetation salinity index was generated based on a ratio between calculated slopes: the $490–665 \text{ nm}$ and $705–1610 \text{ nm}$. This index was denoted as the Sentinel-2-based vegetation salinity index (SVSI) $(\text{band 4} / \text{band 2}) / (\text{band 5} + \text{band 11})$.

Keywords: reflectance spectroscopy; spectral slope; salinity; fresh vegetation; tomato; cotton; Sentinel-2; Sentinel-2-based vegetation salinity index (SVSI)

1. Introduction

Salinity is one of the major ecological factors limiting plant growth and productivity, especially in arid and semi-arid regions with high rates of evapotranspiration and irrigated agriculture [1–3]. Increasing soil salinity leads to degradation of vegetation quality, due to a substantial decrease in water content, chlorophyll, and mineral elements such as K, Ca, Mg, N, and P, for the plants [4–6]. A good example of saline soils can be found in the Jezreel Valley in Northern Israel where over the years, increasing soil salinity has led to deterioration of the soil structure and damage to crops [7]. Importantly, if the saline-prone plot area is recognized on temporal and spatial scales, proper agricultural treatment (e.g., construction of a drainage system) can substantially decrease the crop’s degradation [8].

Conventional methods for monitoring changes in soil salinity are based on field observations and laboratory analyses of both crops and soils. This is mostly done by measuring the electrical
conductivity, exchangeable sodium percentage, and pH of the soil solution. However, this method is
time-consuming, expensive, and restricted to limited areas.

Reflectance spectroscopy in the visible–near infrared–shortwave infrared (VIS–NIR–SWIR)
region combined with multivariate statistical methods is widely used to extract quantitative information
on a material’s composition (e.g., [9–11]). One potential application of such combinations is the
analysis of fresh plants (e.g., leaves, whole plants of undried wheat (Triticum aestivum L.) [12], as
recently discussed by Cozzolino [13]. For example, not only have dry matter, N, oil, and protein been
accurately estimated in fresh samples, but also detection of plant diseases of mandarin in the field
has been reported. In a controlled experiment, the salinity levels of fresh plants were assessed based
on band ratios in the VIS spectral range [14–18]. In another study, the relationship between Na and
Cl contents of tomato plants was correlated with the underlying soil’s spectral reflectance [18]. The
authors found a relatively high accuracy of estimated plant salinity levels. Thus, reflectance spectroscopy
techniques can be used to characterize salinity in growing vegetation.

The spectrum of halite (NaCl) in the VIS–SWIR (400–2500 nm) region is featureless [10].
However, models for assessing salinity based in part on this spectral region have been developed for
vegetation [14,16,17,19], meat [20], Atlantic salmon (Salmo salar) [21], and for mapping salt-affected
soils [8,22–26]. Begley et al. [20] explained that the ability to measure salt in meat by NIRS (near
infrared spectroscopy) is due to the shift in the water spectrum caused by salt-induced changes in the
amount of hydrogen bonding. Ben-Dor and Banin [27] pointed out that featureless spectral properties
(properties without a direct chromophore) may also be predicted via internal correlation with
chromophoric properties. Hence, although NaCl is spectrally featureless, one could determine which
spectral regions might be indicative of the presence of salt features or traces. This knowledge is of
particular interest with the recent launch of the Sentinel-2 multi-spectral instrument (MSI) with its
improved spectral resolution (i.e., narrower bands) that offers significant advantages for capturing
vegetation status [28].

The main goal of our study was to formulate a generic rule for estimating the salinity of different
types of vegetation in the field. We used two populations, tomato and cotton leaves, to study the
relationship between Cl/Na contents and different spectral slopes in the VIS–NIR–SWIR spectral
range. Furthermore, the slope-based model was integrated with conventional PLS analyses. Next, we
investigated the potentially most sensitive spectral channels for vegetation salinity monitoring from a
satellite perspective. We used the recently available Sentinel-2 imagery and different satellite-based
indices were generated and correlated to Cl and Na contents. Finally, a spatial map of vegetation
salinity was generated on a pixel basis for the studied area.

2. Materials and Methods

2.1. Study Area

Three irrigated fields in the Jezreel Valley (Northern Israel) were chosen: two tomato fields in
Ifat (32°37′20.89″N, 35°14′42.02″E) and in Mizra (32°36′30.81″N, 35°14′43.25″E), and one cotton field
in Mizra (Figure 1). The crops in the research area were seasonally affected by soil salinity, due to
groundwater levels rising close to the soil surface and a lack of drainage. The processes leading to
the formation of soil salinity in the upper layer are detailed elsewhere [14,29].

2.2. Field Study

2.2.1. Sample Description

In each field, leaf samples were taken at 20-m intervals along three to five of 120–200-m transects
(Figure 1). Each sample included three leaves taken from the sampled shrub. For tomato, 22 leaf
samples were taken from the Mizra field and 50 from Ifat, and for cotton, 33 samples were taken from
the Mizra field, for a total of 105 plant samples. The leaf samples were collected on 26 June 2008 and
20 May 2009 for tomato samples in Mizra and Ifat, respectively, and on 13 July 2011 for cotton
samples. The leaf samples of tomato were collected at the reproductive stage that includes flowering
and fruit-bearing. In previous studies it was found that yield loss of tomato (*L. esculentum*) was most remarkable in reproductive stage under salinity treatments [30]. Therefore, salinity mapping in early growing stages is important.

