

More detailed instructions can be found in [YCWu-nk/Cell_SAM](#): Large-scale segmentation model facilitates intraoperative histopathology by third harmonic generation microscopy, you can directly copy them

1. We suggest confirming that the basic environment is installed correctly first:

```
(base) [Mon Jul 28 09:59:18 2025] $ nvidia-smi
NVIDIA-SMI 575.64.05    Driver Version: 575.64.05    CUDA Version: 12.9
GPU  Name Persistence-M  Bus-Id Disp.A  Volatile Uncorr. ECC
Fan  Temp  Perf Pwr:Usage/Cap| Memory-Usage GPU-Util Compute M.
Temp MIG M.

0  NVIDIA GeForce RTX 4090   On 00000000:01:00.0 Off
  0% 27C P8      20W / 450W 1MiB / 24564MiB 0% Default N/A
1  NVIDIA GeForce RTX 4090   On 00000000:02:00.0 Off
  0% 28C P8      13W / 450W 1MiB / 24564MiB 0% Default N/A
2  NVIDIA GeForce RTX 4090   On 00000000:C1:00.0 Off
  0% 27C P8      21W / 450W 1MiB / 24564MiB 0% Default N/A
3  NVIDIA GeForce RTX 4090   On 00000000:E1:00.0 Off
  0% 28C P8      11W / 450W 1MiB / 24564MiB 0% Default N/A

Processes:
GPU  GI  CI          PID  Type  Process name          GPU Memory Usage
ID   ID
No running processes found

(base) [ ]$ conda --version
conda 25.5.1
(base) [yuchen@roli-27 yuchen] $ nvcc -V
nvcc: NVIDIA (R) Cuda compiler driver
Copyright (c) 2005-2025 NVIDIA Corporation
Built on Tue_May_27_02:21:03_PDT_2025
Cuda compilation tools, release 12.9, V12.9.86
Build cuda_12.9.r12.9/compiler.36037853_0
(base) [ ]$
```

2. Creating a conda environment, and install *the GPU version* of PyTorch in your preferred way:

Proceed ([y]/n)? y

Downloading and Extracting Packages:

```
Preparing transaction: done
Verifying transaction: done
Executing transaction: done
#
# To activate this environment, use
# $ conda activate cell-sam
#
# To deactivate an active environment, use
# $ conda deactivate
```

```
(base) [cell-sam] $ conda activate cell-sam
```

```
base: ~/.conda/envs/cell-sam
(cell-sam) [ ]$ pip install torch==1.9.0+cu111 torchvision==0.10.0+cu111 torchaudio==0.9.0 -f https://download.pyt
torch.org/whl/torch_stable.html
Looking in indexes: https://pypi.org/simple, https://pypi.ngc.nvidia.com
Looking in links: https://download.pyt
or
Collecting torch==1.9.0+cu111
  Downloading https://download.pyt
or
Collecting torchvision==0.10.0+cu111
  Downloading https://download.pyt
or
Collecting torchaudio==0.9.0
  Downloading https://download.pyt
or
Requirement already satisfied: typing-extensions in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-pa
(fro
Requirement already satisfied: numpy in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-pa
(fro
Requirement already satisfied: pillow>=5.3.0 in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-pa
(fro
Download
  1.9/1.9 MB 94.0 MB/s eta 0:00:00
```

	termcolor	4.4.0	pypi
	terminaltables	3.1.10	pypi
	threadpoolctl	3.5.0	pypi
	tifffile	2023.7.10	pypi
	tk	8.6.13	pypi
	tomli	2.2.1	conda-forge
	torch	1.9.0+cu111	pypi
	torchaudio	0.9.0	pypi
	torchvision	0.10.0+cu111	pypi
	tqdm	4.65.2	pypi
	typing	3.7.4.3	pypi
	typing_extensions	4.12.2	pyha770c72_0
	tzdata	2025.2	pypi
	urllib3	1.26.20	pypi

