

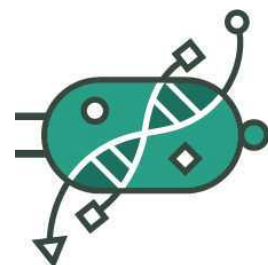
Dec 22, 2023

Version 2

OT-2 Protocol to transfer volume from several plates to a single plate V.2

DOI

<https://dx.doi.org/10.17504/protocols.io.6qpvr4o62gmk/v2>



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Protocol status: Working

We use this protocol and it's working

Created: November 20, 2023

Last Modified: December 22, 2023

Protocol Integer ID: 91172

Keywords: OT-2, Opentrons, Media transfer, 96-well, transposon library variant, throughput workflow for the genotypic characterization, samples of different source plate, description of the lap repository entry lap, fewer final plates than the source number, python script for an opentron, samples from different plate, lap repository entry lap, different source plate, ot2, protocol, fewer final plate, plate, different plate, single plate, opentron, nplatemerging, volumes of transfer, transfer, final plate, laboratory, python, type of plate, throughput workflow, robot, set of instruction, original identities of the sample

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Abstract

This protocol is meant to transfer samples from different plates to a single or fewer final plates than the source number, in other words, merge samples of different source plates in a final plate.

The output of running this script will be the final plate(s) with samples and a reactive (optional), and the corresponding map(s) with the original identities of the samples which will be given by the user in the input file.

This protocol uses a python script for an Opentrons 2 robot and an excel file containing the required variables to set the number of samples, volumes of transfer, type of plates, etc...

In our laboratory, this protocol has been used as part of the "High-throughput workflow for the genotypic characterization of transposon library variants" also available in protocols.io

This protocol is a set of instructions or description of the [LAP repository](#) entry **LAP-NplateMerging-OT2-1.0.0**



Guidelines

This protocol was developed with python 3.7.1, OT App Software Version 6.3.1 and API level version 2.14 in a Linux 4.14.74 system (these are the OT-2 specifications).

In the script several packages are used: pandas (0.25.3), openpyxl (3.1.2), math, random and numpy (1.15.1)

It has been tested with cultures from *Pseudomonas putida* KT2440 as part of the High-throughput workflow for the genotypic characterization of transposon library variants.

The maximum number of 96-well plates per run is 8 source plates to 1 final plates dispensing some media (need of 1 falcon tube rack), 1 pipette is required for all the volumes and the replace of tip racks is allowed (set as True in the input variable file)

Materials

Software

- Python 3.7.1
- opentrons software version 6.3.1
- python packages: pandas (0.25.3), openpyxl (3.1.2), numpy(1.15.1), math, random
- OT App
- Excel

OT-2 Labware

- Opentrons Tip racks

Equipment	
Opentrons 96 Tip Rack 300 µL	NAME
Tip rack	TYPE
Opentrons	BRAND
-	SKU
https://labware.opentrons.com/opentrons_96_tiprack_300ul?category=tipRack ^{LINK}	

Equipment	
Opentrons 96 Tip Rack 20 µL	NAME
Tip rack	TYPE
Opentrons	BRAND
-	SKU
https://labware.opentrons.com/opentrons_96_tiprack_20ul?category=tipRack ^{LINK}	

- 96- well plates

Equipment

96-well plates, flat bottom, non treated

NAME

Cell culture plates

TYPE

VWR

BRAND

734-2781

SKU

https://es.vwr.com/store/catalog/product.jsp?catalog_number=734-2781^{LINK}



- Opentrons Falcon tube rack

Equipment

Opentrons 15 Tube Rack with Falcon 15 mL Conical

NAME

OT Tube Rack

TYPE

Opentrons

BRAND

-

SKU

https://labware.opentrons.com/opentrons_15_tuberack_falcon_15ml_conical/^{LINK}



- 15mL Falcon tubes

Equipment	
Falcon® Conical Centrifuge Tubes 15mL	NAME
Flaocn Tube	TYPE
Falcon	BRAND
352096	SKU
https://ecatalog.corning.com/life-sciences/b2c/US/en/Liquid-Handling/Tubes,-Liquid-Handling/Centrifuge-Tubes/Falcon%C2%AE-Conical-Centrifuge-Tubes/p/falconConicalTubes	LI N K

Equipment:

Equipment	
OT-2	NAME
Liquid handler	TYPE
Opentrons	BRAND
OT-2	SKU

Equipment

Single Channel Electronic Pipette (GEN2) 300uL NAME

Opentrons Pipette TYPE

Opentrons BRAND

- SKU

<https://shop.opentrons.com/single-channel-electronic-pipette-p20/> LINK

Equipment

Single Channel Electronic Pipette (GEN2) 20uL NAME

Opentrons Pipette TYPE

Opentrons BRAND

- SKU

<https://shop.opentrons.com/single-channel-electronic-pipette-p20/> LINK

Before start

It is important to annotate the location of the source plate in the OT-2 slots, the identity of the samples is provided by excel sheets in the input variable and if 2 samples, within a plate or between different source plates has the same name, they will be indistinguishable in the final plate (s)

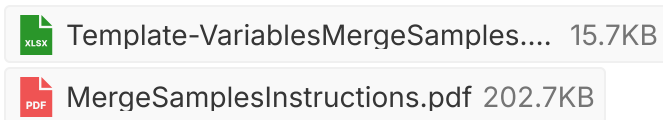
Files Preparation

1 Preparing Customized Template

Preparing the template (a .xlsx) with the specific variables for each experiment.