**Figure 1.** Location of Jezreel Valley on a map of Israel (*left*), with study fields bordering the Genigar Reservoir (*right, top*) and cotton planted in rows in the irrigated field at Mizra (*right, bottom*). Plant sampling locations are marked.

2.2.2. Spectral Reflectance Measurements

The spectral reflectance measurements of the fresh vegetation samples were acquired in the field using an Analytical Spectral Devices (ASD; Boulder, CO, USA) Fieldspec-Pro JR spectrometer furnished with a contact probe. The ASD includes three detectors across the 350–2500 nm spectral region (VIS-NIR between 350 and 1000 nm, SWIR1 between 1001 and 1800 nm and SWIR2 between 1801 and 2500 nm), and measures spectra in 2151 bands at 1-nm intervals [31]. The spectral measurements were acquired by direct contact of the probe to the leaf surface that were placed on the black background. For all experiments, the reflectance, measured relative to a white Halon reflectance panel reference [32], was used to enable conversion of the measurement data into reflectance values. Each spectral measurement represented an average of 40 spectral readings and the average spectra of three leaves per vegetation sample were used in our analyses.

2.2.3. Chemical Analyses

The vegetation samples were oven-dried for 72 h at 60 °C. Three leaves of each sample were ground together to pass through a 1-mm sieve, and subjected to chemical analysis for Na and Cl. Cl anion and Na cation contents, which are indicators of the presence of salinity in vegetation, were selected for correlation with spectral reflectance. Na content weight percent (wt%) was analyzed by flame photometry following digestion with H$_2$SO$_4$ and H$_2$O$_2$ with a measurement error of 5%. Cl content (wt%) was analyzed by chloridimetry following extraction with 0.1 N HNO$_3$ [33] with a measurement error of 10%.
2.2.4. Slope Calculation and Data Analyses

First, the results of all spectral measurements were presented in continuum-removal (CR) spectra [9] to investigate the spectral behavior vs. salinity of the vegetation. To develop a model to assess Cl and Na content in vegetation based on the slope method [34,35], we used the common CR spectral technique [9,36–40]. CR normalizes reflectance spectra to a common baseline, enabling the identification of individual absorption features. The differences in CR spectral behavior as a function of wavelength, which are indicative of changes in vegetation salinity, were identified. Then, different slopes between a pair of wavelengths were calculated and the most correlative were used in our analyses.

In addition, PLS models were generated using the entire spectral range. The original spectra were estimated using CR and first derivative (FD) units. For FD, we used the Savitzky–Golay algorithm [41,42]. The Savitzky–Golay method computes first- or higher-order derivatives, with a smoothing filter of three wavelengths that reduces instrumental noise and determines how many adjacent variables (wavelengths) will be used to estimate the polynomial approximation for the derivation [41].

2.2.5. Calculating the Water-Absorption Area

Water is one of the most prominent factors present in vegetation, which influences the total incident solar energy in the NIR–SWIR region that is absorbed by the vegetation [43]. Hence, calculating the absorbance area of the water bands (e.g., 1200, 1400, 1780, and 1940 nm) [11,43,44] can be a good indicator of water concentration in the vegetation. The spectral absorption features centered at 1940 nm have been found to be good indicators of plant water content [34,45,46]. Therefore, the area between the straight line and the absorption line (Figure 2) was calculated (see [35] for more details).

![Figure 2. Demonstration of total absorption area of two vegetation samples. The larger area (black) is of sample with low Cl and Na contents, whereas the smaller area (gray) is of sample with high Na and Cl contents.](image)

2.2.6. PLS Data Analyses

The calibration equations were created using PLS regression with previously defined spectral slopes in the VIS–NIR–SWIR spectral range of tomato and cotton leaf samples and each chemical reference (i.e., Cl and Na):

\[
Y = A + A_1X_1 + A_2X_2 + A_3X_3 + \ldots + A_nX_n
\]  

(1)

where Y is the chemically-measured Cl or Na content of a leaf sample, A is an empirical coefficient at a specific wavelength/slope, and X is the slope of the spectra in a specific range, or spectral reflectance at a specific wavelength.

Statistical parameters for the calibration model were calculated by leave-one-out cross-validation. The difference between the predicted and measured chemical values was expressed as the root mean square error of prediction (RMSEP) or the root mean square error of cross validation (RMSECV) [47]:

```
Y = A + A_1X_1 + A_2X_2 + A_3X_3 + \ldots + A_nX_n
```
\[
RMSEP = \sqrt{\frac{\sum (X_m - X_p)^2}{n}}
\]

where \(X_m\) and \(X_p\) are the chemically-measured and predicted values, respectively, of a sample based on the spectral analysis, and \(n\) is the number of samples in the calibration stage. In addition, the predictive capability of the model was evaluated by the ratio of prediction to deviation (RPD), which is defined as the ratio of the standard deviation of the reference values (e.g., of Cl) to the RMSECV or RMSEP [47]. For example, RPD values between 2 and 2.5 indicate quite good, and above 2.5, excellent predictive capability of the model [48]. All data management, calculations, PLS analyses, and different spectral pretreatments were performed using Unscrambler version 9.7 (Camo Software, Oslo, Norway).

