

1 **Detection of *Karenia brevis* red tides on the West Florida Shelf using VIIRS observations:**  
2 **Accounting for spatial coherence with artificial intelligence**

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11

12 **Abstract**

13 Harmful algal blooms (HABs) of the toxic dinoflagellate *Karenia brevis* occur annually on the  
14 West Florida Shelf (WFS). Detection of these blooms using satellite observations often suffers  
15 from two problems: lack of accurate algorithms to identify phytoplankton blooms in optically  
16 complex waters and patchiness (i.e., heterogeneity) of *K. brevis* during blooms. Here, using data  
17 collected by the Visible Infrared Imaging Radiometer Suite (VIIRS) on the Suomi National  
18 Polar-orbiting Partnership (SNPP) between 2017 and 2019, we develop a practical approach to  
19 overcome these difficulties despite the lack of a chlorophyll-a fluorescence band on VIIRS. The  
20 approach is based on artificial intelligence (specifically, a deep-learning convolutional neural  
21 network model), which uses spatial coherence of bloom patches to account for the patchiness of  
22 *K. brevis* concentrations. After proper training, the overall performance (i.e., F1 score) of the  
23 deep learning model is 89%. Extracted *K. brevis* patches were consistent with those derived from  
24 the Moderate Resolution Imaging Spectroradiometer (MODIS) on the Aqua satellite that has a  
25 fluorescence band. Furthermore, the wider swath of VIIRS over MODIS (3040-km *versus* 2330-  
26 km) led to more valid observations of bloom extent for improved near-real-time applications.  
27 The results not only demonstrate the capacity of VIIRS in HABs monitoring, but also show the  
28 value of the DL model in extracting *K. brevis* bloom patches for both near real-time applications  
29 and retrospective analysis.

30 **Keywords:** *Karenia brevis*, red tide, bloom detection, deep learning, remote sensing, VIIRS,  
31 MODIS

32 **1. Introduction**

33 Harmful algal blooms (HABs) are a global phenomenon that can negatively impact coastal  
34 ecosystems, economies, and human and wildlife health. Increases in HAB occurrences have been  
35 linked to eutrophication and climate change (Anderson et al., 2021; Fu et al., 2012; Glibert et al.,  
36 2014; Glibert & Burford, 2017). The primary HAB-forming species on the West Florida Shelf  
37 (WFS) is *Karenia brevis*, a toxic dinoflagellate that causes fish, seabird, and marine mammal  
38 mortalities and poses hazards to human and wildlife health (Fleming et al., 2007, 2011;  
39 Flewelling et al., 2005; Kirkpatrick et al., 2004; Steidinger, 2009). *K. brevis* blooms occur near-  
40 annually on the WFS, typically in late summer and fall, although particularly severe blooms have  
41 been reported year-round (e.g., 2005–2007, 2017–2019, and 2020–2021). The spatial scale of *K.*  
42 *brevis* blooms on the WFS also varies considerably from event to event and even over much  
43 shorter (i.e. daily) time scales. Areas with high concentrations of *K. brevis* are often called ‘red  
44 tides’, though these waters can appear in various shades of red, green, brown, or black.

45 The spatial and temporal variability of *K. brevis* blooms and associated impacts on the WFS  
46 require extensive monitoring to inform communication and forecasting. Water samples are  
47 routinely collected during field sampling of WFS coastal waters by the Florida Fish and Wildlife  
48 Conservation Commission (FWC) and dedicated research groups and volunteer networks,  
49 regardless of bloom conditions; additional sampling is also conducted in response to bloom  
50 events. Each year, thousands of samples are enumerated using microscopy to detect and monitor  
51 *K. brevis* and other HABs in Florida’s (U.S.A.) marine and estuarine waters. This information is  
52 compiled within the FWC HAB Monitoring Database and reported by the FWC via regular  
53 updates on HABs. Background *K. brevis* concentrations (< 1,000 cells L<sup>-1</sup>) are often observed in  
54 the non-bloom season (Heil & Steidinger, 2009; Steidinger, 2009). When *K. brevis* cell counts  
55 exceed 1,000 cells L<sup>-1</sup>, commercial shellfish harvesting areas may be closed due to potential  
56 hazards posed by the toxins. The lower limit for satellite detection is 50,000 cells L<sup>-1</sup>, which is  
57 1–2 orders of magnitude less than concentrations at which blooms are visible by the human eye  
58 (Tester & Steidinger, 1997). When cell counts are above 100,000 cells L<sup>-1</sup>, reports of fish

59 mortality and human respiration problems increase (Fleming et al., 2011; Flewelling et al., 2005;  
60 Kirkpatrick et al., 2004).

61 Over the past sixty years, more than a million water samples have been collected on the WFS  
62 and processed for *K. brevis* enumeration by FWC (FWC HAB Monitoring Database, 2021).  
63 However, the majority of samples were collected in nearshore waters, less than 10 km from the  
64 shoreline. This equates to fewer than one sample collected every two months within each 0.05°  
65 grid, although sampling intensity (including offshore) has increased substantially over time (Hu  
66 et al., 2022). Satellite remote sensing can help overcome the bias introduced by the scarcity of  
67 field data (especially offshore) because it provides increased synoptic spatial and temporal  
68 coverage (Amin, Zhou, et al., 2009; Amin et al., 2015; Esaias et al., 1998; Tester & Stumpf,  
69 1998; Tomlinson et al., 2004). At elevated concentrations of *K. brevis*, water discoloration is  
70 captured in satellite images, and such a discoloration is often interpreted as an indication of *K.*  
71 *brevis* blooms (Cannizzaro et al., 2008, 2009; Cullen et al., 1997; Schofield et al., 1999; Tyler &  
72 Stumpf, 1989). Other factors, such as suspended sediments, colored dissolved organic matter  
73 (CDOM), and non-*K. brevis* phytoplankton blooms, can also cause water discoloration and be  
74 misconstrued as red tide (Dierssen et al., 2006). Blooms of *K. brevis* and other species may also  
75 occur at levels that can be observed via satellites but without discoloration observed with the  
76 naked eye. Thus, detection of *K. brevis* blooms using satellite observations requires algorithms,  
77 which are often empirical, to first detect phytoplankton blooms (Amin, Zhou, et al., 2009;  
78 Carvalho et al., 2010, 2011; El-Habashi et al., 2016; Hu & Feng, 2016; Qi et al., 2015; Stumpf et  
79 al., 2003; Tomlinson et al., 2009) and then distinguish the bloom types (e.g., *K. brevis*, diatom,  
80 *Pyrodinium bahamense*, *Tripos hircus*).

81 Satellite sensors equipped with spectral bands to measure solar-stimulated chlorophyll-a  
82 fluorescence, including the Moderate Resolution Imaging Spectroradiometer (MODIS) aboard  
83 Terra (2000–) and Aqua (2002–) and the Ocean and Land Colour Imager (OLCI) aboard  
84 Sentinel-3A (2016–) and Sentinel-3B (2018–), are used for bloom monitoring by U.S. Federal  
85 and State agencies (e.g., the U.S. National Oceanic and Atmospheric Administration (NOAA),  
86 <https://coastalscience.noaa.gov/science-areas/habs/hab-forecasts/gulf-of-mexico/>; FWC,  
87 <https://myfwc.com/research/redtide/statewide/>). Empirical algorithms developed using  
88 concurrent field and satellite data often rely on satellite-field matching pairs (i.e., data collected  
89 from the same location within a short time window), and thus can be regarded as ‘pixel-based’

90 approaches. These include techniques of the chlorophyll-a anomaly (Stumpf, 2001; Stumpf et al.,  
91 2003; Wang et al., 2021), particle backscattering coefficient  $b_{bp}$  ratio (Anderson et al., 2012;  
92 Cannizzaro et al., 2008; Carder et al., 1999; Morel, 1988), normalized water-leaving radiance  
93 ( $nL_w$ ) ratio (Carvalho et al., 2011),  $nL_w$  spectral shape (Tomlinson et al., 2009), normalized  
94 Fluorescence Line Height (nFLH; Hu & Feng, 2016), Red Solar Induced Fluorescence (red SIF,  
95 Luis et al., 2023) and Red-Band Difference (RBD; Amin, Gilerson, et al., 2009; Amin, Zhou, et  
96 al., 2009). However, because *K. brevis* cell concentrations from adjacent waters (i.e., within a  
97 single pixel) can differ by several orders of magnitude (Tomlinson et al., 2009, also see  
98 supplemental Fig. S1), significant data spread is commonly observed when comparing satellite-  
99 estimated *K. brevis* concentration with water sample-determined *K. brevis* concentration (Hu &  
100 Feng, 2016).

101 The Visible Infrared Imaging Radiometer Suite (VIIRS) onboard the Suomi National Polar-  
102 orbiting Partnership (SNPP; 2011–), NOAA-20 (2017–), and NOAA-21 (2022–) exhibits similar  
103 spatial (375/750-m) and temporal (near-daily) resolutions as MODIS (250/500/1000-m) and  
104 OLCI (300-m), but VIIRS does not have the dedicated bands for measuring chlorophyll-a  
105 fluorescence for bloom detection (Hu et al., 2015). Despite this limitation, VIIRS has shown  
106 potential for bloom detection using alternative approaches, including the Red-Green  
107 Chlorophyll-a Index (RGCI; Qi et al., 2015) and neural network models (El-Habashi et al., 2016;  
108 El-Habashi & Ahmed, 2019). While bloom detection may be similar to nFLH and RBD, these  
109 algorithms were developed using pixel-based approaches without the ability to differentiate  
110 between *K. brevis* blooms and other types of blooms. Evaluation of these approaches using a  
111 large field dataset suggested that, although these approaches may work well for case studies, they  
112 are not applicable in a more general sense (supplemental Figs. S2 & S3).

113 Deep learning models are poised to overcome the limitations of pixel-based approaches by  
114 adding an emphasis on the recognition of spatial patterns. Also, unlike empirical approaches that  
115 typically utilize spectral information from only a few wavebands, these models can take  
116 advantage of the spectral information from all bands. Previous studies indicate the potential of  
117 using such approaches to identify HABs of *Magalefidinium polykrikoides* in Korean waters (Kim  
118 et al., 2019; Shin et al., 2022). However, for the *K. brevis* blooms on the WFS, there is no  
119 method for systematic observation using VIIRS data without a fluorescence band. With MODIS  
120 nearing the end of its lifespan (i.e., NASA will cease support of MODIS no later than 2026), the

121 development of robust bloom detection algorithms for VIIRS sensors, scheduled for launch  
122 every five years, is critical to ensure a seamless satellite ocean color data record for monitoring  
123 HABs in near-real-time and documenting long-term trends. In order to address this need, the  
124 objective of this paper is to develop a practical approach to take advantage of computer artificial  
125 intelligence to recognize spatially coherent ocean color patterns in VIIRS imagery associated  
126 with *K. brevis* blooms.

