

v19_220601_no_imputation

June 20, 2022

0.1 Load Data

```
[1]: import pandas as pd
import numpy as np
import os
import glob
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt
import missingno as msno
from sklearn.cluster import KMeans
import seaborn as sns
from matplotlib_venn import venn3
from scipy import stats
from sklearn.metrics import silhouette_score
%matplotlib inline
import datetime
from chord import Chord
```

```
[2]: date = str(datetime.date.today())
print(date)
```

2022-06-07

```
[3]: #set working directory
os.chdir('/Users/kavyasharman/Documents/Projects/microLESA/')

```

```
[4]: #load data as df
pGroups = pd.read_csv('/Users/kavyasharman/Documents/Projects/microLESA/data/
→proteinGroups.txt', delimiter = '\t', low_memory = False)
pGroups.shape
```

```
[4]: (3760, 371)
```

```
[5]: pGroups.head()
```

```
[5]:
```

	Protein IDs \
0	G3UZW7;A0A023T778;Q9CQL1;P61327
1	070589;A0A067XG53;F6Y9I5

```

2 AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AO...
3 AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1
4 AOA075B5P3;AOA0A6YVPO;P01867

```

```

Majority protein IDs Peptide counts (all) \
0 G3UZW7;AOA023T778;Q9CQL1;P61327 1;1;1;1
1 070589;AOA067XG53;F6Y9I5 2;2;1
2 AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AO... 1;1;1;1;1;1;1;1;1
3 AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1 2;1;1;1
4 AOA075B5P3;AOA0A6YVPO;P01867 3;3;3

```

```

Peptide counts (razor+unique) Peptide counts (unique) \
0 1;1;1;1 1;1;1;1
1 2;2;1 2;2;1
2 1;1;1;1;1;1;1;1;1 1;1;1;1;1;1;1;1;1
3 2;1;1;1 2;1;1;1
4 3;3;3 3;3;3

```

```

Protein names \
0 Protein mago nashi homolog 2;Protein mago nash...
1 Peripheral plasma membrane protein CASK
2 Ig kappa chain V-II region 26-10
3 NaN
4 Ig gamma-2B chain C region

```

```

Gene names \
0 Magohb;Magoh
1 Cask
2 Igkv1-110;Igkv1-35;Igkv1-99;Igkv1-115
3 Igkv5-39;Igkv5-43;Igkv5-45
4 Ighg2b;Igh-3

```

```

Fasta headers Number of proteins \
0 tr|G3UZW7|G3UZW7_MOUSE Protein mago nashi homo... 4
1 sp|070589|CSKP_MOUSE Peripheral plasma membran... 3
2 tr|AOA140T8M9|AOA140T8M9_MOUSE Immunoglobulin ... 9
3 tr|AOA075B5M7|AOA075B5M7_MOUSE Immunoglobulin ... 4
4 tr|AOA075B5P3|AOA075B5P3_MOUSE Immunoglobulin ... 3

```

```

Peptides ... Potential contaminant id Peptide IDs \
0 1 ... NaN 0 9939
1 2 ... NaN 1 5371;7221
2 1 ... NaN 2 7144
3 2 ... NaN 3 3664;23181
4 3 ... NaN 4 2087;3907;22401

```

```

Peptide is razor Mod. peptide IDs \

```

0	True	10517
1	True;True	5685;7650
2	True	7567
3	True;True	3880;3881;24763
4	True;True;True	2218;4134;23956

	Evidence IDs \
0	104111;104112;104113;104114
1	57553;57554;57555;74694;74695
2	74067;74068;74069;74070;74071
3	41615;41616;41617;41618;41619;239742
4	24252;24253;24254;24255;24256;24257;24258;4407...

	MS/MS IDs	Best MS/MS \
0	113199;113200;113201;113202;113203	113200
1	62421;62422;62423;81010;81011	62422;81011
2	80341;80342;80343;80344;80345;80346;80347	80344
3	45159;45160;45161;45162;45163;260731	45163;260731
4	26297;26298;26299;26300;26301;26302;26303;2630...	26297;47805;252619

	Oxidation (M) site IDs	Oxidation (M) site positions
0	NaN	NaN
1	NaN	NaN
2	NaN	NaN
3	0	4
4	NaN	NaN

[5 rows x 371 columns]

```
[6]: pGroups['LFQ intensity 4DPI_2_CO_2']
```

```
[6]: 0    0
      1    0
      2    0
      3    0
      4    0
```

```
      ..
3755    0
3756    0
3757    0
3758    0
3759    0
```

Name: LFQ intensity 4DPI_2_CO_2, Length: 3760, dtype: int64

```
[7]: pGroups.tail()
```

```
[7]: Protein IDs \
3755 REV__Q9QZE7
3756 S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5
```

3757	S4R192
3758	V9GWV8;Q3V335;V9G XK3
3759	V9GX06;AOA0R4J0X7;S4R2G5;Q64467

	Majority protein IDs	Peptide counts (all)	\
3755	REV__Q9QZE7	1	
3756	S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	16;16;15;15;15;15	
3757	S4R192	3	
3758	V9GWV8;Q3V335;V9G XK3	4;3;2	
3759	V9GX06	4;1;1;1	

	Peptide counts (razor+unique)	Peptide counts (unique)	Protein names	\
3755	1	1	NaN	
3756	1;1;1;1;1;1	1;1;1;1;1;1	NaN	
3757	1	1	NaN	
3758	1;0;0	1;0;0	NaN	
3759	1;1;1;1	1;1;1;1	NaN	

	Gene names	Fasta headers	\
3755	NaN	sp Q9QZE7 TSNAX_MOUSE Translin-associated prot...	
3756	Ank3	tr S4R2J6 S4R2J6_MOUSE Ankyrin-3 OS=Mus muscul...	
3757	Sec31a	tr S4R192 S4R192_MOUSE Protein transport prote...	
3758	Nedd4	tr V9GWV8 V9GWV8_MOUSE E3 ubiquitin-protein li...	
3759	Gm11214	tr V9GX06 V9GX06_MOUSE Predicted gene 11214 (F...	

	Number of proteins	Peptides	...	Potential contaminant	id	\
3755	1	1	...	NaN	3755	
3756	6	16	...	NaN	3756	
3757	1	3	...	NaN	3757	
3758	3	4	...	NaN	3758	
3759	4	4	...	NaN	3759	

	Peptide IDs	\
3755	13511	
3756	1455;1705;2665;4244;5703;9994;10028;12460;1246...	
3757	10933;16514;22778	
3758	3286;20769;22815;23095	
3759	1045;15252;22879;22880	

	Peptide is razor	\
3755	True	
3756	False;True;False;False;False;False;False;False...	
3757	False;False;True	
3758	False;True;False;False	
3759	True;False;False;False	

Mod. peptide IDs	\
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3755	14418
3756	1533;1796;2831;4501;6036;10573;10608;13162;131...
3757	11562;17664;24344
3758	3485;22231;24385;24386;24676
3759	1093;16340;24452;24453;24454

	Evidence IDs \
3755	136224;136225
3756	16960;16961;16962;16963;16964;16965;16966;1696...
3757	113810;113811;113812;113813;113814;113815;1138...
3758	37878;37879;37880;37881;37882;37883;37884;3788...
3759	12247;12248;154168;154169;154170;154171;154172...

	MS/MS IDs \
3755	147817;147818
3756	18407;18408;18409;18410;18411;18412;18413;1841...
3757	123686;123687;123688;123689;123690;123691;1236...
3758	41143;41144;41145;41146;41147;41148;41149;4115...
3759	13285;13286;13287;167332;167333;167334;167335;...

	Best MS/MS \
3755	147817
3756	18413;21709;33621;51114;66447;113651;113977;13...
3757	123694;178974;256766
3758	41143;230341;257087;259963
3759	13287;167340;257908;257916

	Oxidation (M) site IDs	Oxidation (M) site positions
3755	1946;1947	91;92
3756	552;553	182;203
3757	NaN	NaN
3758	936	111
3759	172;173	176;179

[5 rows x 371 columns]

```
[8]: #remove proteins identified as "reverse", "only identified by site", and
      ↪ "potential contaminant"
pGroups_proc = pGroups[pGroups.Reverse != '+']
pGroups_proc = pGroups_proc[pGroups_proc['Only identified by site'] != '+']
pGroups_proc = pGroups_proc[pGroups_proc['Potential contaminant'] != '+']
pGroups_proc = pGroups_proc.reset_index(drop=True)
pGroups_proc.tail()
```

```
[8]: Protein IDs \
3608 Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2
3609 Q9Z2Z6
3610 Q9Z315;A0A494B9E9
```

3611 S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5
 3612 S4R192

	Majority protein IDs	Peptide counts (all)	\
3608	Q9Z2X1;J3QMT0;J3QM80	10;7;7;3;2;1	
3609	Q9Z2Z6	3	
3610	Q9Z315	7;3	
3611	S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	16;16;15;15;15;15	
3612	S4R192	3	

	Peptide counts (razor+unique)	Peptide counts (unique)	\
3608	8;7;7;3;2;1	8;7;7;3;2;1	
3609	3	3	
3610	7;3	7;3	
3611	1;1;1;1;1;1	1;1;1;1;1;1	
3612	1	1	

	Protein names	Gene names	\
3608	Heterogeneous nuclear ribonucleoprotein F;Hete...	Hnrnpf	
3609	Mitochondrial carnitine/acylcarnitine carrier ...	Slc25a20	
3610	U4/U6.U5 tri-snRNP-associated protein 1	Sart1	
3611	NaN	Ank3	
3612	NaN	Sec31a	

	Fasta headers	Number of proteins	\
3608	sp Q9Z2X1 HNRPF_MOUSE Heterogeneous nuclear ri...	6	
3609	sp Q9Z2Z6 MCAT_MOUSE Mitochondrial carnitine/a...	1	
3610	sp Q9Z315 SNUT1_MOUSE U4/U6.U5 tri-snRNP-assoc...	2	
3611	tr S4R2J6 S4R2J6_MOUSE Ankyrin-3 OS=Mus muscul...	6	
3612	tr S4R192 S4R192_MOUSE Protein transport prote...	1	

	Peptides	...	Potential contaminant	id	\
3608	10	...	NaN	3705	
3609	3	...	NaN	3706	
3610	7	...	NaN	3707	
3611	16	...	NaN	3756	
3612	3	...	NaN	3757	

	Peptide IDs	\
3608	2655;9429;10400;13472;13473;15976;19341;19692;...	
3609	418;5649;19235	
3610	398;2538;2539;4589;4689;13650;15713	
3611	1455;1705;2665;4244;5703;9994;10028;12460;1246...	
3612	10933;16514;22778	

	Peptide is razor	\
3608	False;True;True;True;True;True;True;True;False...	

3609		True;True;True
3610		True;True;True;True;True;True;True
3611	False;True;False;False;False;False;False;False...	
3612		False;False;True

		Mod. peptide IDs \
3608	2820;9982;11005;14366;14367;14368;17104;20682;...	
3609		431;5977;20568
3610		411;2693;2694;4863;4970;4971;14610;16826
3611	1533;1796;2831;4501;6036;10573;10608;13162;131...	
3612		11562;17664;24344

		Evidence IDs \
3608	30975;30976;30977;30978;30979;30980;30981;3098...	
3609	5063;5064;5065;5066;5067;5068;5069;5070;5071;5...	
3610	4882;4883;4884;4885;4886;4887;29644;29645;2964...	
3611	16960;16961;16962;16963;16964;16965;16966;1696...	
3612	113810;113811;113812;113813;113814;113815;1138...	

		MS/MS IDs \
3608	33513;33514;33515;33516;33517;33518;33519;3352...	
3609	5497;5498;5499;5500;5501;5502;5503;5504;5505;5...	
3610	5301;5302;5303;5304;5305;5306;32096;32097;3209...	
3611	18407;18408;18409;18410;18411;18412;18413;1841...	
3612	123686;123687;123688;123689;123690;123691;1236...	

		Best MS/MS \
3608	33513;107092;117607;147535;147570;174314;21329...	
3609		5506;65881;211665
3610		5304;32096;32124;54448;55730;149357;172113
3611	18413;21709;33621;51114;66447;113651;113977;13...	
3612		123694;178974;256766

	Oxidation (M) site IDs	Oxidation (M) site positions
3608	1926;1927	2;345
3609	NaN	NaN
3610	1928	159
3611	552;553	182;203
3612	NaN	NaN

[5 rows x 371 columns]

[9]: pGroups_proc

		Protein IDs \
0		G3UZW7;A0A023T778;Q9CQL1;P61327
1		O70589;A0A067XG53;F6Y9I5
2	A0A140T8M9;A0A140T8M0;A0A0B4J1I0;A0A075B5N0;A0...	

3	AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1
4	AOA075B5P3;AOA0A6YVPO;P01867
...	...
3608	Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2
3609	Q9Z2Z6
3610	Q9Z315;AOA494B9E9
3611	S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5
3612	S4R192

	Majority protein IDs	Peptide counts (all)	\
0	G3UZW7;AOA023T778;Q9CQL1;P61327	1;1;1;1	
1	O70589;AOA067XG53;F6Y9I5	2;2;1	
2	AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AO...	1;1;1;1;1;1;1;1;1	
3	AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	2;1;1;1	
4	AOA075B5P3;AOA0A6YVPO;P01867	3;3;3	
...	
3608	Q9Z2X1;J3QMT0;J3QM80	10;7;7;3;2;1	
3609	Q9Z2Z6	3	
3610	Q9Z315	7;3	
3611	S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	16;16;15;15;15;15	
3612	S4R192	3	

	Peptide counts (razor+unique)	Peptide counts (unique)	\
0	1;1;1;1	1;1;1;1	
1	2;2;1	2;2;1	
2	1;1;1;1;1;1;1;1	1;1;1;1;1;1;1;1	
3	2;1;1;1	2;1;1;1	
4	3;3;3	3;3;3	
...	
3608	8;7;7;3;2;1	8;7;7;3;2;1	
3609	3	3	
3610	7;3	7;3	
3611	1;1;1;1;1;1	1;1;1;1;1;1	
3612	1	1	

	Protein names	\
0	Protein mago nashi homolog 2;Protein mago nash...	
1	Peripheral plasma membrane protein CASK	
2	Ig kappa chain V-II region 26-10	
3	NaN	
4	Ig gamma-2B chain C region	
...	...	
3608	Heterogeneous nuclear ribonucleoprotein F;Hete...	
3609	Mitochondrial carnitine/acylcarnitine carrier ...	
3610	U4/U6.U5 tri-snRNP-associated protein 1	
3611	NaN	
3612	NaN	

	Gene names \
0	Magohb;Magoh
1	Cask
2	Igkv1-110;Igkv1-35;Igkv1-99;Igkv1-115
3	Igkv5-39;Igkv5-43;Igkv5-45
4	Ighg2b;Igh-3
...	...
3608	Hnrnpf
3609	Slc25a20
3610	Sart1
3611	Ank3
3612	Sec31a

	Fasta headers	Number of proteins \
0	tr G3UZW7 G3UZW7_MOUSE Protein mago nashi homo...	4
1	sp O70589 CSKP_MOUSE Peripheral plasma membran...	3
2	tr A0A140T8M9 A0A140T8M9_MOUSE Immunoglobulin ...	9
3	tr A0A075B5M7 A0A075B5M7_MOUSE Immunoglobulin ...	4
4	tr A0A075B5P3 A0A075B5P3_MOUSE Immunoglobulin ...	3
...
3608	sp Q9Z2X1 HNRPF_MOUSE Heterogeneous nuclear ri...	6
3609	sp Q9Z2Z6 MCAT_MOUSE Mitochondrial carnitine/a...	1
3610	sp Q9Z315 SNUT1_MOUSE U4/U6.U5 tri-snRNP-assoc...	2
3611	tr S4R2J6 S4R2J6_MOUSE Ankyrin-3 OS=Mus muscul...	6
3612	tr S4R192 S4R192_MOUSE Protein transport prote...	1

	Peptides	...	Potential contaminant	id \
0	1	...	NaN	0
1	2	...	NaN	1
2	1	...	NaN	2
3	2	...	NaN	3
4	3	...	NaN	4
...
3608	10	...	NaN	3705
3609	3	...	NaN	3706
3610	7	...	NaN	3707
3611	16	...	NaN	3756
3612	3	...	NaN	3757

	Peptide IDs \
0	9939
1	5371;7221
2	7144
3	3664;23181
4	2087;3907;22401
...	...

3608	2655;9429;10400;13472;13473;15976;19341;19692;...
3609	418;5649;19235
3610	398;2538;2539;4589;4689;13650;15713
3611	1455;1705;2665;4244;5703;9994;10028;12460;1246...
3612	10933;16514;22778

	Peptide is razor \
0	True
1	True;True
2	True
3	True;True
4	True;True;True
...	...
3608	False;True;True;True;True;True;True;True;False...
3609	True;True;True
3610	True;True;True;True;True;True;True
3611	False;True;False;False;False;False;False;False...
3612	False;False;True

	Mod. peptide IDs \
0	10517
1	5685;7650
2	7567
3	3880;3881;24763
4	2218;4134;23956
...	...
3608	2820;9982;11005;14366;14367;14368;17104;20682;...
3609	431;5977;20568
3610	411;2693;2694;4863;4970;4971;14610;16826
3611	1533;1796;2831;4501;6036;10573;10608;13162;131...
3612	11562;17664;24344

	Evidence IDs \
0	104111;104112;104113;104114
1	57553;57554;57555;74694;74695
2	74067;74068;74069;74070;74071
3	41615;41616;41617;41618;41619;239742
4	24252;24253;24254;24255;24256;24257;24258;4407...
...	...
3608	30975;30976;30977;30978;30979;30980;30981;3098...
3609	5063;5064;5065;5066;5067;5068;5069;5070;5071;5...
3610	4882;4883;4884;4885;4886;4887;29644;29645;2964...
3611	16960;16961;16962;16963;16964;16965;16966;1696...
3612	113810;113811;113812;113813;113814;113815;1138...