Please confirm again through 'conda list'
'+cu' is right
'cpu' is wrong

3. Install openmim, it's similar to pip

```
(cell-sam) [root@yuchen ~]# pip install -U openmim
Looking in indexes: https://pypi.org/simple, https://pypi.ngc.nv
Collecting openmim
  Downloading openmim-0.3.9-py2.py3-none-any.whl.metadata (16 kB)
Collecting Click (from openmim)
  Downloading click-8.1.8-py3-none-any.whl.metadata (2.3 kB)
Collecting colorama (from openmim)
  Downloading colorama-0.4.6-py2.py3-none-any.whl.metadata (17 kB)
Collecting model-index (from openmim)
  Downloading model_index-0.1.11-py3-none-any.whl.metadata (3.9 kB)
Collecting opendatalab (from openmim)
  Downloading opendatalab-0.0.10-py3-none-any.whl.metadata (6.4 kB)
Collecting pandas (from openmim)
  Downloading pandas-2.0.3-cp38-cp38-manylinux_2_17_x86_64.manylinux2014_x86_64.whl.metadata (1.1 MB)
Requirement already satisfied: pip>=19.3 in /home/yuchen/miniforum (19.3.3)
Collecting requests (from openmim)
  Downloading requests-2.32.4-py3-none-any.whl.metadata (4.9 kB)
Collecting rich (from openmim)
  Downloading rich-14.1.0-py3-none-any.whl.metadata (18 kB)
Collecting tabulate (from openmim)
  Downloading tabulate-0.9.0-py3-none-any.whl.metadata (34 kB)
Collecting pyyaml (from model-index->openmim)
  Downloading PyYAML-6.0.2-cp38-cp38-manylinux_2_17_x86_64.manylinux2014_x86_64.whl.metadata (1.1 MB)
Collecting markdown (from model-index->openmim)
  Downloading Markdown-3.7-py3-none-any.whl.metadata (7.0 kB)
Collecting ordered-set (from model-index->openmim)
```

4. Install mmengine , using mim

```
ch-13.4.2 setuptools-60.2.0 six-1.17.0 tabulate-0.9.0 tqdm-4.65.2
(cell-sam) [ $ mim install mmengine
Looking in indexes: https://pypi.org/simple https://pypi.gc.nv/
Looking in links: https://download.openmmlab.com/mmcv/dist/cu110/
Collecting mmengine
  Downloading mmengine-0.10.7-py3-none-any.whl.metadata (20 kB)
Collecting addict (from mmengine)
  Downloading addict-2.4.0-py3-none-any.whl.metadata (1.0 kB)
Collecting matplotlib (from mmengine)
  Downloading matplotlib-3.7.5-cp38-cp38-manylinux_2_12_x86_64.ma
Requirement already satisfied: numpy in /home/yuchen/miniforge3/e
Requirement already satisfied: pyyaml in /home/yuchen/miniforge3/
Requirement already satisfied: rich in /home/yuchen/miniforge3/en
Collecting termcolor (from mmengine)
  Downloading termcolor-2.4.0-py3-none-any.whl.metadata (6.1 kB)
Collecting yapf (from mmengine)
  Downloading yapf-0.43.0-py3-none-any.whl.metadata (46 kB)
Collecting opencv-python>=3 (from mmengine)
  Downloading opencv_python-4.12.0.88-cp37-abi3-manylinux2014_x86_
Collecting contourpy>=1.0.1 (from matplotlib->mmengine)
  Downloading contourpy-1.1.1-cp38-cp38-manylinux_2_17_x86_64.m
Collecting cycler>=0.10 (from matplotlib->mmengine)
  Downloading cycler-0.12.1-py3-none-any.whl.metadata (3.8 kB)
```

5. Install CUDA Toolkit by hand, if needed (If it is uncertain whether the server has pre configured CUDA, proceed with this step):

```
(cell-sam) [          ]$ conda install cudatoolkit=11.*  
Channels:  
- conda-forge  
Platform: linux-64  
Collecting package metadata (repodata.json): done  
Solving environment: done
```

6. The most important step: configure MMCV:

```
(cell-sam) [1]: mim install mmcv==2.0.*  
Looking in indexes: https://pypi.org/simple, https://pypi.nvidia.com  
Looking in links: https://download.openmmlab.com/mmcv/dist/cu111/torch1.9.0/index.html  
Collecting mmcv==2.0.*  
  Downloading https://download.openmmlab.com/mmcv/dist/cu111/torch1.9.0/mmcv-2.0.1-cp38-cp38-manylinux1_x86_64.whl (77.5 MB)  
    33.0/77.5 MB 14.5 MB/s eta 0:00:04
```