127

128 **2. Data and methods**

129 **2.1 In-situ data**

130 Water samples from near-surface waters were used to determine the in-situ concentration of *K.*  
131 *brevis* at sampling sites. A total of 23,232 field data points for *K. brevis* cell counts were  
132 recorded at sample depths of ~ 0.5 m on the WFS from January 2017 to December 2019 (FWC  
133 HAB Monitoring Database, 2021). Fig. 1a shows the number of field *K. brevis* cell counts data in  
134 each 5-km grid. In this study, *K. brevis* cell counts larger than 100,000 cells L<sup>-1</sup> are considered *K.*  
135 *brevis* blooms.

136 **2.2 VIIRS satellite data**

137 A total of 4,282 level-2 SNPP VIIRS granules covering the WFS from January 2017 to  
138 December 2019 were downloaded from the NOAA CoastWatch data portal  
139 (<https://coastwatch.noaa.gov>). These level-2 products included  $nL_w(\lambda)$  for each band (410, 443,  
140 486, 551, 638, and 671 nm) and quality assurance flag information. Default L3 flags developed  
141 by NOAA (Wang et al., 2017) were applied for quality control to exclude pixels with unreliable  
142 radiance values from further analysis. Fig. 1b shows the increased spatial and temporal coverage  
143 of VIIRS data compared to the *in-situ* cell count data.

144 Remote sensing reflectance ( $R_{rs}(\lambda)$ ) for each band was determined from  $nL_w(\lambda)$  as follows:

145 
$$R_{rs}(\lambda) = nL_w(\lambda)/f_0(\lambda), \quad (1)$$

146 where  $f_0(\lambda)$  is the mean extraterrestrial solar irradiance (Thuillier et al., 2003).

147 A cylindrical equidistant projection was used to map these data within the WFS at 1-km spatial  
148 resolution (Barnes et al., 2021).  $R_{rs}(\lambda)$  data at 671-nm and 551-nm were used to generate the  
149 RGCI (Qi et al., 2015) following the equation:

150 
$$RGCI = R_{rs}(671)/R_{rs}(551), \quad (2)$$

151 In addition,  $R_{rs}(\lambda)$  at 551, 486, and 443 nm were used to generate Enhanced Red-Green-Blue  
152 (ERGB) composite images to show color patterns of coastal waters. The ERGB images  
153 differentiate dark features, caused by high absorption by chlorophyll-a and/or colored dissolved  
154 organic matter (CDOM), from bright features caused by either sediment resuspension or shallow  
155 bottom (Hu et al., 2005).

156 2.3 Deep learning model

157 Deep learning (DL) is a type of artificial intelligence that uses artificial neural networks with  
158 multiple layers to learn from data and make predictions. A Convolution Neural Network (CNN;  
159 Lecun et al., 1998) is a form of deep learning that is widely used in image segmentation for  
160 clustering parts of imagery together that belong to the same object class. Here, a type of CNN  
161 architecture called Res-Unet deep learning model (Diakogiannis et al., 2020; Qi et al., 2021;  
162 Wang & Hu, 2021; Xiao et al., 2018; Yao et al., 2023) is used. This model combines  
163 constructions inherent to both Res-Net (He et al., 2016) and U-net (Ronneberger et al., 2015)  
164 models, thus improving ability to effectively perform image segmentation tasks.

165 The workflow in this study follows three main steps (Fig. 2). First, a set of “ground truth” images  
166 were prepared semi-objectively and combined with satellite  $R_{rs}(\lambda)$  and RGCI for model training.  
167 Here, the term “ground truth” refers to the information determined by integrating ground (i.e.,  
168 field) measurements and image analysis results as opposed to either ground measurements alone  
169 or the theoretical “truth”. The trained model was then validated using a separate set of “ground  
170 truth” images that were reserved for evaluation. Finally, the model was applied to VIIRS data  
171 from 2017–2019, and the model output was used to generate monthly statistics to examine  
172 spatiotemporal variability of *K. brevis* blooms over the course of the bloom event.

173 2.3.1 “Ground truth” image preparation

174 *K. brevis* blooms exhibit high RGCI (Qi et al., 2015) and appear reddish-black in ERGB  
175 composite imagery (Hu et al., 2005), allowing these patches to be differentiated from the

176 surrounding waters. However, not all patches with high RGCI are *K. brevis* blooms because  
177 blooms of other phytoplankton can also lead to high RGCI values, and other factors (e.g.,  
178 CDOM) can cause ERGB images to appear reddish black. Here, field sample data are used to  
179 confirm that patches with high RGCI are *K. brevis* blooms. The patches were identified as *K.*  
180 *brevis* only if the field data showed high *K. brevis* cell counts ( $> 100,000$  cells  $L^{-1}$ ) that  
181 corresponded to high RGCI and reddish-blackish features in ERGB. This practice has been  
182 employed before to delineate *K. brevis* blooms using field sample data and MODIS/Aqua RBD  
183 images (Hu et al., 2022).

184 Based on the criteria above, “ground truth” images were prepared as demonstrated in Fig. 3  
185 using the following steps:

- 186 1. *K. brevis* cell counts data ( $\pm 7$  days) were overlaid on VIIRS daily (i.e., snapshot) RGCI  
187 and ERGB composite imagery (Figs. 3a & 3b).
- 188 2. Patches associated with *K. brevis* cell counts  $> 100,000$  cells  $L^{-1}$  that exhibit high RGCI  
189 and appear reddish black in the ERGB imagery were roughly outlined manually using the  
190 ENVI/IDL region of interest (ROI) tool (Fig. 3b).
- 191 3. Pixels within the outline with  $RGCI \geq$  the threshold of RGCI ( $T_{RGCI}$ ) (e.g., *K. brevis*  
192 bloom) were considered as bloom pixels and extracted objectively (Fig. 3c).  $T_{RGCI}$  was  
193 set as 0.22, which corresponds to a chlorophyll-a concentration of  $1.5 \mu\text{g L}^{-1}$  and  
194 approximately  $150,000$  *K. brevis* cells  $L^{-1}$  (Qi et al., 2015; Stumpf et al., 2003; Tester et  
195 al., 2008). This threshold is consistent with that used for the MODIS RBD by Hu et al.  
196 (2022).
- 197 4. The bloom pixels were assigned a value of 1. All remaining pixels, including those  
198 outside the outline or with  $RGCI < T_{RGCI}$ , were assigned a value of 0 (e.g., non-*K. brevis*  
199 bloom) or NaN (not a number, due to no observation or invalid pixels) (Fig. 3d).

200 A total of 100 VIIRS images were delineated following the above steps. Twenty three of these  
201 images contained high VIIRS RGCI with *K. brevis* cell counts equal to zero and are intended to  
202 help prevent false positives. Through random selection, 47 of these images were designated for  
203 training, and the remaining 53 images were reserved for validation. Here, although the cell  
204 counts data were likely collected not in the same day of the image acquisition and water could  
205 have moved within  $\pm 7$  days to cause a mismatch between the locations of the *in situ* data and  
206 image feature, as long as there were high cell counts within or near an image feature, the feature

207 is delineated as a *K. brevis* bloom patch. This is also one reason why a patch-wise approach  
208 should work better than a pixel-wise approach.

209 *2.3.2 Model training*

210 A total of 47 “ground truth” images and their corresponding  $R_{rs}(\lambda)$  and RGCI data were used as a  
211 training dataset for developing and training the DL model. To balance the weight of the input  
212 data and make the deep learning model training converge smoothly, each  $R_{rs}(\lambda)$  band (410, 443,  
213 486, 551, 638, 671 nm) was normalized by:

214 
$$nR_{rs}(\lambda) = (\log(R_{rs}(\lambda)) - \log(R_{rs}(\lambda)_{min})) / (\log(R_{rs}(\lambda)_{max}) - \log(R_{rs}(\lambda)_{min})), \quad (3)$$

215 where  $R_{rs}(\lambda)_{min}$  and  $R_{rs}(\lambda)_{max}$  were determined to be 0.0001 and 0.02 by trial and error,  
216 respectively. If  $R_{rs}(\lambda)$  was less than 0.0001, it was set to 0.0001; and if  $R_{rs}(\lambda)$  was great than  
217 0.02, it was set to 0.02.

218 Likewise, RGCI was normalized as follows:

219 
$$nRGCI = (RGCI - RGCI_{min}) / (RGCI_{max} - RGCI_{min}), \quad (4)$$

220 where  $RGCI_{min}$  and  $RGCI_{max}$  were determined to be 0.1 and 1.5 by trial and error, respectively.  
221 If RGCI was less than 0.1, it was set to 0.1; and if RGCI was greater than 1.5, it was set to 1.5.

222 Squared convolution kernels were applied in this training network. Thus, each input training  
223 image was divided into several spatially non-overlapping sub-images of  $256 \times 256$  pixels, with  
224 the sub-image size determined by computing power. Each sub-image was then used to train the  
225 DL model, and the Jaccard distance index was used to assess the model convergence. After  
226 passing through the deep convolutional layers, the model can recognize the characteristic  $R_{rs}(\lambda)$   
227 spectral shapes of *K. brevis* bloom patch and the coherent spatial relationships among the  $R_{rs}(\lambda)$   
228 spectral features that help identify the *K. brevis* bloom patches.

229 *2.3.3 Model validation*

230 A total of 53 delineated VIIRS “ground truth” images were reserved for validation to evaluate  
231 the model performance. The morphology of each patch in the model extracted images was  
232 visually compared with the semi-objectively delineated patches of the “ground truth” images and  
233 field *K. brevis* cell concentration data to determine whether the model extracted results matched  
234 those from the “ground truth” images. A confusion matrix (Stehman, 1997) was used to report

235 the number of true-positives (TP), true-negatives (TN), false-positives (FP), and false-negatives  
236 (FN), as well as the F1 score to evaluate the overall accuracy. The F1 score was calculated as:

237 
$$F1 = 2TP / (2TP + FP + FN) \times 100\%. \quad (5)$$

238 The above statistics is based on the evaluation of the 53 image pairs, each containing a “ground  
239 truth” image (or truth image) and an image of model results (or model image). There are 41 *K.*  
240 *brevis* bloom truth images and 12 non-bloom truth images. Each model image was compared to  
241 its corresponding truth image to determine whether the model image is a TP, TN, FP, or FN. A  
242 model image is a TP if 1) the morphology of each bloom patch in the model image matches that  
243 in the truth image and 2) the overlapping bloom area (as measured by the number of pixels)  
244 between the model image and the truth image is > 75% of the bloom area in the truth image,  
245 otherwise the model image is an FN. A model image is an FP if any patch is classified as a  
246 bloom patch, but the corresponding truth image shows no bloom, otherwise the model image is  
247 an TN.