	MS/MS IDs \
0	113199;113200;113201;113202;113203

```

1          62421;62422;62423;81010;81011
2      80341;80342;80343;80344;80345;80346;80347
3          45159;45160;45161;45162;45163;260731
4      26297;26298;26299;26300;26301;26302;26303;2630...
...
3608  33513;33514;33515;33516;33517;33518;33519;3352...
3609  5497;5498;5499;5500;5501;5502;5503;5504;5505;5...
3610  5301;5302;5303;5304;5305;5306;32096;32097;3209...
3611  18407;18408;18409;18410;18411;18412;18413;1841...
3612  123686;123687;123688;123689;123690;123691;1236...

```

```

Best MS/MS \
0          113200
1      62422;81011
2          80344
3      45163;260731
4      26297;47805;252619
...
3608  33513;107092;117607;147535;147570;174314;21329...
3609          5506;65881;211665
3610      5304;32096;32124;54448;55730;149357;172113
3611  18413;21709;33621;51114;66447;113651;113977;13...
3612          123694;178974;256766

```

```

Oxidation (M) site IDs  Oxidation (M) site positions
0          NaN          NaN
1          NaN          NaN
2          NaN          NaN
3              0              4
4          NaN          NaN
...
3608      1926;1927      2;345
3609          NaN          NaN
3610      1928          159
3611      552;553      182;203
3612          NaN          NaN

```

[3613 rows x 371 columns]

```

[10]: #create database of all proteins in protein groups
proteins = pGroups_proc[' Protein IDs'].str.split(";", expand=True)
proteins = proteins.stack()
proteins = list(set(proteins))
proteins = pd.DataFrame(proteins)
proteins.to_csv('outs/' + date + '_allProteinsFromPGroups.csv')

```

```

[215]: #extract LFQ intensity and create new df with expt classifiers
#instead of group by region or timepoint, add region+timepoint as factors

```

```

def make_lfq(dataframe, species):
    temp = dataframe.columns
    lfq = pd.DataFrame()
    filename = str('/Users/kavyasharman/Documents/Projects/microLESA/outs/' +
    →date + '_' + species + '_lfq.csv')
    string = "LFQ"
    for i in temp:
        if string in i:
            lfq = lfq.append(dataframe[i])
    lfq.columns = dataframe[' Protein IDs']
    lfq = lfq.reset_index()
    lfq['index'] = lfq['index'].str.strip("LFQ intensity ")
    lfq[['TimePoint', 'BioRep', 'Region', 'TechRep']] = lfq['index'].str.
    →split("_", expand=True)
    lfq['Region'] = lfq['Region'].replace({'LE': 'interface', 'AB': 'SAC', 'CO':
    → 'cortex'})
    lfq['DPI_Region_BioRep'] = lfq['TimePoint'] + "_" + lfq['Region'] + "_" +
    →lfq['BioRep']
    lfq['BioRep_TechRep'] = lfq['BioRep'] + "_" + lfq['TechRep']
    lfq['DPI_Region'] = lfq['TimePoint'] + "_" + lfq['Region']
    lfq['index'] = lfq['TimePoint'] + "_" + lfq['Region'] + "_" + lfq['BioRep']
    →+ "_" + lfq['TechRep']
    lfq = lfq.set_index('index')
    lfq = lfq.sort_values(by=['TimePoint', 'Region', 'BioRep', 'TechRep'],
    →ascending=False)
    return lfq

#extract LFQ intensity for all proteins and create new df 'lfq' with expt
    →classifiers
lfq = make_lfq(pGroups_proc, 'combined')
lfq.shape

```

[215]: (42, 3620)

[222]: lfq.head()

```

[222]: Protein IDs      G3UZW7;A0A023T778;Q9CQL1;P61327  070589;A0A067XG53;F6Y9I5  \
index
4DPI_interface_2_4      0.0      0.0
4DPI_interface_2_2      0.0      0.0
4DPI_interface_2_1      0.0      0.0
4DPI_interface_1_2      0.0      0.0
4DPI_interface_1_1      0.0      0.0

Protein IDs      A0A140T8M9;A0A140T8M0;A0A0B4J1I0;A0A075B5N0;A0A075B5K8;F6XWB
2;A0A140T8N1;A0A0G2JDE5;P01631  \
index

```

4DPI_interface_2_4	0.0
4DPI_interface_2_2	0.0
4DPI_interface_2_1	0.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	0.0

Protein IDs	AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	\
index		
4DPI_interface_2_4	0.0	
4DPI_interface_2_2	0.0	
4DPI_interface_2_1	0.0	
4DPI_interface_1_2	0.0	
4DPI_interface_1_1	0.0	

Protein IDs	AOA075B5P3;AOA0A6YVP0;P01867	\
index		
4DPI_interface_2_4	0.0	
4DPI_interface_2_2	0.0	
4DPI_interface_2_1	0.0	
4DPI_interface_1_2	0.0	
4DPI_interface_1_1	0.0	

Protein IDs	AOA075B5P5;AOA1Y7VJN6;P03987	\
index		
4DPI_interface_2_4	0.0	
4DPI_interface_2_2	0.0	
4DPI_interface_2_1	0.0	
4DPI_interface_1_2	0.0	
4DPI_interface_1_1	0.0	

Protein IDs	AOA075B6A0;AOA075B5P6;P01872	\
index		
4DPI_interface_2_4	11850000.0	
4DPI_interface_2_2	13198000.0	
4DPI_interface_2_1	14041000.0	
4DPI_interface_1_2	8327300.0	
4DPI_interface_1_1	9088300.0	

Protein IDs	AOA075B5R2;AOA075B5R3;AOA0A6YX91;P01786	\
index		
4DPI_interface_2_4	489990.0	
4DPI_interface_2_2	0.0	
4DPI_interface_2_1	1280300.0	
4DPI_interface_1_2	0.0	
4DPI_interface_1_1	0.0	

Protein IDs	AOA075B5S5;AOA0A6YXL5	AOA075B5T3;J3QK03;AOA0A6YWS9	... \
-------------	-----------------------	------------------------------	-------

index			...
4DPI_interface_2_4	0.0	0.0	...
4DPI_interface_2_2	0.0	0.0	...
4DPI_interface_2_1	0.0	0.0	...
4DPI_interface_1_2	0.0	0.0	...
4DPI_interface_1_1	0.0	0.0	...

Protein IDs Q9Z315;A0A494B9E9 \

index	
4DPI_interface_2_4	954220.0
4DPI_interface_2_2	1377400.0
4DPI_interface_2_1	922440.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	1117600.0

Protein IDs S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5 S4R192 \

index		
4DPI_interface_2_4	0.0	0.0
4DPI_interface_2_2	0.0	0.0
4DPI_interface_2_1	0.0	0.0
4DPI_interface_1_2	0.0	0.0
4DPI_interface_1_1	0.0	0.0

Protein IDs TimePoint BioRep Region TechRep DPI_Region_BioRep \

index					
4DPI_interface_2_4	4DPI	2	interface	4	4DPI_interface_2
4DPI_interface_2_2	4DPI	2	interface	2	4DPI_interface_2
4DPI_interface_2_1	4DPI	2	interface	1	4DPI_interface_2
4DPI_interface_1_2	4DPI	1	interface	2	4DPI_interface_1
4DPI_interface_1_1	4DPI	1	interface	1	4DPI_interface_1

Protein IDs BioRep_TechRep DPI_Region

index		
4DPI_interface_2_4	2_4	4DPI_interface
4DPI_interface_2_2	2_2	4DPI_interface
4DPI_interface_2_1	2_1	4DPI_interface
4DPI_interface_1_2	1_2	4DPI_interface
4DPI_interface_1_1	1_1	4DPI_interface

[5 rows x 3620 columns]

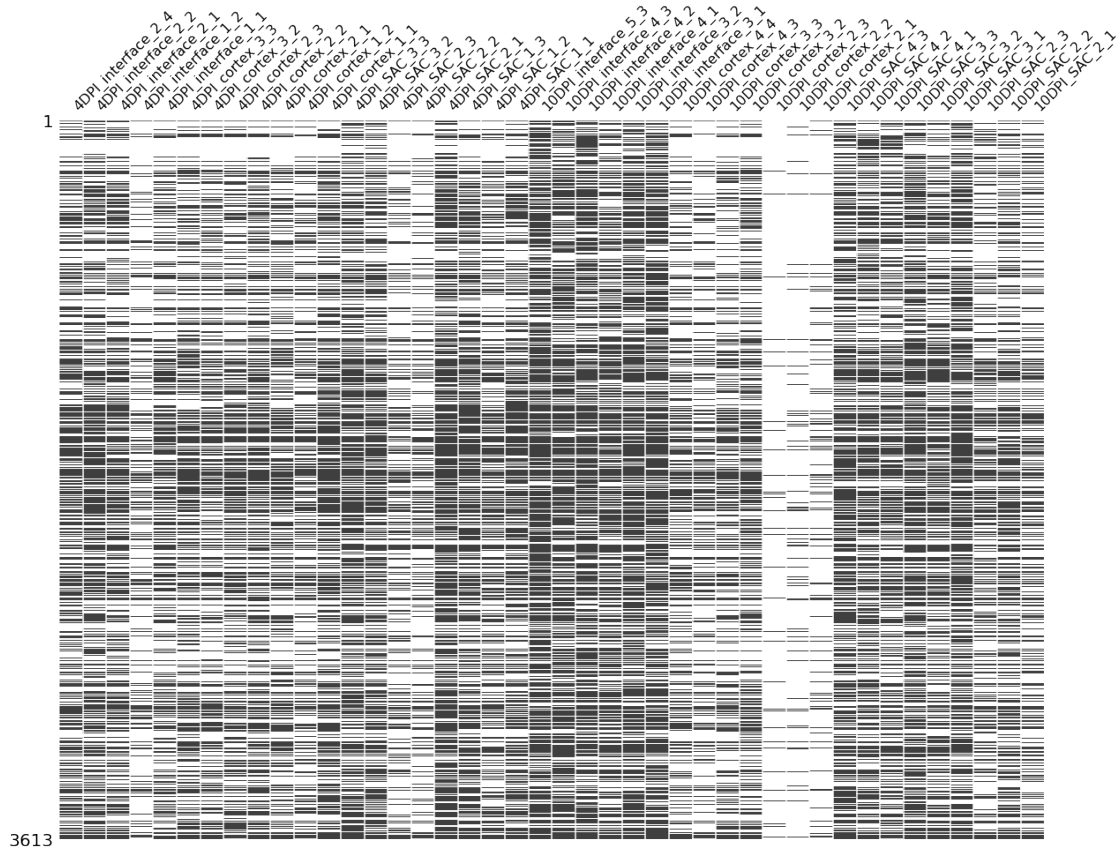
```
[219]: # calculate missing values (i.e. LFQ = 0)
lfq_nan = lfq.drop(['TimePoint', 'BioRep', 'Region', 'TechRep'],
→ 'DPI_Region_BioRep', 'BioRep_TechRep', 'DPI_Region'], axis=1)
lfq_nan = lfq_nan.replace(0, np.nan)
number_missing_values = lfq_nan.transpose().isnull().sum()
percent_missing_values = lfq_nan.transpose().isnull().mean() * 100
```

```
percent_missing_values
```

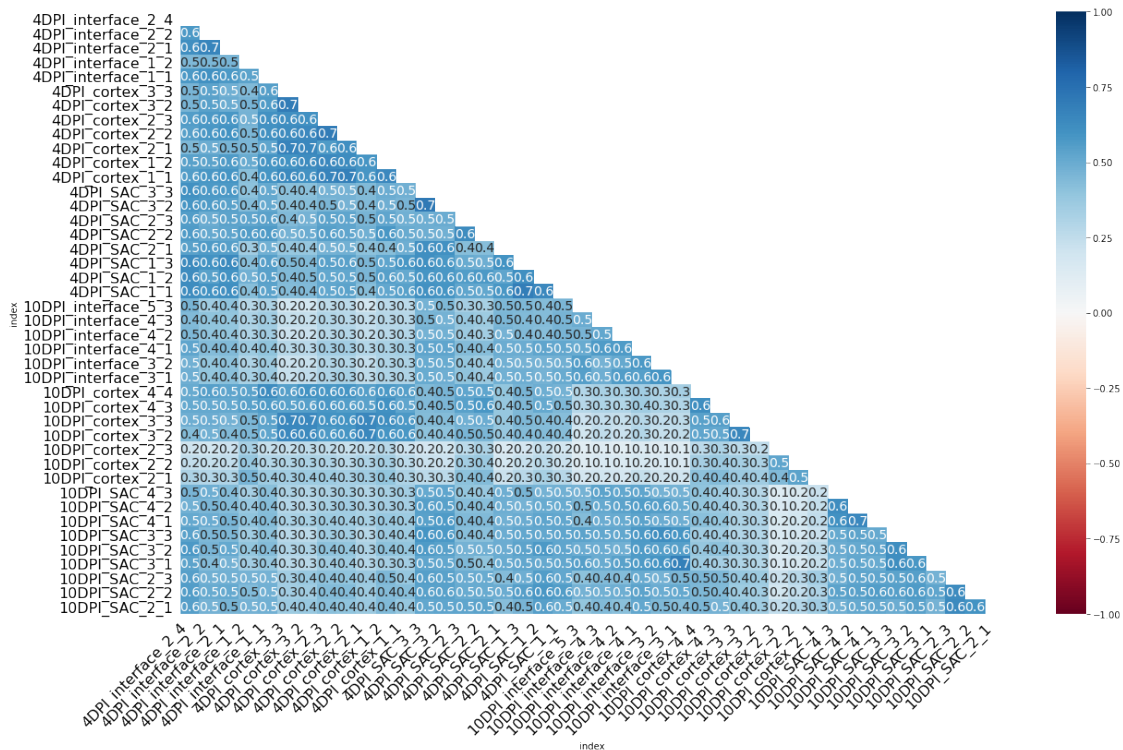
```
[219]: index
4DPI_interface_2_4      52.809300
4DPI_interface_2_2      47.329089
4DPI_interface_2_1      52.421810
4DPI_interface_1_2      77.359535
4DPI_interface_1_1      59.811791
4DPI_cortex_3_3         56.545807
4DPI_cortex_3_2         60.891226
4DPI_cortex_2_3         57.846665
4DPI_cortex_2_2         53.390534
4DPI_cortex_2_1         58.815389
4DPI_cortex_1_2         65.292001
4DPI_cortex_1_1         50.429006
4DPI_SAC_3_3            46.277332
4DPI_SAC_3_2            48.879048
4DPI_SAC_2_3            71.851647
4DPI_SAC_2_2            73.124827
4DPI_SAC_2_1            37.032937
4DPI_SAC_1_3            46.000554
4DPI_SAC_1_2            59.562690
4DPI_SAC_1_1            48.851370
10DPI_interface_5_3     31.082203
10DPI_interface_4_3     38.998063
10DPI_interface_4_2     36.424024
10DPI_interface_4_1     46.360365
10DPI_interface_3_2     36.534736
10DPI_interface_3_1     37.337393
10DPI_cortex_4_4        64.738445
10DPI_cortex_4_3        72.349848
10DPI_cortex_3_3        63.963465
10DPI_cortex_3_2        59.009134
10DPI_cortex_2_3        95.599225
10DPI_cortex_2_2        93.052865
10DPI_cortex_2_1        89.593136
10DPI_SAC_4_3           41.821201
10DPI_SAC_4_2           47.993357
10DPI_SAC_4_1           51.868254
10DPI_SAC_3_3           42.402436
10DPI_SAC_3_2           52.920011
10DPI_SAC_3_1           41.129255
10DPI_SAC_2_3           63.520620
10DPI_SAC_2_2           55.438694
10DPI_SAC_2_1           66.343759
dtype: float64
```

```
[220]: pd.DataFrame(percent_missing_values).to_csv('outs/' + date + '_missingValues.  
→csv')
```

```
[221]: msno.matrix(lfq_nan.transpose(), sparkline=False, figsize=(20,15))  
plt.savefig('outs/' + date + '_missingValues_matrix.png')
```



```
[223]: msno.heatmap(lfq_nan.transpose())  
plt.savefig('outs/' + date + '_missingValues_heatmap.svg')
```

```
[45]: #z-score calculation to drop outliers
```

```
z = np.abs(stats.zscore(percent_missing_values))
print(z)
print(np.where(z > 2))
```

```
[0.21484596 0.58111471 0.24074375 1.42596406 0.25316412 0.03488274
 0.32530796 0.12182532 0.17599927 0.1865698 0.61943287 0.37393238
 0.65140871 0.47752355 1.05784547 1.14293821 1.269256 0.66990713
 0.23651554 0.47937339 1.66697207 1.13791721 1.30995253 0.64585919
 1.30255316 1.24890774 0.58243603 1.09114263 0.53064044 0.19951869
 2.64501006 2.47482458 2.24359431 0.9492333 0.5367185 0.27774059
 0.91038662 0.20744659 0.99547936 0.50104297 0.03911095 0.68972687]
(array([30, 31, 32]),)
```

```
[46]: #z-score calculation to drop outliers
```

```
z = np.abs(stats.zscore(percent_missing_values))
z>2
```

```
[46]: array([False, False, False, False, False, False, False, False, False,
        False, False, False, False, False, False, False, False, False,
        False, False, False, False, False, False, False, False, False,
        False, False, False, True,  True,  True, False, False, False,
        False, False, False, False, False, False])
```

```
[47]: #list of outlier samples
outliers = percent_missing_values.sort_values(ascending=False)[0:3].index
outliers
```

```
[47]: Index(['10DPI_cortex_2_3', '10DPI_cortex_2_2', '10DPI_cortex_2_1'],
dtype='object', name='index')
```

```
[48]: #drop outliers
combined_lfq = make_lfq(pGroups_proc, 'combined').drop(outliers, axis=0)
combined_lfq.to_csv('outs/' + date + '_combined_lfq.csv')
```

