MicrographCleaner: a python package for cryo-EM micrograph segmentation using deep learning

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Abstract

Cryo-EM workflows require from tens of thousands of high-quality particle projections to unveil the three-dimensional structure of macromolecules. Current methods for automatic particle picking tend to suffer from high false-positive rates, hurdling the reconstruction process. One common cause of this problem is the presence of carbon and different types of high-contrast contaminations that, in many cases, affect large areas of micrographs. In order to overcome this limitation, we have developed MicrographCleaner, a deep learning approach designed to discriminate which regions of micrographs are suitable for particle picking and which are not, that we will refer to as "contaminated". MicrographCleaner implements a U-net-like model trained on a manually curated dataset compiled from over five hundred micrographs. The benchmarking, carried out on about one hundred independent micrographs, shows that MicrographCleaner is a very efficient approach for micrograph preprocessing.

Availability and implementation
MicrographCleaner package is available at PyPI and Anaconda Cloud repositories as micrograph_cleaner_em. Source code is available at https://github.com/rsanchezgarc/micrograph_cleaner_em. Integration with the cryo-EM software Scipion/Xmipp is also provided through the deepMicrographScreen protocol.

1. Introduction.

Cryogenic-Electron Microscopy (cryo-EM) Single Particle Analysis (SPA) has recently become a powerful technique for the determination of macromolecular structures achieving, in many cases, atomic resolutions. SPA consists of a set of complex and variable operations that, departing from thousands of particle projections, leads to the synthesis of electronic density maps of macromolecules. The massive number of particles that are needed for SPA has made of automatic particle picking one of the most influential steps in virtually all reconstruction workflows. Nevertheless, some problems intrinsic to the cryo-EM pipelines, such as low signal-to-noise ratio and the presence of high contrast artifacts and contaminants in the micrographs, degrades the performance of particle picking algorithms (Zhu *et al.*, 2004; Vargas *et al.*, 2013) and leads to the addition of false positive particles in SPA workflows. This problem can be mitigated trough different algorithms that clean and remove incorrectly selected particles after automatic picking (Sanchez-Garcia et al., 2018; Vargas et al., 2013).

One of the most common shortcomings observed during automatic picking is the attraction of these methods to select grid carbon spots, especially at the holes edges. Due to its relevance, some algorithms

have been designed to prevent particle selection in those regions. For example, the em_hole_finder program, included in the Appion package (Lander et al., 2009) is based on morphological image processing operations to compute masks around carbon holes. More recently, EMHP (Berndsen et al., 2017) was designed to perform a similar task through image filtering and thresholding operations followed by a circle fitting procedure. Although very useful when grid edges are clearly visible, both approaches struggle in those cases where high contrast contaminations are present in micrographs. Moreover, both of them require human supervision to determine the presence of carbon in the micrographs and to set some user-defined parameters. As a result, its applicability is limited to supervised scenarios.

In order to overcome these limitations, we have developed MicrographCleaner, a fully automatic deep learning solution that, instead of trying to find the regions of micrographs where carbon is located, it performs a pixel-wise classification – segmentation – of micrographs into two categories, desirable and undesirable regions. Moreover, our model, trained for pixel-wise classification, is not only able to provide a more robust and accurate solution for carbon detection, but it is also able to remove other types of contaminations such as ice crystals or ethane.

2. Material and methods

2.1 Algorithm

MicrographCleaner computes binary semantic segmentation of micrographs with the aim of delineating optimal regions for particle picking and isolating those areas containing high-contrast contaminants and other artifacts. To that end, MicrographCleaner implements a U-net-like model (Ronneberger et al., 2015) trained on a dataset of 539 manually segmented micrographs collected from 16 different EMPIAR (Iudin et al., 2016) entries. The evaluation was performed on an independent set of 97 micrographs compiled from two EMPIAR projects and another two in-home projects (see Supplementary Material S4). Both training and testing sets of micrographs include examples of clean, carbon-containing and contamination-containing as well as mixed cases. Neural network training was carried out using the Adam optimizer and a combination of perceptual loss (Johnson et al., 2016) and weighted binary cross-entropy (see Supplementary Material S1 and S2). A previous normalization step is required to adjust the different intensity scales of micrographs. Thus, all micrographs are normalized using a robust scaling strategy and donwsampled (see Supplementary Material S3). Finally, overlapping patches of 256x256 pixels with strides of 128 pixels are extracted from the micrograph and fed to the network.