248 In addition to the confusion matrix, the bloom areas of all image pairs (i.e., the truth images and  
249 the model images) were compared using a linear fitting with the coefficient of determination ( $R^2$ )  
250 and the root mean square error (RMSE). RMSE was calculated as follows:

251 
$$RMSE = \sqrt{\frac{1}{n} \sum_{i=1}^n \left( \frac{x_i - \hat{x}_i}{\hat{x}_i} \right)^2} \times 100\%, \quad (6)$$

252 where  $x_i$  is the bloom area (i.e., number of pixels) of the model extracted result of the  $i$ -th model  
253 image, and  $\hat{x}_i$  is the area of the  $i$ -th truth image.

254 Spectral similarities between  $R_{rs}(\lambda)$  measured in *K. brevis* bloom and non-*K. brevis* bloom  
255 patches were examined using the Spectral Angle Mapper (SAM) index (Kruse et al., 1993). The  
256 SAM (in degrees) indicates the spectral similarity between two  $R_{rs}(\lambda)$  spectra by calculating the  
257 angle between them. The closer SAM is to 0-degree, the greater the similarity between the two  
258 spectral shapes.

259 *2.3.4 Statistics of K. brevis coverage*

260 After the model was trained and validated, it was applied to a 3-year series of VIIRS data (2017–  
261 2019) that encompassed a long-lasting *K. brevis* bloom event. Each pixel was classified into one  
262 of three classes: ‘*K. brevis* bloom’ with a value of 1, ‘non-*K. brevis* bloom’ with a value of 0, or

263 ‘no valid observation’. The ‘no valid observation’ class was determined using the default L3  
264 quality assurance flag information processed by NOAA (Wang et al., 2017), and these pixels  
265 were excluded from the following statistics.

266 Monthly maps of bloom occurrence frequency (BOF) were generated from the individual  
267 classified images and for a given location were calculated as follows:

268 
$$BOF = (N_{kb}/(N_{kb} + N_{nkb})) \times 100\%, \quad (7)$$

269 where  $N_{kb}$  is the total number of ‘*K. brevis* bloom’ pixels and  $N_{nkb}$  is the total number of ‘non-*K.*  
270 *brevis* bloom’ pixels. To make the statistics more meaningful, pixels with fewer than five valid  
271 observations (see definition above) in any given month were excluded. Bloom areal footprints  
272 were then calculated from all > 0% pixels in the monthly BOF maps.

273 Monthly BOF maps were also generated from individual classified maps derived based on RGCI >  
274 TRGCI and chlorophyll-a > 1.5  $\mu\text{g L}^{-1}$  derived using a neural network model (El-Habashi et al.,  
275 2016) to compare with model results.

276 **2.4 MODIS-based *K. brevis* bloom occurrence frequency maps**

277 MODIS/Aqua data were used to qualitatively assess the VIIRS deep learning model results.  
278 Specifically, *K. brevis* blooms were classified following the work of Hu et al. (2022) by  
279 integrating water sample data and MODIS data. Briefly, field *K. brevis* cell counts were first  
280 overlaid on MODIS RBD (Amin et al., 2015; Amin, Zhou, et al., 2009). Patches with high RBD  
281 and high field *K. brevis* cell concentrations were delineated semi-objectively: a crude outline was  
282 manually drawn over each patch, and pixels within the outline with RBD > 0.15  $\text{mW cm}^{-2} \mu\text{m}^{-1}$   
283  $\text{sr}^{-1}$  (which corresponds to 150,000 cells  $\text{L}^{-1}$ , Hu & Feng, 2016) were identified as *K. brevis*  
284 bloom pixels. Monthly BOF maps were generated from the daily imagery similarly to VIIRS  
285 (Section 2.3.4).

286

287 **3. Results**

288 **3.1 Model validation**

289 A confusion matrix for assessing the performance of the DL model is shown in Table 1. The  
290 overall F1 score was 89% with an accuracy of 81% and precision of 83%. Four sets of example

291 images from the validation dataset are presented in Fig. 4, showcasing TP, TN, FP, and FN  
 292 results. The first set of images in the top row displays the TP results where the DL model  
 293 successfully extracted *K. brevis* patches that match the “ground truth” image. Here, each TP case  
 294 meets the criteria of both morphological evaluation and 75% threshold of bloom area as  
 295 described in the methodology above. Of the 39 TP cases, the ratio of the overlapping bloom area  
 296 to the true bloom area for each image pair ranged between 77.7% and 96.3%, with an overall  
 297 ratio of 82.2% when all image pairs were combined. The second set of images illustrates the TN  
 298 results where the DL model correctly identified non-*K. brevis* bloom (i.e., not a single patch was  
 299 a bloom patch in both the “ground truth” image and the model image). The third set of images  
 300 presents the FP results where the DL model extracted incorrect bloom patches, and the fourth set  
 301 of images displays the FN results where the DL model failed to identify > 25% of the  
 302 overlapping bloom areas between the “ground truth” image and the model image.

303 Table 1. Performance evaluation of the DL model where P and PP are the number of “true” and predicted *K. brevis*  
 304 blooms, respectively; N and PN are the number of “true” and predicted non-*K. brevis* bloom, respectively. TN and  
 305 TP are the number of true negatives and positives, respectively; FN and FP are the number of false negatives and  
 306 positives, respectively.

Total n = 53		Predicted		F1 score	2TP/(2TP+FP+FN)	88.6%
		PP	PN			
“Ground truth”	P	TP 39	FN 2	True positive rate (TPR)	TP/(TP+FN)	95.1%
	N	FP 8	TN 4	False positive rate (FPR)	FP/(FP+TN)	66.7%
Precision		TP/(TP+FP)	83.0%	Accuracy	(TP+TN)/n	81.1%

307  
 308 A comparison between bloom areas (in number of pixels) determined from the “ground truth”  
 309 images and the corresponding model images is presented in Fig. 5, wherein the overall RMSE  
 310 was found to be 31.5%, and the coefficient of determination  $R^2$  was calculated to be 0.92. Bloom  
 311 areal extent was underestimated by the model in the Panhandle region in late-2018. These results  
 312 will be discussed in detail below.

313 3.2 Model performance

314 The DL model, trained and verified based on the VIIRS spectral information and image coherent  
315 context, can identify the *K. brevis* patches and distinguish them from non-*K. brevis* bloom  
316 patches.

317 Fig. 6 shows a VIIRS scene collected on 16 September 2018 that contains two separate patches  
318 of high RGCI water that appear darkish red in the ERGB imagery. *K. brevis* cell counts  $>$   
319 100,000 cells L $^{-1}$  confirm that the northern patch offshore of Charlotte Harbor was a true bloom,  
320 and the model correctly identified this bloom patch. Background cell counts ( $<1,000$  cells L $^{-1}$ )  
321 were collected in the more southerly patch located south of Cape Romano ( $\sim 26^{\circ}$ N), and the  
322 model correctly identified this patch as a non-*K. brevis* bloom. VIIRS  $R_{rs}(\lambda)$  spectral shapes were  
323 examined in Fig. 6a at locations within these patches. The high similarity in spectral shape (SAM  
324 = 4.94 $^{\circ}$ ) indicates that both patches would be categorized as *K. brevis* blooms based on RGCI  
325 alone. The DL model accurately differentiated between the *K. brevis* bloom patch and the non-*K.*  
326 *brevis* bloom patches.

327 Fig. 7 further demonstrates how the VIIRS DL model generates fewer false-positive  
328 classifications compared to both RGCI (Qi et al., 2015) and the neural network model (El-  
329 Habashi et al., 2016). Monthly BOF maps using all three techniques were generated during a *K.*  
330 *brevis* bloom event (August 2018) and non-*K. brevis* bloom event (June 2019), and are compared  
331 to monthly FWC cell abundance data. While the neural network and RGCI retrieval results  
332 accurately detect the *K. brevis* bloom in the central WFS in August 2018, there are some false-  
333 positive results in nearshore waters to the north in the Panhandle/Big Bend regions and south of  
334 Cape Romano ( $\sim 26^{\circ}$ N). False positive classifications were also prevalent in these regions during  
335 the non-*K. brevis* bloom event in June 2019. The VIIRS DL model, on the other hand, shows  
336 strong consistency with *K. brevis* cell abundance, indicating improved performance in accurately  
337 identifying both *K. brevis* blooms and non-blooms.

338 3.3 Comparisons between VIIRS and MODIS

339 Monthly MODIS BOF maps generated by semi-objective delineation for May 2018 to January  
340 2019 were previously presented by Hu et al. (2022). In Fig. 8, comparisons are made between  
341 bloom footprints generated from these maps and those derived using the VIIRS DL model. *K.*

342 *brevis* blooms were detected by both methods from June 2018 to January 2019, but several  
343 differences were observed in the footprint areas. The VIIRS DL model often estimated *K. brevis*  
344 blooms extending further into shallow coastal waters than the MODIS BOF, resulting in larger  
345 footprint areal estimates than those observed with MODIS, except for cases during October and  
346 November 2018 when MODIS derived BOF footprints were larger than those from VIIRS in the  
347 Panhandle and Big Bend regions.

348 Fig. 9 provides a detailed visualization of the differences in bloom footprint observed between  
349 VIIRS and MODIS for imagery acquired approximately one hour apart on October 30, 2018. The  
350 VIIRS bloom footprint was 44% lower than that from MODIS. While MODIS RBD was well  
351 above the threshold used by Hu et al. (2022) for identifying bloom patches, VIIRS RGCI was  
352 close to the bloom threshold used when training the model. Residual increased suspended  
353 sediment following the recent passage of a winter frontal system is evident in the VIIRS ERGB  
354 and may explain why VIIRS failed to detect this patch. VIIRS and MODIS  $R_s(\lambda)$  spectra  
355 extracted from within the bloom patch are similar (SAM = 9.07°).

356

#### 357 **4. Discussion**

##### 358 4.1. Strengths and limitations

359 *K. brevis* blooms on the WFS pose threats to coastal ecosystems and public health and can  
360 negatively impact local economies. An accurate means for near-real-time monitoring is required  
361 to help protect public health, and long-term monitoring is needed to better understand the  
362 underlying causes of blooms and identify bloom trends. Field measurements of *K. brevis* cell  
363 counts are highly precise; however, their limited spatial and temporal resolutions restrict their  
364 overall efficacy in consistently monitoring blooms with accuracy. Remote sensing may serve as a  
365 valuable tool to complement field-based monitoring programs. However, previous remote  
366 sensing algorithms often rely on pixel-based approaches with pre-determined thresholds applied  
367 to identify blooms for each pixel with an image (Amin, Zhou, et al., 2009; Cannizzaro et al.,  
368 2008, 2009; Carvalho et al., 2010, 2011; Hu & Feng, 2016; Qi et al., 2015; Soto et al., 2015;  
369 Stumpf et al., 2003; Tomlinson et al., 2009). These had limited success due to the problems  
370 associated with sub-pixel variability (Hu & Feng, 2016; Fig. S1). While neural network models  
371 (El-Habashi et al., 2016; El-Habashi & Ahmed, 2019) offer several advantages over threshold-

372 based empirical approaches, systemic testing showed unsatisfactory performance (Figs. S2 & S3).  
373 Here, we developed a deep learning model for detecting *K. brevis* blooms on the WFS using  
374 VIIRS imagery that outperforms these other methods.