To validate our model, several datasets were constructed: slopes, reflectance, CR, and FD. Each set was divided into two subsets: (i) a calibration set, comprised of 70–76 representative tomato and cotton samples, which was calculated by the cross-validation method; and (ii) an external test set, comprised of 14–17 tomato and cotton samples, which was used to examine the model’s predictive ability. As the number of samples was limited, the calculation of statistical parameters for the calibration model was done by leave-one-out cross-validation [41]. Cl and Na were log-transformed to obtain a normal distribution of reference values and were subjected to PLS analysis.

2.3. Satellite-Based Estimation of Salinity

As an example of future applications, we estimated a currently available system, Sentinel-2. First, we compared salinity indices that were previously generated for Landsat. The Landsat-based indices were matched to the Sentinel-2 configuration. Next, the results of all field measurements were spectrally resampled to match this configuration. The spectral resampling to match between field spectral measurements to Sentinel-2 was done by using “spectral resampling” procedure implemented in ENVI software (Version 5.2) [49]. Briefly, it uses a Gaussian model with FWHM (full width at half maximum) to match the desired sensor configuration. We calculated the following: salinity index (SI), normalized differential salinity index (NDSI), vegetation soil salinity index (VSSI) and salinity ratio [50,51]. The calculated indices were correlated to Cl and Na contents. For example, SI (\(\sqrt{b_1 \times b_3}\)) with band 1, centered at 485 nm and band 3 centered at 660 nm, of Landsat 8 was equivalent to band 2 centered at 490 nm and band 4 centered at 665 nm of Sentinel-2.

To generate a map of vegetation salinity levels, we similarly correlated the Na/Cl contents and slopes of all possible pairs of channels using simple linear regression analysis. In addition, we estimated the different ratio indices based on calculated slopes. The spectral slope or ratio with the highest regression coefficient was chosen to represent vegetation salinity levels on a pixel base.

To obtain a spatial representation of vegetation salinity levels using Sentinel-2, a cloud-free image (3 July 2016) of the study area was downloaded from the archives of the USGS Earth Explorer website [52]. We used a level-1C product which is a geo-coded top of the atmosphere reflectance with a subpixel multispectral registration (European Space Agency (ESA, [53]). Sentinel-2 carries a MSI that records in 13 spectral channels, covering a range of wavelengths from 440 to 2200 nm with spatial resolutions of 10, 20, and 60 m. The Sen2Cor atmospheric correction processor was used to retrieve surface reflectance (ESA, [53]. To generate a salinity map, we first masked all non-vegetative pixels using the Sentinel-2 image: NDVI (normalized difference vegetation index) values were calculated and all values lower than 0.5 were removed from our analyses. Finally, a vegetation salinity map was generated using the slope or ratio with the highest regression coefficient.

3. Results

3.1. Chemical Reference: Cl and Na

Table 1 shows the statistical parameters of Cl and Na in fresh vegetation samples. The range of Cl values in the tomato samples was wider than in the cotton samples, whereas the range of Na values in cotton was wider than in tomato. The range of Cl and Na values in tomato samples was 0.75%–3.86%
(70 samples, with an average of 1.45% and STD = 0.64) and 0.13%–0.53% (average 0.22% and STD = 0.07), respectively. The range of Cl and Na values in cotton samples was 1.24%–2.86% (33 samples, average 1.94% and STD = 0.43) and 0.21%–1.21% (average 0.44% and STD = 0.23), respectively.

Table 1. Average, standard deviation, median, minimum and maximum values of Cl and Na concentrations (wt%) in tomato and cotton.

<table>
<thead>
<tr>
<th></th>
<th>Tomato + Cotton (103 Samples)</th>
<th>Tomato (70 Samples)</th>
<th>Cotton (33 Samples)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Cl</td>
<td>Na</td>
<td>Cl</td>
</tr>
<tr>
<td>Average</td>
<td>1.60</td>
<td>0.29</td>
<td>1.45</td>
</tr>
<tr>
<td>Standard Deviation</td>
<td>0.62</td>
<td>0.17</td>
<td>0.64</td>
</tr>
<tr>
<td>Median</td>
<td>1.46</td>
<td>0.23</td>
<td>1.25</td>
</tr>
<tr>
<td>Minimum</td>
<td>0.75</td>
<td>0.13</td>
<td>0.75</td>
</tr>
<tr>
<td>Maximum</td>
<td>3.86</td>
<td>1.12</td>
<td>3.86</td>
</tr>
</tbody>
</table>

3.2. Spectral Slope Analysis: Spectral Regions Sensitive to NaCl

Differences in 14 spectral ranges indicating changes in salinity levels were identified across the VIS–NIR–SWIR region (350–2500 nm) (Table 2). Figure 3a,b shows changes in the slope values as a function of different Cl and Na contents across nine of these spectral ranges. The slopes increased/decreased with increasing/decreasing Cl and Na contents. Figure 3c demonstrates these relationships by zooming in on the 2400–2440 nm spectral range, where the slope is seen to increase with increasing Cl and Na contents. Figure 4 shows the box plot of the slopes for the 14 spectral ranges. In general, the slopes can be divided into two groups, with large and small variability in their values between tomato and cotton.