Attached there is a template of the variable file with several sheets and a PDF file explaining each variable:

1. **GeneralVariables:** variables related to the labware to be used
2. **PipetteVariables:** variables related to the pipettes to be used
3. **PerPlateVariables:** variables related to the specifications of each source plate
4. **Maps (Optional):** sheet(s) with the names of the samples in the source plates. These will be reflected in the final plate map → *not included in the template but needs to be included and have the same names as established in the variable **Name Sheet Map Identifiers** from the PerPlateVariablesSheet*



1.1 *Fill the template with the corresponding values*

1.2 *Save it with the name VariablesMergeSamples.xlsx*

Note

The file should be spelt **exactly** *VariablesMergeSamples.xlsx* or the Python script won't work correctly

2 Transferring *VariablesMergeSamples.xlsx* file to Robot

Transfer the *VariablesMergeSamples.xlsx* to the directory */data/user_storage* of the OT robot that we are going to use to perform the protocol.



Note

Before transferring any file to the OT, we need to know the **IP of the robot**.

This can be obtained in the Networking section of the Device that we will use. To obtain this info go to **OT-App -> Devices -> Chosen Robot (three dots) -> Robot Settings -> Networking**

In this tab, you can see 2 types of IP; one is shown if both the robot and you are connected to the same Wifi, and the other is shown if the computer and the robot are connected via USB. Both connections can be used for this step

Note

To connect to the robot an **OT-key** should have been previously generated, and it is done with the `ssh-keygen` command and **transferring the public key to the OT**.

For more information about how to generate and set the connection between your computer and the Opentrons robot, visit <https://support.opentrons.com/s/article/Setting-up-SSH-access-to-your-OT-2>

Here, we present a summary of how to transfer the files in 3 Operative Systems:
Windows, MacOS and Linux

MacOS/Linux

We will use the command line with `scp` to transfer the file `VariablesMergeSamples.xlsx` to the OT system.

We need to perform the following command

Command

File passing from our computer to robot's linux (OT raspberry)

Passing Files to OT

```
scp -i [ot_key] [file] root@[IP_OT]:/data/user_storage
```



Note

You could face difficulties transferring files in MacOS Ventura (13) and Sonoma (14). These problems can be solved by adding the argument -O (uppercase o) to the command

Command

Transferring files to OT (MacOS 13 and 14)

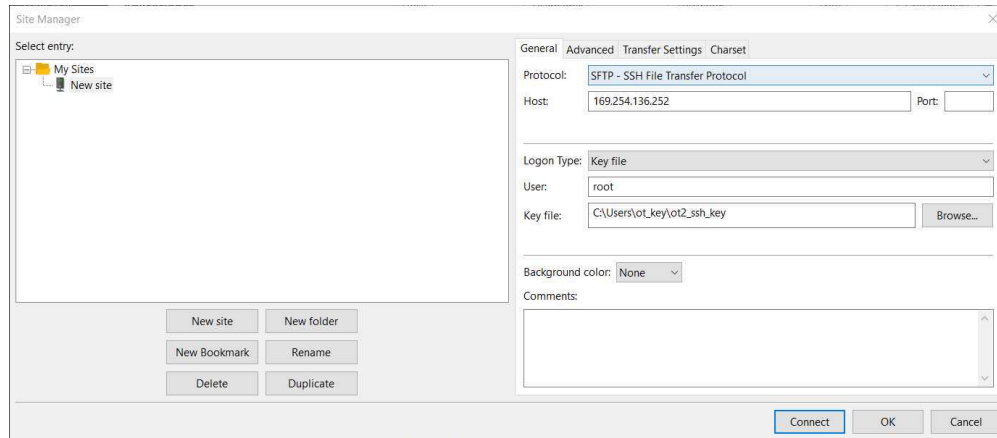
```
scp -Oi [ot_keypath] [file path]  
root@[OP_robot]:/data/user_storage
```

Windows

There are several ways to send files from a Windows to a Linux (for example, with a virtual machine or Windows Powershell in the latest versions of Windows).

Here, we will use **FileZilla** (<https://filezilla-project.org/download.php?type=client>).

Go to **File → Site Manager → New Site → Change Protocol to SFTP**. Then, introduce in Host the OT IP, change the *Logon Type* to key file, change *User* to root and give the directory where the ot key is. It should look something like this



Example of setting the FileZilla to transfer files from Windows (our computer) to Linux (OT)

Then press *Connect*, and we will have a connection between our computer and the robot.

After this connection, we should be able to move the file *VariablesMergeSamples.xlsx* (in our computer) to the directory */data/user_storage* in the robot.

This method can be used as well in some Linux and MacOS

Note

Take into account that the IP of the robot could change, so it is possible that it will be needed to change the host in these connections from time to time.

3 Adding the custom labware



There is only a need to do this step when the labware that you are using is not OT official or is not included in the OT app

3.1 Creation of .json file

The description file can be obtained by describing the labware dimensions at <https://labware.opentrons.com/create/>

3.2 Uploading files to the OT App

In the OT app, we need to perform the following route: **Labware → Import → Choose File → Select file we have created in step 3.1**



Expected result

After uploading the labware you should be able to see the new labware in the Labware tab of the OT App, all custom labware can be found more easily in the category *Custom Labware*

3.3 Transfer labware files to the robot

If you are using the entry **LAP-NplateMerging-OT2-1.0.0 and custom labware**, an additional step is needed, which is transferring a folder with the custom labware

We need to create for our custom labware a folder with the API name containing the description file (.json) called 1.json and then transfer that folder to the robot's folder `/data/labware/v2/custom_definitions/custom_beta` in a similar way as in the Step 2 but with the difference that is a directory that needs to be transferred and not a file.

