

Infection rate determination

DAPI and NP segmentation



Name your images correctly from the start

Please label your data as followed:

date_condition_descriptionYouWouldlike

Info: Please use „_“ for separating different informations, otherwise you have to change it in the code.

Example:

3.11.2023_Female_H7N7_MOI0.05_6h_IBA1,CD68,Ki67_20x.czi

If you code your condition like this:

3.11.2023_A1_IBA1,CD68,Ki67_20x.czi

Please decode your data prior to the following analysis (you can use another code for this or do it manual).

```

27 # =====
28 # Please specify your channel and pixel size
29 # =====
30
31 DAPI = 0
32 NP = 2
33 pixels_to_um = 0.2930000 #size of pixel
34
35 # =====
36 # Please specify your conditions for plotting
37 # =====
38
39 # Define condition names for labels and actual condition strings
40 # if you do not have 2 belonging conditions (e.g. sex), you can name them here
41 # Important: Make sure that the value is exactly like this in your filename!
42
43 condition_definitions = {
44     'Female 6h': 'Female_H7N7_MOI0.05_6h',
45     'Male 6h': 'Male_H7N7_MOI0.05_6h',
46     'Female 24h': 'Female_H7N7_MOI0.05_24h',
47     'Male 24h': 'Male_H7N7_MOI0.05_24h',
48 }
49
50
51 #####
52 # Functions
53 #####
54
55 # =====
56 # The following functions need to be altered for different users as the naming
57 # of your files is important here
58 # =====
59
60 def ConditionFinder (filename):
61     condition = filename.split("_")[1:5]
62     condition = '_'.join(condition[0:])
63
64     return condition
65
66
67 def RoundFinder(filename):
68     cell_round = filename.split("_")[0]
69
70     return cell_round
71

```

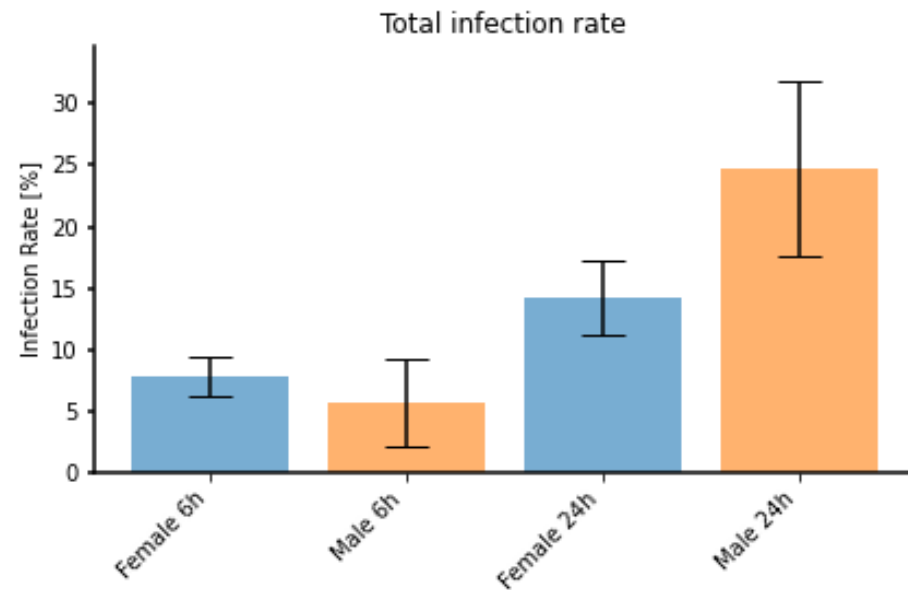
Please select the channel correctly! **Important**, otherwise you will create wrong data!!! And please add your pixel size, for size filtering.

Please enter here your condition names. Only important for plotting, if you do not want your data to be plotted, you do not need this here.

If you did the naming differently then recommended, please change it here accordingly!

Run the code

- You need: your data in one folder and the correct labelling
- Press on run and select the folder with your image data
- You get:



Infection rates per condition

	A	B	C	D	E
1	round	condition	DAPI_count	NP_count	NP_%
2	9.8.23	Female_H7N	143.5	21.5	14.2185877
3	9.8.23	Female_H7N	145	11.5	7.77041625
4	9.8.23	Male_H7N7_I	61.5	17	24.5538665
5	9.8.23	Male_H7N7_I	142.5	8	5.62641584

Example segmentation