2.1 Package

MicrographCleaner has been implemented as an easy-to-install and easy-to-employ Python 3.x package. Thus, the command line tool can be automatically installed from Anaconda Cloud and PyPI repositories whereas the GUI version can be installed through the Scipion (de la Rosa-Trevín et al., 2016) plugin manager. The neural network was implemented using the Keras (Chollet, 2015) package and the Tensorflow (Abadi et al., 2016) backend. Micrograph preprocessing is carried out using the scikit-image (van der Walt et al., 2014) package.

3. Results

The evaluation of MicrographCleaner was performed comparing the predicted masks with the ground truth of testing micrographs. To that end, the mean Intersection over Union (mIoU) metric was

calculated considering predicted and manually curated micrograph regions (see Supplementary Material S2). MicrographCleaner achieved a mIoU value of 0.544. This score implies a good agreement between ground truth and predicted masks, especially when taking into account that the testing set contains clean micrographs examples together with carbon-containing and contaminated micrographs. Figure 1 shows the predictions for four different micrographs, illustrating that MicrographCleaner is capable of successfully detecting both contaminations and carbon.

We also have compared MicrographCleaner with two state-of-the-art carbon finder programs: em_hole_finder and EMHP. Before entering into these comparisons, it is important to highlight that MicrographCleaner, contrary to the others, is a parameter-free method and it does not require manual intervention in order to determine whether or not carbon is present in a micrograph. It is also quite fast, in the order of 3 seconds per micrograph. Consequently, MicrographCleaner can be employed in automatic pipelines and thus, it is suitable for automatic Cryo-EM analysis at facilities. Yet, with the aim of strictly compare carbon detection efficacy, we have taken the subset of the testing set in which all micrographs contain some carbon and executed the three algorithms. As it can be appreciated in Table 1, MicrographCleaner achieves a much better performance than previously published methods (see Supplementary Material S5 for more details).

Table 1. MicrographCleaner performance for carbon detection compared to other methods.

Algorithm	**mIoU**	**stdIoU**	**Failure percentage**
MicrographCleaner	0.788335	0.229398	3.33%
EMHP	0.198055	0.211471	45.00%
em_hole_finder	0.056918	0.046918	63.00%

Notes: mIoU: mean Intersection over Union; stdIoT standard deviation Intersection over Union; Failure percentage: percentage of the testing set for which the IoU was equal to 0.

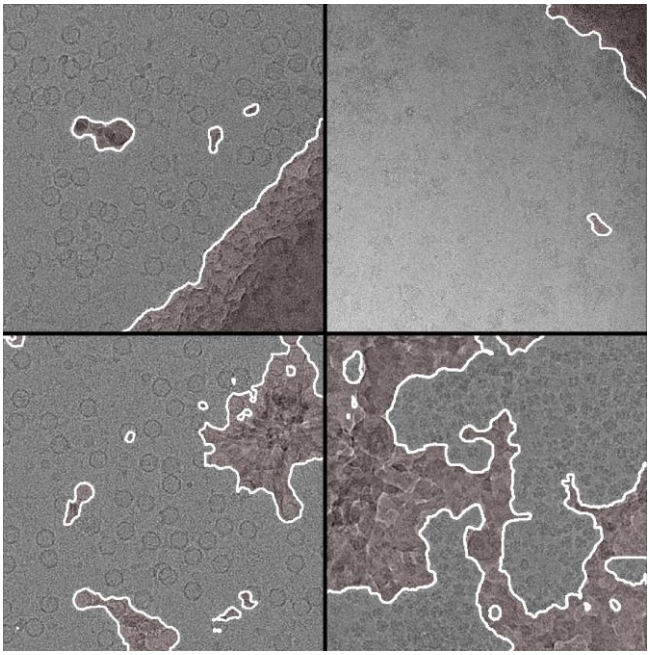

Fig 1. MicrographCleaner predictions. Red shadowed regions correspond to micrograph areas labeled as “non suitable” with 50% or more confidence. Top images show MicrographCleaner capability to detect carbon in the presence of contaminants. Bottom images show MicrographCleaner capability to detect a wide variety of different contaminants.