Command

Transferring the custom labware to OT (Linux)

```
scp -i [ot_key] -r [directory_custom_labware]  
root@[IP_OT]:/data/labware/v2/custom_definitions/custom_beta
```

Note

We do not need to execute this part every time the protocol is used, only when that labware is not included in the OT official labware and these directories are not in the robot

Prepare RobotOS

4 Install needed packages



This script needs the package *openpyxl*, which is not installed by default in the OT-2 robots

Note

This step is only needed if the package is not installed in the robot, not every run of the protocol

If the package is not installed, an error will appear when running the script in the robot. While simulating the script in the app, this error will appear, but you can ignore it

4.1 *Connect to the robot*



to find the IP of the robot in which you want to run the script

To connect to the robot, you can do it via ssh with the following command

Command

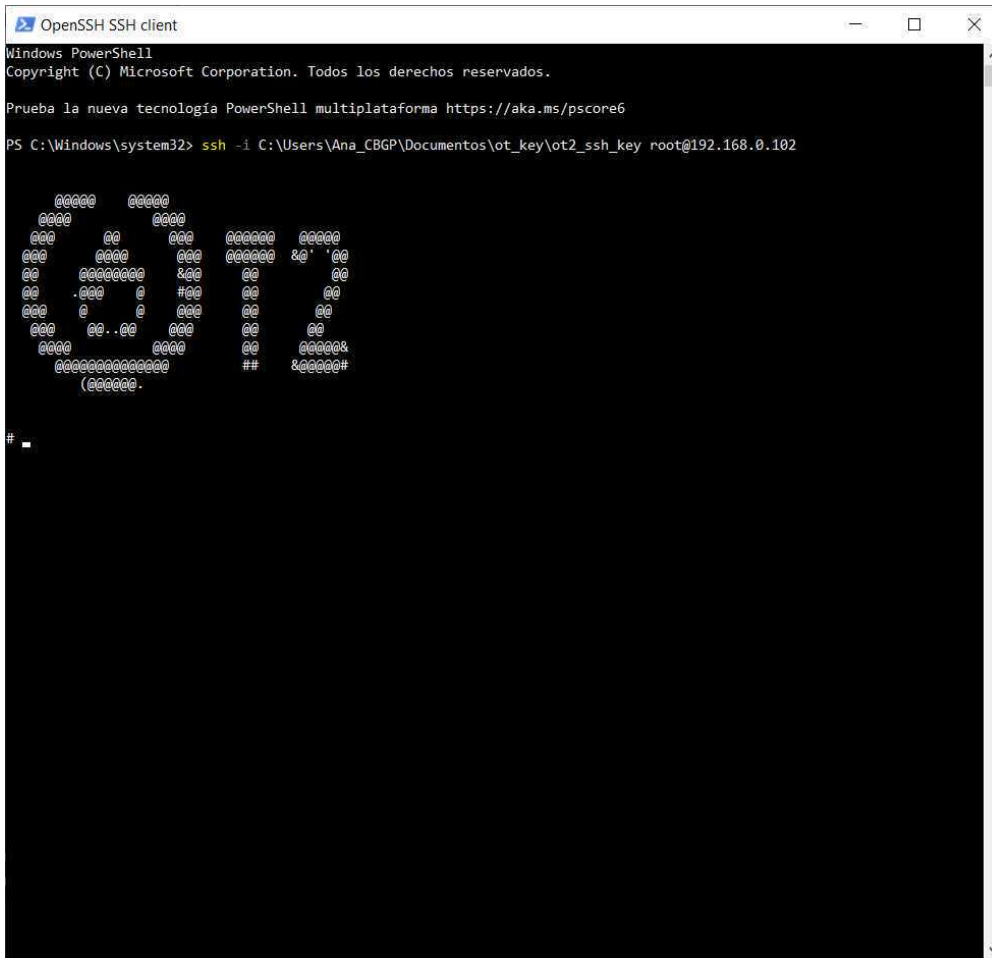
Connect to Linux based OT via ssh

```
ssh -i [path ot_key] root@[Robot_IP]
```

In Windows you can do this command in Windows Powershell

Expected result

If the connection has been successful you should obtain a screen similar to the following image



Drawing obtained when entering an OT-2 system

4.2 Install the package

Once inside the robot's system, you need to run the following command



Command

Install openpyxl package (Linux 4.14.74-v7)

```
pip install openpyxl
```

Note

For more information about installing packages in the opentrons robots, check the following Opentrons page: <https://support.opentrons.com/s/article/Using-Python-packages-in-Python-API-protocols>

Running Protocol

5 Load script in OT-App

Now that we have transferred the variable files to the robot, we can import the script and run it in the selected robot

Note

This whole step has been developed with version 6.3.1 of the OT-App and has been tested until version 7.0.2

Indications may vary from version to version of the opentrons App and the version of the script.



Software

Opentrons App

NAME

Windows >=10, Mac >=10 , Ubuntu >=12.04

OS

Opentrons

DEVELOPER

<https://opentrons.com/ot-app/>

REPOSITORY

5.1 Load the script in the App

Protocols → Import → Drag Python script

Note

The last script version can be found at

<https://github.com/BiocomputationLab/LAPrepository/tree/main/LAPEntries> (the name of this file is the user's choice). The name of the directory should be **LAP-NplateMerging-OT2** followed by the version.

As well we can find the latest version of the script at

<https://www.laprepo.cbgp.upm.es/repository/> with the same name as in GitHub

Software

LAP Repository

NAME

<https://biocomputationlab.com/>

DEVELOPER

www.laprepo.com

REPOSITORY



Note

The App with version 6.3.1 analyzes your protocol before setting a robot to run, so the labware will not be shown before assigning the protocol to a specific robot when you import it into the App

