Combing through cell counts

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1 Combing through cell counts

1.0.1 Jonathan Ramos 3/6/2024

Looks like there were still a few subtle discrepancies in the mean cell ns. Let's just check once more to enusre that we are counting only completed colocalized groupings.

The idea here is to consider each image in our set an independent network of nodes and vertices (as in from graph theory) where each stain type (row of data) represents a node. If any number (up to 4) of unique staintypes are colocalized they should can be represented as adjacent nodes in a directed graph. If a colocalization is "true" or "real" then each node in a colocalized group should point to every other node in the grouping, i.e. if a PV is colocalized with a WFA, then that WFA is also colocalized with that PV. In graph theory, this is called a complete subgraph (or clique in an undirected graph). This way, we expect that each colocalized grouping (or complete subgraph) is therefore also disjoint from any other colocalized grouping; that is, our complete subgraphs do not overlap with each other. This means that a given roi_id can only ever be a part of a single colocalized grouping.

[81]:
```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import ast
import sys
import statsmodels.api as sm
from statsmodels.formula.api import ols

# loading some functions we wrote before
sys.path.append("/Users/jonathanramos/Desktop/LRI/Image ROI Data Wrangling/")
from clean import *
from norm import *
from count import *
```

2 Load in sets

[82]:
```
df_coloc = pd.read_csv('KET-VR5_FULL_SET.csv').drop('Unnamed: 0', axis=1)

# literal eval for true_grouping tuples
df_coloc['true_grouping_literal'] = df_coloc.true_grouping.apply(ast.
 ↪literal_eval)

print(df_coloc.columns)
print(df_coloc.shape)
df_coloc.head()
```

```
Index(['index', 'filename', 'image_name', 'roi_id', 'true_grouping',
       'dummy_PV', 'dummy_cFos', 'dummy_Npas4', 'dummy_WFA', 'CoM_x', 'CoM_y',
       'background', 'mean_intensity', 'stain_type', 'filename.1', 'rat_n',
       'treatment', 'group_name', 'snr', 'mean-background',
       'adjusted_mean-background', 'true_grouping_literal'],
      dtype='object')
(18632, 22)
```

[82]:

	index	filename	image_name	roi_id	true_grouping	dummy_PV	dummy_cFos	dummy_Npas4	dummy_WFA	CoM_x	…	mean_intensity	stain_type	filename.1	rat_n	treatment	group_name	snr
0	0	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00000_PV	('0-000-00000_PV', '0-FFF-00045_Npas4', '0-FFF…	True	False	True	True	297.86	…	536.8331	PV	KET-10-12_PFC_3.7_A_2.tif	KET-10-12	FR1_KET	KET-10	2.247372
1	1	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00001_PV	('0-000-00001_PV', '0-FFF-00070_cFos', '0-FFF-…	True	True	True	False	340.47	…	314.9278	PV	KET-10-12_PFC_3.7_A_2.tif	KET-10-12	FR1_KET	KET-10	1.318398
2	2	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00002_PV	('0-000-00002_PV', '0-FFF-00044_Npas4', '0-FFF…	True	False	True	True	154.85	…	324.0556	PV	KET-10-12_PFC_3.7_A_2.tif	KET-10-12	FR1_KET	KET-10	1.356610
3	3	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00003_PV	('0-000-00003_PV', '0-FFF-00082_Npas4', '0-FFF…	True	False	True	True	310.10	…	346.0313	PV	KET-10-12_PFC_3.7_A_2.tif	KET-10-12	FR1_KET	KET-10	1.448608
4	4	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00004_PV	('0-000-00004_PV',)	True	False	False	False	44.35	…	429.6127	PV	KET-10-12_PFC_3.7_A_2.tif	KET-10-12	FR1_KET	KET-10	1.798510

```
  mean-background  adjusted_mean-background  \
0      297.961600                 382.72460
1       76.056290                 160.81929
2       85.184100                 169.94710
3      107.159805                 191.92280
4      190.741200                 275.50420

                                true_grouping_literal
0  (0-000-00000_PV, 0-FFF-00045_Npas4, 0-FFF-0000…
1  (0-000-00001_PV, 0-FFF-00070_cFos, 0-FFF-00012…
2  (0-000-00002_PV, 0-FFF-00044_Npas4, 0-FFF-0000…
3  (0-000-00003_PV, 0-FFF-00082_Npas4, 0-FFF-0000…
4                                (0-000-00004_PV,)

[5 rows x 22 columns]
```