Table 2. Coefficient of determination between slopes in selected spectral regions and Cl and Na contents of vegetation. The bold numbers are the selected range for Cl and/or for Na

<table>
<thead>
<tr>
<th>Slope Spectral Range (nm)</th>
<th>Cl</th>
<th>Na</th>
<th>R^2 (N = 103)</th>
</tr>
</thead>
<tbody>
<tr>
<td>410–430</td>
<td>0.4596</td>
<td>0.3863</td>
<td></td>
</tr>
<tr>
<td>430–452</td>
<td>0.4364</td>
<td>0.0918</td>
<td></td>
</tr>
<tr>
<td>500–536</td>
<td>0.3095</td>
<td>0.0811</td>
<td></td>
</tr>
<tr>
<td>550–585</td>
<td>0.3145</td>
<td>0.1151</td>
<td></td>
</tr>
<tr>
<td>585–675</td>
<td>0.1558</td>
<td>0.045</td>
<td></td>
</tr>
<tr>
<td>675–750</td>
<td>0.2045</td>
<td>0.0085</td>
<td></td>
</tr>
<tr>
<td>1124–1161</td>
<td>0.4184</td>
<td>0.4376</td>
<td></td>
</tr>
<tr>
<td>1162–1200</td>
<td>0.0384</td>
<td>0.2317</td>
<td></td>
</tr>
<tr>
<td>1200–1275</td>
<td>0.3456</td>
<td>0.3254</td>
<td></td>
</tr>
<tr>
<td>1294–1364</td>
<td>0.3575</td>
<td>0.4099</td>
<td></td>
</tr>
<tr>
<td>1915–2235</td>
<td>0.5371</td>
<td>0.2719</td>
<td></td>
</tr>
<tr>
<td>2340–2400</td>
<td>0.5964</td>
<td>0.2989</td>
<td></td>
</tr>
<tr>
<td>2400–2440</td>
<td>0.5872</td>
<td>0.376</td>
<td></td>
</tr>
<tr>
<td>2450–2480</td>
<td>0.047</td>
<td>0.1914</td>
<td></td>
</tr>
</tbody>
</table>

To test differences between the slopes of the tomato and cotton populations we used a paired t-test. The p-value between the two is presented in Figure 4. It can be seen that in four spectral ranges (430–452 nm, 500–536 nm, 585–675 nm, and 675–750 nm), slopes were similar between tomato and cotton with p > 0.05 and in 10 spectral ranges (410–430 nm, 550–585 nm, 1124–1161 nm, 1162–1200 nm, 1200–1275 nm, 1294–1364 nm, 1915–2235 nm, 2340–2400 nm, 2400–2440 nm, and 2450–2480 nm), the slope populations differed (p < 0.05). It is interesting to note that two of the four, and eight of the 10 spectral ranges (excluding 500–536 nm, 585–675 nm, 1162–1200 nm, and 2450–2480 nm) were found significant for predicting Cl and Na, respectively, and appear in the prediction model (Table 3).
Namely, similarities or dissimilarities between spectral slopes of tomato and cotton enabled prediction of Cl and Na contents. This also indicates the potential for developing a generic model for predicting salinity in other types of vegetation.

Figure 3. (a) Continuum removal (CR) reflectance spectra of vegetation samples with different percentages of Cl/Na. Note the variability in the slopes across the different spectral ranges: 410–430 nm, 430–452 nm, 550–585 nm, 1124–1161 nm, 1200–1275 nm, 1294–1364 nm, 2340–2400 nm, and 2400–2440 nm. Note that both cotton and tomato leaves exhibit spectral variability across similar spectral ranges as a function of changes in salinity levels. Therefore, both populations can be combined in the analyses; (b) visualization of the slopes’ tendency to increase or decrease as a function of different Cl/Na contents (the right Y axis is the scale for the slope between 1915 and 2235 nm). Up-pointing arrow indicates a slope increase with Cl/Na content increase; down-pointing arrow indicates a slope decrease with Cl/Na content increase; and (c) zooming in on the 2400–2440 nm spectral range to demonstrate the changes in slope with changes in Cl/Na concentration.
Table 3. Prediction models for Cl and Na using the slope-based method.

<table>
<thead>
<tr>
<th>Model Description</th>
<th>Cl</th>
<th>Na</th>
</tr>
</thead>
<tbody>
<tr>
<td>$2.539 \times \text{Slope} (410 - 430) + 1.255 \times \text{Slope} (675 - 750) + 0.542 \times \text{Slope} (1124 - 1161) + 0.27 \times \text{Slope} (1200 - 1275) - 1.036 \times \text{Slope} (1294 - 1364) - 2.896 \times \text{Slope} (1915 - 2235) + 0.854 \times \text{Slope} (2340 - 2400) + 0.963 \times \text{Slope} (2400 - 2440)$</td>
<td>$-1.248 + 3.209 \times \text{Slope} (410 - 430) - 2.506 \times \text{Slope} (430 - 452) - 1.781 \times \text{Slope} (550 - 585) - 1.025 \times \text{Slope} (1124 - 1161) - 1.619 \times \text{Slope} (1294 - 1364) + 1.457 \times \text{Slope} (2340 - 2400) + 1.003 \times \text{Slope} (2400 - 2440)$</td>
<td></td>
</tr>
</tbody>
</table>

3.3. PLS Analysis

Table 2 displays the coefficient of determination between slopes in specific spectral ranges and the Cl and Na contents of the plant samples. The values ranged from 0.1558 to 0.5964 for Cl and from 0.0918 to 0.4099 for Na. Fourteen spectral ranges were found to be correlated with Cl and/or Na, whereas 12 spectral ranges were selected to be correlated with Cl and 10 with Na. Note, that eight spectral ranges were found to be correlated with both Cl and Na (Table 2).