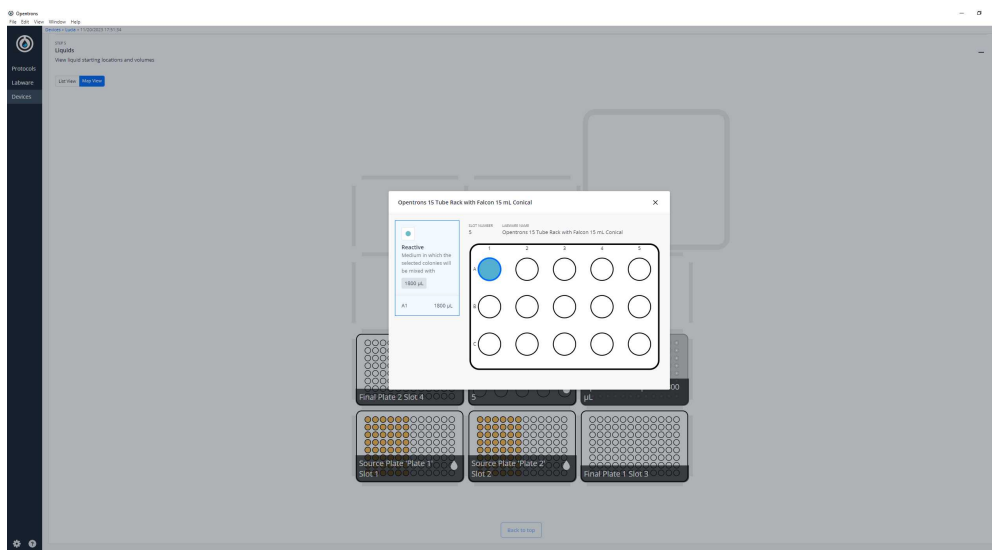
5.2 *Select Robot to Perform Script*

Click in the protocol → Start setup → Choose the OT where the file *VariablesMergeSamples.xlsx* is → Proceed To Setup

After clicking on Proceed to Setup, you should obtain, the positions of the labware in the *Labware* tab and in the *Liquids* tab, the reagents and their corresponding volume.

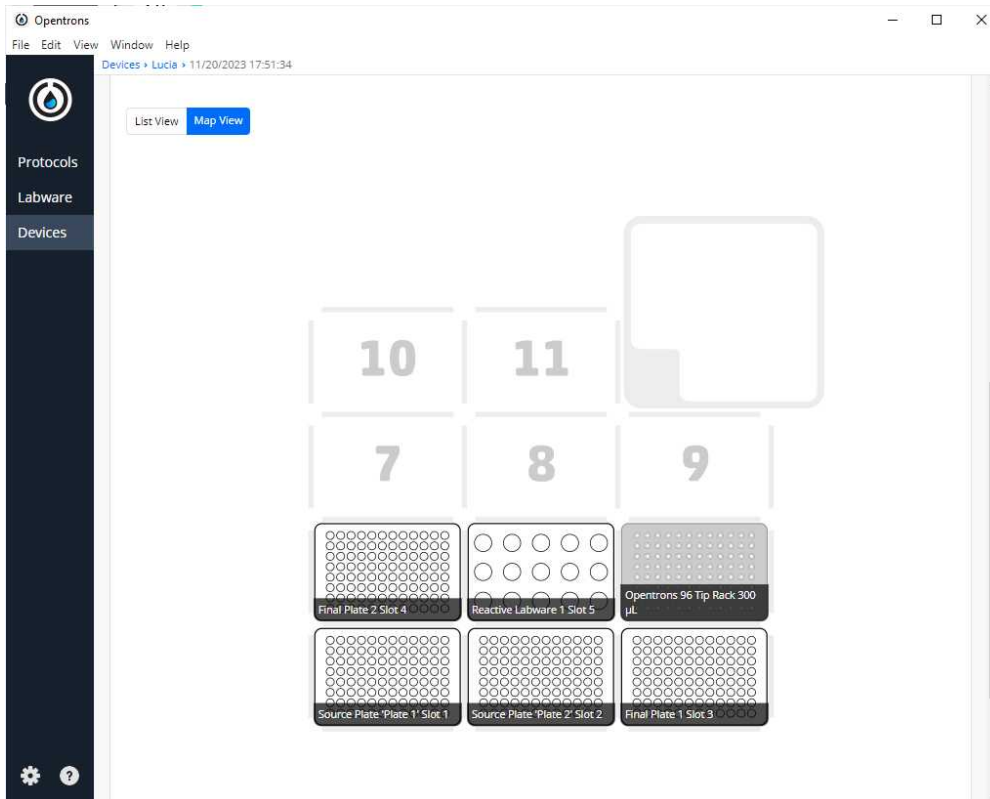
Expected result

A **liquid setup** should look like the following image, where you can find the samples in the initial plates and the reactives in the Falcon labware



Liquid Set-up Example of a Samples Merge Protocol

A **labware setup** should look like the following image, where you can find the initial and final plates, the falcon labware (if needed) to store the reagent and the corresponding tip rack(s)



Labware Set-up example of a Samples Merge Protocol

Note

The volume of the initial samples is established to be 90% of the max volume of the well, but it is only a recommendation. **Just make sure that there is enough volume to transfer to the final plate(s).**

The volume of the reagents is exactly what is needed, so it is **suggested to pour always more to take in account the error of pipetting**

Note

It is recommended that you perform a labware position check.

You can do it with test plates after loading the script but before cleaning the surface. That way, you reduce the probability of contamination (using the test plates and labware) and pipetting errors (position check).

6 Run Protocol in OT



6.1 *Make sure the needed calibrations are done*

Pipettes, tip racks and tip length calibrations need to be done for the items used in this run

6.2 *Labware position check is performed (if required)*



6.3 *Clean the surface of the robot with 70% ethanol to clean and disinfect the surfaces*

Note

Check the Opentrons page <https://support.opentrons.com/s/article/Cleaning-your-OT-2?> for more information about cleaning the OT-2 robot with the proper materials

6.4 *Set the labware and reagents as shown in the OT-App*

6.5 *Start Run*

The procedure that the robot is going to do is mainly divided into 3 parts:

1. Select which samples, taking into account the variable *Type of Sample Selection*
2. (Optional) Distribute reactive(s)
3. Distribute selected samples

Expected result

One or more plates, with a reactive mixed with the samples from different sources or only the samples. In addition to an **Excel file where every sheet is the map of a final plate located in the folder /data/user_storage** that will give the position in these plates with their identifiers (the name given in the maps of each source plate in the input variable file)

After-Running

7 **Retrieve labware from the OT**

8 Import map from robot

There will be a map with the name set in the variable *Name File Final Map* set in the sheet *GeneralVariables* followed by the extension .xlsx: *[Name File Final Map].xlsx*

This file will be located in the directory */data/user_storage* in the robot where the script has been run.