0.1.1 Remove proteins with missing values

```
[49]: combined_lfq
```

```
[49]: Protein IDs      G3UZW7;A0A023T778;Q9CQL1;P61327  \
index
4DPI_interface_2_4      0.0
4DPI_interface_2_2      0.0
4DPI_interface_2_1      0.0
4DPI_interface_1_2      0.0
4DPI_interface_1_1      0.0
4DPI_cortex_3_3          0.0
4DPI_cortex_3_2          0.0
4DPI_cortex_2_3          0.0
4DPI_cortex_2_2          0.0
4DPI_cortex_2_1          0.0
4DPI_cortex_1_2          0.0
4DPI_cortex_1_1          0.0
4DPI_SAC_3_3             0.0
4DPI_SAC_3_2             0.0
4DPI_SAC_2_3             0.0
4DPI_SAC_2_2             0.0
4DPI_SAC_2_1            189280.0
4DPI_SAC_1_3            561840.0
4DPI_SAC_1_2            779590.0
4DPI_SAC_1_1             0.0
10DPI_interface_5_3      0.0
10DPI_interface_4_3      0.0
10DPI_interface_4_2     222210.0
10DPI_interface_4_1      0.0
10DPI_interface_3_2      0.0
10DPI_interface_3_1      0.0
10DPI_cortex_4_4         0.0
10DPI_cortex_4_3         0.0
10DPI_cortex_3_3         0.0
10DPI_cortex_3_2         0.0
10DPI_SAC_4_3            0.0
```

10DPI_SAC_4_2	0.0
10DPI_SAC_4_1	0.0
10DPI_SAC_3_3	0.0
10DPI_SAC_3_2	0.0
10DPI_SAC_3_1	0.0
10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	0.0
10DPI_SAC_2_1	0.0

Protein IDs 070589;A0A067XG53;F6Y9I5 \

index

4DPI_interface_2_4	0.0
4DPI_interface_2_2	0.0
4DPI_interface_2_1	0.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	0.0
4DPI_cortex_3_3	0.0
4DPI_cortex_3_2	0.0
4DPI_cortex_2_3	0.0
4DPI_cortex_2_2	0.0
4DPI_cortex_2_1	497950.0
4DPI_cortex_1_2	0.0
4DPI_cortex_1_1	0.0
4DPI_SAC_3_3	0.0
4DPI_SAC_3_2	0.0
4DPI_SAC_2_3	0.0
4DPI_SAC_2_2	0.0
4DPI_SAC_2_1	235580.0
4DPI_SAC_1_3	0.0
4DPI_SAC_1_2	0.0
4DPI_SAC_1_1	0.0
10DPI_interface_5_3	0.0
10DPI_interface_4_3	0.0
10DPI_interface_4_2	0.0
10DPI_interface_4_1	0.0
10DPI_interface_3_2	0.0
10DPI_interface_3_1	0.0
10DPI_cortex_4_4	0.0
10DPI_cortex_4_3	0.0
10DPI_cortex_3_3	0.0
10DPI_cortex_3_2	434700.0
10DPI_SAC_4_3	0.0
10DPI_SAC_4_2	0.0
10DPI_SAC_4_1	0.0
10DPI_SAC_3_3	0.0
10DPI_SAC_3_2	0.0
10DPI_SAC_3_1	0.0

10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	0.0
10DPI_SAC_2_1	0.0

Protein IDs AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA075B5K8;F6XW
B2;AOA140T8N1;AOA0G2JDE5;P01631 \

index

4DPI_interface_2_4	0.0
4DPI_interface_2_2	0.0
4DPI_interface_2_1	0.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	0.0
4DPI_cortex_3_3	0.0
4DPI_cortex_3_2	0.0
4DPI_cortex_2_3	0.0
4DPI_cortex_2_2	0.0
4DPI_cortex_2_1	0.0
4DPI_cortex_1_2	0.0
4DPI_cortex_1_1	0.0
4DPI_SAC_3_3	0.0
4DPI_SAC_3_2	0.0
4DPI_SAC_2_3	0.0
4DPI_SAC_2_2	0.0
4DPI_SAC_2_1	0.0
4DPI_SAC_1_3	0.0
4DPI_SAC_1_2	0.0
4DPI_SAC_1_1	0.0
10DPI_interface_5_3	0.0
10DPI_interface_4_3	0.0
10DPI_interface_4_2	0.0
10DPI_interface_4_1	0.0
10DPI_interface_3_2	0.0
10DPI_interface_3_1	0.0
10DPI_cortex_4_4	0.0
10DPI_cortex_4_3	0.0
10DPI_cortex_3_3	855310.0
10DPI_cortex_3_2	498950.0
10DPI_SAC_4_3	1674100.0
10DPI_SAC_4_2	431080.0
10DPI_SAC_4_1	548100.0
10DPI_SAC_3_3	0.0
10DPI_SAC_3_2	0.0
10DPI_SAC_3_1	0.0
10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	0.0
10DPI_SAC_2_1	0.0

Protein IDs	A0A075B5M7;A0A0G2JDV4;A0A0B4J1J2;A0A0B4J1J1	\
index		
4DPI_interface_2_4		0.0
4DPI_interface_2_2		0.0
4DPI_interface_2_1		0.0
4DPI_interface_1_2		0.0
4DPI_interface_1_1		0.0
4DPI_cortex_3_3		0.0
4DPI_cortex_3_2		0.0
4DPI_cortex_2_3		0.0
4DPI_cortex_2_2		0.0
4DPI_cortex_2_1		0.0
4DPI_cortex_1_2		0.0
4DPI_cortex_1_1		0.0
4DPI_SAC_3_3		0.0
4DPI_SAC_3_2		0.0
4DPI_SAC_2_3		0.0
4DPI_SAC_2_2		0.0
4DPI_SAC_2_1		0.0
4DPI_SAC_1_3		0.0
4DPI_SAC_1_2		0.0
4DPI_SAC_1_1		0.0
10DPI_interface_5_3		81477.0
10DPI_interface_4_3		207140.0
10DPI_interface_4_2		0.0
10DPI_interface_4_1		0.0
10DPI_interface_3_2		0.0
10DPI_interface_3_1		0.0
10DPI_cortex_4_4		0.0
10DPI_cortex_4_3		0.0
10DPI_cortex_3_3		0.0
10DPI_cortex_3_2		0.0
10DPI_SAC_4_3		202930.0
10DPI_SAC_4_2		290690.0
10DPI_SAC_4_1		0.0
10DPI_SAC_3_3		0.0
10DPI_SAC_3_2		0.0
10DPI_SAC_3_1		0.0
10DPI_SAC_2_3		0.0
10DPI_SAC_2_2		0.0
10DPI_SAC_2_1		0.0

Protein IDs	A0A075B5P3;A0A0A6YVP0;P01867	\
index		
4DPI_interface_2_4	0.0	
4DPI_interface_2_2	0.0	
4DPI_interface_2_1	0.0	

4DPI_interface_1_2	0.0
4DPI_interface_1_1	0.0
4DPI_cortex_3_3	0.0
4DPI_cortex_3_2	0.0
4DPI_cortex_2_3	0.0
4DPI_cortex_2_2	0.0
4DPI_cortex_2_1	0.0
4DPI_cortex_1_2	0.0
4DPI_cortex_1_1	0.0
4DPI_SAC_3_3	0.0
4DPI_SAC_3_2	0.0
4DPI_SAC_2_3	0.0
4DPI_SAC_2_2	0.0
4DPI_SAC_2_1	0.0
4DPI_SAC_1_3	0.0
4DPI_SAC_1_2	0.0
4DPI_SAC_1_1	0.0
10DPI_interface_5_3	0.0
10DPI_interface_4_3	1183700.0
10DPI_interface_4_2	2295600.0
10DPI_interface_4_1	1475700.0
10DPI_interface_3_2	869990.0
10DPI_interface_3_1	0.0
10DPI_cortex_4_4	0.0
10DPI_cortex_4_3	0.0
10DPI_cortex_3_3	0.0
10DPI_cortex_3_2	0.0
10DPI_SAC_4_3	1207600.0
10DPI_SAC_4_2	1526100.0
10DPI_SAC_4_1	1165600.0
10DPI_SAC_3_3	0.0
10DPI_SAC_3_2	0.0
10DPI_SAC_3_1	724260.0
10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	0.0
10DPI_SAC_2_1	0.0

Protein IDs AOA075B5P5;AOA1Y7VJN6;P03987 \

index

4DPI_interface_2_4	0.0
4DPI_interface_2_2	0.0
4DPI_interface_2_1	0.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	0.0
4DPI_cortex_3_3	0.0
4DPI_cortex_3_2	0.0
4DPI_cortex_2_3	0.0

4DPI_cortex_2_2	0.0
4DPI_cortex_2_1	0.0
4DPI_cortex_1_2	0.0
4DPI_cortex_1_1	0.0
4DPI_SAC_3_3	0.0
4DPI_SAC_3_2	0.0
4DPI_SAC_2_3	0.0
4DPI_SAC_2_2	0.0
4DPI_SAC_2_1	0.0
4DPI_SAC_1_3	0.0
4DPI_SAC_1_2	0.0
4DPI_SAC_1_1	0.0
10DPI_interface_5_3	1109600.0
10DPI_interface_4_3	12624000.0
10DPI_interface_4_2	14386000.0
10DPI_interface_4_1	19824000.0
10DPI_interface_3_2	3005900.0
10DPI_interface_3_1	447980.0
10DPI_cortex_4_4	21601000.0
10DPI_cortex_4_3	526710.0
10DPI_cortex_3_3	1102700.0
10DPI_cortex_3_2	654170.0
10DPI_SAC_4_3	29513000.0
10DPI_SAC_4_2	13752000.0
10DPI_SAC_4_1	17690000.0
10DPI_SAC_3_3	848950.0
10DPI_SAC_3_2	997060.0
10DPI_SAC_3_1	295760.0
10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	1941400.0
10DPI_SAC_2_1	0.0

Protein IDs	AOA075B6A0;AOA075B5P6;P01872 \
index	
4DPI_interface_2_4	11850000.0
4DPI_interface_2_2	13198000.0
4DPI_interface_2_1	14041000.0
4DPI_interface_1_2	8327300.0
4DPI_interface_1_1	9088300.0
4DPI_cortex_3_3	28479000.0
4DPI_cortex_3_2	9475600.0
4DPI_cortex_2_3	12548000.0
4DPI_cortex_2_2	17613000.0
4DPI_cortex_2_1	18376000.0
4DPI_cortex_1_2	13749000.0
4DPI_cortex_1_1	19070000.0
4DPI_SAC_3_3	4038600.0

4DPI_SAC_3_2	1051500.0
4DPI_SAC_2_3	10500000.0
4DPI_SAC_2_2	4068800.0
4DPI_SAC_2_1	6212200.0
4DPI_SAC_1_3	7820100.0
4DPI_SAC_1_2	3859700.0
4DPI_SAC_1_1	4501600.0
10DPI_interface_5_3	7111900.0
10DPI_interface_4_3	15394000.0
10DPI_interface_4_2	20355000.0
10DPI_interface_4_1	24453000.0
10DPI_interface_3_2	6101300.0
10DPI_interface_3_1	9250300.0
10DPI_cortex_4_4	49938000.0
10DPI_cortex_4_3	16417000.0
10DPI_cortex_3_3	16583000.0
10DPI_cortex_3_2	14249000.0
10DPI_SAC_4_3	36836000.0
10DPI_SAC_4_2	21015000.0
10DPI_SAC_4_1	20078000.0
10DPI_SAC_3_3	5357600.0
10DPI_SAC_3_2	2988700.0
10DPI_SAC_3_1	6025900.0
10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	6863300.0
10DPI_SAC_2_1	2568900.0

Protein IDs	A0A075B5R2;A0A075B5R3;A0A0A6YX91;P01786 \
index	
4DPI_interface_2_4	489990.0
4DPI_interface_2_2	0.0
4DPI_interface_2_1	1280300.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	0.0
4DPI_cortex_3_3	0.0
4DPI_cortex_3_2	0.0
4DPI_cortex_2_3	537080.0
4DPI_cortex_2_2	1492000.0
4DPI_cortex_2_1	247720.0
4DPI_cortex_1_2	0.0
4DPI_cortex_1_1	0.0
4DPI_SAC_3_3	0.0
4DPI_SAC_3_2	0.0
4DPI_SAC_2_3	0.0
4DPI_SAC_2_2	0.0
4DPI_SAC_2_1	0.0
4DPI_SAC_1_3	0.0

4DPI_SAC_1_2	0.0
4DPI_SAC_1_1	0.0
10DPI_interface_5_3	923480.0
10DPI_interface_4_3	2742000.0
10DPI_interface_4_2	2199900.0
10DPI_interface_4_1	2193900.0
10DPI_interface_3_2	875140.0
10DPI_interface_3_1	1436900.0
10DPI_cortex_4_4	1717800.0
10DPI_cortex_4_3	1395700.0
10DPI_cortex_3_3	880010.0
10DPI_cortex_3_2	708140.0
10DPI_SAC_4_3	1401500.0
10DPI_SAC_4_2	0.0
10DPI_SAC_4_1	1453700.0
10DPI_SAC_3_3	536050.0
10DPI_SAC_3_2	2143600.0
10DPI_SAC_3_1	944670.0
10DPI_SAC_2_3	0.0
10DPI_SAC_2_2	533920.0
10DPI_SAC_2_1	0.0

Protein IDs	AOA075B5S5;AOA0A6YXL5	AOA075B5T3;J3QK03;AOA0A6YWS9	...	\
index			...	
4DPI_interface_2_4	0.0	0.0	...	
4DPI_interface_2_2	0.0	0.0	...	
4DPI_interface_2_1	0.0	0.0	...	
4DPI_interface_1_2	0.0	0.0	...	
4DPI_interface_1_1	0.0	0.0	...	
4DPI_cortex_3_3	0.0	0.0	...	
4DPI_cortex_3_2	0.0	0.0	...	
4DPI_cortex_2_3	0.0	0.0	...	
4DPI_cortex_2_2	0.0	0.0	...	
4DPI_cortex_2_1	0.0	0.0	...	
4DPI_cortex_1_2	0.0	0.0	...	
4DPI_cortex_1_1	0.0	0.0	...	
4DPI_SAC_3_3	0.0	0.0	...	
4DPI_SAC_3_2	0.0	0.0	...	
4DPI_SAC_2_3	0.0	0.0	...	
4DPI_SAC_2_2	0.0	0.0	...	
4DPI_SAC_2_1	0.0	0.0	...	
4DPI_SAC_1_3	0.0	0.0	...	
4DPI_SAC_1_2	0.0	0.0	...	
4DPI_SAC_1_1	0.0	0.0	...	
10DPI_interface_5_3	582000.0	624130.0	...	
10DPI_interface_4_3	10474000.0	1520300.0	...	
10DPI_interface_4_2	4267600.0	3319600.0	...	

10DPI_interface_4_1	1899800.0	1112100.0	...
10DPI_interface_3_2	521140.0	826650.0	...
10DPI_interface_3_1	554440.0	0.0	...
10DPI_cortex_4_4	3546300.0	2818100.0	...
10DPI_cortex_4_3	0.0	2965700.0	...
10DPI_cortex_3_3	0.0	992900.0	...
10DPI_cortex_3_2	739180.0	0.0	...
10DPI_SAC_4_3	2447600.0	5502600.0	...
10DPI_SAC_4_2	3972000.0	2063300.0	...
10DPI_SAC_4_1	8814000.0	2556300.0	...
10DPI_SAC_3_3	0.0	1867700.0	...
10DPI_SAC_3_2	0.0	766080.0	...
10DPI_SAC_3_1	0.0	0.0	...
10DPI_SAC_2_3	0.0	0.0	...
10DPI_SAC_2_2	0.0	0.0	...
10DPI_SAC_2_1	0.0	0.0	...