Table 4 presents the PLS regression analyses for the slope method and the various data preprocessing models. In general, the Cl-prediction models were better than the Na-prediction models. Specifically, the $R^2$ and RPD of the Cl-prediction models were: 0.84/2.56, 0.78/2.22, 0.77/1.99, and 0.62/1.68 for CR, reflectance, slope and FD, respectively. On the other hand, the $R^2$ and RPD of the Na-prediction models were: 0.52/1.49, 0.50/1.47, 0.63/1.69, and 0.62/1.67 for CR, reflectance, slope and FD, respectively.
Table 4. Partial least squares (PLS) regression model results of slope method and of various data pre-processing techniques to assess Cl and Na in fresh vegetation samples.

<table>
<thead>
<tr>
<th></th>
<th>Cl Model Statistics Characteristical</th>
<th>Log(Cl) Model Statistics Characteristical</th>
<th>Na Model Statistics Characteristical</th>
<th>Log(Na) Model Statistics Characteristical</th>
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<tbody>
<tr>
<td></td>
<td>Calibration</td>
<td>Validation</td>
<td>Prediction</td>
<td>Calibration</td>
</tr>
<tr>
<td>Total number of samples (Tomato/Cotton)</td>
<td>74 (50/24)</td>
<td>17 (12/5)</td>
<td>74 (50/24)</td>
<td>17 (12/5)</td>
</tr>
<tr>
<td>Slope</td>
<td>0.78</td>
<td>0.76</td>
<td>0.82</td>
<td>0.77</td>
</tr>
<tr>
<td>Offset</td>
<td>0.34</td>
<td>0.37</td>
<td>0.20</td>
<td>0.04</td>
</tr>
<tr>
<td>RMSE</td>
<td>0.25</td>
<td>0.27</td>
<td>0.22</td>
<td>0.07</td>
</tr>
<tr>
<td>RPD</td>
<td>2.13</td>
<td>2.00</td>
<td>1.99</td>
<td>2.10</td>
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<tr>
<td>R²</td>
<td>0.78</td>
<td>0.76</td>
<td>0.77</td>
<td>0.77</td>
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Continuum Removal

<table>
<thead>
<tr>
<th></th>
<th>Cl Model Statistics Characteristical</th>
<th>Log(Cl) Model Statistics Characteristical</th>
<th>Na Model Statistics Characteristical</th>
<th>Log(Na) Model Statistics Characteristical</th>
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<tr>
<td></td>
<td>Calibration</td>
<td>Validation</td>
<td>Prediction</td>
<td>Calibration</td>
</tr>
<tr>
<td>Total number of samples (Tomato/Cotton)</td>
<td>75 (52/23)</td>
<td>15 (11/4)</td>
<td>75 (52/23)</td>
<td>15 (11/4)</td>
</tr>
<tr>
<td>Slope</td>
<td>0.76</td>
<td>0.74</td>
<td>0.87</td>
<td>0.72</td>
</tr>
<tr>
<td>Offset</td>
<td>0.36</td>
<td>0.39</td>
<td>0.24</td>
<td>0.04</td>
</tr>
<tr>
<td>RMSE</td>
<td>0.24</td>
<td>0.26</td>
<td>0.22</td>
<td>0.07</td>
</tr>
<tr>
<td>RPD</td>
<td>2.06</td>
<td>1.92</td>
<td>2.56</td>
<td>1.91</td>
</tr>
<tr>
<td>R²</td>
<td>0.76</td>
<td>0.74</td>
<td>0.84</td>
<td>0.72</td>
</tr>
</tbody>
</table>

Reflectance

<table>
<thead>
<tr>
<th></th>
<th>Cl Model Statistics Characteristical</th>
<th>Log(Cl) Model Statistics Characteristical</th>
<th>Na Model Statistics Characteristical</th>
<th>Log(Na) Model Statistics Characteristical</th>
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<tbody>
<tr>
<td></td>
<td>Calibration</td>
<td>Validation</td>
<td>Prediction</td>
<td>Calibration</td>
</tr>
<tr>
<td>Total number of samples (Tomato/Cotton)</td>
<td>76 (54/22)</td>
<td>15 (10/5)</td>
<td>76 (54/22)</td>
<td>15 (10/5)</td>
</tr>
<tr>
<td>Slope</td>
<td>0.74</td>
<td>0.71</td>
<td>0.77</td>
<td>0.71</td>
</tr>
<tr>
<td>Offset</td>
<td>0.40</td>
<td>0.44</td>
<td>0.38</td>
<td>0.05</td>
</tr>
<tr>
<td>RMSE</td>
<td>0.26</td>
<td>0.28</td>
<td>0.27</td>
<td>0.08</td>
</tr>
<tr>
<td>RPD</td>
<td>1.97</td>
<td>1.81</td>
<td>2.18</td>
<td>1.88</td>
</tr>
<tr>
<td>R²</td>
<td>0.74</td>
<td>0.70</td>
<td>0.78</td>
<td>0.71</td>
</tr>
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</table>