To retrieve the file, we can  [go to step #2](#) and reproduce it by transferring the files from the robot to the computer.

Command

Transferring files from OT to computer (Linux, macOS)

```
scp -i [path_ot_key]  
root@[IP_robot]:/data/user_storage/[name_map].xlsx  
[ultimate_path_computer]
```

Expected result

The sheets in the final map contain the identity of the samples selected in the places where they have been placed/distributed

These identities are going to be the names given in the map Excel sheets from the input file

Example

1h 5m

- 9 **We want to merge 16 samples from 5 source plates in a different way, 1 of them selecting samples from back to front, 2 in a random way and 2 from front to back. Our final plate will have half the plate already occupied by other samples, and we want to add to all the samples water, so we need to take into account those details**

We will use a computer with a Windows 10 system

9.1 Prepare variable file

10m

Excel template that we can find [go to step #1](#) filled and saved with the name *VariablesMergeSamples.xlsx*



VariablesMergeSamples.xlsx 22.6KB

9.2 Upload custom labware to app

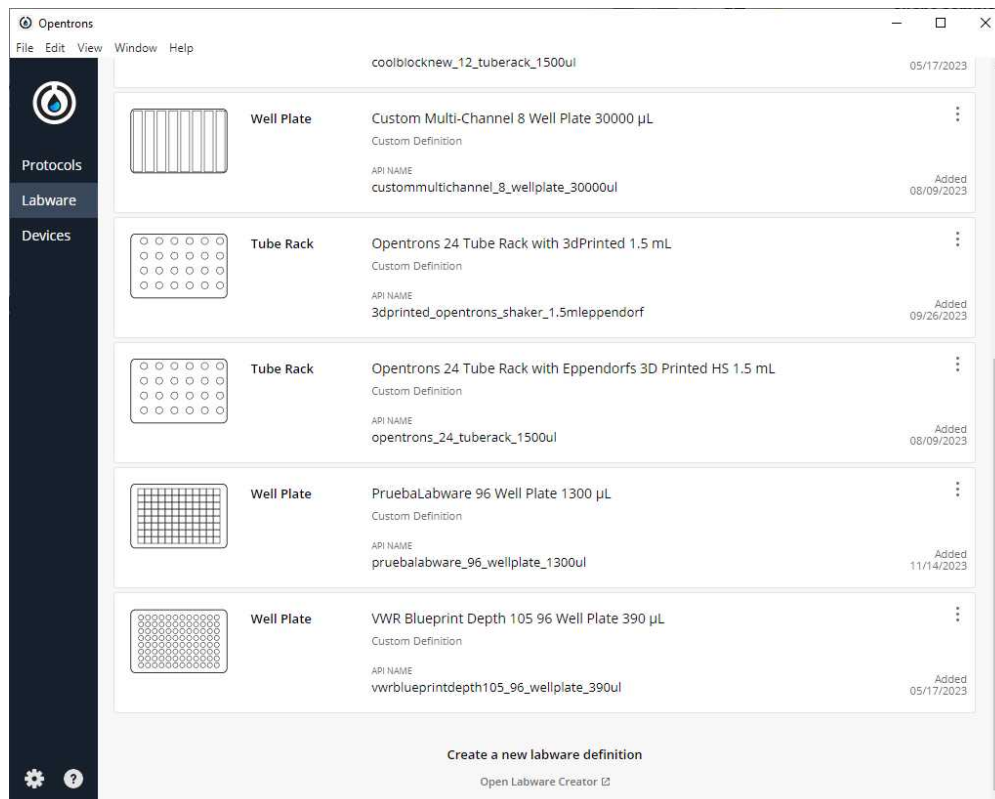
2m

We are using a custom labware called *vwrblueprintdepth105_96_wellplate_390ul* that has been created with the labware creator that opentrons offers (<https://labware.opentrons.com/create/>)