Protein IDs Q9Z315;A0A494B9E9 \

index

4DPI_interface_2_4	954220.0
4DPI_interface_2_2	1377400.0
4DPI_interface_2_1	922440.0
4DPI_interface_1_2	0.0
4DPI_interface_1_1	1117600.0
4DPI_cortex_3_3	638210.0
4DPI_cortex_3_2	661230.0
4DPI_cortex_2_3	0.0
4DPI_cortex_2_2	494860.0
4DPI_cortex_2_1	674540.0
4DPI_cortex_1_2	0.0
4DPI_cortex_1_1	334760.0
4DPI_SAC_3_3	1396800.0
4DPI_SAC_3_2	1242700.0
4DPI_SAC_2_3	1401200.0
4DPI_SAC_2_2	0.0
4DPI_SAC_2_1	458240.0
4DPI_SAC_1_3	780230.0
4DPI_SAC_1_2	1219000.0
4DPI_SAC_1_1	600830.0
10DPI_interface_5_3	1281000.0
10DPI_interface_4_3	2218300.0
10DPI_interface_4_2	1495700.0
10DPI_interface_4_1	927020.0
10DPI_interface_3_2	1865800.0
10DPI_interface_3_1	1160000.0
10DPI_cortex_4_4	946380.0
10DPI_cortex_4_3	0.0

10DPI_cortex_3_3	633750.0
10DPI_cortex_3_2	556170.0
10DPI_SAC_4_3	1137800.0
10DPI_SAC_4_2	1302200.0
10DPI_SAC_4_1	0.0
10DPI_SAC_3_3	1731200.0
10DPI_SAC_3_2	1444700.0
10DPI_SAC_3_1	1059500.0
10DPI_SAC_2_3	3343300.0
10DPI_SAC_2_2	800330.0
10DPI_SAC_2_1	2338200.0

Protein IDs	S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	S4R192 \
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4DPI_interface_2_4	0.0	0.0
4DPI_interface_2_2	0.0	0.0
4DPI_interface_2_1	0.0	0.0
4DPI_interface_1_2	0.0	0.0
4DPI_interface_1_1	0.0	0.0
4DPI_cortex_3_3	0.0	0.0
4DPI_cortex_3_2	0.0	0.0
4DPI_cortex_2_3	662690.0	0.0
4DPI_cortex_2_2	0.0	0.0
4DPI_cortex_2_1	0.0	0.0
4DPI_cortex_1_2	714480.0	0.0
4DPI_cortex_1_1	716000.0	0.0
4DPI_SAC_3_3	407450.0	0.0
4DPI_SAC_3_2	0.0	357930.0
4DPI_SAC_2_3	0.0	0.0
4DPI_SAC_2_2	0.0	0.0
4DPI_SAC_2_1	360500.0	169050.0
4DPI_SAC_1_3	610920.0	0.0
4DPI_SAC_1_2	326340.0	0.0
4DPI_SAC_1_1	0.0	251670.0
10DPI_interface_5_3	126670.0	0.0
10DPI_interface_4_3	0.0	305120.0
10DPI_interface_4_2	0.0	549550.0
10DPI_interface_4_1	0.0	0.0
10DPI_interface_3_2	0.0	364880.0
10DPI_interface_3_1	0.0	781480.0
10DPI_cortex_4_4	0.0	0.0
10DPI_cortex_4_3	0.0	0.0
10DPI_cortex_3_3	0.0	0.0
10DPI_cortex_3_2	0.0	0.0
10DPI_SAC_4_3	0.0	0.0
10DPI_SAC_4_2	0.0	0.0
10DPI_SAC_4_1	0.0	326950.0

10DPI_SAC_3_3	0.0	423330.0
10DPI_SAC_3_2	0.0	0.0
10DPI_SAC_3_1	0.0	0.0
10DPI_SAC_2_3	0.0	0.0
10DPI_SAC_2_2	0.0	0.0
10DPI_SAC_2_1	0.0	0.0

Protein IDs index	TimePoint	BioRep	Region	TechRep	DPI_Region_BioRep \
4DPI_interface_2_4	4DPI	2	interface	4	4DPI_interface_2
4DPI_interface_2_2	4DPI	2	interface	2	4DPI_interface_2
4DPI_interface_2_1	4DPI	2	interface	1	4DPI_interface_2
4DPI_interface_1_2	4DPI	1	interface	2	4DPI_interface_1
4DPI_interface_1_1	4DPI	1	interface	1	4DPI_interface_1
4DPI_cortex_3_3	4DPI	3	cortex	3	4DPI_cortex_3
4DPI_cortex_3_2	4DPI	3	cortex	2	4DPI_cortex_3
4DPI_cortex_2_3	4DPI	2	cortex	3	4DPI_cortex_2
4DPI_cortex_2_2	4DPI	2	cortex	2	4DPI_cortex_2
4DPI_cortex_2_1	4DPI	2	cortex	1	4DPI_cortex_2
4DPI_cortex_1_2	4DPI	1	cortex	2	4DPI_cortex_1
4DPI_cortex_1_1	4DPI	1	cortex	1	4DPI_cortex_1
4DPI_SAC_3_3	4DPI	3	SAC	3	4DPI_SAC_3
4DPI_SAC_3_2	4DPI	3	SAC	2	4DPI_SAC_3
4DPI_SAC_2_3	4DPI	2	SAC	3	4DPI_SAC_2
4DPI_SAC_2_2	4DPI	2	SAC	2	4DPI_SAC_2
4DPI_SAC_2_1	4DPI	2	SAC	1	4DPI_SAC_2
4DPI_SAC_1_3	4DPI	1	SAC	3	4DPI_SAC_1
4DPI_SAC_1_2	4DPI	1	SAC	2	4DPI_SAC_1
4DPI_SAC_1_1	4DPI	1	SAC	1	4DPI_SAC_1
10DPI_interface_5_3	10DPI	5	interface	3	10DPI_interface_5
10DPI_interface_4_3	10DPI	4	interface	3	10DPI_interface_4
10DPI_interface_4_2	10DPI	4	interface	2	10DPI_interface_4
10DPI_interface_4_1	10DPI	4	interface	1	10DPI_interface_4
10DPI_interface_3_2	10DPI	3	interface	2	10DPI_interface_3
10DPI_interface_3_1	10DPI	3	interface	1	10DPI_interface_3
10DPI_cortex_4_4	10DPI	4	cortex	4	10DPI_cortex_4
10DPI_cortex_4_3	10DPI	4	cortex	3	10DPI_cortex_4
10DPI_cortex_3_3	10DPI	3	cortex	3	10DPI_cortex_3
10DPI_cortex_3_2	10DPI	3	cortex	2	10DPI_cortex_3
10DPI_SAC_4_3	10DPI	4	SAC	3	10DPI_SAC_4
10DPI_SAC_4_2	10DPI	4	SAC	2	10DPI_SAC_4
10DPI_SAC_4_1	10DPI	4	SAC	1	10DPI_SAC_4
10DPI_SAC_3_3	10DPI	3	SAC	3	10DPI_SAC_3
10DPI_SAC_3_2	10DPI	3	SAC	2	10DPI_SAC_3
10DPI_SAC_3_1	10DPI	3	SAC	1	10DPI_SAC_3
10DPI_SAC_2_3	10DPI	2	SAC	3	10DPI_SAC_2
10DPI_SAC_2_2	10DPI	2	SAC	2	10DPI_SAC_2

10DPI_SAC_2_1	10DPI	2	SAC	1	10DPI_SAC_2
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Protein IDs	BioRep_TechRep	DPI_Region
index		
4DPI_interface_2_4	2_4	4DPI_interface
4DPI_interface_2_2	2_2	4DPI_interface
4DPI_interface_2_1	2_1	4DPI_interface
4DPI_interface_1_2	1_2	4DPI_interface
4DPI_interface_1_1	1_1	4DPI_interface
4DPI_cortex_3_3	3_3	4DPI_cortex
4DPI_cortex_3_2	3_2	4DPI_cortex
4DPI_cortex_2_3	2_3	4DPI_cortex
4DPI_cortex_2_2	2_2	4DPI_cortex
4DPI_cortex_2_1	2_1	4DPI_cortex
4DPI_cortex_1_2	1_2	4DPI_cortex
4DPI_cortex_1_1	1_1	4DPI_cortex
4DPI_SAC_3_3	3_3	4DPI_SAC
4DPI_SAC_3_2	3_2	4DPI_SAC
4DPI_SAC_2_3	2_3	4DPI_SAC
4DPI_SAC_2_2	2_2	4DPI_SAC
4DPI_SAC_2_1	2_1	4DPI_SAC
4DPI_SAC_1_3	1_3	4DPI_SAC
4DPI_SAC_1_2	1_2	4DPI_SAC
4DPI_SAC_1_1	1_1	4DPI_SAC
10DPI_interface_5_3	5_3	10DPI_interface
10DPI_interface_4_3	4_3	10DPI_interface
10DPI_interface_4_2	4_2	10DPI_interface
10DPI_interface_4_1	4_1	10DPI_interface
10DPI_interface_3_2	3_2	10DPI_interface
10DPI_interface_3_1	3_1	10DPI_interface
10DPI_cortex_4_4	4_4	10DPI_cortex
10DPI_cortex_4_3	4_3	10DPI_cortex
10DPI_cortex_3_3	3_3	10DPI_cortex
10DPI_cortex_3_2	3_2	10DPI_cortex
10DPI_SAC_4_3	4_3	10DPI_SAC
10DPI_SAC_4_2	4_2	10DPI_SAC
10DPI_SAC_4_1	4_1	10DPI_SAC
10DPI_SAC_3_3	3_3	10DPI_SAC
10DPI_SAC_3_2	3_2	10DPI_SAC
10DPI_SAC_3_1	3_1	10DPI_SAC
10DPI_SAC_2_3	2_3	10DPI_SAC
10DPI_SAC_2_2	2_2	10DPI_SAC
10DPI_SAC_2_1	2_1	10DPI_SAC

[39 rows x 3620 columns]

```
[50]: combined_lfq = combined_lfq.replace(0, np.nan)
combined_lfq
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```
[50]: Protein IDs      G3UZW7;A0A023T778;Q9CQL1;P61327  \
index
4DPI_interface_2_4      NaN
4DPI_interface_2_2      NaN
4DPI_interface_2_1      NaN
4DPI_interface_1_2      NaN
4DPI_interface_1_1      NaN
4DPI_cortex_3_3         NaN
4DPI_cortex_3_2         NaN
4DPI_cortex_2_3         NaN
4DPI_cortex_2_2         NaN
4DPI_cortex_2_1         NaN
4DPI_cortex_1_2         NaN
4DPI_cortex_1_1         NaN
4DPI_SAC_3_3           NaN
4DPI_SAC_3_2           NaN
4DPI_SAC_2_3           NaN
4DPI_SAC_2_2           NaN
4DPI_SAC_2_1           189280.0
4DPI_SAC_1_3           561840.0
4DPI_SAC_1_2           779590.0
4DPI_SAC_1_1           NaN
10DPI_interface_5_3     NaN
10DPI_interface_4_3     NaN
10DPI_interface_4_2     222210.0
10DPI_interface_4_1     NaN
10DPI_interface_3_2     NaN
10DPI_interface_3_1     NaN
10DPI_cortex_4_4        NaN
10DPI_cortex_4_3        NaN
10DPI_cortex_3_3        NaN
10DPI_cortex_3_2        NaN
10DPI_SAC_4_3           NaN
10DPI_SAC_4_2           NaN
10DPI_SAC_4_1           NaN
10DPI_SAC_3_3           NaN
10DPI_SAC_3_2           NaN
10DPI_SAC_3_1           NaN
10DPI_SAC_2_3           NaN
10DPI_SAC_2_2           NaN
10DPI_SAC_2_1           NaN
```

```
Protein IDs      070589;A0A067XG53;F6Y9I5  \
index
```

4DPI_interface_2_4	NaN
4DPI_interface_2_2	NaN
4DPI_interface_2_1	NaN
4DPI_interface_1_2	NaN
4DPI_interface_1_1	NaN
4DPI_cortex_3_3	NaN
4DPI_cortex_3_2	NaN
4DPI_cortex_2_3	NaN
4DPI_cortex_2_2	NaN
4DPI_cortex_2_1	497950.0
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	NaN
4DPI_SAC_3_3	NaN
4DPI_SAC_3_2	NaN
4DPI_SAC_2_3	NaN
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	235580.0
4DPI_SAC_1_3	NaN
4DPI_SAC_1_2	NaN
4DPI_SAC_1_1	NaN
10DPI_interface_5_3	NaN
10DPI_interface_4_3	NaN
10DPI_interface_4_2	NaN
10DPI_interface_4_1	NaN
10DPI_interface_3_2	NaN
10DPI_interface_3_1	NaN
10DPI_cortex_4_4	NaN
10DPI_cortex_4_3	NaN
10DPI_cortex_3_3	NaN
10DPI_cortex_3_2	434700.0
10DPI_SAC_4_3	NaN
10DPI_SAC_4_2	NaN
10DPI_SAC_4_1	NaN
10DPI_SAC_3_3	NaN
10DPI_SAC_3_2	NaN
10DPI_SAC_3_1	NaN
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	NaN
10DPI_SAC_2_1	NaN

Protein IDs AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA075B5K8;F6XW
B2;AOA140T8N1;AOA0G2JDE5;P01631 \

index

4DPI_interface_2_4	NaN
4DPI_interface_2_2	NaN
4DPI_interface_2_1	NaN
4DPI_interface_1_2	NaN

4DPI_interface_1_1	NaN
4DPI_cortex_3_3	NaN
4DPI_cortex_3_2	NaN
4DPI_cortex_2_3	NaN
4DPI_cortex_2_2	NaN
4DPI_cortex_2_1	NaN
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	NaN
4DPI_SAC_3_3	NaN
4DPI_SAC_3_2	NaN
4DPI_SAC_2_3	NaN
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	NaN
4DPI_SAC_1_3	NaN
4DPI_SAC_1_2	NaN
4DPI_SAC_1_1	NaN
10DPI_interface_5_3	NaN
10DPI_interface_4_3	NaN
10DPI_interface_4_2	NaN
10DPI_interface_4_1	NaN
10DPI_interface_3_2	NaN
10DPI_interface_3_1	NaN
10DPI_cortex_4_4	NaN
10DPI_cortex_4_3	NaN
10DPI_cortex_3_3	855310.0
10DPI_cortex_3_2	498950.0
10DPI_SAC_4_3	1674100.0
10DPI_SAC_4_2	431080.0
10DPI_SAC_4_1	548100.0
10DPI_SAC_3_3	NaN
10DPI_SAC_3_2	NaN
10DPI_SAC_3_1	NaN
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	NaN
10DPI_SAC_2_1	NaN

Protein IDs AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1 \

index

4DPI_interface_2_4	NaN
4DPI_interface_2_2	NaN
4DPI_interface_2_1	NaN
4DPI_interface_1_2	NaN
4DPI_interface_1_1	NaN
4DPI_cortex_3_3	NaN
4DPI_cortex_3_2	NaN
4DPI_cortex_2_3	NaN
4DPI_cortex_2_2	NaN

4DPI_cortex_2_1	NaN
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	NaN
4DPI_SAC_3_3	NaN
4DPI_SAC_3_2	NaN
4DPI_SAC_2_3	NaN
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	NaN
4DPI_SAC_1_3	NaN
4DPI_SAC_1_2	NaN
4DPI_SAC_1_1	NaN
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10DPI_interface_4_3	207140.0
10DPI_interface_4_2	NaN
10DPI_interface_4_1	NaN
10DPI_interface_3_2	NaN
10DPI_interface_3_1	NaN
10DPI_cortex_4_4	NaN
10DPI_cortex_4_3	NaN
10DPI_cortex_3_3	NaN
10DPI_cortex_3_2	NaN
10DPI_SAC_4_3	202930.0
10DPI_SAC_4_2	290690.0
10DPI_SAC_4_1	NaN
10DPI_SAC_3_3	NaN
10DPI_SAC_3_2	NaN
10DPI_SAC_3_1	NaN
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	NaN
10DPI_SAC_2_1	NaN

Protein IDs A0A075B5P3;A0A0A6YVP0;P01867 \

index

4DPI_interface_2_4	NaN
4DPI_interface_2_2	NaN
4DPI_interface_2_1	NaN
4DPI_interface_1_2	NaN
4DPI_interface_1_1	NaN
4DPI_cortex_3_3	NaN
4DPI_cortex_3_2	NaN
4DPI_cortex_2_3	NaN
4DPI_cortex_2_2	NaN
4DPI_cortex_2_1	NaN
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	NaN
4DPI_SAC_3_3	NaN
4DPI_SAC_3_2	NaN

4DPI_SAC_2_3	NaN
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	NaN
4DPI_SAC_1_3	NaN
4DPI_SAC_1_2	NaN
4DPI_SAC_1_1	NaN
10DPI_interface_5_3	NaN
10DPI_interface_4_3	1183700.0
10DPI_interface_4_2	2295600.0
10DPI_interface_4_1	1475700.0
10DPI_interface_3_2	869990.0
10DPI_interface_3_1	NaN
10DPI_cortex_4_4	NaN
10DPI_cortex_4_3	NaN
10DPI_cortex_3_3	NaN
10DPI_cortex_3_2	NaN
10DPI_SAC_4_3	1207600.0
10DPI_SAC_4_2	1526100.0
10DPI_SAC_4_1	1165600.0
10DPI_SAC_3_3	NaN
10DPI_SAC_3_2	NaN
10DPI_SAC_3_1	724260.0
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	NaN
10DPI_SAC_2_1	NaN