Wait for a few minutes. If there are no errors, the installation could be successful.

```
Requirement already satisfied: typing-extensions<5.0,>=4.0.0 in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-packages (from rich->mmengine)>=0.3.0->mmcv==2.0.*) (4.12.2)
Requirement already satisfied: zipp>=3.1.0 in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-packages (from importlib-resources>=3.2.0->matplotlib->mmengine)>=0.3.0->mmcv==2.0.*) (3.20.2)
Requirement already satisfied: mdurl=0.1 in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-packages (from markdown-it-py>=2.2.0->rich->mmengine)>=0.3.0->mmcv==2.0.*) (0.1.2)
Requirement already satisfied: six>=1.5 in /home/yuchen/miniforge3/envs/cell-sam/lib/python3.8/site-packages (from python-dateutil)=2.7>matplotlib>mmengine)>=0.3.0->mmcv==2.0.*) (1.17.0)
Installing collected packages: mmcv
Successfully installed mmcv-2.0.1
```

If there is an error, please check the above steps, such as whether the CUDA Toolkit is installed correctly, and try different versions of PyTorch (GPU version as a prerequisite), which can be fixed after a simple one or two version changes. Or find more solutions from [Issues · open-mmlab/mmdetection](#) (We have deployed it on more than 6 servers, including different CPUs and graphics cards (1080Ti, 3060, V100, 3080, 3090Ti, 4090), and different operating systems (Windows, Ubuntu, Arch Linux). Please report any bugs on your server to us.).

6. Install other used packages and SAM :

```
cd prompt_SAM; pip install -v -e .; pip install -r requirements.txt
```

```
Requirement already satisfied: cell2img==0.1.11 in /home/yuchen/mmlab/cell_SAM/Cell_SAM/requirements/optional.t  
s>cityscapesScripts->r /home/drew/Dropbox/Notebooks/yuchen/Cell_SAM/prompt_SAM/requirements/optional.t  
Downloading cython-3.1.2-cp38-cp38-manylinux_2_17_x86_64.manylinux2014_x86_64.whl (3.3 MB)  
 3.3/3.3 MB 82.9 MB/s eta 0:00:00  
Downloading cityscapesScripts-2.2.4-py3-none-any.whl (473 kB)  
Downloading imagecorruptions-1.1.2-py3-none-any.whl (2.1 kB)  
 2.1/2.1 MB 117.8 MB/s eta 0:00:00  
Downloading scikit_learn-1.3.2-cp38-cp38-manylinux_2_17_x86_64.manylinux2014_x86_64.whl (11.1 MB)  
 11.1/11.1 MB 117.1 MB/s eta 0:00:00  
Downloading joblib-1.4.2-py3-none-any.whl (301 kB)  
Downloading scikit_image-0.21.0-cp38-cp38-manylinux_2_17_x86_64.manylinux2014_x86_64.whl (13.9 MB)  
 13.9/13.9 MB 116.8 MB/s eta 0:00:00  
Downloading threadpoolctl-3.5.0-py3-none-any.whl (18 kB)  
Downloading appdirs-1.4.4-py2.py3-none-any.whl (9.6 kB)  
Downloading coloredlogs-15.0.1-py2.py3-none-any.whl (46 kB)  
Downloading quaternion-0.9.9-py3-none-any.whl (14 kB)  
Downloading humanfriendly-10.0-py2.py3-none-any.whl (86 kB)  
Downloading imageio-2.35.1-py3-none-any.whl (315 kB)  
Downloading lazy_loader-0.4-py3-none-any.whl (12 kB)  
Downloading networkx-3.1-py3-none-any.whl (2.1 MB)  
 2.1/2.1 MB 119.6 MB/s eta 0:00:00  
Downloading PyWavelets-1.4.1-cp38-cp38-manylinux_2_17_x86_64.manylinux2014_x86_64.whl (6.9 MB)  
 6.9/6.9 MB 117.4 MB/s eta 0:00:00  
Downloading tifffile-2023.7.10-py3-none-any.whl (220 kB)  
Building wheels for collected packages: typing  
 Building wheel for typing (setup.py) ... done
```

```
(cell-SAM) [1]: $ pip install git+https://github.com/facebookresearch/segment-anything.git  
Looking in indexes: https://pypi.org/simple, https://pypi.python.org/pypi  
Collecting git+https://github.com/facebookresearch/segment-anything.git  
  Cloning https://github.com/facebookresearch/segment-anything.git to /tmp/pip-req-build-ph5ikcxh  
    Running command git clone --filter=blob:none --quiet https://github.com/facebookresearch/segment-anything.git  
h5ikcxh  
  Resolved https://github.com/facebookresearch/segment-anything.git to commit dca509fe793f601edb92606367a655c15a  
  Preparing metadata (setup.py) ... done  
Building wheels for collected packages: segment-anything  
  Building wheel for segment-anything (setup.py) ... done  
  Created wheel for segment-anything: filename=segment Anything-1.0-py3-none-any.whl size=36623 sha256=99624b97ff4ee61b812dd4be546831e43d96f99140b  
  Stored in directory: /tmp/pip-ephem-wheel-cache-amnfwa_0/wheels/b0/7e/40/20f0ble23280cc4a66dc8009c29f42cb4afcl  
Successfully built segment-anything  
Installing collected packages: segment-anything  
Successfully installed segment-anything-1.0
```