3 Update true groupings

[122]:
```
images = [df_coloc.query(f'image_name == "{im}"') for im in df_coloc.image_name.
↪unique()]

updated_true = []
for i, df_img in enumerate(images):
    # group by true_grouping_literal, then get arr of unique roi_ids that had␣
↪that grouping
    df_grouped = df_img.groupby('true_grouping_literal')['roi_id'].unique().
↪reset_index(name='roi_ids')

    # check if the length of the true grouping equals the length of the arr of␣
↪roi ids we just found
    df_grouped['matching_len'] = df_grouped.apply(lambda x: len(x.
↪true_grouping_literal) == len(x.roi_ids), axis=1)

    # update true grouping, cast arr of roi_ids with a given grouping to a tuple
    df_grouped['updated_true_grouping'] = df_grouped.roi_ids.apply(lambda x:␣
↪tuple(x))

    # rename for merge
    df_grouped = df_grouped.rename(columns= {'roi_ids': 'roi_id'})

    # select, then join
    updated_true.append(df_img.merge(df_grouped.explode('roi_id')[['roi_id',␣
↪'matching_len', 'updated_true_grouping']]))
```

```
# concat updated true groupings
df_coloc_updated = pd.concat(updated_true)
```

[125]:
```
# quickly check our result
images = [df_coloc_updated.query(f'image_name == "{im}"') for im in␣
 ↪df_coloc_updated.image_name.unique()]

updated_true = []
for i, df_img in enumerate(images):
    df_grouped = df_img.groupby('updated_true_grouping')['roi_id'].unique().
 ↪reset_index(name='roi_ids')
    df_grouped['matching_len'] = df_grouped.apply(lambda x: len(x.
 ↪updated_true_grouping) == len(x.roi_ids), axis=1)

    # no assertion fails.. whew :-)
    assert df_grouped.matching_len.all()
```

4 Building new dummies

[126]:
```
def get_dummies(x):
    groupings = [rid.split('_')[-1] for rid in x]

    dummy_PV = False
    dummy_cFos = False
    dummy_Npas4 = False
    dummy_WFA = False

    if 'PV' in groupings:
        dummy_PV = True
    if 'cFos' in groupings:
        dummy_cFos = True
    if 'Npas4' in groupings:
        dummy_Npas4 = True
    if 'WFA' in groupings:
        dummy_WFA = True

    return dummy_PV, dummy_cFos, dummy_Npas4, dummy_WFA

df_coloc_updated['dummy'] = df_coloc_updated.updated_true_grouping.
 ↪apply(get_dummies)
df_coloc_updated['dummy_PV'], df_coloc_updated['dummy_cFos'],␣
 ↪df_coloc_updated['dummy_Npas4'], df_coloc_updated['dummy_WFA'] =␣
 ↪zip(*df_coloc_updated['dummy'])

df_coloc_updated
```