Protein IDs AOA075B5P5;AOA1Y7VJN6;P03987 \

index

4DPI_interface_2_4	NaN
4DPI_interface_2_2	NaN
4DPI_interface_2_1	NaN
4DPI_interface_1_2	NaN
4DPI_interface_1_1	NaN
4DPI_cortex_3_3	NaN
4DPI_cortex_3_2	NaN
4DPI_cortex_2_3	NaN
4DPI_cortex_2_2	NaN
4DPI_cortex_2_1	NaN
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	NaN
4DPI_SAC_3_3	NaN
4DPI_SAC_3_2	NaN
4DPI_SAC_2_3	NaN
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	NaN
4DPI_SAC_1_3	NaN
4DPI_SAC_1_2	NaN

4DPI_SAC_1_1	NaN
10DPI_interface_5_3	1109600.0
10DPI_interface_4_3	12624000.0
10DPI_interface_4_2	14386000.0
10DPI_interface_4_1	19824000.0
10DPI_interface_3_2	3005900.0
10DPI_interface_3_1	447980.0
10DPI_cortex_4_4	21601000.0
10DPI_cortex_4_3	526710.0
10DPI_cortex_3_3	1102700.0
10DPI_cortex_3_2	654170.0
10DPI_SAC_4_3	29513000.0
10DPI_SAC_4_2	13752000.0
10DPI_SAC_4_1	17690000.0
10DPI_SAC_3_3	848950.0
10DPI_SAC_3_2	997060.0
10DPI_SAC_3_1	295760.0
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	1941400.0
10DPI_SAC_2_1	NaN

Protein IDs AOA075B6A0;AOA075B5P6;P01872 \

index

4DPI_interface_2_4	11850000.0
4DPI_interface_2_2	13198000.0
4DPI_interface_2_1	14041000.0
4DPI_interface_1_2	8327300.0
4DPI_interface_1_1	9088300.0
4DPI_cortex_3_3	28479000.0
4DPI_cortex_3_2	9475600.0
4DPI_cortex_2_3	12548000.0
4DPI_cortex_2_2	17613000.0
4DPI_cortex_2_1	18376000.0
4DPI_cortex_1_2	13749000.0
4DPI_cortex_1_1	19070000.0
4DPI_SAC_3_3	4038600.0
4DPI_SAC_3_2	1051500.0
4DPI_SAC_2_3	10500000.0
4DPI_SAC_2_2	4068800.0
4DPI_SAC_2_1	6212200.0
4DPI_SAC_1_3	7820100.0
4DPI_SAC_1_2	3859700.0
4DPI_SAC_1_1	4501600.0
10DPI_interface_5_3	7111900.0
10DPI_interface_4_3	15394000.0
10DPI_interface_4_2	20355000.0
10DPI_interface_4_1	24453000.0

10DPI_interface_3_2	6101300.0
10DPI_interface_3_1	9250300.0
10DPI_cortex_4_4	49938000.0
10DPI_cortex_4_3	16417000.0
10DPI_cortex_3_3	16583000.0
10DPI_cortex_3_2	14249000.0
10DPI_SAC_4_3	36836000.0
10DPI_SAC_4_2	21015000.0
10DPI_SAC_4_1	20078000.0
10DPI_SAC_3_3	5357600.0
10DPI_SAC_3_2	2988700.0
10DPI_SAC_3_1	6025900.0
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	6863300.0
10DPI_SAC_2_1	2568900.0

Protein IDs	AOA075B5R2;AOA075B5R3;AOA0A6YX91;P01786 \
index	
4DPI_interface_2_4	489990.0
4DPI_interface_2_2	NaN
4DPI_interface_2_1	1280300.0
4DPI_interface_1_2	NaN
4DPI_interface_1_1	NaN
4DPI_cortex_3_3	NaN
4DPI_cortex_3_2	NaN
4DPI_cortex_2_3	537080.0
4DPI_cortex_2_2	1492000.0
4DPI_cortex_2_1	247720.0
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	NaN
4DPI_SAC_3_3	NaN
4DPI_SAC_3_2	NaN
4DPI_SAC_2_3	NaN
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	NaN
4DPI_SAC_1_3	NaN
4DPI_SAC_1_2	NaN
4DPI_SAC_1_1	NaN
10DPI_interface_5_3	923480.0
10DPI_interface_4_3	2742000.0
10DPI_interface_4_2	2199900.0
10DPI_interface_4_1	2193900.0
10DPI_interface_3_2	875140.0
10DPI_interface_3_1	1436900.0
10DPI_cortex_4_4	1717800.0
10DPI_cortex_4_3	1395700.0
10DPI_cortex_3_3	880010.0

10DPI_cortex_3_2	708140.0
10DPI_SAC_4_3	1401500.0
10DPI_SAC_4_2	NaN
10DPI_SAC_4_1	1453700.0
10DPI_SAC_3_3	536050.0
10DPI_SAC_3_2	2143600.0
10DPI_SAC_3_1	944670.0
10DPI_SAC_2_3	NaN
10DPI_SAC_2_2	533920.0
10DPI_SAC_2_1	NaN

Protein IDs	AOA075B5S5;AOA0A6YXL5	AOA075B5T3;J3QK03;AOA0A6YWS9	...	\
index			...	
4DPI_interface_2_4	NaN	NaN	...	
4DPI_interface_2_2	NaN	NaN	...	
4DPI_interface_2_1	NaN	NaN	...	
4DPI_interface_1_2	NaN	NaN	...	
4DPI_interface_1_1	NaN	NaN	...	
4DPI_cortex_3_3	NaN	NaN	...	
4DPI_cortex_3_2	NaN	NaN	...	
4DPI_cortex_2_3	NaN	NaN	...	
4DPI_cortex_2_2	NaN	NaN	...	
4DPI_cortex_2_1	NaN	NaN	...	
4DPI_cortex_1_2	NaN	NaN	...	
4DPI_cortex_1_1	NaN	NaN	...	
4DPI_SAC_3_3	NaN	NaN	...	
4DPI_SAC_3_2	NaN	NaN	...	
4DPI_SAC_2_3	NaN	NaN	...	
4DPI_SAC_2_2	NaN	NaN	...	
4DPI_SAC_2_1	NaN	NaN	...	
4DPI_SAC_1_3	NaN	NaN	...	
4DPI_SAC_1_2	NaN	NaN	...	
4DPI_SAC_1_1	NaN	NaN	...	
10DPI_interface_5_3	582000.0	624130.0	...	
10DPI_interface_4_3	10474000.0	1520300.0	...	
10DPI_interface_4_2	4267600.0	3319600.0	...	
10DPI_interface_4_1	1899800.0	1112100.0	...	
10DPI_interface_3_2	521140.0	826650.0	...	
10DPI_interface_3_1	554440.0	NaN	...	
10DPI_cortex_4_4	3546300.0	2818100.0	...	
10DPI_cortex_4_3	NaN	2965700.0	...	
10DPI_cortex_3_3	NaN	992900.0	...	
10DPI_cortex_3_2	739180.0	NaN	...	
10DPI_SAC_4_3	2447600.0	5502600.0	...	
10DPI_SAC_4_2	3972000.0	2063300.0	...	
10DPI_SAC_4_1	8814000.0	2556300.0	...	
10DPI_SAC_3_3	NaN	1867700.0	...	

10DPI_SAC_3_2	NaN	766080.0	...
10DPI_SAC_3_1	NaN	NaN	...
10DPI_SAC_2_3	NaN	NaN	...
10DPI_SAC_2_2	NaN	NaN	...
10DPI_SAC_2_1	NaN	NaN	...

Protein IDs Q9Z315;A0A494B9E9 \

index

4DPI_interface_2_4	954220.0
4DPI_interface_2_2	1377400.0
4DPI_interface_2_1	922440.0
4DPI_interface_1_2	NaN
4DPI_interface_1_1	1117600.0
4DPI_cortex_3_3	638210.0
4DPI_cortex_3_2	661230.0
4DPI_cortex_2_3	NaN
4DPI_cortex_2_2	494860.0
4DPI_cortex_2_1	674540.0
4DPI_cortex_1_2	NaN
4DPI_cortex_1_1	334760.0
4DPI_SAC_3_3	1396800.0
4DPI_SAC_3_2	1242700.0
4DPI_SAC_2_3	1401200.0
4DPI_SAC_2_2	NaN
4DPI_SAC_2_1	458240.0
4DPI_SAC_1_3	780230.0
4DPI_SAC_1_2	1219000.0
4DPI_SAC_1_1	600830.0
10DPI_interface_5_3	1281000.0
10DPI_interface_4_3	2218300.0
10DPI_interface_4_2	1495700.0
10DPI_interface_4_1	927020.0
10DPI_interface_3_2	1865800.0
10DPI_interface_3_1	1160000.0
10DPI_cortex_4_4	946380.0
10DPI_cortex_4_3	NaN
10DPI_cortex_3_3	633750.0
10DPI_cortex_3_2	556170.0
10DPI_SAC_4_3	1137800.0
10DPI_SAC_4_2	1302200.0
10DPI_SAC_4_1	NaN
10DPI_SAC_3_3	1731200.0
10DPI_SAC_3_2	1444700.0
10DPI_SAC_3_1	1059500.0
10DPI_SAC_2_3	3343300.0
10DPI_SAC_2_2	800330.0
10DPI_SAC_2_1	2338200.0

Protein IDs	S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	S4R192	\
index			
4DPI_interface_2_4	NaN	NaN	
4DPI_interface_2_2	NaN	NaN	
4DPI_interface_2_1	NaN	NaN	
4DPI_interface_1_2	NaN	NaN	
4DPI_interface_1_1	NaN	NaN	
4DPI_cortex_3_3	NaN	NaN	
4DPI_cortex_3_2	NaN	NaN	
4DPI_cortex_2_3	662690.0	NaN	
4DPI_cortex_2_2	NaN	NaN	
4DPI_cortex_2_1	NaN	NaN	
4DPI_cortex_1_2	714480.0	NaN	
4DPI_cortex_1_1	716000.0	NaN	
4DPI_SAC_3_3	407450.0	NaN	
4DPI_SAC_3_2	NaN	357930.0	
4DPI_SAC_2_3	NaN	NaN	
4DPI_SAC_2_2	NaN	NaN	
4DPI_SAC_2_1	360500.0	169050.0	
4DPI_SAC_1_3	610920.0	NaN	
4DPI_SAC_1_2	326340.0	NaN	
4DPI_SAC_1_1	NaN	251670.0	
10DPI_interface_5_3	126670.0	NaN	
10DPI_interface_4_3	NaN	305120.0	
10DPI_interface_4_2	NaN	549550.0	
10DPI_interface_4_1	NaN	NaN	
10DPI_interface_3_2	NaN	364880.0	
10DPI_interface_3_1	NaN	781480.0	
10DPI_cortex_4_4	NaN	NaN	
10DPI_cortex_4_3	NaN	NaN	
10DPI_cortex_3_3	NaN	NaN	
10DPI_cortex_3_2	NaN	NaN	
10DPI_SAC_4_3	NaN	NaN	
10DPI_SAC_4_2	NaN	NaN	
10DPI_SAC_4_1	NaN	326950.0	
10DPI_SAC_3_3	NaN	423330.0	
10DPI_SAC_3_2	NaN	NaN	
10DPI_SAC_3_1	NaN	NaN	
10DPI_SAC_2_3	NaN	NaN	
10DPI_SAC_2_2	NaN	NaN	
10DPI_SAC_2_1	NaN	NaN	

Protein IDs	TimePoint	BioRep	Region	TechRep	DPI_Region_BioRep	\
index						
4DPI_interface_2_4	4DPI	2	interface	4	4DPI_interface_2	
4DPI_interface_2_2	4DPI	2	interface	2	4DPI_interface_2	

4DPI_interface_2_1	4DPI	2	interface	1	4DPI_interface_2
4DPI_interface_1_2	4DPI	1	interface	2	4DPI_interface_1
4DPI_interface_1_1	4DPI	1	interface	1	4DPI_interface_1
4DPI_cortex_3_3	4DPI	3	cortex	3	4DPI_cortex_3
4DPI_cortex_3_2	4DPI	3	cortex	2	4DPI_cortex_3
4DPI_cortex_2_3	4DPI	2	cortex	3	4DPI_cortex_2
4DPI_cortex_2_2	4DPI	2	cortex	2	4DPI_cortex_2
4DPI_cortex_2_1	4DPI	2	cortex	1	4DPI_cortex_2
4DPI_cortex_1_2	4DPI	1	cortex	2	4DPI_cortex_1
4DPI_cortex_1_1	4DPI	1	cortex	1	4DPI_cortex_1
4DPI_SAC_3_3	4DPI	3	SAC	3	4DPI_SAC_3
4DPI_SAC_3_2	4DPI	3	SAC	2	4DPI_SAC_3
4DPI_SAC_2_3	4DPI	2	SAC	3	4DPI_SAC_2
4DPI_SAC_2_2	4DPI	2	SAC	2	4DPI_SAC_2
4DPI_SAC_2_1	4DPI	2	SAC	1	4DPI_SAC_2
4DPI_SAC_1_3	4DPI	1	SAC	3	4DPI_SAC_1
4DPI_SAC_1_2	4DPI	1	SAC	2	4DPI_SAC_1
4DPI_SAC_1_1	4DPI	1	SAC	1	4DPI_SAC_1
10DPI_interface_5_3	10DPI	5	interface	3	10DPI_interface_5
10DPI_interface_4_3	10DPI	4	interface	3	10DPI_interface_4
10DPI_interface_4_2	10DPI	4	interface	2	10DPI_interface_4
10DPI_interface_4_1	10DPI	4	interface	1	10DPI_interface_4
10DPI_interface_3_2	10DPI	3	interface	2	10DPI_interface_3
10DPI_interface_3_1	10DPI	3	interface	1	10DPI_interface_3
10DPI_cortex_4_4	10DPI	4	cortex	4	10DPI_cortex_4
10DPI_cortex_4_3	10DPI	4	cortex	3	10DPI_cortex_4
10DPI_cortex_3_3	10DPI	3	cortex	3	10DPI_cortex_3
10DPI_cortex_3_2	10DPI	3	cortex	2	10DPI_cortex_3
10DPI_SAC_4_3	10DPI	4	SAC	3	10DPI_SAC_4
10DPI_SAC_4_2	10DPI	4	SAC	2	10DPI_SAC_4
10DPI_SAC_4_1	10DPI	4	SAC	1	10DPI_SAC_4
10DPI_SAC_3_3	10DPI	3	SAC	3	10DPI_SAC_3
10DPI_SAC_3_2	10DPI	3	SAC	2	10DPI_SAC_3
10DPI_SAC_3_1	10DPI	3	SAC	1	10DPI_SAC_3
10DPI_SAC_2_3	10DPI	2	SAC	3	10DPI_SAC_2
10DPI_SAC_2_2	10DPI	2	SAC	2	10DPI_SAC_2
10DPI_SAC_2_1	10DPI	2	SAC	1	10DPI_SAC_2

Protein IDs index	BioRep_TechRep	DPI_Region
4DPI_interface_2_4	2_4	4DPI_interface
4DPI_interface_2_2	2_2	4DPI_interface
4DPI_interface_2_1	2_1	4DPI_interface
4DPI_interface_1_2	1_2	4DPI_interface
4DPI_interface_1_1	1_1	4DPI_interface
4DPI_cortex_3_3	3_3	4DPI_cortex
4DPI_cortex_3_2	3_2	4DPI_cortex

4DPI_cortex_2_3	2_3	4DPI_cortex
4DPI_cortex_2_2	2_2	4DPI_cortex
4DPI_cortex_2_1	2_1	4DPI_cortex
4DPI_cortex_1_2	1_2	4DPI_cortex
4DPI_cortex_1_1	1_1	4DPI_cortex
4DPI_SAC_3_3	3_3	4DPI_SAC
4DPI_SAC_3_2	3_2	4DPI_SAC
4DPI_SAC_2_3	2_3	4DPI_SAC
4DPI_SAC_2_2	2_2	4DPI_SAC
4DPI_SAC_2_1	2_1	4DPI_SAC
4DPI_SAC_1_3	1_3	4DPI_SAC
4DPI_SAC_1_2	1_2	4DPI_SAC
4DPI_SAC_1_1	1_1	4DPI_SAC
10DPI_interface_5_3	5_3	10DPI_interface
10DPI_interface_4_3	4_3	10DPI_interface
10DPI_interface_4_2	4_2	10DPI_interface
10DPI_interface_4_1	4_1	10DPI_interface
10DPI_interface_3_2	3_2	10DPI_interface
10DPI_interface_3_1	3_1	10DPI_interface
10DPI_cortex_4_4	4_4	10DPI_cortex
10DPI_cortex_4_3	4_3	10DPI_cortex
10DPI_cortex_3_3	3_3	10DPI_cortex
10DPI_cortex_3_2	3_2	10DPI_cortex
10DPI_SAC_4_3	4_3	10DPI_SAC
10DPI_SAC_4_2	4_2	10DPI_SAC
10DPI_SAC_4_1	4_1	10DPI_SAC
10DPI_SAC_3_3	3_3	10DPI_SAC
10DPI_SAC_3_2	3_2	10DPI_SAC
10DPI_SAC_3_1	3_1	10DPI_SAC
10DPI_SAC_2_3	2_3	10DPI_SAC
10DPI_SAC_2_2	2_2	10DPI_SAC
10DPI_SAC_2_1	2_1	10DPI_SAC