375 By adopting a patch-wise approach that considers spatial information (He et al., 2016), the  
376 VIIRS DL model can overcome limitations associated with subpixel variability that are inherent  
377 in pixel-wise approaches (Hu & Feng, 2016; Fig. S1). Furthermore, unlike empirical algorithms  
378 using a few bands, such as the RGCI (Qi et al., 2015) and RBD (Amin, Zhou, et al., 2009)  
379 algorithms, the VIIRS DL model utilizes all VIIRS bands as data input and relies on  $R_{rs}(\lambda)$   
380 spectral shapes for bloom identification. This spectral data from all bands can provide more  
381 comprehensive information compared to the limited utilization of just two or three bands in other  
382 empirical algorithms, therefore improving the accuracy of the deep learning model (Krizhevsky  
383 et al., 2017).

384 As an automated patch-wise approach, the VIIRS DL model reduces false positives and  
385 improves *K. brevis* bloom patch delineation, thus reducing the need for secondary verification by  
386 *in situ* data and/or human interpretation. In contrast, most traditional approaches first determine  
387 chlorophyll-a concentrations or a bloom patch (El-Habashi et al., 2016; Hu et al., 2005; Soto et  
388 al., 2015), and then use in-situ sampling and/or human interpretation to confirm whether the  
389 bloom patch is due to *K. brevis* or other phytoplankton. However, this does not indicate that the  
390 DL model does not require *in situ* data for verification, particularly because *K. brevis* is not the  
391 only dinoflagellate and blooms of other dinoflagellates may have similar optical properties to be  
392 detected by the same DL model. If this is the case, what the DL model detects are blooms of  
393 dinoflagellates. Yet because *K. brevis* is the dominant dinoflagellate to cause red tides, one can  
394 assume that most of the detected blooms are likely due to *K. brevis*.

395 Furthermore, VIIRS has a wider swath width (3040-km) compared with MODIS (2330-km),  
396 which means VIIRS has a greater number of observations to compare. Fig. 10 compares the  
397 monthly coverage and the number of valid observations for MODIS and VIIRS during the latter  
398 part of the 2017–2019 HABs bloom event (July 2018–December 2018). Although MODIS and  
399 VIIRS had similar *K. brevis* bloom trends, there were differences in the number of valid  
400 observation numbers. VIIRS had an average of ten or more valid observations per pixel per  
401 month in the WFS region, while MODIS had only around five. Under good observation

402 conditions (e.g., cloud-free and optimal solar/sensor zenith angles), as in October 2018, VIIRS  
403 could achieve more than 25 valid observations in the offshore area of Florida, while MODIS had  
404 only about 15.

405 Additionally, unlike MODIS that saturates its fluorescence band (678-nm) over moderate to high  
406 sun glint (Hu et al., 2012), VIIRS does not saturate under such conditions. Fig. 11 displays a case  
407 study where a 5-day period of VIIRS and MODIS observations showed the advantage of using  
408 VIIRS data for near-real-time monitoring. Due to the land adjacency effect, the saturation of the  
409 fluorescence band, and the narrower swatch (than VIIRS), MODIS had only three images  
410 showing scattered *K. brevis* patches during the 5-day period, with none of them capturing the full  
411 extent of the bloom. In contrast, VIIRS had at least one image per day in this 5-day period, with  
412 each of them showing near-complete bloom extent, which has extensive value toward guiding  
413 timely and targeted resource management and public health communications during *K. brevis*  
414 blooms.

415 Despite the advantages of using a deep learning model with VIIRS observation to detect *K.*  
416 *brevis* blooms, there are several limitations. One is the definition of “bloom”. Here, the dataset  
417 used to train the VIIRS DL model was prepared based on the RGCI threshold corresponding to  
418 150,000 cells L<sup>-1</sup> of *K. brevis* (Amin et al., 2015; Amin, Zhou, et al., 2009; Hu et al., 2022; Hu &  
419 Feng, 2016; Qi et al., 2015; Soto et al., 2015) if the phytoplankton population is dominated by *K.*  
420 *brevis*. Because of the significant bloom patchiness (Fig. S1) and because of mixed  
421 phytoplankton assemblage, this definition does not indicate that within a delineated bloom patch,  
422 *K. brevis* cell concentration is always > 150,000 cells L<sup>-1</sup>. As shown in Fig. 3, cell concentration  
423 within the bloom patch can be much lower than this threshold, and sometimes can be 0 – 1,000.  
424 This certainly does not mean that a *K. brevis* bloom patch with maximum concentration of  
425 ~10,000 cells L<sup>-1</sup> (or even ~5,000 cells L<sup>-1</sup>) can be detected by the DL model. What it means is  
426 that an image feature with maximum cell counts lower than this threshold is considered as “non-  
427 bloom” in the training and validation datasets. However, this threshold is higher than the  
428 threshold of 5,000 cells L<sup>-1</sup> when the commercial shellfish harvesting areas were previously  
429 required to be closed. It is also higher than the cell count threshold above which fish mortality  
430 and human respiration irritation often occur (Fleming et al., 2011; Flewelling et al., 2005;  
431 Kirkpatrick et al., 2004). Correspondingly, the *K. brevis* blooms detected here are rather

432 conservative, i.e., without including bloom patches (or other image features) with maximum  $K$ .  
433 *brevis* concentrations lower than this threshold, although these waters are also harmful to marine  
434 animals. Also, bloom detection is currently a binary classification that only allows for a  
435 distinction between the presence and absence of blooms without quantifying the intensity of  
436 them, although such a quantification may be possible when taking account of the RGCI values of  
437 the delineated bloom patches. Clearly, future efforts are required to detect blooms at lower  $K$ .  
438 *brevis* concentrations and to quantify the concentrations beyond presence/absence detections.  
439 This would improve the utility of the tool for tracking bloom transport, evolution, and  
440 appearance/disappearance.

441 The second limitation relates to the VIIRS DL model's applicability even under cloud free  
442 conditions. Similar to other satellite sensors, the DL model is not applicable to image pixels  
443 immediately adjacent to land because these pixels may be mixed pixels (between water and land)  
444 or contaminated by land adjacency effect. In this study, a 2-pixel buffer was applied immediately  
445 adjacent to land, effectively masking those areas. Estuaries were also masked to eliminate the  
446 influence of land adjacency effects. Furthermore, the strength of avoiding false-positive  
447 detection in sediment-rich waters (because of the use of the full  $R_{rs}(\lambda)$  spectral information  
448 together with spatial context) can become a weakness in some special cases. For example, if  
449 sediment resuspension, due to the passage of cold fronts or storms, occurs during a *K. brevis*  
450 bloom, the high concentrations of sediment particles can obscure the *K. brevis* signals, leading to  
451 no bloom detection. Fig. 12 shows such a case. The ERGB images in Fig. 12a and Fig. 12c,  
452 overlaid with field-measured *K. brevis* cell counts, reveal that in the Epicenter region, there was  
453 a persistent and expansive *K. brevis* bloom during mid-November 2018, and the bloom patches  
454 were correctly extracted by the VIIRS DL model (Figs. 12d and 12f). During this period and on  
455 16 November 2018, a cold front passed through the Epicenter region, causing high  
456 concentrations of resuspended sediment particles (bright features in Fig. 12b), which led to no  
457 bloom detection (Fig. 12e). However, such a false-negative detection can be easily remedied by  
458 inspecting sequential images: if similar bloom patches are detected in t1 and t3 but not in t2  
459 when sediment resuspension occurs, one can safely assume that similar bloom patches still exist  
460 in t2. Likewise, for near-real-time applications, if bloom patches are found in t1, the lack of  
461 detected bloom patches in t2 due to sediment resuspension does not indicate the end of the bloom.

462 Therefore, the lack of ability to detect *K. brevis* blooms in sediment-rich waters is much less of a  
463 problem than the false-positive detections in sediment-rich waters by other methods.

464

465 4.2. Future perspective

466 VIIRS measurements used in this study are from the SNPP satellite, yet the same sensor is also  
467 carried by the NOAA-20 (2017–present) and NOAA-21 (2022–present) satellites. Future  
468 satellites carrying the same VIIRS are expected to be launched about every 5 years. A  
469 combination of these sensors, each with a different equatorial crossing time, can provide multiple  
470 observations of *K. brevis* blooms in a single day. This will not only improve the cloud-free data  
471 coverage, but also may provide more than one observation per day to capture the diel vertical  
472 migration of *K. brevis* cells (Arnone et al., 2017; Hu, Barnes, et al., 2016; Qi et al., 2017;  
473 Schofield et al., 2006). Likewise, the multi-sensor observations can not only help to study the  
474 timing, intensity, and short-term dynamics of *K. brevis* blooms, but also improve near-real-time  
475 observations to alert the public on bloom situations (e.g., NOAA's HAB Forecast System, or the  
476 Integrated Redtide Information System (IRIS), Hu, Murch, et al., 2016;  
477 <https://optics.marine.usf.edu/projects/iris.html>). The same logic can be extended to other sensors  
478 such as the Ocean and Land Colour Instrument (OLCI) on the Sentinel-3A (2016–present) and  
479 Sentinel-3B (2018–present) satellites. While cross-sensor consistency is yet to be determined, the  
480 integration of these different satellite sensors can provide more comprehensive and accurate  
481 observations of *K. brevis* blooms than being offered by any single sensor, thus facilitating both  
482 research on bloom dynamics and near-real-time monitoring.

483 The findings here demonstrate the success of combining VIIRS observations and computer  
484 artificial intelligence to detect HABs, while near-real-time applications require implementation  
485 of this approach to generate *K. brevis* bloom maps automatically, so these maps can be  
486 incorporated in the current IRIS. We expect to implement this approach in IRIS to monitor *K.*  
487 *brevis* blooms in near-real-time in the next step.

488 Finally, the demonstration is for *K. brevis* blooms on the WFS between 2017 and 2019. Can the  
489 same DL model be applied to other years for the same WFS and to other regions in the Gulf of  
490 Mexico (GoM) where *K. brevis* have also been reported (e.g., coastal waters off Texas)? Because  
491 the DL model is strictly data driven, if the training used here for the period of 2017 – 2019 does

492 not encompass all observing scenarios (e.g., solar/viewing geometry, weak-moderate sun glint,  
493 different aerosol types and thicknesses) and all optical complexity (e.g., optically shallow bottom,  
494 non-algal water constituents) for other years or for other GoM regions, then the DL model needs  
495 to be retrained to include those scenarios. Otherwise, there is no need for retraining. For these  
496 reasons, because a 3-year observing period is believed to be long enough to cover all possible  
497 observing scenarios, application of the DL model for the WFS but to other years is unlikely to  
498 require retraining. In contrast, for other regions of the GoM, because the reasons leading to  
499 optical complexity may be different, a retraining is very likely needed. For the same reason,  
500 because HABs are a global phenomenon (Anderson et al., 2021) and because of the global  
501 coverage of VIIRS and other satellite data, we expect that such a machine learning approach may  
502 find more applications in other regions where HABs also occur, once field data are available for  
503 training and validation. These HABs are not necessarily caused by *K. brevis*, but can be caused  
504 by other dinoflagellates. In particular, future satellite missions will have the capacity to collect  
505 hyperspectral data on both sun-synchronous and geostationary satellite platforms. These include  
506 NASA's Plankton, Aerosol, Cloud, ocean Ecosystem (PACE) mission, NASA's Geostationary  
507 Littoral Imaging and Monitoring Radiometer (GLIMR) mission, and NOAA's Geostationary  
508 Extended Observations (GeoXO) mission. These missions will provide unprecedented ocean  
509 color data to bolster the ability to detect HABs by accounting for spatial coherence, spectral  
510 contrasts, and short-term changes.