First derivative

<table>
<thead>
<tr>
<th></th>
<th>Cl Model Statistics Characteristical</th>
<th>Log(Cl) Model Statistics Characteristical</th>
<th>Na Model Statistics Characteristical</th>
<th>Log(Na) Model Statistics Characteristical</th>
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</thead>
<tbody>
<tr>
<td></td>
<td>Calibration</td>
<td>Validation</td>
<td>Prediction</td>
<td>Calibration</td>
</tr>
<tr>
<td>Total number of samples (Tomato/Cotton)</td>
<td>76 (53/23)</td>
<td>15 (11/4)</td>
<td>76 (53/23)</td>
<td>15 (11/4)</td>
</tr>
<tr>
<td>Slope</td>
<td>0.77</td>
<td>0.71</td>
<td>0.71</td>
<td>0.75</td>
</tr>
<tr>
<td>Offset</td>
<td>0.36</td>
<td>0.44</td>
<td>0.47</td>
<td>0.04</td>
</tr>
<tr>
<td>RMSE</td>
<td>0.25</td>
<td>0.29</td>
<td>0.31</td>
<td>0.07</td>
</tr>
<tr>
<td>RPD</td>
<td>2.09</td>
<td>1.83</td>
<td>1.62</td>
<td>2.03</td>
</tr>
<tr>
<td>R²</td>
<td>0.77</td>
<td>0.71</td>
<td>0.61</td>
<td>0.75</td>
</tr>
</tbody>
</table>
As can be seen, the best model for the prediction of Cl content was based on CR reflectance. However, the CR model for Na estimation was the least precise (Table 4). Compared to CR, FD, and reflectance models, the slope-based model was quite good at predicting Cl content and best at predicting Na content (Table 4). The slope-based prediction models for Cl and Na are presented in Table 3, where in the Cl model, eight of the 12 selected spectral ranges were found significant, seven of 10 for the Na model. Moreover, when we compared the percentage of X and Y variability explained by the first two LV (latent variables) components in the PLS model, we found it to be higher for the slope method. Specifically, the first two LVs of the slope-based model explained 92% and 86% of the X variance (spectra) and 78% and 66% of the Y variance (Cl and Na, respectively), whereas in the CR model, they explained 82% and 82% of the X variance (spectra) and 74% and 52% of the Y variance (Cl and Na, respectively). A higher percentage of variance explained by a lower number of components indicates model stability [47].

3.4. Estimating the Potential of Vegetation Salinity Mapping Using Sentinel-2 Image

Table 5 shows the coefficient of determination between known salinity indices based on the Landsat configuration [50,51] and the Cl/Na contents of the plant samples. As can be seen, when these indices are modified to the Sentinel-2 bands, the correlation ranges from 0.128 to 0.235 for Cl and less than 0.03 for Na.

<table>
<thead>
<tr>
<th>Salinity Indices</th>
<th>Landsat bands</th>
<th>Sentinel-2 bands</th>
<th>$R^2$ (n=103)</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>SI</td>
<td>$\sqrt{(b_1 \times b_3)}$</td>
<td>$\sqrt{(b_2 \times b_4)}$</td>
<td>0.205</td>
<td>0.0163</td>
</tr>
<tr>
<td>NDSI</td>
<td>$(b_3 - b_4)/(b_3 + b_4)$</td>
<td>$(b_4 - b_8)/(b_4 + b_8)$</td>
<td>0.189</td>
<td>0.0121</td>
</tr>
<tr>
<td>VSSI</td>
<td>$2 \times b_2 - 5(b_3 + b_4)$</td>
<td>$2 \times b_3 - 5(b_4 + b_8)$</td>
<td>0.235</td>
<td>0.0245</td>
</tr>
<tr>
<td>SI</td>
<td>$\sqrt{(b_3 \times b_4)}$</td>
<td>$\sqrt{(b_4 \times b_8)}$</td>
<td>0.189</td>
<td>0.012</td>
</tr>
<tr>
<td>Salinity Ratio</td>
<td>$(b_3 - b_4)/(b_2 + b_4)$</td>
<td>$(b_4 - b_8)/(b_3 + b_8)$</td>
<td>0.128</td>
<td>0.025</td>
</tr>
</tbody>
</table>

Table 6 shows the coefficient of determination between the slopes calculated in the current study, again, based on the Sentinel-2 configuration. As can be seen, the higher spectral resolution provides better estimates of salinity. For Cl content, the correlation ranges from 0.138 to 0.434 and for Na from 0.140 to 0.368. Note that informative spectral ranges are between 490–1610 nm, 665–1610 nm, 490–2190 nm, and 665–2190 nm (values of $R^2$ above 0.4 were highlighted). Furthermore, in Table 6, two additional indices based on the slope logic are also presented, with $R^2$ between 0.464 and 0.488. The index with the highest $R^2$ is denoted as the Sentinel-2-based vegetation salinity index (SVSI) ((b_4 - b_2)/(b_5 + b_11), Table 6).