[126]:

	index	filename	image_name	roi_id	\
0	0	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00000_PV	
1	1	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00001_PV	
2	2	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00002_PV	
3	3	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00003_PV	
4	4	KET-10-12_PFC_3.7_A_2.tif	KET-10-12_PFC_3.7_A	0-000-00004_PV	
..	…	…	…	…	
118	898	PE-13-9_PFC_4.0_B_5.tif	PE-13-9_PFC_4.0_B	0-FFF-00008_WFA	
119	899	PE-13-9_PFC_4.0_B_5.tif	PE-13-9_PFC_4.0_B	0-FFF-00009_WFA	
120	900	PE-13-9_PFC_4.0_B_5.tif	PE-13-9_PFC_4.0_B	0-FFF-00010_WFA	
121	901	PE-13-9_PFC_4.0_B_5.tif	PE-13-9_PFC_4.0_B	0-FFF-00011_WFA	
122	902	PE-13-9_PFC_4.0_B_5.tif	PE-13-9_PFC_4.0_B	0-FFF-00012_WFA	

	true_grouping	dummy_PV	dummy_cFos	\
0	('0-000-00000_PV', '0-FFF-00045_Npas4', '0-FFF…	True	False	
1	('0-000-00001_PV', '0-FFF-00070_cFos', '0-FFF-…	True	True	
2	('0-000-00002_PV', '0-FFF-00044_Npas4', '0-FFF…	True	False	
3	('0-000-00003_PV', '0-FFF-00082_Npas4', '0-FFF…	True	False	
4	('0-000-00004_PV',)	True	False	
..	…	…	…	
118	('0-FFF-00008_WFA',)	False	False	
119	('0-000-00002_PV', '0-FFF-00009_WFA')	True	False	
120	('0-FFF-00010_WFA',)	False	False	
121	('0-000-00005_PV', '0-FFF-00011_WFA')	True	False	
122	('0-000-00000_PV', '0-FFF-00002_Npas4', '0-FFF…	True	False	

	dummy_Npas4	dummy_WFA	CoM_x	…	rat_n	treatment	group_name	\
0	True	True	297.86	…	KET-10-12	FR1_KET	KET-10	
1	True	False	340.47	…	KET-10-12	FR1_KET	KET-10	
2	True	True	154.85	…	KET-10-12	FR1_KET	KET-10	
3	True	True	310.10	…	KET-10-12	FR1_KET	KET-10	
4	False	False	44.35	…	KET-10-12	FR1_KET	KET-10	
..	…	…	…	…	…	…	…	
118	False	True	385.12	…	PE-13-9	VR5_SAL	PE-13	
119	False	True	409.79	…	PE-13-9	VR5_SAL	PE-13	
120	False	True	300.24	…	PE-13-9	VR5_SAL	PE-13	
121	False	True	414.17	…	PE-13-9	VR5_SAL	PE-13	
122	True	True	203.68	…	PE-13-9	VR5_SAL	PE-13	

	snr	mean-background	adjusted_mean-background	\
0	2.247372	297.961600	382.724600	
1	1.318398	76.056290	160.819290	
2	1.356610	85.184100	169.947100	
3	1.448608	107.159805	191.922800	
4	1.798510	190.741200	275.504200	
..	…	…	…	
118	1.087687	9.611397	26.986496	

```
119  1.175773       19.266396                36.641495
120  1.129718       14.218300                31.593400
121  1.099090       10.861198                28.236298
122  1.028254        3.096893                20.471992

                                true_grouping_literal matching_len  \
0    (0-000-00000_PV, 0-FFF-00045_Npas4, 0-FFF-0000…         True
1    (0-000-00001_PV, 0-FFF-00070_cFos, 0-FFF-00012…         True
2    (0-000-00002_PV, 0-FFF-00044_Npas4, 0-FFF-0000…         True
3    (0-000-00003_PV, 0-FFF-00082_Npas4, 0-FFF-0000…         True
4                                    (0-000-00004_PV,)         True
..                                                 …          …
118                                (0-FFF-00008_WFA,)         True
119                 (0-000-00002_PV, 0-FFF-00009_WFA)         True
120                                (0-FFF-00010_WFA,)         True
121                 (0-000-00005_PV, 0-FFF-00011_WFA)         True
122  (0-000-00000_PV, 0-FFF-00002_Npas4, 0-FFF-0001…         True

                                updated_true_grouping  \
0    (0-000-00000_PV, 0-FFF-00045_Npas4, 0-FFF-0000…
1    (0-000-00001_PV, 0-FFF-00070_cFos, 0-FFF-00012…
2    (0-000-00002_PV, 0-FFF-00044_Npas4, 0-FFF-0000…
3    (0-000-00003_PV, 0-FFF-00082_Npas4, 0-FFF-0000…
4                                    (0-000-00004_PV,)
..                                                 …
118                                (0-FFF-00008_WFA,)
119                 (0-000-00002_PV, 0-FFF-00009_WFA)
120                                (0-FFF-00010_WFA,)
121                 (0-000-00005_PV, 0-FFF-00011_WFA)
122  (0-000-00000_PV, 0-FFF-00002_Npas4, 0-FFF-0001…

                          dummy
0      (True, False, True, True)
1      (True, True, True, False)
2      (True, False, True, True)
3      (True, False, True, True)
4    (True, False, False, False)
..                           …
118  (False, False, False, True)
119   (True, False, False, True)
120  (False, False, False, True)
121   (True, False, False, True)
122    (True, False, True, True)

[18632 rows x 25 columns]
```