511

## 512 **5. Conclusion**

513 To date, compared with MODIS or OLCI, the use of VIIRS in detecting HABs in the Gulf of  
514 Mexico is limited, possibly due to its lack of a fluorescence band. This technical challenge is  
515 circumvented here through the use of full spectral information from each VIIRS image pixel and  
516 a deep learning model to account for the spatial context of bloom pixels. Such an approach  
517 detects *K. brevis* blooms on the West Florida Shelf as spatially coherent features, thus avoiding  
518 typical problems of *K. brevis* patchiness (i.e., heterogeneity) encountered by traditional pixel-  
519 based methods. The approach led to detected *K. brevis* bloom patterns that are consistent with  
520 those derived from MODIS and microscopy observations and, meanwhile, the wide swath makes  
521 VIIRS particularly useful in both retrospective analyses of bloom dynamics and near-real-time

522 monitoring of bloom occurrence. We expect to implement such an approach for near-real-time  
523 data production in the current IRIS.

524

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537

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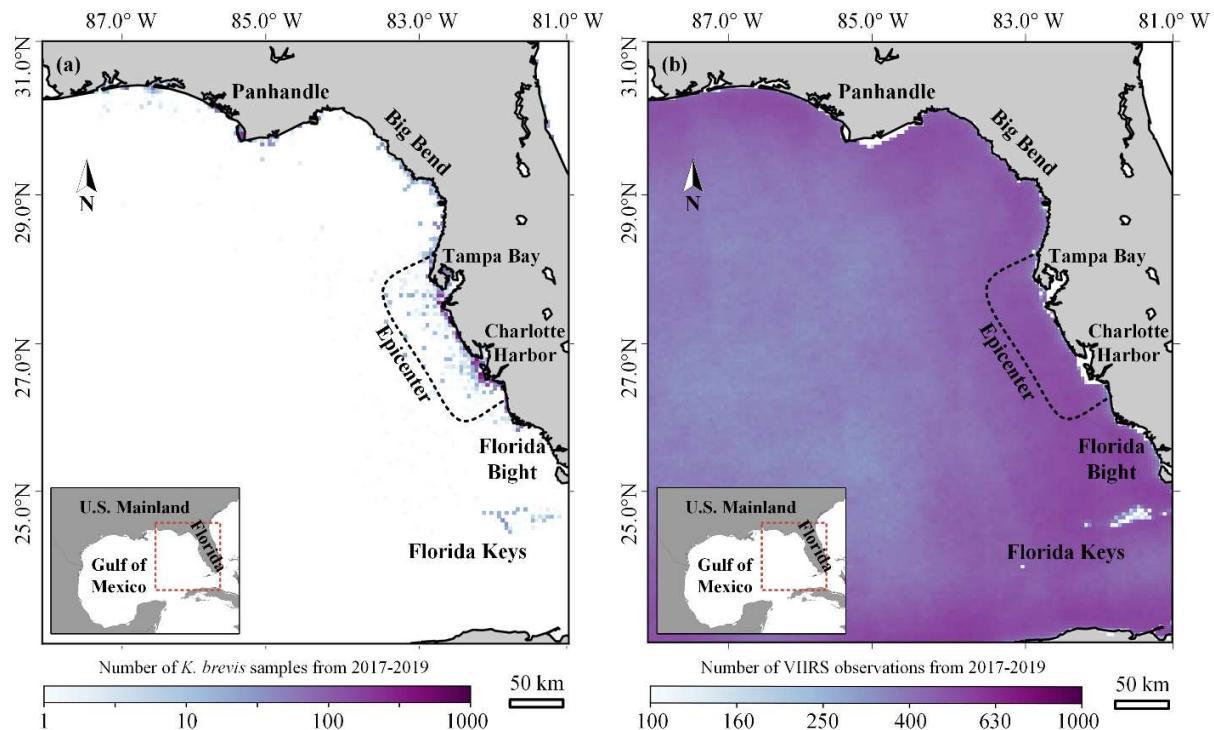
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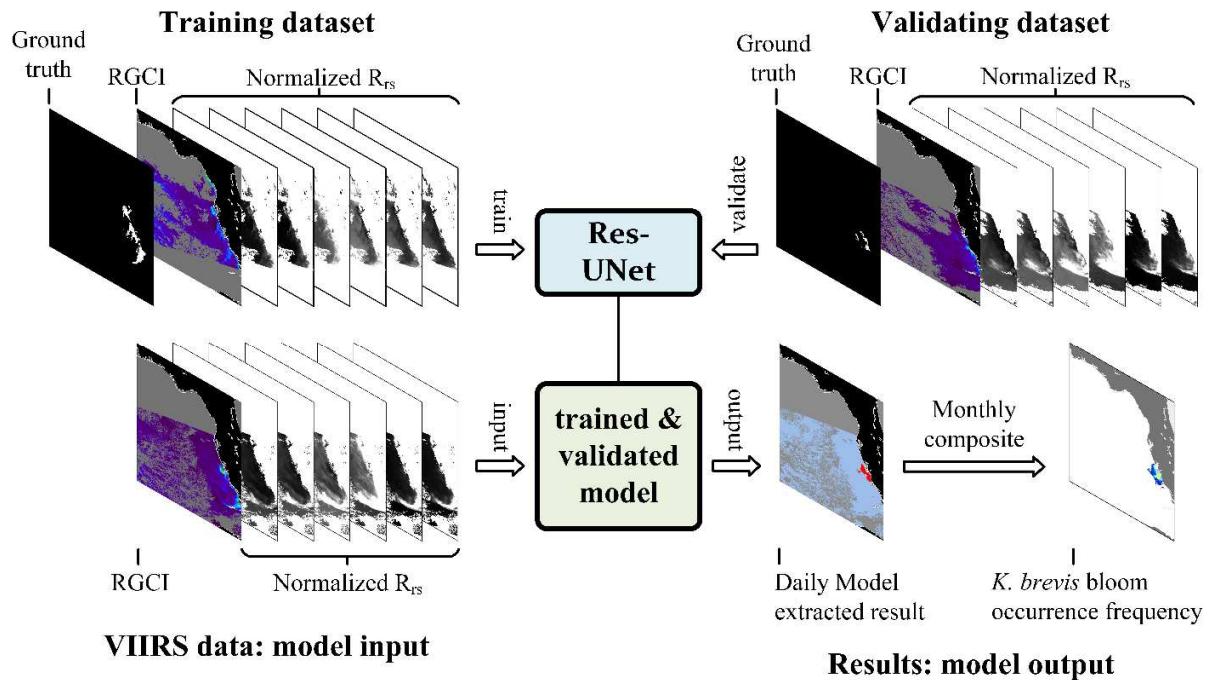
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776 **Figures**



777  
778 Fig. 1. The West Florida Shelf is located in the eastern Gulf of Mexico (inset map), spanning the  
779 region west of the Florida peninsula, encompassing the Panhandle, Big Bend, Central West  
780 Florida Shelf, including Tampa Bay and Charlotte Harbor, and the Florida Keys. The number of  
781 (a) discrete *in situ* *K. brevis* cellular abundance observations with scale from 1–1000 and (b)  
782 valid VIIRS observations in each 5-km grid in 2017–2019 with scale from 100–1000 are shown.  
783 The number of valid MODIS observations has been shown in Hu et al. (2022). Following  
784 [Weisberg et al. \(2019\)](#), the region from the north of Tampa Bay to the south of Charlotte Harbor  
785 is outlined as the *K. brevis* bloom “epicenter”, i.e., where most *K. brevis* blooms were found and  
786 most water samples were collected.

