

¹ Deep Learning for Single Molecule Localization ² Microscopy

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⁴ Summary

⁵ Single Molecule Localization Microscopy (SMLM) enables researchers to interrogate nanoscale
⁶ spatial details in a range of systems. Biological investigations benefit greatly from SMLM
⁷ due to its ability to quantitatively investigate details of great importance such as protein
⁸ distribution in the cell membrane or protein-protein interactions([Baddeley & Bewersdorf, 2018](#)).
⁹ However, SMLM set-ups can be quite expensive, imaging times can be lengthy and data
¹⁰ analysis can require expert knowledge. Recently, Deep Learning (DL) algorithms have been
¹¹ developed to reduce imaging time and automate data analysis. Naturally each subsequent
¹² model aims to address a different shortcoming of the prior work and so there is a family of
¹³ model architectures. However, there is not a singular location for researchers to have access
¹⁴ to these models dedicated to SMLM. We developed the Deep Learning for Single Molecule
¹⁵ Localization Microscopy (DL4SMLM) in Python using the PyTorch framework to democratize
¹⁶ access to these models and lower the barrier of entry to SMLM.

¹⁸ Statement of Need

¹⁹ DL4SMLM is a Python package that uses the Pytorch DL framework to implement Convolu-
²⁰ tional Neural Nets (CNN) dedicated to SMLM. We created it for researchers and engineers who
²¹ wish to use DL for SMLM in whichever field they practice granted they have sufficient data.
²² Currently, there are six models: Super Resolution Convolutional Neural Network (SRCNN)([Dong](#)
²³ et al., 2015), Deep-Stochastic Optical Reconstruction Microscopy (Deep-STORM)([Nehme](#) et
²⁴ al., 2018), Skip-STORM, Unet for STORM (U-STORM)([W. Yao](#) et al., 2018), Deep Residual
²⁵ STORM (DRL-STORM)([B. Yao](#) et al., 2020), and Fast Dense Image Reconstruction based
²⁶ Deep Learning STORM (FID-STORM)([Zhou](#) et al., 2023). SRCNN and Skip-STORM are
²⁷ novel architectures to the SMLM field. While SRCNN has been applied to perform the super
²⁸ resolution task in more traditional image processing tasks, its utility in SMLM is unknown.
²⁹ Skip-STORM's architecture is similar to Deep-STORM but a skip connection between the
³⁰ initial image and final layer is introduced to enable it to have a spatial context during its
³¹ reconstruction phase which could assist it in emitter dense images. U-STORM is inspired
³² by U-Net and adopts a similar biphasic architecture where in the first half the initial image
³³ is max pooled while simultaneously increasing channel number and the second half involves
³⁴ the up sampling and reduction in channel. It also contains skip connections that connect the
³⁵ features of the first half to the those of second half during image reconstruction. These models
³⁶ are implemented using the object-oriented programming paradigm enabling researchers to
³⁷ instantiate multiple models for a single data set to allow downstream comparison of inference
³⁸ performance. Additionally, we have implemented the L1L2 Loss metric first introduced in
³⁹ ([Nehme](#) et al., 2018) and used for all subsequent architectures implemented in this library.
⁴⁰ Other features include functions that automate the Training and Validation process for a
⁴¹ user-set number of epochs in addition to a function that automates the inference procedure
⁴² of a trained model on a test data set and reports the Structured Similarity Index Measure
⁴³ (SSIM) and the Normalized Mean Square Error (NMSE). To assist researchers in visualizing
the emitter localization performance of their model, we have implemented a function that

44 visualizes the diffraction limited image of the emitters, its super-resolved version, and the
45 predicted super-resolved image according to the trained model. This empowers the researcher
46 to troubleshoot the performance of the model and determine if the model should be retrained
47 with different parameters or to proceed with the experiment. To facilitate the development
48 of powerful yet accurate models we have implemented two functions that enable knowledge
49 distillation between a teacher model and a student model: Hint Learning and Knowledge
50 Transfer. Hint training automates the learning process of a student network to have its
51 intermediate representation mirror that of a teacher network. Knowledge Transfer uses the
52 attentive imitation loss function from ([Saputra et al., 2019](#)), the teacher model, and the ground
53 truth data set to optimize the student network. Altogether, we aim for this package to serve
54 as a widely used resource for any experimenter in the SMLM field.

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60 References

- 61 Baddeley, D., & Bewersdorf, J. (2018). Biological insight from super-resolution microscopy:
62 What we can learn from localization-based images. *Annual Review of Biochemistry*, 87(1),
63 965–989.
- 64 Dong, C., Loy, C. C., He, K., & Tang, X. (2015). Image super-resolution using deep
65 convolutional networks. *IEEE Transactions on Pattern Analysis and Machine Intelligence*,
66 38(2), 295–307.
- 67 Nehme, E., Weiss, L. E., Michaeli, T., & Shechtman, Y. (2018). Deep-STORM: Super-
68 resolution single-molecule microscopy by deep learning. *Optica*, 5(4), 458–464. <https://doi.org/10.1364/OPTICA.5.000458>
- 69 Saputra, M. R. U., De Gusmao, P. P., Almalioglu, Y., Markham, A., & Trigoni, N. (2019).
70 Distilling knowledge from a deep pose regressor network. *Proceedings of the IEEE/CVF
71 International Conference on Computer Vision*, 263–272.
- 72 Yao, B., Li, W., Pan, W., Yang, Z., Chen, D., Li, J., & Qu, J. (2020). Image reconstruction
73 with a deep convolutional neural network in high-density super-resolution microscopy. *Opt.
74 Express*, 28(10), 15432–15446. <https://doi.org/10.1364/OE.392358>
- 75 Yao, W., Zeng, Z., Lian, C., & Tang, H. (2018). Pixel-wise regression using u-net and its
76 application on pansharpening. *Neurocomputing*, 312, 364–371. <https://doi.org/10.1016/j.neucom.2018.05.103>
- 77 Zhou, Z., Wu, J., Wang, Z., & Huang, Z.-L. (2023). Deep learning using a residual decon-
78 volutional network enables real-time high-density single-molecule localization microscopy.
79 *Biomed. Opt. Express*, 14(4), 1833–1847. <https://doi.org/10.1364/BOE.484540>