5 Do we match?

[177]:
```
import itertools

# do our doubles agree?
print('double labeled differences: ')
for stain_x, stain_y in itertools.combinations(['PV', 'cFos', 'Npas4',␣
↪'WFA'],2):
    x_on_y = df_coloc_updated.query(f'dummy_{stain_x} == True and␣
↪dummy_{stain_y} == True and stain_type == "{stain_x}"')
    y_on_x = df_coloc_updated.query(f'dummy_{stain_x} == True and␣
↪dummy_{stain_y} == True and stain_type == "{stain_y}"')
    diff = x_on_y.__len__() - y_on_x.__len__()
    print(f'{stain_x}, {stain_y}:    {diff}')

# do our quads agree?
print('\n\nquad labeled ns: ')
quads = df_coloc_updated.query('dummy_PV == True and dummy_cFos == True and␣
↪dummy_Npas4 == True and dummy_WFA == True')
for stain in ['PV', 'cFos', 'Npas4', 'WFA']:
    q = quads.query(f'stain_type == "{stain}"')
    print(stain,': ', q.__len__())

# looks like cFos has all the issues here.
```

```
double labeled differences:
PV, cFos:    -3
PV, Npas4:    0
PV, WFA:    0
cFos, Npas4:    4
cFos, WFA:    3
Npas4, WFA:    0


quad labeled ns:
PV :  173
cFos :  175
Npas4 :  173
WFA :  173
```

6 Investigating cFos

[201]:
```
df_PV_cFos = df_coloc_updated.query('dummy_PV == True and dummy_cFos == True␣
↪and (stain_type == "PV" or stain_type == "cFos")')
df_PV_cFos_paired = df_PV_cFos.groupby(['image_name',␣
↪'updated_true_grouping'])['roi_id'].unique().reset_index(name='paired')
```

```
df_PV_cFos_paired['n'] = df_PV_cFos_paired.paired.apply(lambda x: len(x))

print(np.array(df_PV_cFos_paired[df_PV_cFos_paired.n != 2].paired.to_list()))
df_PV_cFos_paired[df_PV_cFos_paired.n != 2]
```

```
[['0-000-00006_PV' '0-005-00057_cFos' '0-005-00069_cFos']
 ['0-000-00005_PV' '0-005-00019_cFos' '0-FFF-00045_cFos']
 ['0-000-00008_PV' '0-005-00024_cFos' '0-005-00026_cFos']]
```