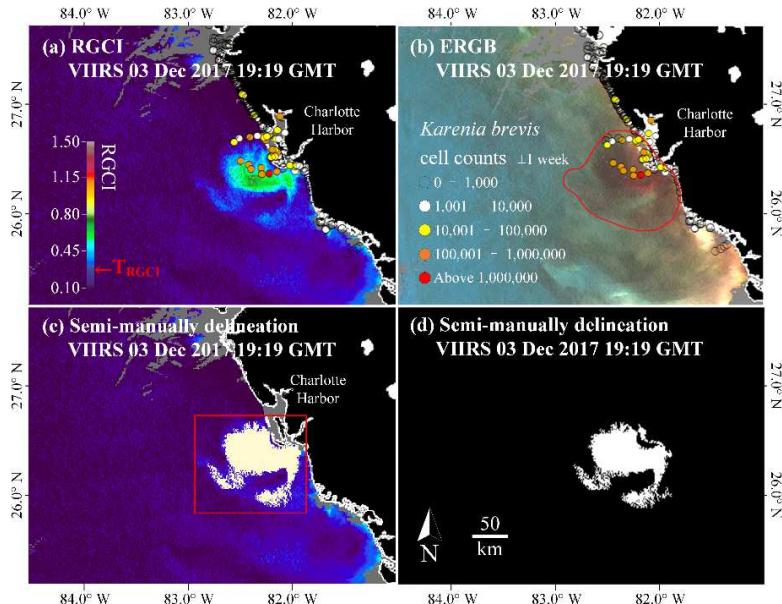
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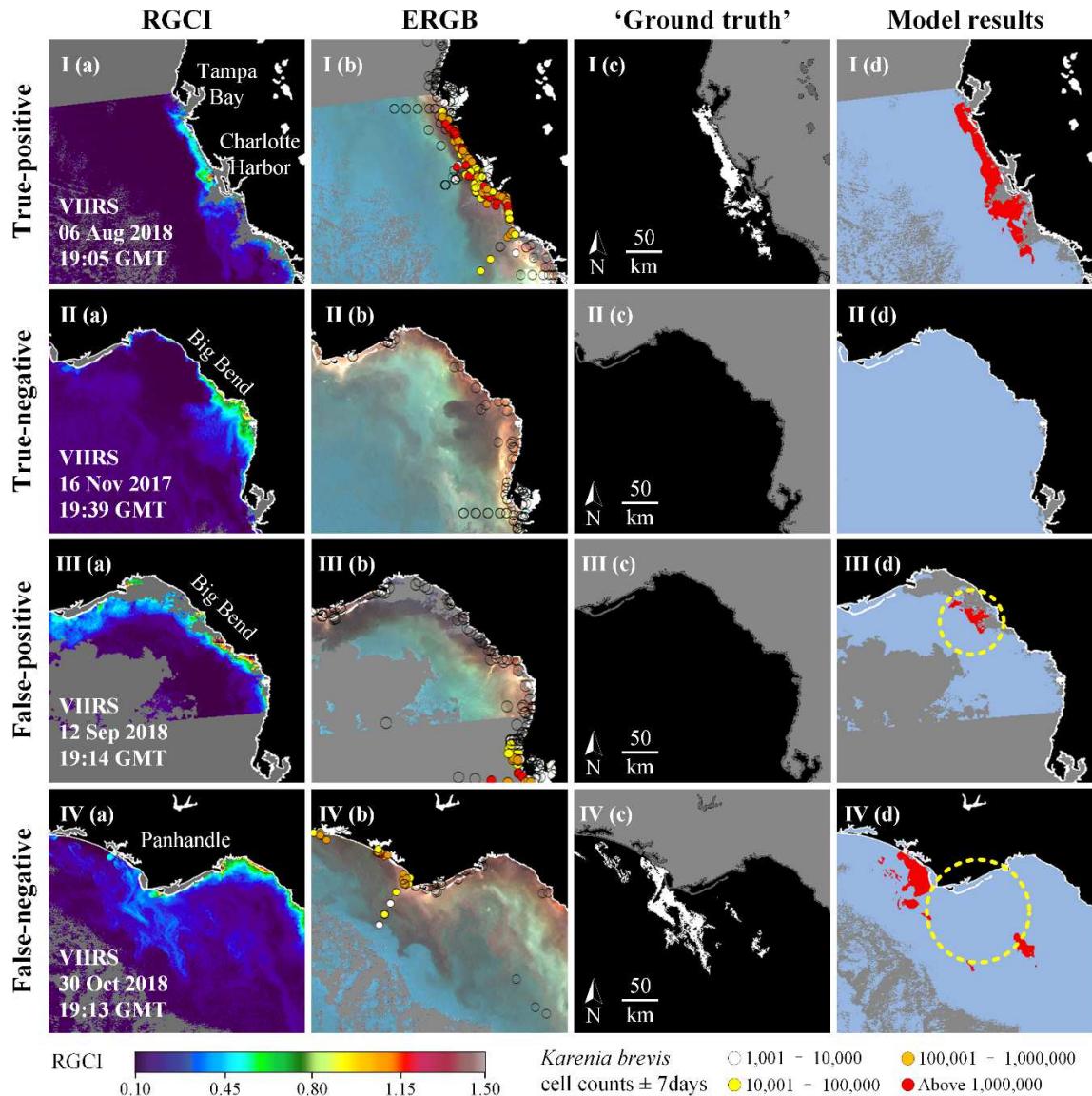
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790 Fig. 2. Conceptual illustration of the model training, validation, and application activities  
 791 conducted in this study for classifying *K. brevis* blooms in VIIRS imagery using a deep learning  
 792 approach. In the top row, VIIRS “ground truth”,  $R_{rs}(\lambda)$ , and RGCI images are used for training  
 793 and validating the deep learning model. In the bottom row, the validated model was applied to  
 794 VIIRS  $R_{rs}(\lambda)$  and RGCI data to delineate *K. brevis* bloom patches. The pixels were classified as  
 795 ‘*K. brevis* bloom’ (red), ‘non-*K. brevis* bloom’ (blue), and ‘no valid observation’ (grey). Monthly  
 796 bloom occurrence frequency maps were generated from the individual (near-daily) model  
 797 extracted results.

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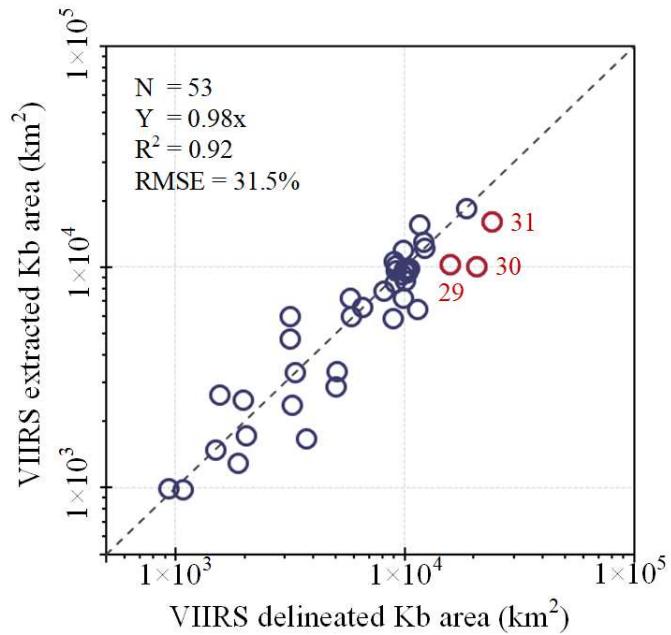


801 Fig. 3. “Ground truth” image preparation steps for the training the DL model. In (a), VIIIRS  
 802 RGCI is overlaid with *K. brevis* field sample data ( $\pm 1$  week) (shown in colored circles). The  
 803 spatially coherent high-RGCI patch in (a) and reddish-dark patch in the VIIIRS ERGB composite  
 804 image (b) that is collocated with high *K. brevis* cell counts ( $> 100,000$  cells  $L^{-1}$ ) suggest that this  
 805 is a *K. brevis* bloom patch. A crude outline is manually drawn over the patch. In (c), pixels  
 806 within the manual outline with RGCI values greater than the RGCI threshold ( $T_{RGCI}$  is marked  
 807 on the color bar) are considered as bloom pixels. In (d), pixels in the delineated patch (*K. brevis*  
 808 bloom) are marked as 1 (white), and all other pixels (non-*K. brevis* bloom) are marked as 0  
 809 (black).



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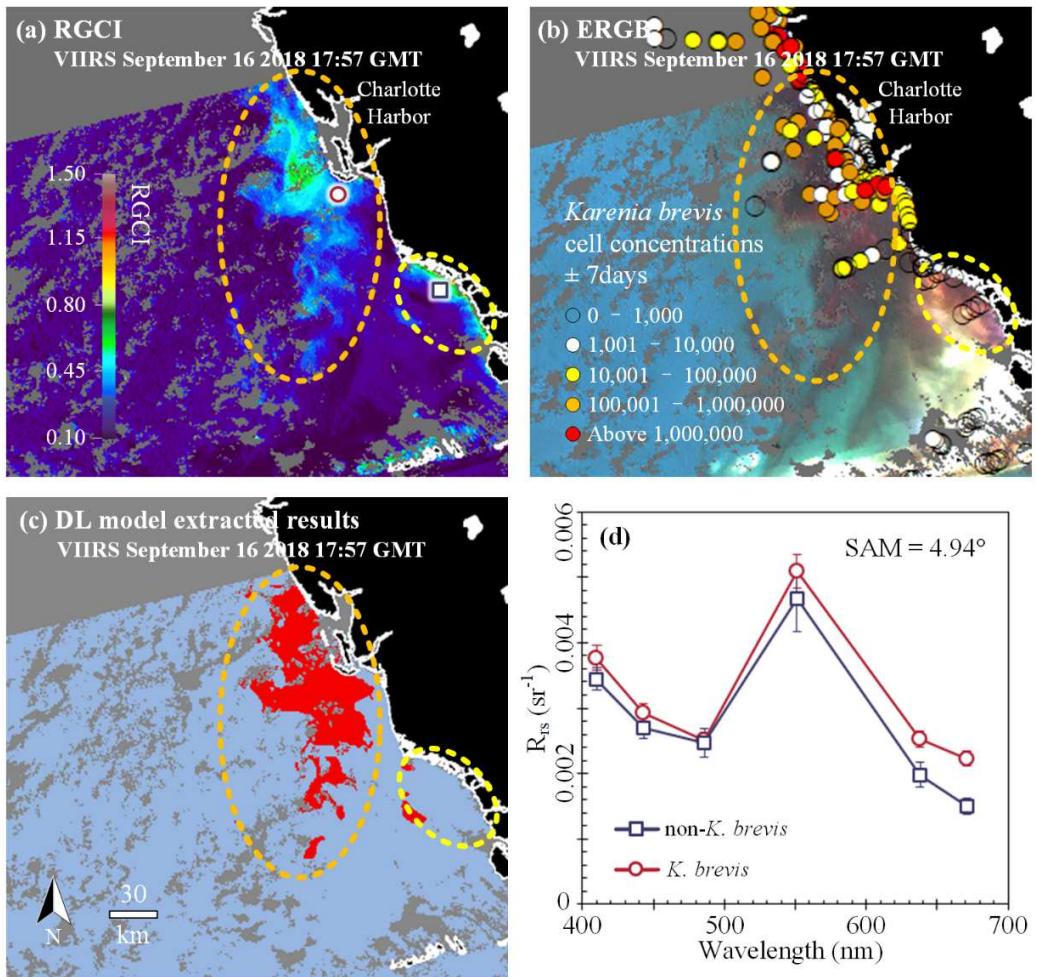
811 Fig. 4. Four sets of example imagery from the validation dataset. The first row (I) shows a case  
 812 with true-positive results. The second row (II) shows a case with true-negative results. The third  
 813 row (III) shows the false-positive results, and the fourth row (IV) shows the false-negative  
 814 results (poor results are circled in dashed lines) as compared with the “ground truth” delineation  
 815 in (c). Imagery shown include: (a) VIIRS RGCI; (b) VIIRS ERGB-composite images with field  
 816 *K. brevis* cell count data overlaid, cell counts less than 1000 show as transparent circles; (c)  
 817 VIIRS semi-objectively delineated results, and (d) VIIRS DL model extracted results with red,  
 818 blue, and grey representing *K. brevis* blooms, non-*K. brevis* bloom, and no valid observation or  
 819 no satellite data, respectively.



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821 Fig. 5. Scatter plot of *K. brevis* bloom area ( $\text{km}^2$ ) determined from the VIIRS delineated “ground  
 822 truth” dataset (based on visual inspection of co-located VIIRS imagery and in-situ data) and  
 823 trained VIIRS DL model. Red circles represent the image pairs that the bloom detected in  
 824 Panhandle region on 29, 30 and 31 October 2018. Cases where neither method identified a *K.*  
 825 *brevis* bloom are not shown.