vwrblueprintdepth105_96_wellplate... 11.7KB

Upload it to the labware and make sure it is loaded in the app

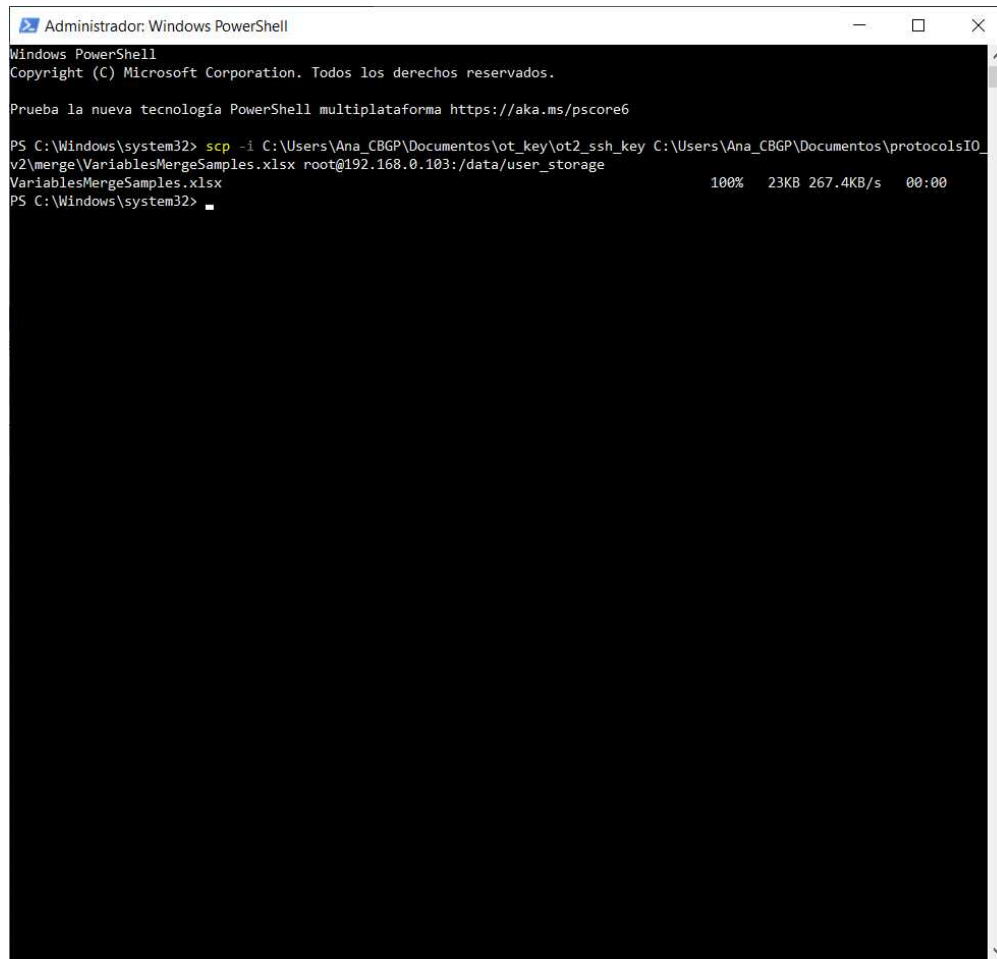


List of custom labware recorded in the Opentrons App



9.3 Export the variable file to the */data/user_storage* folder in the robot

2m



```
Administrador: Windows PowerShell
Windows PowerShell
Copyright (C) Microsoft Corporation. Todos los derechos reservados.

Prueba la nueva tecnología PowerShell multiplataforma https://aka.ms/pscore6

PS C:\Windows\system32> scp -i C:\Users\Ana_CBG\Documents\ot_key\ot2_ssh_key C:\Users\Ana_CBG\Documents\protocolsIO_v2\merge\VariablesMergeSamples.xlsx root@192.168.0.103:/data/user_storage
VariablesMergeSamples.xlsx                               100% 23KB 267.4KB/s 00:00
PS C:\Windows\system32>
```

command line window with scp commands to transfer the variables .xlsx from our computer to the OT-2

9.4 Because we are using version 1.0.0 of the script in this example, we will transfer the directory of the labware as well (here we have attached a zip, but it is the folder that must be transferred, not the zip)

1m

 vwrblueprintdepth105_96_wellplate... 1.8KB

Command

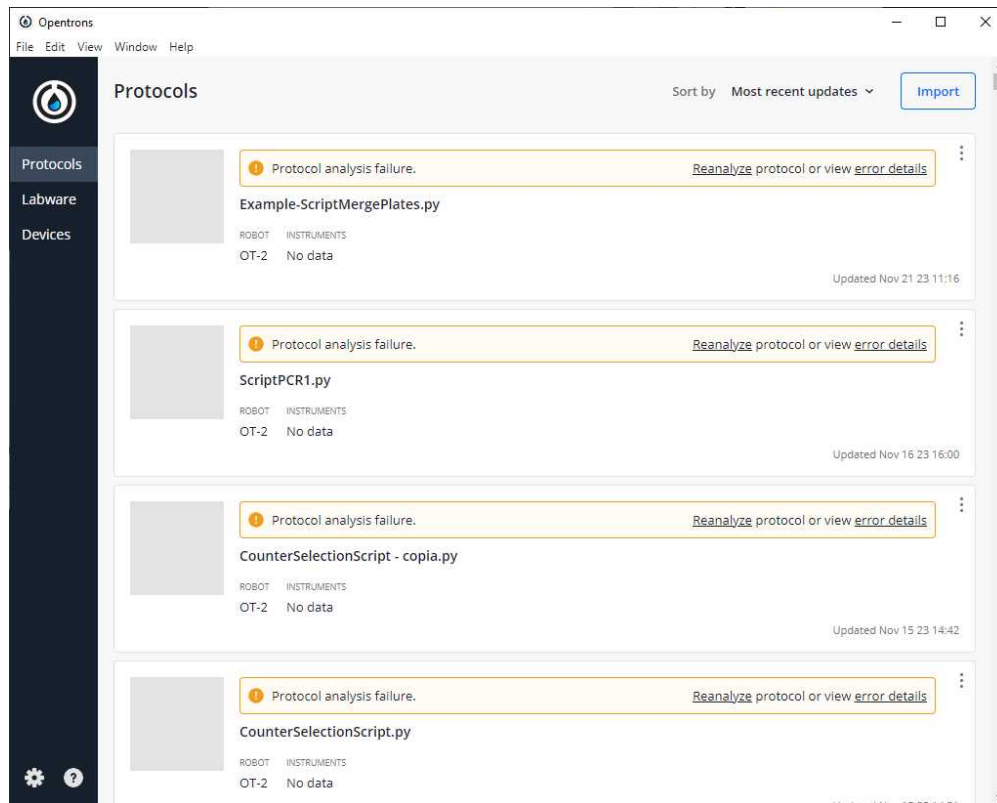
Transferring the used custom labware to OT (Linux)

```
scp -i [ot_key] -r vwrblueprintdepth105_96_wellplate_390ul
root@[IP_OT]:/data/labware/v2/custom_definitions/custom_beta
```

- 9.5 Import the script that we have downloaded from the step [go to step #5.1](#) (I named it *Example-ScriptMergePlates.py*) to the OT-App