Normally, SAM will automatically download the weight file called, but you can also manually download it, if SAM cannot access its website. We use 'sam_vit_b_01ec64.pth'

7. Test demo:

Download the dataset and weight files as instructed by GitHub, and place them in the required location. (prompt SAM)

-  **datasam** Put the data here
-  **outputs** Wait for the results here
-  **det4sam_spark_8xb32_r50-300e.py** Config file of the detection model
-  **detector_sam_demo.py** Python files combining SAM/SAM2 and detection model
-  **detector_sam2_demo.py**
-  **epoch_300.pth** Weight of the detection model
-  **sam_vit_b_01ec64.pth** Weight of SAM
-  **utils.py**

7. Test demo:

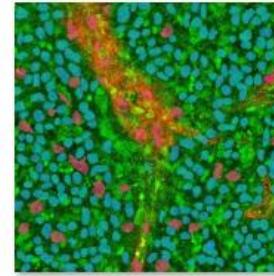
Download the dataset and weight files as instructed by GitHub, and place them in the required location. (finetune SAM)

- 📁 configs Config file of the model
 - 📁 datasets Put the data here →
 - 📁 load
 - 📁 models SAM/SAM2 models
 - 📁 visualizations Wait for the results here
 - 📄 __init__.py
 - 📄 LICENSE
 - 📄 model_finetune.pth Weight of the trained model
 - 📄 save.py
 - 📄 sod_metric.py
 - 📄 test.py
 - 📄 train.py
 - 📄 utils.py
- 📁 Ima Test data path
 - 📁 images Train data path
 - 📁 ma Test mask path
 - 📁 masks Train mask path

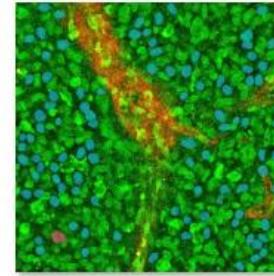
8. Get results:

Enter instructions, wait for the program to run, and obtain visible results.

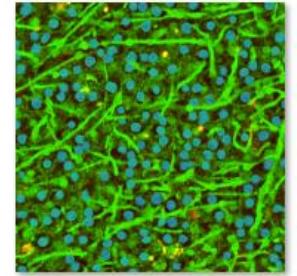
Prompt-SAM like these, in outputs



img3_slic2.jpg

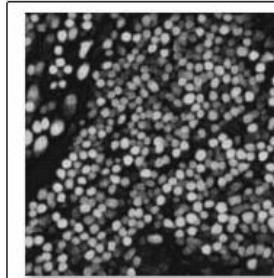


imgtest.jpg

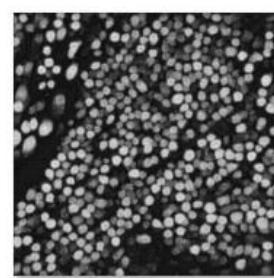


our.png

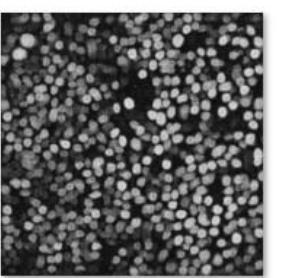
Finetuned-SAM like these , in visualizations



pred_0_0.png



pred_0_0-1.tif



pred_1_0.png