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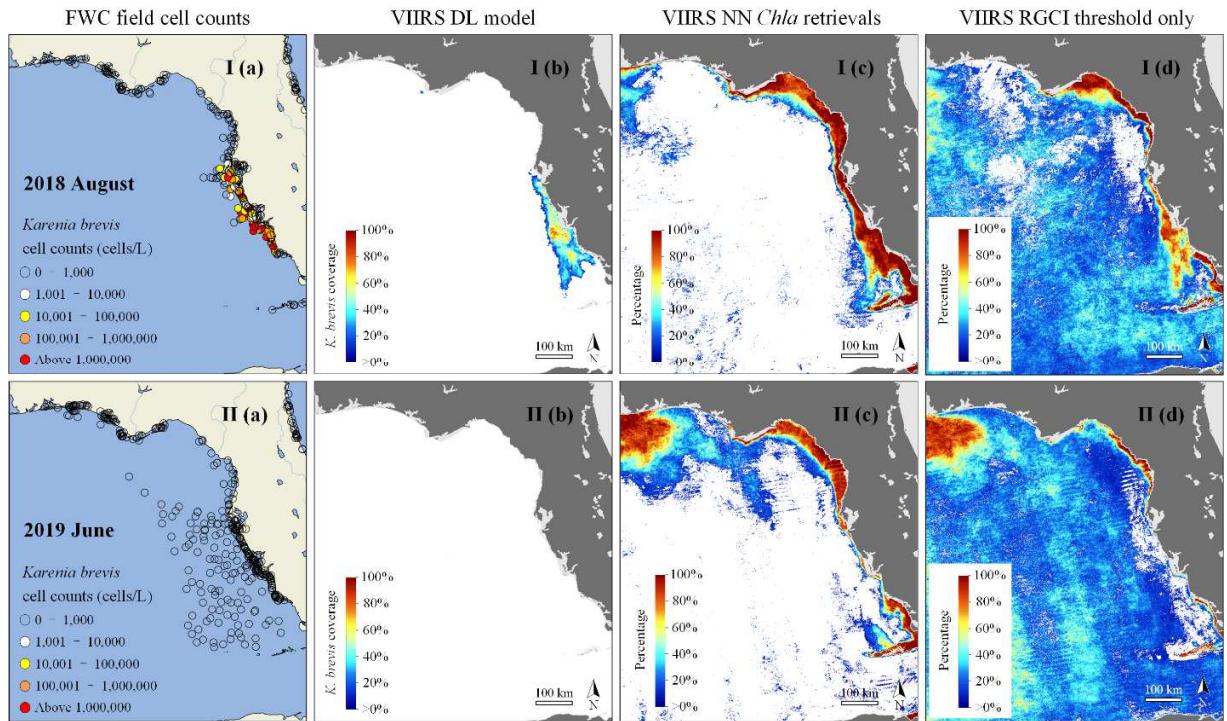


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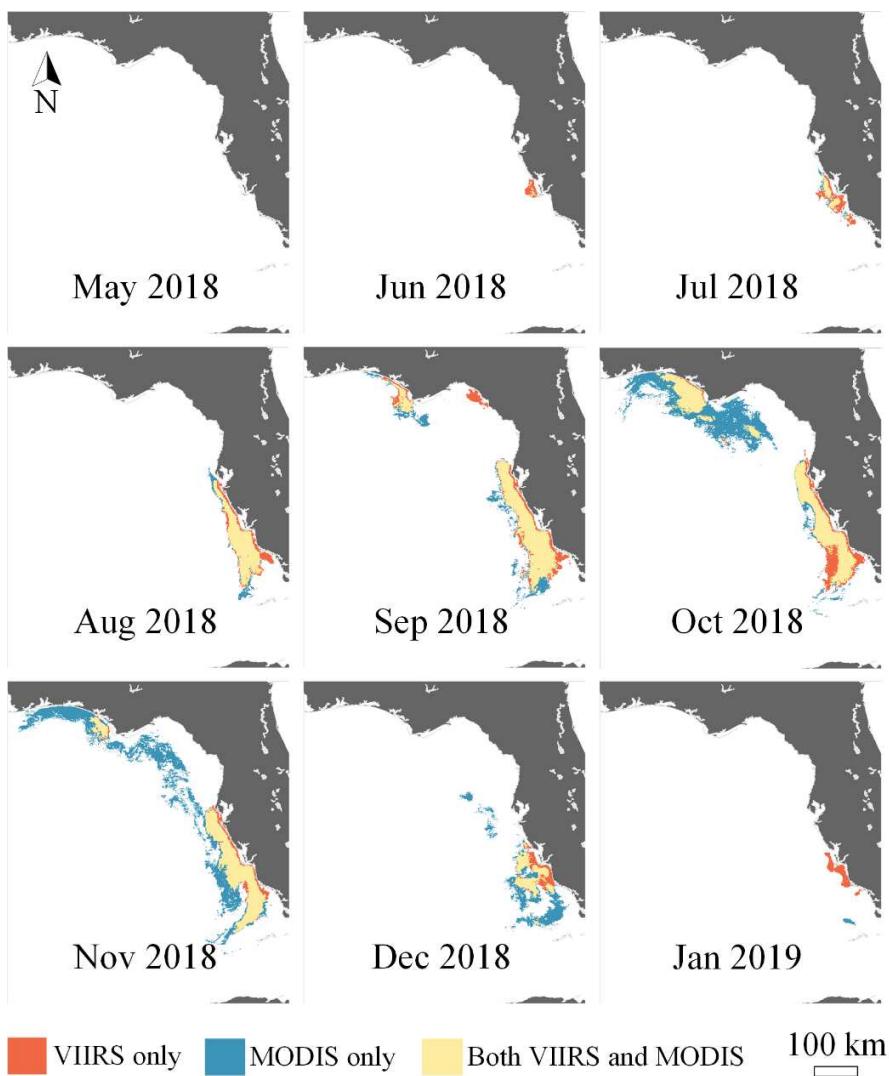
828 Fig. 6. The capacity of the DL model in separating *K. brevis* and non-*K. brevis* bloom patches in  
 829 optically complex nearshore waters is demonstrated here using a VIIRS scene of the WFS  
 830 collected on 16 September 2018 for cases of (a) VIIRS RGCI, (b) VIIRS ERGB composite  
 831 image overlaid with field *K. brevis* cellular abundance data, where high RGCI values are shown  
 832 to correspond to dark waters, and (c) *K. brevis* bloom patch determined from the VIIRS DL  
 833 model, showing that the DL model correctly classified a high-RGCI patch (yellow dashed circle)  
 834 south of the *K. bloom* patch (orange dashed circle) as non-*K. brevis* bloom. Panel (d) shows that  
 835 the spectral shapes from the two locations (one in the *K. brevis* bloom and the other in the non-*K.*  
 836 *brevis* bloom, see locations marked in (a)) are similar, yet the DL model could differentiate them.

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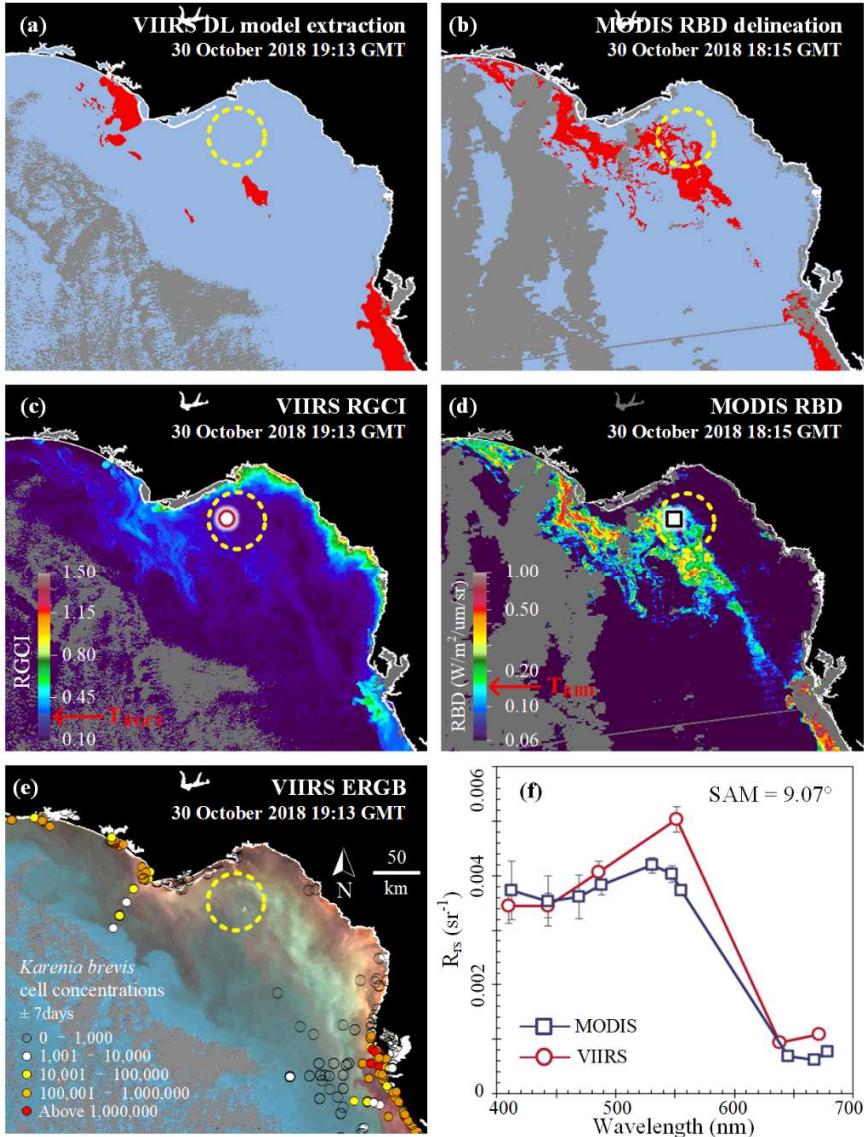
840 Fig. 7. Comparisons between (a) field *K. brevis* cellular abundance data ( $\text{cells L}^{-1}$ ) from FWC  
 841 and monthly VIIRS bloom occurrence frequency maps generated in I(a) August 2018 and II(a)  
 842 June 2019 using the approaches of I(b) & II(b) DL model, I(c) & II(c) neural network model (El-  
 843 Habashi et al., 2016), and I(d) & II(d) RGCI during *K. brevis* bloom (I) and non-*K. brevis* bloom  
 844 (II) events. Only the DL model can correctly identify the bloom and non-bloom events.



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846 Fig. 8. Monthly *K. brevis* bloom footprint maps derived from MODIS by semi-objective  
 847 delineation (Hu et al., 2022) and VIIRS using the DL model for the bloom event between mid-  
 848 2018 and early-2019.