[39 rows x 3620 columns]

```
[51]: combined_lfq = combined_lfq.dropna(axis=1)
combined_lfq.shape
```

[51]: (39, 294)

0.2 Combined analysis

```
[52]: combined_lfq.head()
```

```
[52]: Protein IDs      AOA0J9YTY0;AOA0J9YUL3;Q8C1B7;AOA0J9YUV6;AOA0J9YVA6  \
index
4DPI_interface_2_4      8884400.0
4DPI_interface_2_2      4805100.0
```

4DPI_interface_2_1	4679400.0
4DPI_interface_1_2	2752000.0
4DPI_interface_1_1	6446300.0

Protein IDs A0A0J9YUZ4;P63158;A0A0J9YUD8;D3YVC6;D3YZ18 \

index

4DPI_interface_2_4	13795000.0
4DPI_interface_2_2	16043000.0
4DPI_interface_2_1	14321000.0
4DPI_interface_1_2	2559000.0
4DPI_interface_1_1	18740000.0

Protein IDs A0A0R4J093;Q9DBP5 A0A0R4J0I1;P07759;E9Q499;Q80X76 \

index

4DPI_interface_2_4	5749000.0	73676000.0
4DPI_interface_2_2	10405000.0	50598000.0
4DPI_interface_2_1	9228800.0	105190000.0
4DPI_interface_1_2	6292300.0	22129000.0
4DPI_interface_1_1	18973000.0	38062000.0

Protein IDs 070404;A0A0R4J0R1;A0A0U1RPE8 A0A0R4J0Z1;P08003 \

index

4DPI_interface_2_4	4863000.0	20661000.0
4DPI_interface_2_2	5048700.0	20709000.0
4DPI_interface_2_1	4421600.0	17844000.0
4DPI_interface_1_2	1316400.0	22085000.0
4DPI_interface_1_1	2682100.0	24030000.0

Protein IDs

E9QL31;A0A0R4J104;P98078;E9PX84;Q3TRE6;F6TQN9;Q9DCE6;E0CXT5;E0CZ53;E0CYJ2 \

index

4DPI_interface_2_4	10187000.0
4DPI_interface_2_2	10975000.0
4DPI_interface_2_1	8142400.0
4DPI_interface_1_2	704720.0
4DPI_interface_1_1	4963900.0

Protein IDs A0A0R4J195;Q9CSU0;F6YB25;F7DAY5;Q8VDS4 \

index

4DPI_interface_2_4	1394900.0
4DPI_interface_2_2	959390.0
4DPI_interface_2_1	549190.0
4DPI_interface_1_2	1506600.0
4DPI_interface_1_1	1400400.0

Protein IDs

A0A1B0GQU8;P35980;A0A1B0GSS8;A0A1B0GSF7;A0A1B0GSA8;A0A1B0GS28;A0A1B0GRZ3 \

```

index
4DPI_interface_2_4      11981000.0
4DPI_interface_2_2      22897000.0
4DPI_interface_2_1      20121000.0
4DPI_interface_1_2      36610000.0
4DPI_interface_1_1      23111000.0

Protein IDs      P06151;Q564E2;AOA1BOGSR9;AOA1BOGSX0;D3YZQ9;AOA1BOGSL7;AOA1BO
GT41;AOA1BOGQX5;AOA1BOGS79;AOA1BOGRW9;AOA1BOGRC1;AOA1BOGRS2;D3YVR7;D3YZE4;AOA1BO
GRE9;AOA1BOGSR2;P00342 \
index
4DPI_interface_2_4      48765000.0
4DPI_interface_2_2      84281000.0
4DPI_interface_2_1      62405000.0
4DPI_interface_1_2      61114000.0
4DPI_interface_1_1      36589000.0

Protein IDs      ...      Q9Z1Q5      Q9Z2D6      Q9Z2I8;AOA0N4SWD1      TimePoint \
index      ...
4DPI_interface_2_4      ...      15999000.0      3568300.0      12254000.0      4DPI
4DPI_interface_2_2      ...      21039000.0      1709200.0      17425000.0      4DPI
4DPI_interface_2_1      ...      27600000.0      1083700.0      14862000.0      4DPI
4DPI_interface_1_2      ...      22611000.0      2757000.0      7584200.0      4DPI
4DPI_interface_1_1      ...      23281000.0      5819700.0      16155000.0      4DPI

Protein IDs      BioRep      Region      TechRep      DPI_Region_BioRep \
index
4DPI_interface_2_4      2      interface      4      4DPI_interface_2
4DPI_interface_2_2      2      interface      2      4DPI_interface_2
4DPI_interface_2_1      2      interface      1      4DPI_interface_2
4DPI_interface_1_2      1      interface      2      4DPI_interface_1
4DPI_interface_1_1      1      interface      1      4DPI_interface_1

Protein IDs      BioRep_TechRep      DPI_Region
index
4DPI_interface_2_4      2_4      4DPI_interface
4DPI_interface_2_2      2_2      4DPI_interface
4DPI_interface_2_1      2_1      4DPI_interface
4DPI_interface_1_2      1_2      4DPI_interface
4DPI_interface_1_1      1_1      4DPI_interface

```

[5 rows x 294 columns]

```

[60]: ### Function to separate data and labels
def df_sep(label, dataframe):
    X = dataframe.drop(['TimePoint', 'TechRep', 'BioRep', 'Region'],
        → 'DPI_Region_BioRep', 'DPI_Region', 'BioRep_TechRep'], axis=1)

```

```
y = dataframe[label]
return X,y
```

```
[61]: X,Y = df_sep('DPI_Region', combined_lfq)
Y
y = pd.factorize(Y)[0]
```

```
[62]: X.to_csv('/Users/kavyasharman/Documents/Projects/microLESA/outs/' + date +
→ '_combined_X.csv')
```

0.3 PCA

```
[63]: pca = PCA()
pca.fit_transform(X)
```

```
[63]: array([[ -1.28844218e+09, -7.11460826e+08,  6.31811418e+08, ...,
          1.03164582e+07, -1.17846874e+07,  2.92942842e-07],
 [ 5.28450418e+08, -5.99037940e+08,  1.94400410e+08, ...,
          2.28226796e+07, -5.41262642e+04,  2.92942842e-07],
 [ 2.67146200e+08, -1.33471004e+09, -8.64781484e+06, ...,
          -2.71817551e+07,  2.19963656e+06,  2.92942842e-07],
 ...,
 [-2.75694213e+09, -1.98017577e+08,  4.88713329e+08, ...,
          -5.89531193e+06, -4.21840665e+06,  2.92942842e-07],
 [-2.09023427e+09, -1.40884868e+08,  1.02851665e+09, ...,
          -3.57478427e+06,  3.89483396e+06,  2.92942842e-07],
 [-2.01507561e+09, -1.92089267e+08, -3.76911129e+08, ...,
          9.41837935e+06, -6.70903517e+06,  2.92942842e-07]])
```

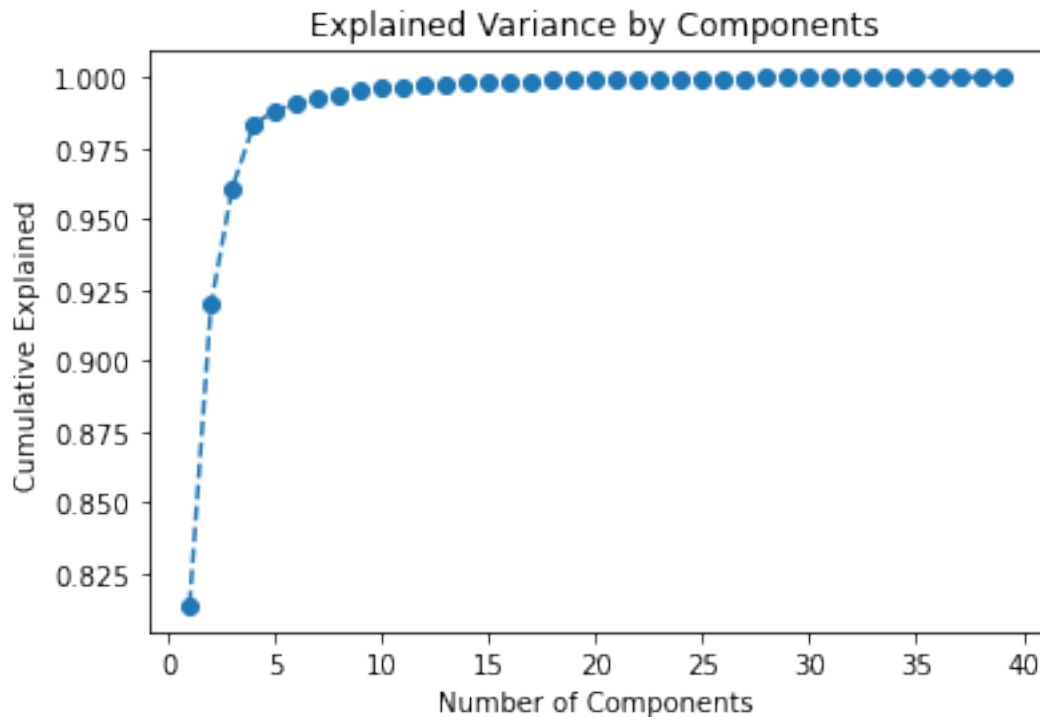
```
[64]: components = pca.components_
components.shape
```

```
[64]: (39, 287)
```

```
[65]: X.shape
```

```
[65]: (39, 287)
```

```
[66]: plt.figure()
plt.plot(range(1,40), pca.explained_variance_ratio_.cumsum(), marker = 'o',
→ linestyle = '--')
plt.title('Explained Variance by Components')
plt.xlabel('Number of Components')
plt.ylabel('Cumulative Explained')
plt.savefig('outs/' + date + '_combined_PCA_explainedVariance.svg')
```



```
[67]: pca_expl_var = pca.explained_variance_ratio_
      pca_expl_var
```

```
[67]: array([8.13496140e-01, 1.06088298e-01, 4.13704385e-02, 2.22253702e-02,
          5.13907045e-03, 2.23608178e-03, 1.84227069e-03, 1.69922876e-03,
          1.17444745e-03, 8.60060940e-04, 7.28495022e-04, 4.43397100e-04,
          3.70086472e-04, 3.25457317e-04, 3.03346162e-04, 2.37529489e-04,
          2.18973177e-04, 1.92780927e-04, 1.65274961e-04, 1.14038533e-04,
          9.70533536e-05, 9.03254288e-05, 7.95687166e-05, 7.23250045e-05,
          5.98391318e-05, 5.79711467e-05, 5.17241307e-05, 4.12563898e-05,
          3.54151933e-05, 3.43962530e-05, 2.82922579e-05, 2.28373465e-05,
          2.19429014e-05, 1.94594840e-05, 1.89470482e-05, 1.44635243e-05,
          1.38293277e-05, 9.56707717e-06, 5.19960053e-06])
```

```
[68]: scores_pca = pca.fit_transform(X)

      #pca loadings
      loadings = pd.DataFrame(pca.components_.T, index=X.columns)
```

```
[69]: scores_pca.shape
```

```
[69]: (39, 39)
```

```
[70]: pd.DataFrame(scores_pca).to_csv('/Users/kavyasharman/Documents/Projects/
      ↳microLESA/outs/' + date + '_combined_scores_pca.csv')
```

```
[71]: loadings.to_csv('/Users/kavyasharman/Documents/Projects/microLESA/outs/' + date_
      →+ '_combined_loadings_pca.csv')
```

0.3.1 Silhouette scores to determine k

```
[72]: from sklearn.datasets import make_blobs
      from sklearn.cluster import KMeans
      from sklearn.metrics import silhouette_samples, silhouette_score

      import matplotlib.pyplot as plt
      import matplotlib.cm as cm
      import numpy as np

      range_n_clusters = [2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15]
      silhouette_scores_euc = pd.DataFrame(columns = ['k', 'score'])

      for n_clusters in range_n_clusters:
          # Create a subplot with 1 row and 2 columns
          fig, (ax1, ax2) = plt.subplots(1, 2)
          fig.set_size_inches(18, 7)

          # The 1st subplot is the silhouette plot
          # The silhouette coefficient can range from -1, 1 but in this example all
          # lie within [-0.1, 1]
          ax1.set_xlim([-0.1, 1])
          # The (n_clusters+1)*10 is for inserting blank space between silhouette
          # plots of individual clusters, to demarcate them clearly.
          ax1.set_ylim([0, len(scores_pca) + (n_clusters + 1) * 10])

          # Initialize the clusterer with n_clusters value and a random generator
          # seed of 10 for reproducibility.
          clusterer = KMeans(n_clusters=n_clusters, random_state=42)
          cluster_labels = clusterer.fit_predict(scores_pca)

          # The silhouette_score gives the average value for all the samples.
          # This gives a perspective into the density and separation of the formed
          # clusters
          silhouette_avg = silhouette_score(scores_pca, cluster_labels)
          # print("For n_clusters =", n_clusters,
          #       "The average silhouette_score is :", silhouette_avg)
          silhouette_scores_euc = silhouette_scores_euc.append(pd.DataFrame({'k':
      →n_clusters, 'score': silhouette_avg}, index=[0]), ignore_index=True)

          # Compute the silhouette scores for each sample
          sample_silhouette_values = silhouette_samples(scores_pca, cluster_labels)
```

```

y_lower = 10
for i in range(n_clusters):
    # Aggregate the silhouette scores for samples belonging to
    # cluster i, and sort them
    ith_cluster_silhouette_values = \
        sample_silhouette_values[cluster_labels == i]

    ith_cluster_silhouette_values.sort()

    size_cluster_i = ith_cluster_silhouette_values.shape[0]
    y_upper = y_lower + size_cluster_i

    color = cm.nipy_spectral(float(i) / n_clusters)
    ax1.fill_betweenx(np.arange(y_lower, y_upper),
                      0, ith_cluster_silhouette_values,
                      facecolor=color, edgecolor=color, alpha=0.7)

    # Label the silhouette plots with their cluster numbers at the middle
    ax1.text(-0.05, y_lower + 0.5 * size_cluster_i, str(i))

    # Compute the new y_lower for next plot
    y_lower = y_upper + 10  # 10 for the 0 samples

ax1.set_title("The silhouette plot for the various clusters.")
ax1.set_xlabel("The silhouette coefficient values")
ax1.set_ylabel("Cluster label")

# The vertical line for average silhouette score of all the values
ax1.axvline(x=silhouette_avg, color="red", linestyle="--")

ax1.set_yticks([])  # Clear the yaxis labels / ticks
ax1.set_xticks([-0.1, 0, 0.2, 0.4, 0.6, 0.8, 1])

# 2nd Plot showing the actual clusters formed
colors = cm.nipy_spectral(cluster_labels.astype(float) / n_clusters)
ax2.scatter(scores_pca[:, 0], scores_pca[:, 1], marker='.', s=300, lw=0,
→alpha=0.7,
           c=colors, edgecolor='k')

# Labeling the clusters
centers = clusterer.cluster_centers_
# Draw white circles at cluster centers
ax2.scatter(centers[:, 0], centers[:, 1], marker='o',
           c="white", alpha=1, s=200, edgecolor='k')

for i, c in enumerate(centers):
    ax2.scatter(c[0], c[1], marker='$%d$' % i, alpha=1,