[201]:

	image_name	updated_true_grouping	\
159	KET-10-3_PFC_3.8_B	(0-000-00006_PV, 0-005-00057_cFos, 0-005-00069…	
177	KET-10-4_PFC_3.7_D	(0-000-00005_PV, 0-005-00019_cFos, 0-FFF-00045…	
281	KET-8-7_PFC_3.7_C	(0-000-00008_PV, 0-005-00024_cFos, 0-005-00026…	

	paired	n
159	[0-000-00006_PV, 0-005-00057_cFos, 0-005-00069…	3
177	[0-000-00005_PV, 0-005-00019_cFos, 0-FFF-00045…	3
281	[0-000-00008_PV, 0-005-00024_cFos, 0-005-00026…	3

[202]:
```
df_WFA_cFos = df_coloc_updated.query('dummy_WFA == True and dummy_cFos == True␣
↪and (stain_type == "WFA" or stain_type == "cFos")')
df_WFA_cFos_paired = df_WFA_cFos.groupby(['image_name',␣
↪'updated_true_grouping'])['roi_id'].unique().reset_index(name='paired')
df_WFA_cFos_paired['n'] = df_WFA_cFos_paired.paired.apply(lambda x: len(x))

print(np.array(df_WFA_cFos_paired[df_WFA_cFos_paired.n != 2].paired.to_list()))
df_WFA_cFos_paired[df_WFA_cFos_paired.n != 2]
```

```
[['0-005-00057_cFos' '0-005-00069_cFos' '0-FFF-00003_WFA']
 ['0-005-00024_cFos' '0-005-00026_cFos' '0-FFF-00010_WFA']
 ['0-FFF-00062_cFos' '0-FFF-00063_cFos' '0-FFF-00007_WFA']]
```

[202]:

	image_name	updated_true_grouping	\
85	KET-10-3_PFC_3.8_B	(0-000-00006_PV, 0-005-00057_cFos, 0-005-00069…	
144	KET-8-7_PFC_3.7_C	(0-000-00008_PV, 0-005-00024_cFos, 0-005-00026…	
242	KET-9-6_PFC_3.6_E	(0-FFF-00062_cFos, 0-FFF-00063_cFos, 0-200-000…	

	paired	n
85	[0-005-00057_cFos, 0-005-00069_cFos, 0-FFF-000…	3
144	[0-005-00024_cFos, 0-005-00026_cFos, 0-FFF-000…	3
242	[0-FFF-00062_cFos, 0-FFF-00063_cFos, 0-FFF-000…	3

[203]:
```
df_Npas4_cFos = df_coloc_updated.query('dummy_Npas4 == True and dummy_cFos ==␣
↪True and (stain_type == "Npas4" or stain_type == "cFos")')
df_Npas4_cFos_paired = df_Npas4_cFos.groupby(['image_name',␣
↪'updated_true_grouping'])['roi_id'].unique().reset_index(name='paired')
df_Npas4_cFos_paired['n'] = df_Npas4_cFos_paired.paired.apply(lambda x: len(x))
```

```
print(np.array(df_Npas4_cFos_paired[df_Npas4_cFos_paired.n != 2].paired.
 ↪to_list()))
df_Npas4_cFos_paired[df_Npas4_cFos_paired.n != 2]
```

```
[['0-005-00057_cFos' '0-005-00069_cFos' '0-FFF-00051_Npas4']
 ['0-005-00019_cFos' '0-FFF-00045_cFos' '0-200-00000_Npas4']
 ['0-005-00024_cFos' '0-005-00026_cFos' '0-FFF-00088_Npas4']
 ['0-FFF-00062_cFos' '0-FFF-00063_cFos' '0-200-00003_Npas4']]
```

[203]:

	image_name	updated_true_grouping	\
1185	KET-10-3_PFC_3.8_B	(0-000-00006_PV, 0-005-00057_cFos, 0-005-00069…	
1295	KET-10-4_PFC_3.7_D	(0-000-00005_PV, 0-005-00019_cFos, 0-FFF-00045…	
1989	KET-8-7_PFC_3.7_C	(0-000-00008_PV, 0-005-00024_cFos, 0-005-00026…	
2587	KET-9-6_PFC_3.6_E	(0-FFF-00062_cFos, 0-FFF-00063_cFos, 0-200-000…	

	paired	n
1185	[0-005-00057_cFos, 0-005-00069_cFos, 0-FFF-000…	3
1295	[0-005-00019_cFos, 0-FFF-00045_cFos, 0-200-000…	3
1989	[0-005-00024_cFos, 0-005-00026_cFos, 0-FFF-000…	3
2587	[0-FFF-00062_cFos, 0-FFF-00063_cFos, 0-200-000…	3