30s

 Example-ScriptMergePlates.py 47.3KB



Result of importing the Python script into the OT-App

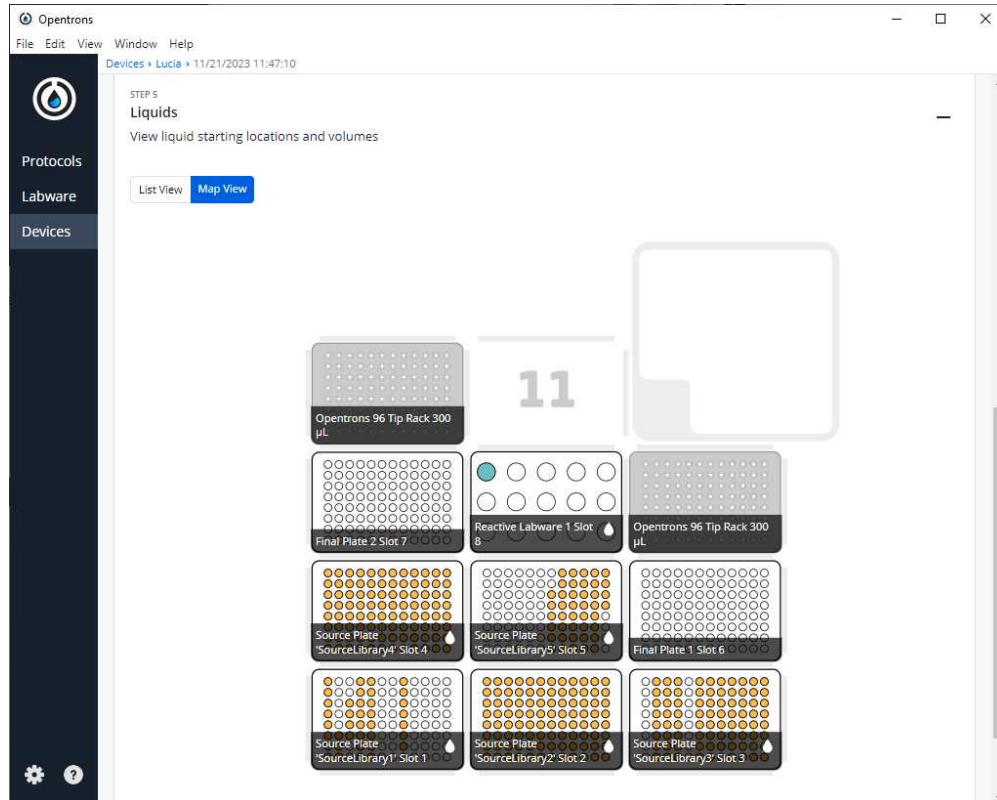
As we can see, we have an error, but that is programmed because the script is meant to work in the robot but not in your computer

9.6 Run the protocol in the robot that we have transferred the Excel file

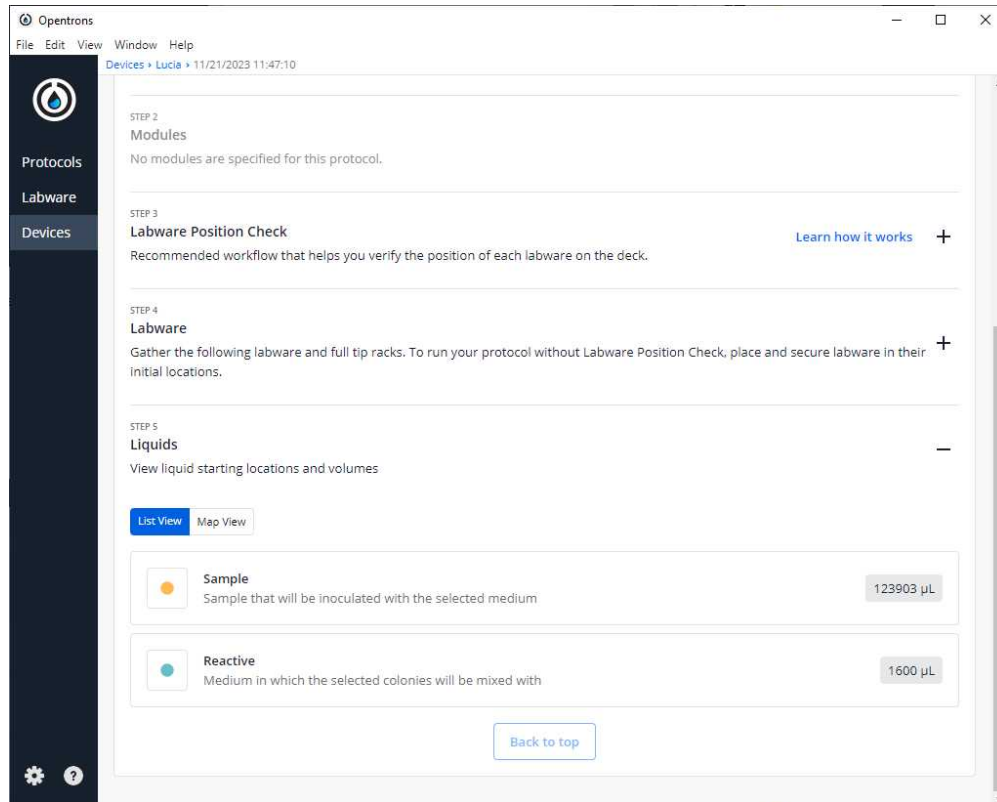
1m

Example-ScriptMergePlates.py → **Start setup** → **Select the robot** in which we are going to run the protocol

If we do not have any errors, the output should look similar to the following pictures