```

```

s=50, edgecolor='k')

ax2.set_title("The visualization of the clustered data.")
ax2.set_xlabel("Feature space for the 1st feature")
ax2.set_ylabel("Feature space for the 2nd feature")

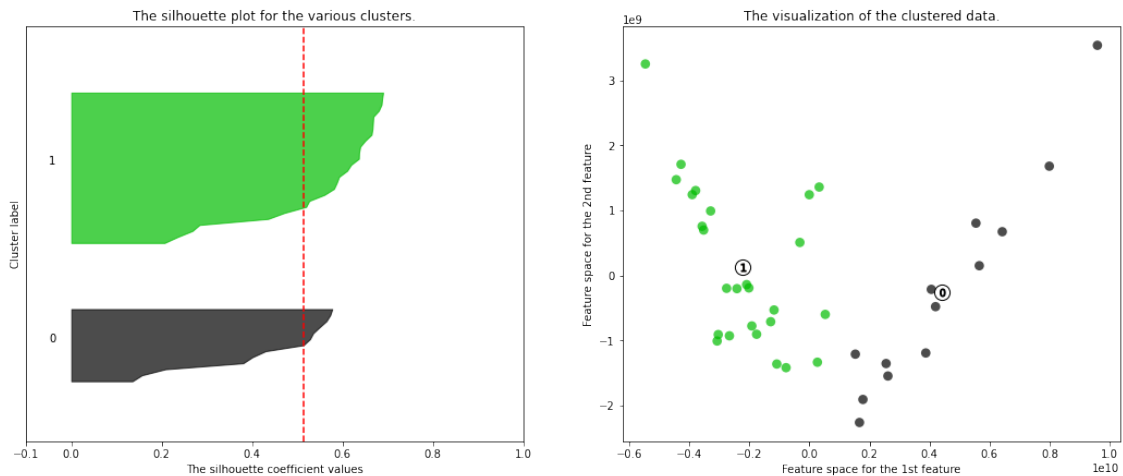
plt.suptitle(("Silhouette analysis for KMeans clustering on sample data "
              "with n_clusters = %d" % n_clusters),
             fontsize=14, fontweight='bold')

plt.savefig('outs/' + date + '_combined_silhouette_scores_euc_' +
            str(n_clusters) + '.svg')

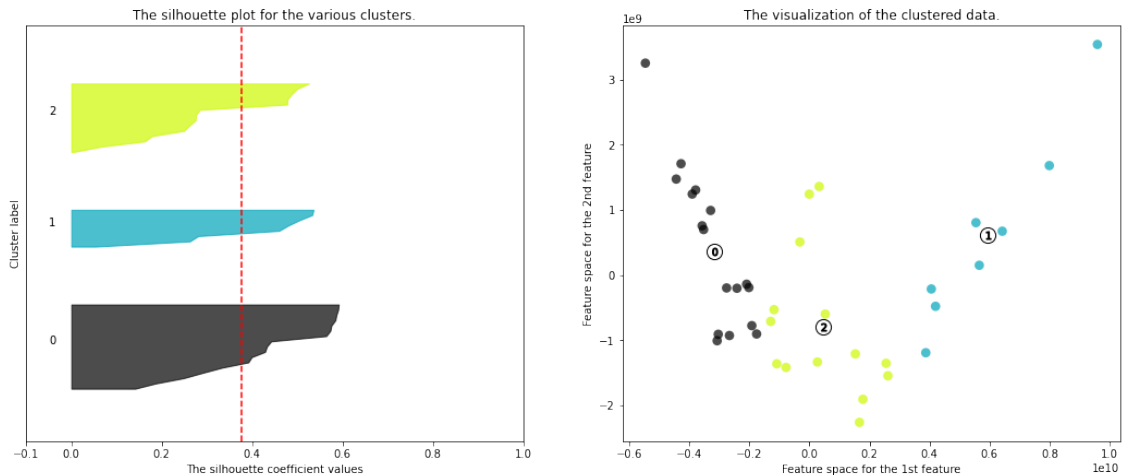
silhouette_scores_euc.to_csv('outs/' + date + '_combined_silhouette_scores_euc.'
                             + csv')

```

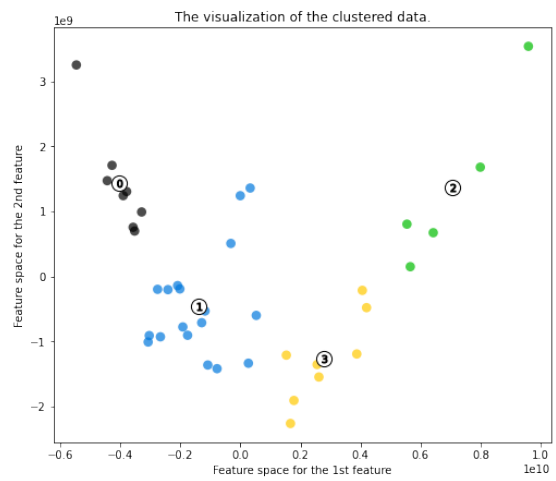
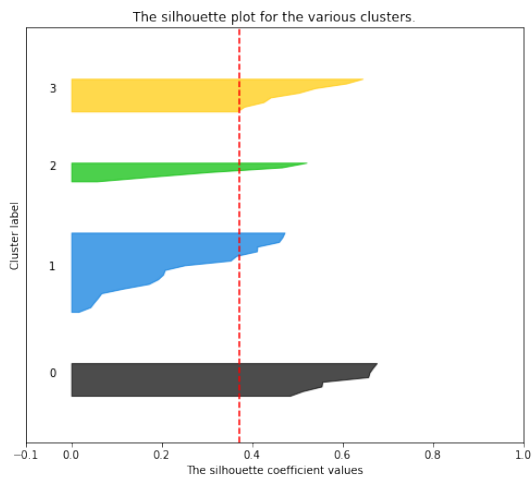
Silhouette analysis for KMeans clustering on sample data with n_clusters = 2



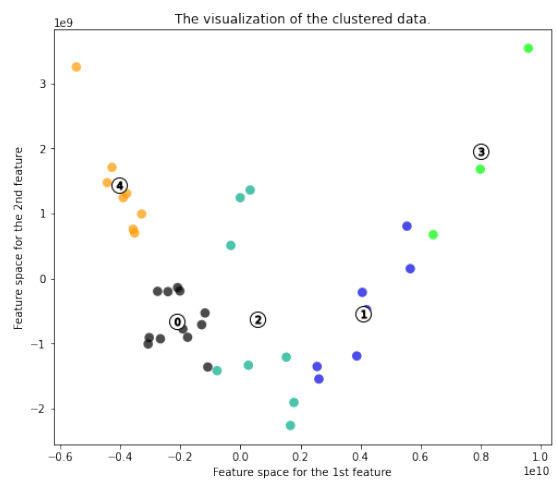
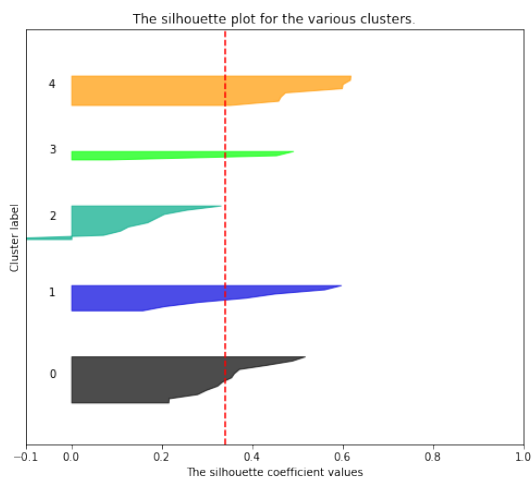
Silhouette analysis for KMeans clustering on sample data with n_clusters = 3



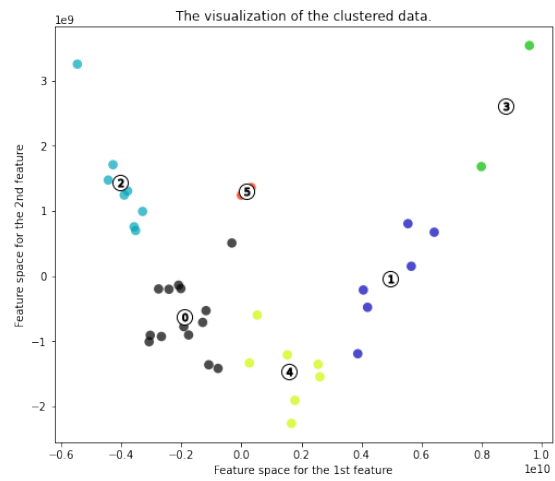
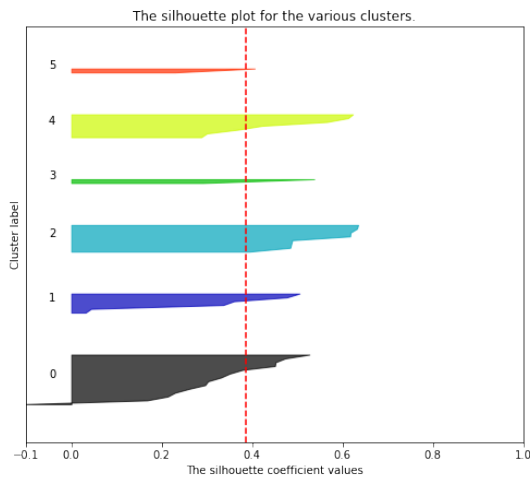
Silhouette analysis for KMeans clustering on sample data with $n_clusters = 4$



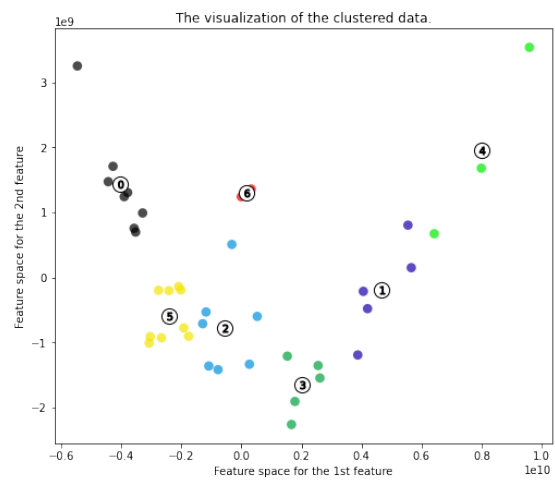
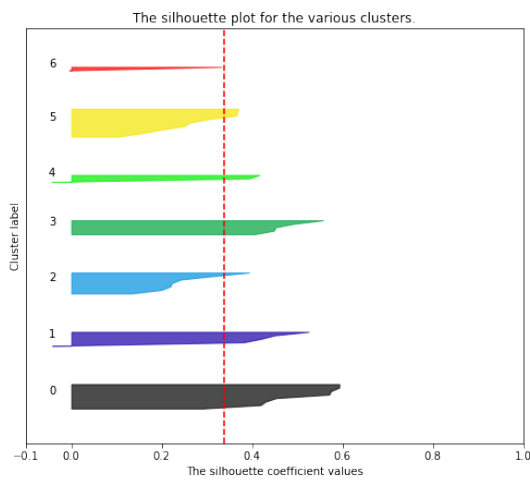
Silhouette analysis for KMeans clustering on sample data with $n_clusters = 5$



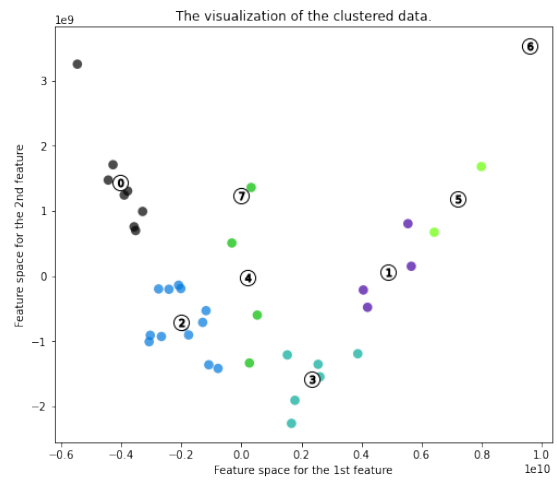
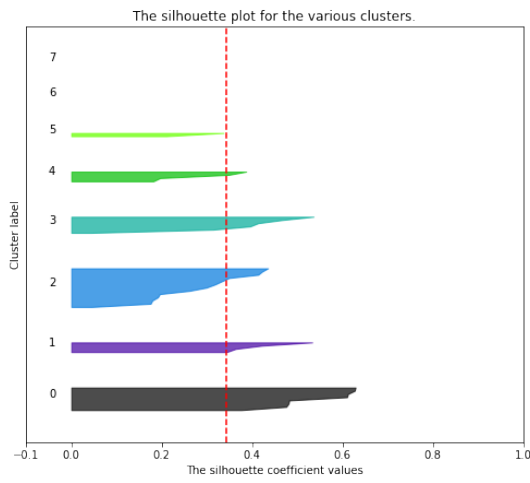
Silhouette analysis for KMeans clustering on sample data with $n_clusters = 6$



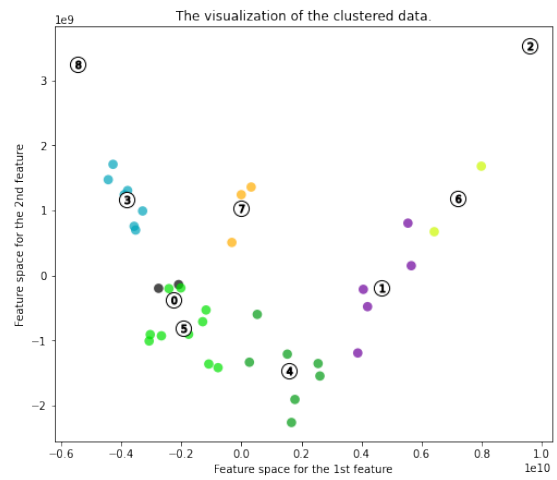
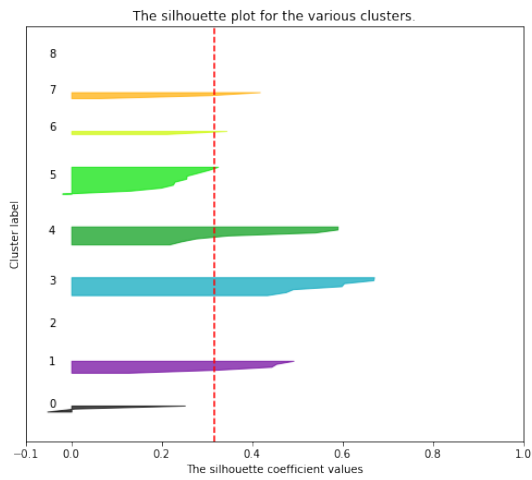
Silhouette analysis for KMeans clustering on sample data with $n_clusters = 7$



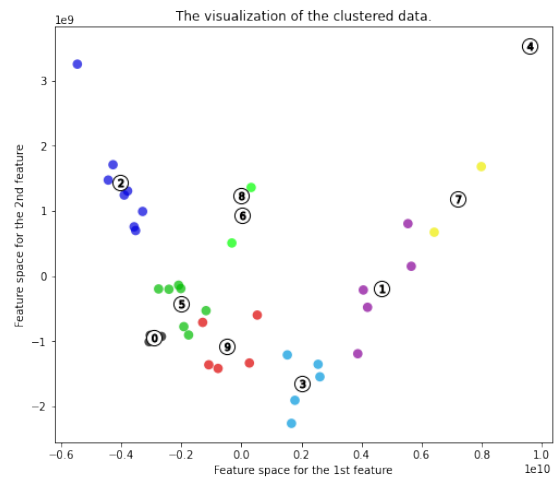
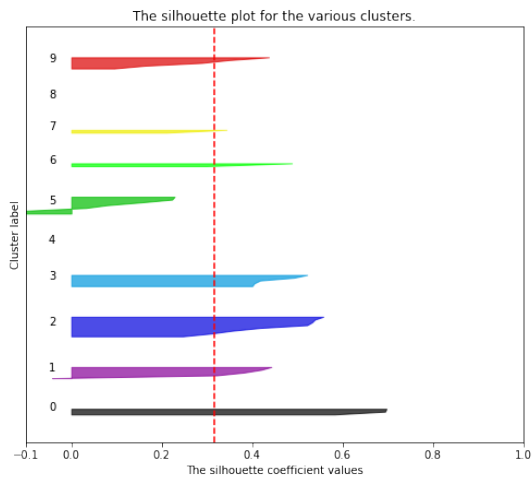
Silhouette analysis for KMeans clustering on sample data with n_clusters = 8



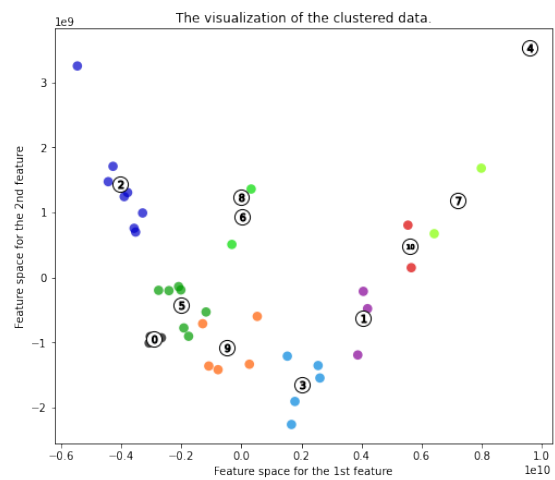
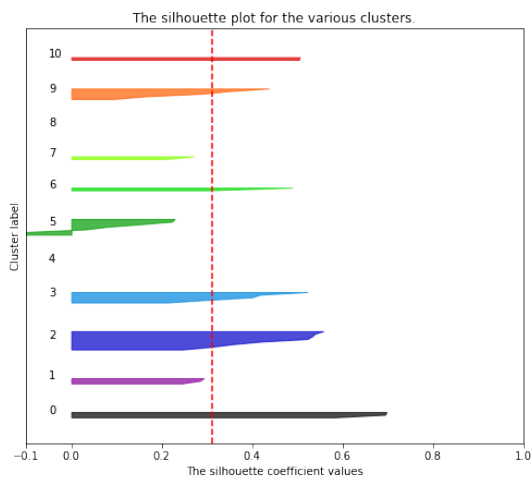
Silhouette analysis for KMeans clustering on sample data with n_clusters = 9