<table>
<thead>
<tr>
<th>Slope Spectral Range (nm)</th>
<th>Sentinel-2 Bands</th>
<th>$R^2$ (N=103)</th>
<th>$Cl^*$</th>
<th>$Na^*$</th>
</tr>
</thead>
<tbody>
<tr>
<td>490–560</td>
<td>b_2–b_3</td>
<td>0.299</td>
<td>0.086</td>
<td></td>
</tr>
<tr>
<td>490–665</td>
<td>b_2–b_4</td>
<td>0.232</td>
<td>0.062</td>
<td></td>
</tr>
<tr>
<td>560–665</td>
<td>b_3–b_4</td>
<td>0.138</td>
<td>0.040</td>
<td></td>
</tr>
<tr>
<td>490–705</td>
<td>b_2–b_5</td>
<td>0.392</td>
<td>0.113</td>
<td></td>
</tr>
<tr>
<td>560–705</td>
<td>b_3–b_5</td>
<td>0.337</td>
<td>0.097</td>
<td></td>
</tr>
<tr>
<td>665–705</td>
<td>b_4–b_5</td>
<td>0.331</td>
<td>0.095</td>
<td></td>
</tr>
<tr>
<td>490–740</td>
<td>b_2–b_6</td>
<td>0.273</td>
<td>0.045</td>
<td></td>
</tr>
<tr>
<td>490–1610</td>
<td>b_2–b_11</td>
<td>0.404</td>
<td>0.168</td>
<td></td>
</tr>
</tbody>
</table>
Table 6. Cont.

<table>
<thead>
<tr>
<th>Slope Spectral Range (nm)</th>
<th>Sentinel-2 Bands</th>
<th>Cl* Na* (R^2 (N = 103))</th>
</tr>
</thead>
<tbody>
<tr>
<td>560–1610</td>
<td>b3–b11</td>
<td>0.327 0.157</td>
</tr>
<tr>
<td>665–1610</td>
<td>b4–b11</td>
<td>0.408 0.179</td>
</tr>
<tr>
<td>705–1610</td>
<td>b5–b11</td>
<td>0.222 0.140</td>
</tr>
<tr>
<td>783–1610</td>
<td>b7–b11</td>
<td>0.343 0.321</td>
</tr>
<tr>
<td>490–2190</td>
<td>b2–b12</td>
<td>0.434 0.230</td>
</tr>
<tr>
<td>560–2190</td>
<td>b3–b12</td>
<td>0.325 0.226</td>
</tr>
<tr>
<td>665–2190</td>
<td>b4–b12</td>
<td>0.425 0.245</td>
</tr>
<tr>
<td>705–2190</td>
<td>b5–b12</td>
<td>0.238 0.225</td>
</tr>
<tr>
<td>783–2190</td>
<td>b7–b12</td>
<td>0.291 0.368</td>
</tr>
<tr>
<td>((490 – 665) / (560 + 2190))</td>
<td>((b4 – b2) / (b3 + b12))</td>
<td>0.464 0.364</td>
</tr>
<tr>
<td>((490 – 665) / (705 + 1610))</td>
<td>((b4 – b2) / (b5 + b11))</td>
<td>0.488 0.284</td>
</tr>
</tbody>
</table>

In Figure 5, we map the spatial variability in salinity levels between different vegetative areas using the SVSI. The salinity level increases with decreasing values of SVSI (Figure 5d). In the study area (white rectangle in Figure 5a,d), the vegetation salinity levels typically range between low and medium. A few plots show high salinity values. Importantly, a few plots (e.g., numbered 1) exhibit high variability in salinity, which was captured by the SVSI index. For example, at 120 m distance (numbered 1.1 in Figure 5d), the SVSI ranges from ~0.07 to ~0.25 (~0.85%–2.5% Cl). Note that the northwestern part of the generated map shows high vegetation salinity levels (white circle in Figure 5d, ~2.5%–4%).

Figure 5. Vegetation salinity map of the study area based on slope ratio index (SVSI) using Sentinel-2 imagery. (a) Sentinel-2 image of the study area; (b) NDVI; (c) masked NDVI < 0.5; and (d) vegetation salinity map based on slope ratio index.
4. Discussion

Previous studies highlighted that increasing salinity levels significantly decrease stomatal conductance, produce more negative water potentials, and decrease root hydraulic conductance, resulting in decreased water content in the plant [4,5,18,54]. Begley et al. [20] detected differences in the absorption bands of water at 1806 nm that were well correlated with different concentrations of NaCl in meat. A similar relationship between the total area of water absorption and crude protein content was found by Lugassi et al. [35].

Our results also indicate that water absorption bands can be used as tracers for salinity monitoring. Note that the slopes in the spectral range of 1915–2235 nm and also the total area of water absorption between 1840 and 2235 nm (Figure 3a) decreased with increasing Cl and Na contents. In addition, the highest coefficient in the prediction equation of the slope-based CI model (Table 3) was for the slope between 1915 and 2235 nm. Furthermore, when analyzing score plots of our PLS model, there is an increase in Cl values (Figure 6) and a decrease in water content of the vegetation samples (Figure 7) when moving from left to right. We can speculate that an increase in salinity decreases the total amount of water in the plant. To further investigate this point, we calculated the total area between the shoulders at 1840 and 2235 nm (of the absorption peak of water at 1940 nm) (Figure 2). The results were uploaded into the score plot of the slope method (Figure 6) and are presented in Figure 7.

![Figure 6](image1.png)

**Figure 6.** Score plot for partial least squares (PLS) of the Cl model using the slope method. PC, principal component.