6.1 Narrowing down our search

By inspecting the dataframes and print outs shown above I've narrowed my search down to 8 suspect cFos cells. We can see that these groupings imply 4 cells (2 quads, and two triples) that are each colocalized with 2 cFos cells. The dataframes above overlap on the following cFos cells, suggesting one of each of the following pairs: - KET-10-3_PFC_3.8_B : '0-005-00057_cFos' or '0-005-00069_cFos' - KET-10-4_PFC_3.7_D : '0-005-00019_cFos' or '0-FFF-00045_cFos' - KET-8-7_PFC_3.7_C : '0-005-00024_cFos' or '0-005-00026_cFos' - KET-9-6_PFC_3.6_E : '0-FFF-00062_cFos' or '0-FFF-00063_cFos'

The above search also implies that PV, WFA and Npas4 all agree with each other and since Npas4 is paired with each of our suspect cFos hits, we can just use the Npas4 roi_id to determine which cFos roi_id truly belongs.

[264]:
```
def tie_breaker(im, offending_rid):
    df_image = df_coloc_updated.query(f'image_name == "{im}"')
    target = df_image.query(f'roi_id == "{offending_rid}"').
 ↪updated_true_grouping.to_list()[0]
    df_extra = df_image[df_image.updated_true_grouping == target][['roi_id',␣
 ↪'stain_type', 'CoM_x', 'CoM_y']]
    df_extra['coord'] = list(zip(df_extra.CoM_x, df_extra.CoM_y))

    sus_rids = [rid for rid in target if 'cFos' in rid]
    sus_coords = [(rid, df_extra.query(f'roi_id == "{rid}"').coord.item()) for␣
 ↪rid in sus_rids]
```

```
    for rid, sus_coord in sus_coords:
        df_extra[f'dist_{rid}'] = df_extra.apply(lambda x: distance(x.coord,␣
↪sus_coord), axis=1)

    df_extra = df_extra[(df_extra != 0).all(1)]
    df_extra['image_name'] = im

    return df_extra

extra_cfos1 = tie_breaker('KET-10-3_PFC_3.8_B', '0-005-00057_cFos')
extra_cfos2 = tie_breaker('KET-10-4_PFC_3.7_D', '0-005-00019_cFos')
extra_cfos3 = tie_breaker('KET-8-7_PFC_3.7_C', '0-005-00026_cFos')
extra_cfos4 = tie_breaker('KET-9-6_PFC_3.6_E', '0-FFF-00063_cFos')

extra_cfos = pd.concat([extra_cfos1, extra_cfos2, extra_cfos3, extra_cfos4])
extra_cfos
```

[264]:

	roi_id	stain_type	CoM_x	CoM_y	coord	\
5	0-000-00006_PV	PV	329.96	397.15	(329.96, 397.15)	
138	0-FFF-00051_Npas4	Npas4	324.83	400.60	(324.83, 400.6)	
166	0-FFF-00003_WFA	WFA	332.21	390.56	(332.21, 390.56)	
5	0-000-00005_PV	PV	253.98	469.11	(253.98, 469.11)	
73	0-200-00000_Npas4	Npas4	252.09	461.34	(252.09, 461.34)	
8	0-000-00008_PV	PV	428.22	444.75	(428.22, 444.75)	
185	0-FFF-00088_Npas4	Npas4	429.76	440.65	(429.76, 440.65)	
203	0-FFF-00010_WFA	WFA	431.42	443.67	(431.42, 443.67)	
102	0-200-00003_Npas4	Npas4	332.80	358.93	(332.8, 358.93)	
203	0-FFF-00007_WFA	WFA	336.03	361.95	(336.03, 361.95)	