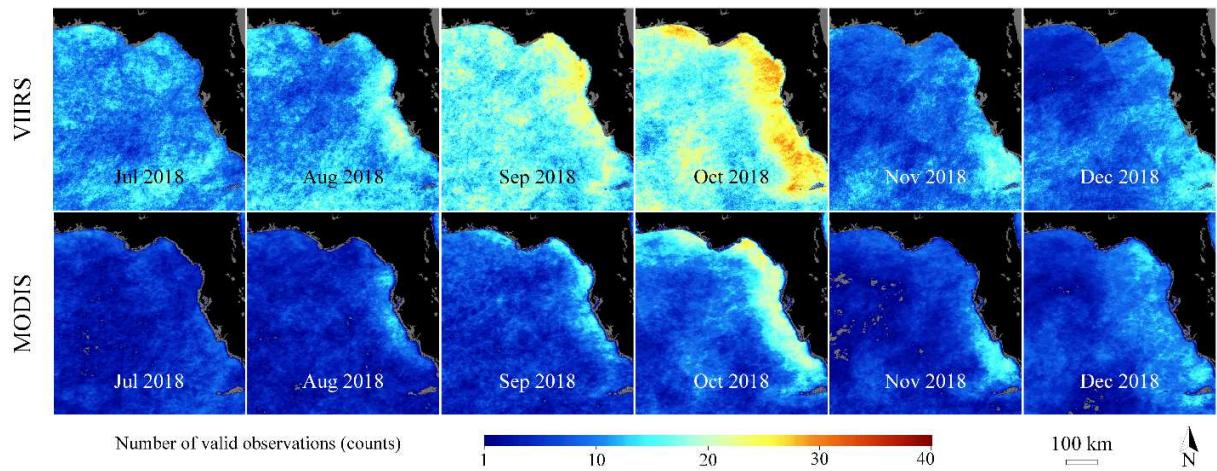
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851 Fig. 9. The detection difference between VIIRS and MODIS in the Panhandle region on 30  
 852 October 2018 for (a) VIIRS DL model extraction results, (b) MODIS RBD delineation results  
 853 based on the RBD threshold in Hu et al. (2022), (c)VIIRS RGCI image, (d) MODIS RBD image,  
 854 (e) VIIRS ERGB images overlaid with *K. brevis* cell counts, and (f) VIIRS and MODIS  $R_{rs}(\lambda)$   
 855 spectra from dash circled locations as noted in (d) and (e). Note that red, blue, and grey in panels  
 856 (a) and (b) represent *K. brevis* blooms, non-*K. brevis* bloom, and no valid observation or no  
 857 satellite data, respectively.

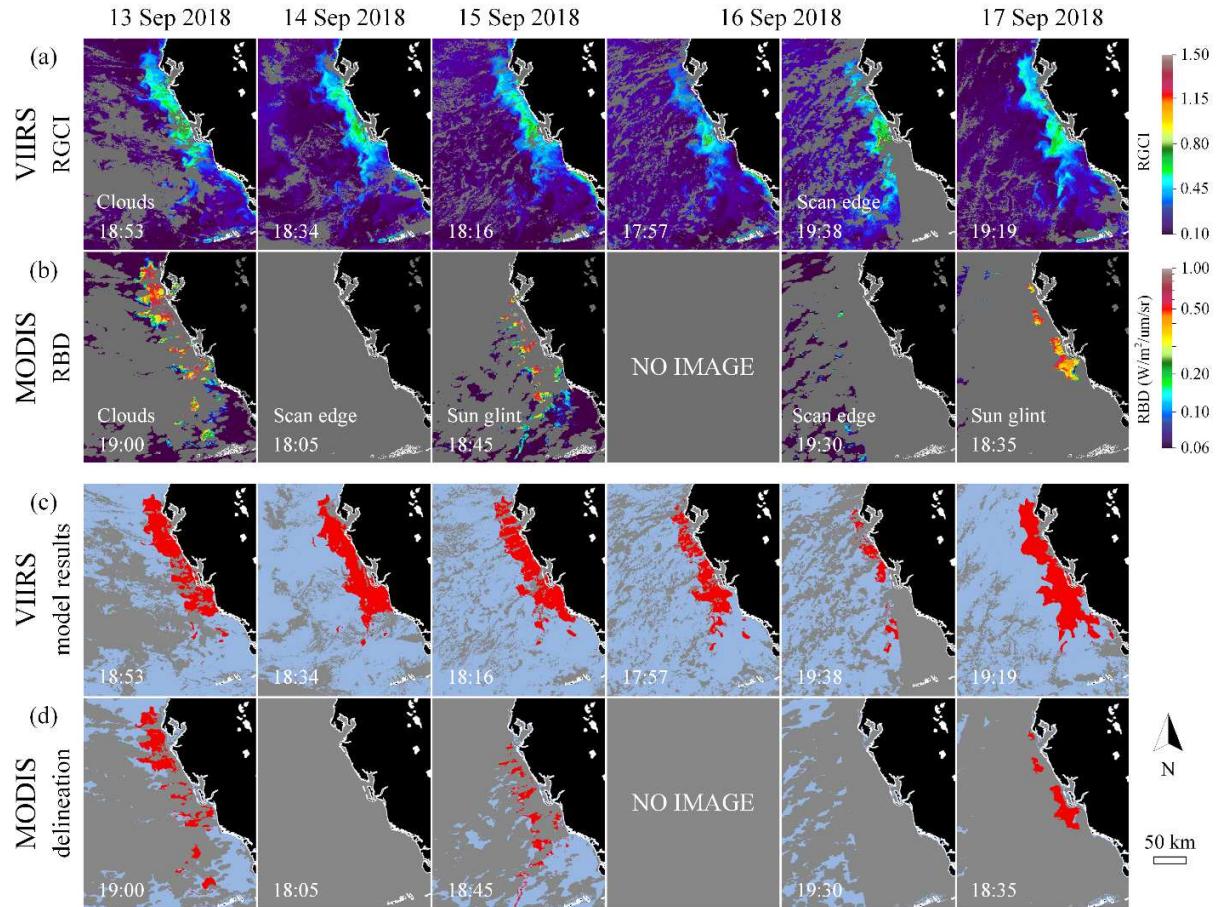
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860 Fig. 10. Monthly images of VIIRS (top row) and MODIS (bottom row) showing the spatial  
 861 distributions of number of valid observations at each 1-km location in the eastern Gulf of Mexico  
 862 for each month from July to December 2018.

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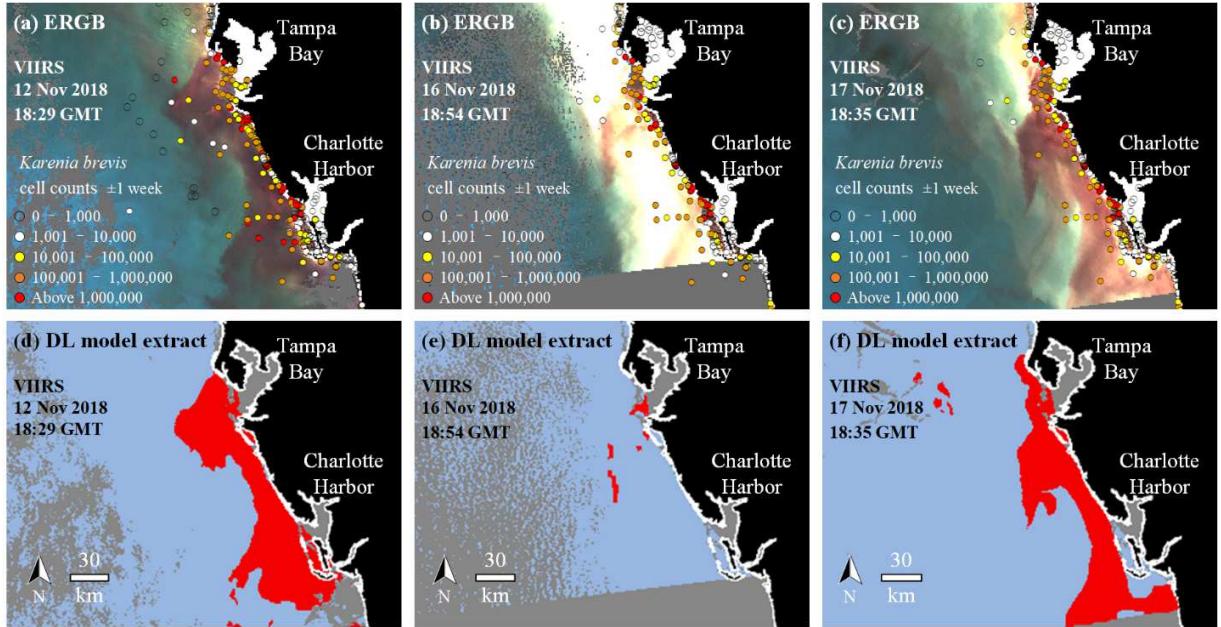
865 Fig. 11. An example showing the difference between VIIRS and MODIS valid observations and  
 866 *K. brevis* bloom detections. The rows in (a) and (b) show the VIIRS RGCI and MODIS RBD  
 867 images, respectively, over the central WFS for five consecutive days. The grey color represents  
 868 no valid observation or no satellite data. The rows in (c) and (d) show their corresponding *K.*  
 869 *brevis* bloom detection results determined with VIIRS using the DL model and MODIS by semi-  
 870 objective delineation (Hu et al., 2022). Red, blue, and grey represent *K. brevis* blooms, non-*K.*  
 871 *brevis* bloom, no valid observation or no satellite data, respectively.

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877 Fig. 12. An example showing how the strength of the VIIRS DL model (i.e., avoid false-positive  
 878 bloom detection from sediment-rich waters) can turn into a weakness, and how such a weakness  
 879 can be overcome by inspecting sequential images. Panels (a-c) are the VIIRS ERGB images in  
 880 the Epicenter region overlaid with *K. brevis* cell counts, collected on 12, 16, and 17 November  
 881 2018, respectively. Panels (d-f) are the corresponding VIIRS DL model extraction results, with  
 882 red, blue, and grey representing *K. brevis* bloom patches, non-*K. brevis* bloom waters, and no  
 883 valid observation or no satellite data, respectively. The field data and ERGB images suggest a  
 884 continuous *K. brevis* bloom in the Epicenter region in mid-November 2018. During the bloom  
 885 period, the sediment resuspension event on November 16 (b) led to no bloom detection (e), but  
 886 one can still safely assume the existence of similar bloom patches as in (d) and (f).

887

## Graphical Abstract

Harmful algal blooms (HABs) of the toxic dinoflagellate *Karenia brevis*, often called red tides, occur annually on the West Florida Shelf (WFS). Detection of these HABs using satellite observations often suffers from two problems: lack of accurate algorithms to identify phytoplankton blooms in optically complex waters and patchiness (i.e., heterogeneity) of *K. brevis* cellular abundance in bloom waters. Here, to take advantage of the wide swath (3040 km) and non-saturation of the Visible Infrared Imaging Radiometer Suite (VIIRS) while realizing its disadvantage due to the lack of a fluorescence band, we develop a deep-learning (DL) convolutional neural network model to overcome the above technical challenges, especially on the spatial coherence of bloom patches. After proper training, the overall performance (i.e., F1 score) of the DL model is 89%. The results for the period of 2017 – 2019 not only demonstrate the capacity of VIIRS in HABs monitoring, but also show the value of the DL model in extracting *K. brevis* bloom patches for both near real-time applications and retrospective analysis.