Labware and liquid set-up layout



Volumes of the needed liquids to perform the protocol

9.7 Turn the HEPA filter module

30s



9.8 Clean platform of the robot that we are going to perform the protocol

2m

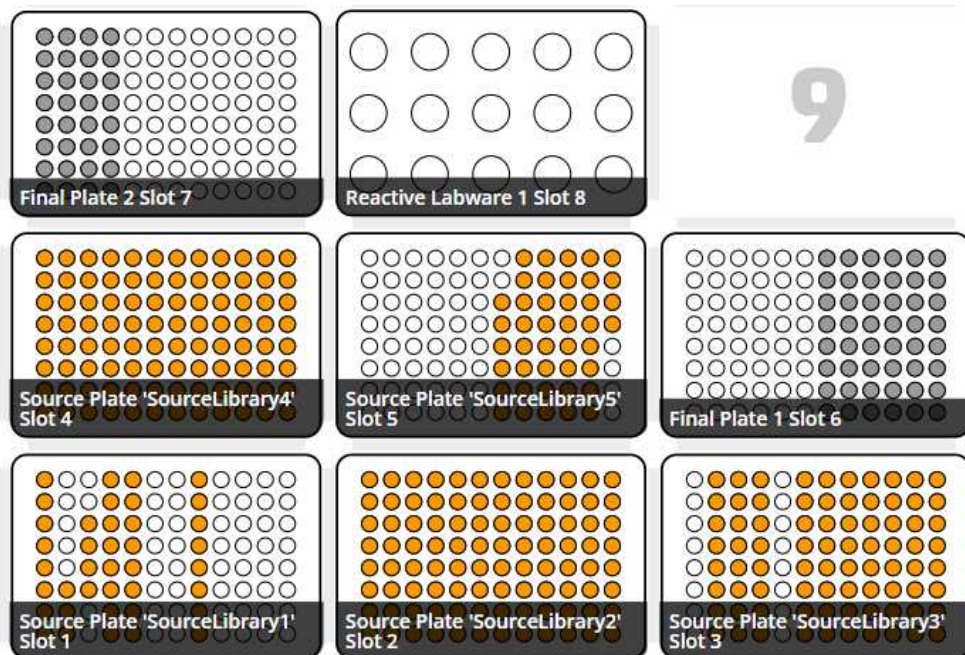
9.9 Prepare all reagents and labware in the places the App is showing and take into account the notes in step  Notes

5m

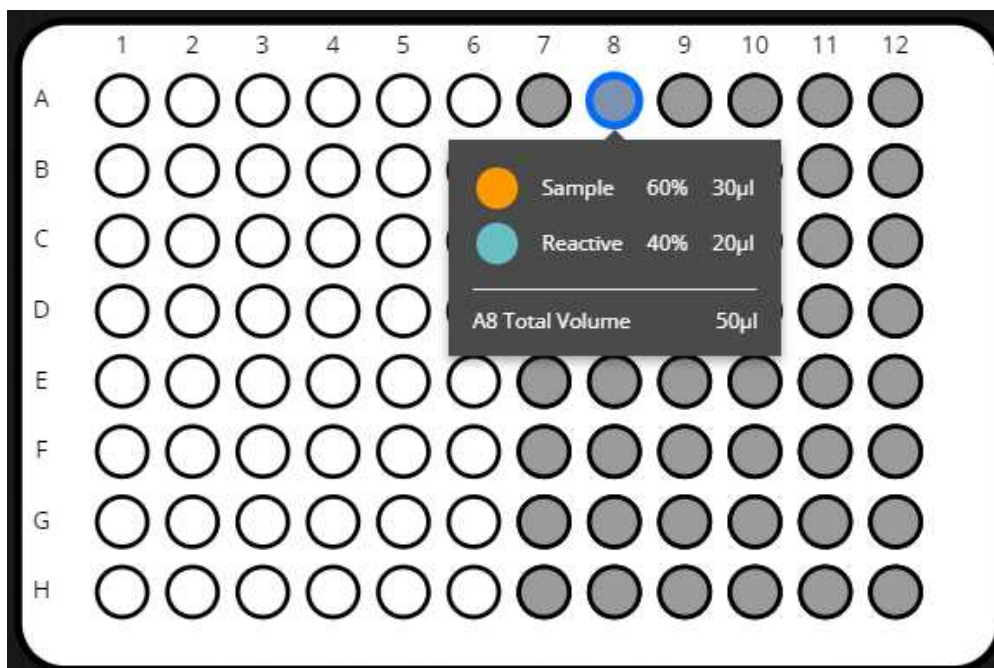
9.10 Start run

35m

Expected result



Final layout of source and final plates in the run



Example of the content of A8 in the labware *Final Plate 1 Slot 6*

Here we will obtain the mix between the volume of the reactive, in this case, water, and the samples selected by the values in the variable file. These positions are seen in the image by the grey wells, and we can see the info of the plate and reactive in the plate on slot 6 in the second picture



9.11 Retrieve labwares from the OT

5m

9.12 Retrieve the final maps, in this case, the map file will be called *example-mapMerge.xlsx* (name that is stated in the variable file in the variable *Name File Final Map*)

1m

```
Administrator: Windows PowerShell
PS C:\Windows\system32> scp -i C:\Users\Ana_CBG\Documents\ot_key\ot2_ssh_key root@192.168.0.103:/data/user_storage/example-mapMerge.xlsx C:\Users\Ana_CBG\Documents\protocolsIO_v2\merge
example-mapMerge.xlsx 100% 6455 905.7KB/s 00:00
PS C:\Windows\system32>
```

Command line windows with the transfer command of the final plate(s) map(s) file from the OT to our computer

 example-mapMerge.xlsx 10KB



Protocol references

pBLAM1-x: standardized transposon tools for high-throughput screening (*Synthetic Biology*)

<https://doi.org/10.1093/synbio/ysad012>

The Laboratory Automation Protocol (LAP) Format and Repository: A Platform for Enhancing Workflow Efficiency in Synthetic Biology (*ACS Synth. Biol.*) <https://doi.org/10.1021/acssynbio.3c00397>