	dist_0-005-00057_cFos	dist_0-005-00069_cFos	image_name	\
5	1.989271	7.724066	KET-10-3_PFC_3.8_B	
138	5.016393	2.750455	KET-10-3_PFC_3.8_B	
166	8.933678	14.502090	KET-10-3_PFC_3.8_B	
5	NaN	NaN	KET-10-4_PFC_3.7_D	
73	NaN	NaN	KET-10-4_PFC_3.7_D	
8	NaN	NaN	KET-8-7_PFC_3.7_C	
185	NaN	NaN	KET-8-7_PFC_3.7_C	
203	NaN	NaN	KET-8-7_PFC_3.7_C	
102	NaN	NaN	KET-9-6_PFC_3.6_E	
203	NaN	NaN	KET-9-6_PFC_3.6_E	

	dist_0-005-00019_cFos	dist_0-FFF-00045_cFos	dist_0-005-00024_cFos	\
5	NaN	NaN	NaN	
138	NaN	NaN	NaN	
166	NaN	NaN	NaN	
5	2.539016	13.877684	NaN	
73	5.596713	5.970301	NaN	

```
8                      NaN                     NaN                1.721046
185                    NaN                     NaN                6.005231
203                    NaN                     NaN                4.207089
102                    NaN                     NaN                     NaN
203                    NaN                     NaN                     NaN

     dist_0-005-00026_cFos  dist_0-FFF-00062_cFos  dist_0-FFF-00063_cFos
5                      NaN                     NaN                     NaN
138                    NaN                     NaN                     NaN
166                    NaN                     NaN                     NaN
5                      NaN                     NaN                     NaN
73                     NaN                     NaN                     NaN
8                 3.950215                     NaN                     NaN
185               0.430116                     NaN                     NaN
203               3.164838                     NaN                     NaN
102                    NaN                6.865712                9.485383
203                    NaN                3.513702               12.939876
```

6.1.1 Success!

By computing distances, I've computationally decided tie breakers. No eye balling required. in summary, we have the following coloc cFos cells: - KET-10-3_PFC_3.8_B : '0-005-00057_cFos' - KET-10-4_PFC_3.7_D : '0-005-00019_cFos' - KET-8-7_PFC_3.7_C : '0-005-00026_cFos' - KET-9-6_PFC_3.6_E : '0-FFF-00062_cFos'

and the following cFos cells will be relabeled as single-labeled cFos only (lonely cfos) - KET-10-3_PFC_3.8_B : '0-005-00069_cFos' - KET-10-4_PFC_3.7_D : '0-FFF-00045_cFos' - KET-8-7_PFC_3.7_C : '0-005-00024_cFos' - KET-9-6_PFC_3.6_E : '0-FFF-00063_cFos'

[353]:
```
df_coloc_updated = df_coloc_updated.drop('index', axis=1).reset_index()

target_images = df_Npas4_cFos_paired[df_Npas4_cFos_paired.n != 2].image_name.
 ↪values
target_cfos = [[rid for rid in x if 'cFos' in rid] for x in␣
 ↪df_Npas4_cFos_paired[df_Npas4_cFos_paired.n != 2].paired.values]

true_coloc_cfos = ['0-005-00057_cFos', '0-005-00019_cFos', '0-005-00026_cFos',␣
 ↪'0-FFF-00062_cFos']
single_cfos = [rid for rid in np.ravel(target_cfos) if not rid in␣
 ↪true_coloc_cfos]

targets = zip(target_images, single_cfos)

for im, single_rid in targets:
    single_i = df_coloc_updated.query(f'image_name == "{im}" and roi_id ==␣
 ↪"{single_rid}"').index
    grouping = df_coloc_updated.iloc[single_i,:].updated_true_grouping.item()
```

```
    updated_grouping = tuple(roi_id for roi_id in grouping if roi_id !=␣
↪single_rid)

    # update lonely cfos
    df_coloc_updated.at[single_i.item(), 'updated_true_grouping'] =␣
↪tuple([single_rid])
    df_coloc_updated.at[single_i.item(), 'dummy_PV'] = False
    df_coloc_updated.at[single_i.item(), 'dummy_Npas4'] = False
    df_coloc_updated.at[single_i.item(), 'dummy_WFA'] = False

    # update grouping of all other true coloc stain types
    for rid in updated_grouping:
        coloc_i = df_coloc_updated.query(f'image_name == "{im}" and roi_id ==␣
↪"{rid}"').index
        df_coloc_updated.at[coloc_i.item(), 'updated_true_grouping'] =␣
↪updated_grouping
```