4. Conclusions

MicrographCleaner is an easy-to-install and easy-to-use python package that allows efficient and automatic micrograph segmentation with the aim of preventing particle pickers from selecting inappropriate regions. To that end, MicrographCleaner relays on a U-net-like model that has being trained on about 500 micrographs. When compared to other methodologies, MicrographCleaner has proven more robust, achieving results closer to the human criterion than other state-of-the-art methods. As a result, we consider that MicrographCleaner is a powerful approach to be applied at the very beginning of cryo-EM workflows, even within on-the-fly processing pipelines, leading to cleaner sets of input particle and, consequently to a better processing performance.

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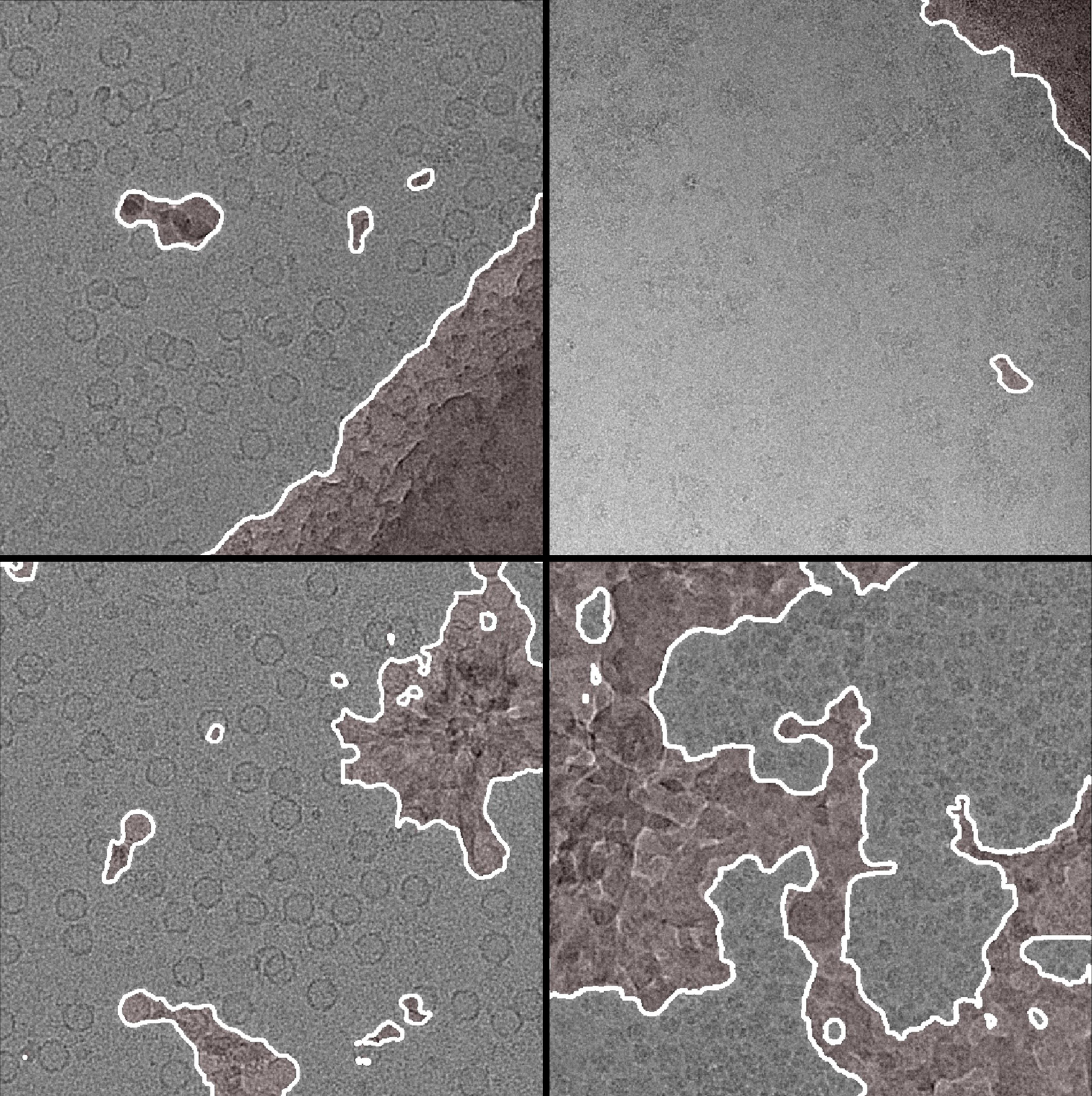
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