![Figure 7](image2.png)

**Figure 7.** Uploading the total absorption area into the score plot of partial least squares (PLS) of the Cl model using the slope method. Numbers next to gray circles represent the total area of water absorption.
In this regard, we can speculate that although the wide water absorption masks the spectral features of the plant, this absorption is still diagnostic for salinity, probably due to the internal connection between water and salinity.

Generally, spectral ranges that were found informative using field measurements were also found informative when using satellite-based Sentinel-2 data, but using much coarser spectral resolution. Currently, there is only one available hyperspectral satellite (HYPERION) with few more planned missions (EnMap, expected launch in 2018, and HyspIRI in 2022), which will be able to generate calculations of narrow bands at high spectral resolution. However, in contrast to hyperspectral imagery, multispectral data is easier to pre-process [55]. These include, for example, less complicated atmospheric correction algorithms, as well as obtaining high temporal coverage data (by means of orbital sensors), therefore, using relatively coarse spectral ranges and only four spectral bands of Sentinel-2 to estimate salinity is already encouraging result. Furthermore, although samples were collected in 2008–2011, whereas the Sentinel-2 image was available from 2015, we expect that similar trends will be derived in future studies that combine in situ and satellite-based analyses.

Previous studies have also shown that by using spectral indices, it is possible to determine salinity levels on a pixel base using coarse spectral resolution imagery (Landsat and IKONOS) [56,57]. These studies reported coefficients of determination between salinity and spectral indices ranging from $R^2 = 0.23$ to $R^2 = 0.78$. In this regard, our results showed that Sentinel-2, with its improved spectral and spatial resolution, has a potential for more accurate mapping of vegetation salinity status ($R^2 = 0.488$). Although the coefficient of determination might be interpreted as somewhat low, the relationship between both parameters (e.g., salinity vs. slopes) was found to be significant ($p$ value < 0.05). Furthermore, we also calculated the confidence interval for a given confidence level (95%) to achieve the needed accuracy. For example, to obtain 50% accuracy, we need a sample size of 103 spectral measurements, whereas to obtain 90% accuracy, 286 measurements are needed. Taking into account all of the above, we can argue that in order to achieve a higher accuracy of salinity estimation, future studies should be expanded and include the following steps: (1) using the dataset with the much wider range of chemical reference values and vegetation types; (2) larger calibration and validation dataset samples; and (3) incorporating multiple slopes into the analyses (similarly to Lugassi et al. [35]).

Our results indicate that by using either field spectral measurements or satellite imagery, many spectral indices indicative of chemical composition can be generated and matched to Sentinel-2 or any other satellite system. Indeed, both field and satellite measurements indicated similar estimations of salinity when using SVSI index (field $R^2 = 0.486$, satellite-based $R^2 = 0.488$). In this regard we can conclude that vegetation indices can be customized for generating salinity values, as well as for any other particulate application.

5. Conclusions

We hypothesized that a generic rule can be formulated for estimating salinity in different types of vegetation. The spectral regions that are most sensitive to changes in salinity were identified using both field and satellite (Sentinel-2) observations. Importantly, both cotton and tomato leaves exhibited similar spectral variability across similar spectral ranges as a function of changes in salinity levels. Therefore, both populations were combined in the analyses. Although salts lack spectral fingerprints across the VIS–NIR–SWIR region, by using spectral slope calculations and PLS regression analyses, we were able to quantitatively assess, with relatively good estimation ($R^2 = 0.77$ for Cl and $R^2 = 0.63$ for Na), the salinity of tomato and cotton plants using field measurements.

Next, the results of our field measurements were expanded to satellite-based observations. We used similar analyses of spectral slopes run on different bands, and different spectral indices were generated. The best estimate of vegetation salinity when using Sentinel-2-imagery ($R^2 = 0.488$) takes into account the following spectral ranges: 490–665 nm and 705–1610 nm. This index was denoted as the Sentinel-2-based vegetation salinity index (SVSI) (band 4 – band 2)/(band 5 + band 11).

In practice, more samples are needed to produce a more robust model. Nevertheless, our results suggest that vegetation reflectance spectroscopy is potentially useful for predicting plant salinity. Since the method is non-destructive and can measure several constituents quickly, inexpensively,
accurately, and simultaneously, it could become a valuable tool for the quality control of processed salinity in plants on site and, hence, enable monitoring of the salinity status of plants throughout their growth. The potential of using Sentinel-2 data is encouraging due to its high spatial resolution of 20 m, which enables high-resolution salinity mapping.

Future study is no doubt required to estimate the salinity of different types of vegetation over areas with a salinity problem in Israel using Sentinel-2 data.

Acknowledgments: This research was supported by the Ministry of Science Technology & Space, Israel (grant 3-13140). Additional funding was provided by the Samaria and Jordan Rift Regional R&D Center, Israel. Authors greatly acknowledge three anonymous reviewers for their important and significant comments.

Author Contributions: Rachel Lugassi analyzed the data. Rachel Lugassi, Alexandra Chudnovsky performed the statistical analysis and modeling. All authors were involved in paper writing. Alexandra Chudnovsky and Naftaly Goldshleger, designed, performed and supported the experiments and project PI’s.

Conflicts of Interest: The authors declare no conflict of interest.

References


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