6.1.2 now let's see if all our cell ns match

[361]:
```
# do our doubles agree?
print('double labeled differences: ')
for stain_x, stain_y in itertools.combinations(['PV', 'cFos', 'Npas4', 'WFA'],␣
↪r=2):
    x_on_y = df_coloc_updated.query(f'dummy_{stain_x} == True and␣
↪dummy_{stain_y} == True and stain_type == "{stain_x}"')
    y_on_x = df_coloc_updated.query(f'dummy_{stain_x} == True and␣
↪dummy_{stain_y} == True and stain_type == "{stain_y}"')
    diff = x_on_y.__len__() - y_on_x.__len__()
    print(f'{stain_x}, {stain_y}:    {diff}')

# do our triples agree?
def check_triple_ns(comb):
    stain_x, stain_y, stain_z = comb
    q = df_coloc_updated.query(
        f'dummy_{stain_x} == True and dummy_{stain_y} == True and␣
↪dummy_{stain_z} == True and\
         (stain_type == "{stain_x}" or stain_type == "{stain_y}" or stain_type␣
↪== "{stain_z}")'
    )

    q_x = q.query(f'stain_type == "{stain_x}"')
    q_y = q.query(f'stain_type == "{stain_y}"')
    q_z = q.query(f'stain_type == "{stain_z}"')

    print(f'\ntriple {stain_x},{stain_y},{stain_z} ns:')
    print(stain_x, ' :', q_x.__len__())
```

```
    print(stain_y, ' :', q_y.__len__())
    print(stain_z, ' :', q_z.__len__())

for comb in itertools.combinations(['PV', 'cFos', 'Npas4', 'WFA'], r=3):
    check_triple_ns(comb)

# do our quads agree?
print('\n\nquad labeled ns: ')
quads = df_coloc_updated.query('dummy_PV == True and dummy_cFos == True and␣
 ↪dummy_Npas4 == True and dummy_WFA == True')
for stain in ['PV', 'cFos', 'Npas4', 'WFA']:
    q = quads.query(f'stain_type == "{stain}"')
    print(stain,': ', q.__len__())

# looks like cFos has all the issues here.
```

```
double labeled differences:
PV, cFos:    0
PV, Npas4:    0
PV, WFA:    0
cFos, Npas4:    0
cFos, WFA:    0
Npas4, WFA:    0

triple PV,cFos,Npas4 ns:
PV  : 353
cFos  : 353
Npas4  : 353

triple PV,cFos,WFA ns:
PV  : 270
cFos  : 270
WFA  : 270

triple PV,Npas4,WFA ns:
PV  : 252
Npas4  : 252
WFA  : 252

triple cFos,Npas4,WFA ns:
cFos  : 224
Npas4  : 224
WFA  : 224


quad labeled ns:
PV :  173
```

```
cFos :  173
Npas4 :  173
WFA :  173
```

6.2 GREAT. now we can write to disk and proceed with our analyses

[362]:
```
df_coloc_updated.to_csv('KET-VR5_FINAL.csv')
```