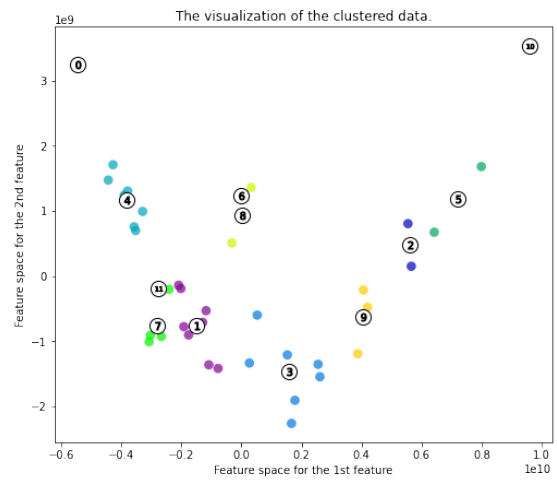
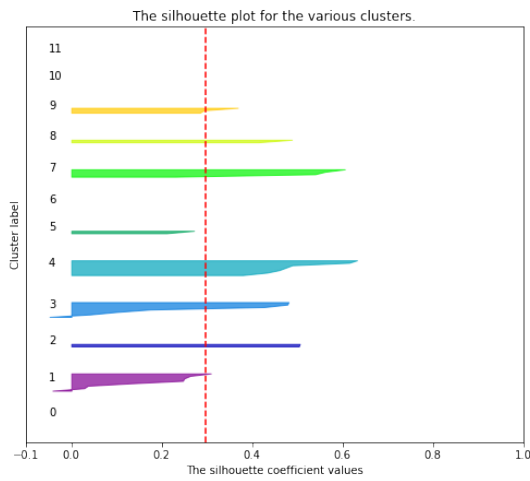
Silhouette analysis for KMeans clustering on sample data with n_clusters = 10



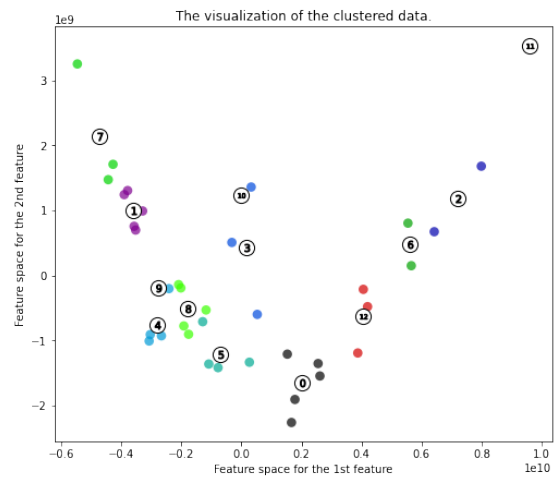
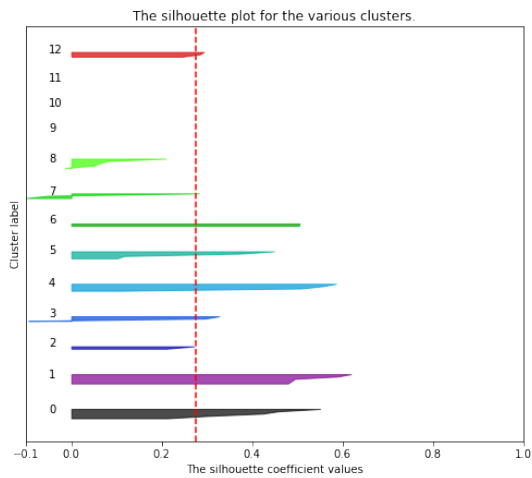
Silhouette analysis for KMeans clustering on sample data with n_clusters = 11



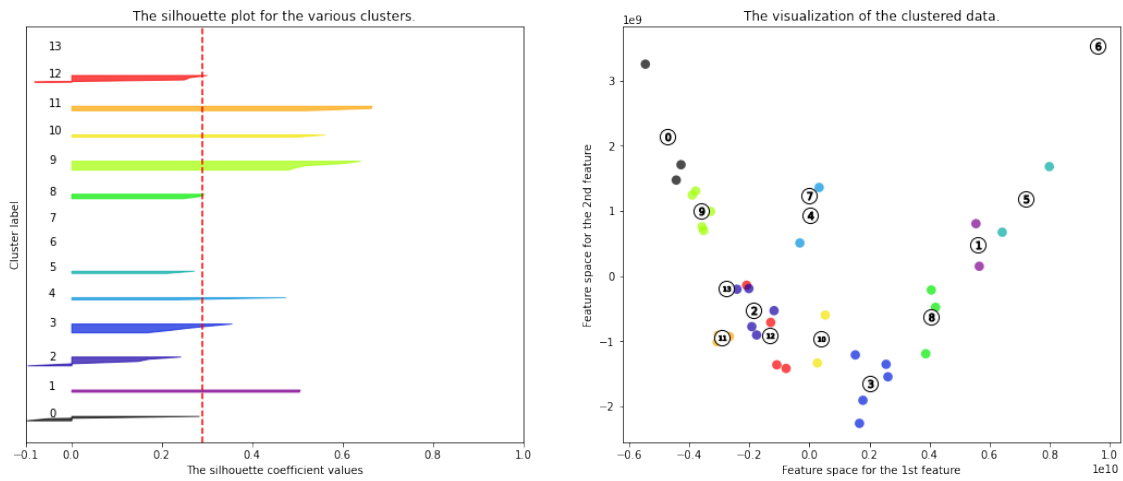
Silhouette analysis for KMeans clustering on sample data with n_clusters = 12



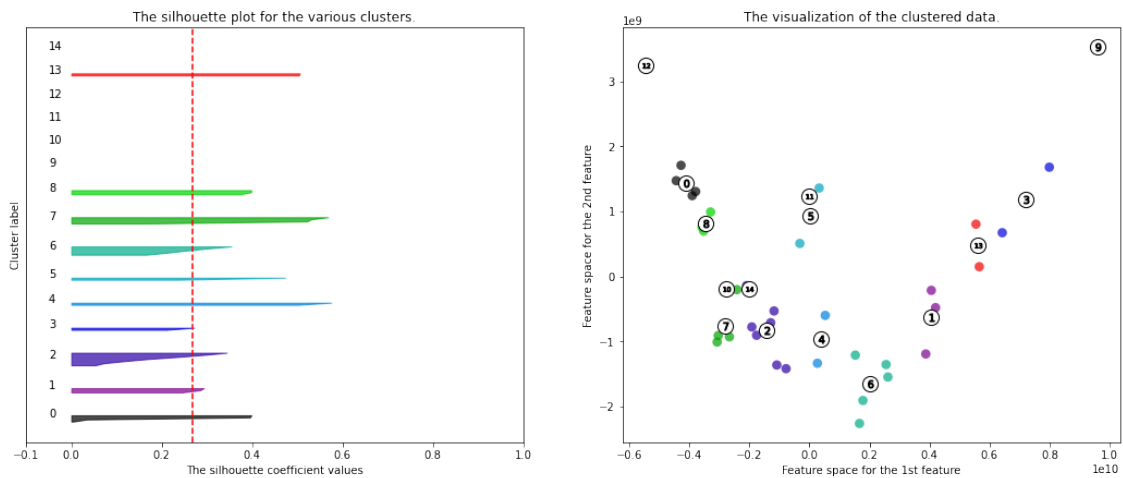
Silhouette analysis for KMeans clustering on sample data with n_clusters = 13



Silhouette analysis for KMeans clustering on sample data with n_clusters = 14



Silhouette analysis for KMeans clustering on sample data with n_clusters = 15



[73]: silhouette_scores_euc

[73]:

	k	score
0	2	0.513197
1	3	0.376735
2	4	0.371904
3	5	0.340121
4	6	0.386350
5	7	0.338519
6	8	0.343562
7	9	0.315147
8	10	0.315310
9	11	0.311548

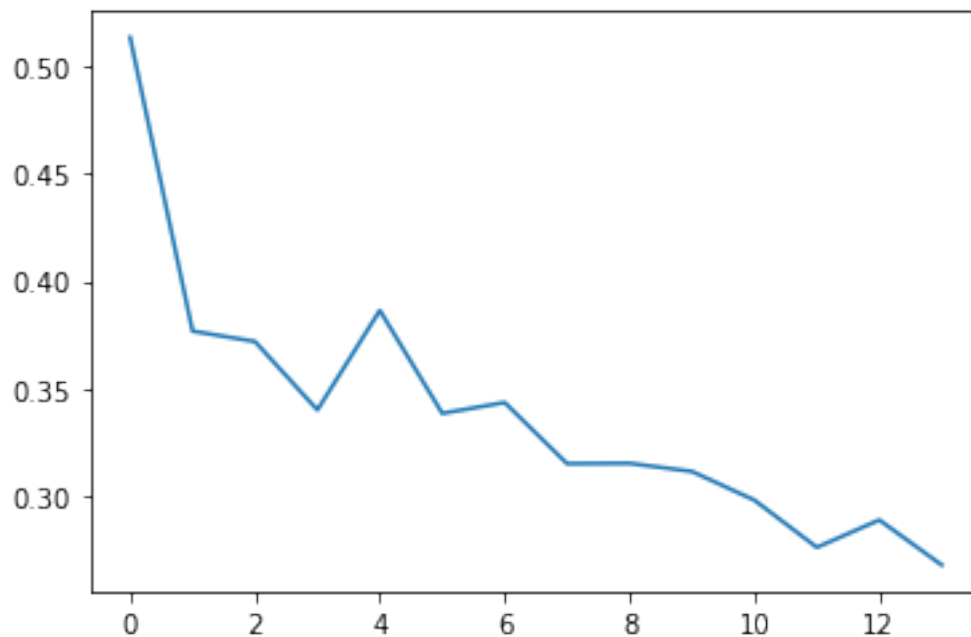
```

10 12 0.298222
11 13 0.276279
12 14 0.289012
13 15 0.268083

```

```
[74]: plt.plot(silhouette_scores_euc['score'])
```

```
[74]: [<matplotlib.lines.Line2D at 0x7fdc0f199940>]
```



0.3.2 k=4

```
[77]: kmeans_pca = KMeans(n_clusters=4, init='k-means++', random_state=42)
kmeans_pca.fit(scores_pca)
kmeans_labels = kmeans_pca.labels_
```

```
[78]: kmeans_labels_df = pd.DataFrame(kmeans_labels)
```

```
[79]: cluster_centers = kmeans_pca.cluster_centers_
cluster_centers_df = pd.DataFrame(cluster_centers).to_csv('outs/' + date + '_'
↳ '_combined_kmeans_clusterCenters.csv')
```

```
[81]: cluster_centers = kmeans_pca.cluster_centers_
df_segk_pca_kmeans = pd.concat([X.reset_index(drop=True), pd.
↳ DataFrame(scores_pca)], axis=1)
df_segk_pca_kmeans.columns.values[-34]
df_segk_pca_kmeans.columns.values[-39:-34] = ['Component 1', 'Component 2',
↳ 'Component 3', 'Component 4', 'Component 5']
df_segk_pca_kmeans['Segment K-means PCA'] = kmeans_pca.labels_
```

```
df_segm_pca_kmeans.index = X.index
df_segm_pca_kmeans[['TimePoint', 'BioRep', 'TechRep', 'Region', 'DPI_Region',
    ↳ 'BioRep_TechRep']] = combined_lfq[['TimePoint', 'BioRep', 'TechRep',
    ↳ 'Region', 'DPI_Region', 'BioRep_TechRep']]
df_segm_pca_kmeans.to_csv('outs/' + date + '_combined_df_segm_pca_kmeans.csv')
df_segm_pca_kmeans
```

[81]: AOA0J9YTY0;AOA0J9YUL3;Q8C1B7;AOA0J9YUV6;AOA0J9YVA6 \

index	
4DPI_interface_2_4	8884400.0
4DPI_interface_2_2	4805100.0
4DPI_interface_2_1	4679400.0
4DPI_interface_1_2	2752000.0
4DPI_interface_1_1	6446300.0
4DPI_cortex_3_3	3966100.0
4DPI_cortex_3_2	3174000.0
4DPI_cortex_2_3	3505600.0
4DPI_cortex_2_2	3623400.0
4DPI_cortex_2_1	5408800.0
4DPI_cortex_1_2	3500500.0
4DPI_cortex_1_1	4027400.0
4DPI_SAC_3_3	10933000.0
4DPI_SAC_3_2	9909200.0
4DPI_SAC_2_3	7513200.0
4DPI_SAC_2_2	5452600.0
4DPI_SAC_2_1	11385000.0
4DPI_SAC_1_3	8059300.0
4DPI_SAC_1_2	5600200.0
4DPI_SAC_1_1	6698000.0
10DPI_interface_5_3	14004000.0
10DPI_interface_4_3	11083000.0
10DPI_interface_4_2	10662000.0
10DPI_interface_4_1	10168000.0
10DPI_interface_3_2	10654000.0
10DPI_interface_3_1	13143000.0
10DPI_cortex_4_4	4613200.0
10DPI_cortex_4_3	7330400.0
10DPI_cortex_3_3	2525300.0
10DPI_cortex_3_2	4286400.0
10DPI_SAC_4_3	9362000.0
10DPI_SAC_4_2	10750000.0
10DPI_SAC_4_1	10526000.0
10DPI_SAC_3_3	22060000.0
10DPI_SAC_3_2	16201000.0
10DPI_SAC_3_1	11879000.0
10DPI_SAC_2_3	10519000.0
10DPI_SAC_2_2	11053000.0

10DPI_SAC_2_1 12673000.0

AOA0J9YUZ4;P63158;AOA0J9YUD8;D3YVC6;D3YZ18 \

index

4DPI_interface_2_4	13795000.0
4DPI_interface_2_2	16043000.0
4DPI_interface_2_1	14321000.0
4DPI_interface_1_2	2559000.0
4DPI_interface_1_1	18740000.0
4DPI_cortex_3_3	10538000.0
4DPI_cortex_3_2	5800300.0
4DPI_cortex_2_3	7019000.0
4DPI_cortex_2_2	9811200.0
4DPI_cortex_2_1	7080900.0
4DPI_cortex_1_2	13352000.0
4DPI_cortex_1_1	15997000.0
4DPI_SAC_3_3	24605000.0
4DPI_SAC_3_2	18164000.0
4DPI_SAC_2_3	11153000.0
4DPI_SAC_2_2	7476200.0
4DPI_SAC_2_1	24354000.0
4DPI_SAC_1_3	15659000.0
4DPI_SAC_1_2	18231000.0
4DPI_SAC_1_1	17122000.0
10DPI_interface_5_3	19714000.0
10DPI_interface_4_3	41455000.0
10DPI_interface_4_2	18782000.0
10DPI_interface_4_1	26015000.0
10DPI_interface_3_2	28269000.0
10DPI_interface_3_1	18962000.0
10DPI_cortex_4_4	6571500.0
10DPI_cortex_4_3	12236000.0
10DPI_cortex_3_3	2883700.0
10DPI_cortex_3_2	3060800.0
10DPI_SAC_4_3	19296000.0
10DPI_SAC_4_2	26852000.0
10DPI_SAC_4_1	17329000.0
10DPI_SAC_3_3	20678000.0
10DPI_SAC_3_2	18474000.0
10DPI_SAC_3_1	22727000.0
10DPI_SAC_2_3	21473000.0
10DPI_SAC_2_2	33776000.0
10DPI_SAC_2_1	13833000.0

AOA0R4J093;Q9DBP5 AOA0R4J0I1;P07759;E9Q499;Q80X76 \

index

4DPI_interface_2_4	5749000.0	73676000.0
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4DPI_interface_2_1	9228800.0	105190000.0
4DPI_interface_1_2	6292300.0	22129000.0
4DPI_interface_1_1	18973000.0	38062000.0
4DPI_cortex_3_3	14700000.0	32753000.0
4DPI_cortex_3_2	12382000.0	24879000.0
4DPI_cortex_2_3	15484000.0	63258000.0
4DPI_cortex_2_2	11188000.0	47764000.0
4DPI_cortex_2_1	7899100.0	44351000.0
4DPI_cortex_1_2	15600000.0	60586000.0
4DPI_cortex_1_1	14481000.0	84161000.0
4DPI_SAC_3_3	10172000.0	33907000.0
4DPI_SAC_3_2	11552000.0	28193000.0
4DPI_SAC_2_3	13766000.0	56852000.0
4DPI_SAC_2_2	9574300.0	37292000.0
4DPI_SAC_2_1	12936000.0	60285000.0
4DPI_SAC_1_3	6236600.0	79677000.0
4DPI_SAC_1_2	4541100.0	63271000.0
4DPI_SAC_1_1	5351400.0	53311000.0
10DPI_interface_5_3	12167000.0	16442000.0
10DPI_interface_4_3	11927000.0	20993000.0
10DPI_interface_4_2	12228000.0	37156000.0
10DPI_interface_4_1	9103100.0	37811000.0
10DPI_interface_3_2	12785000.0	17304000.0
10DPI_interface_3_1	13679000.0	23913000.0
10DPI_cortex_4_4	15905000.0	21923000.0
10DPI_cortex_4_3	3591100.0	28778000.0
10DPI_cortex_3_3	2846000.0	17751000.0
10DPI_cortex_3_2	8662600.0	16765000.0
10DPI_SAC_4_3	12920000.0	26820000.0
10DPI_SAC_4_2	12945000.0	18110000.0
10DPI_SAC_4_1	11294000.0	17675000.0
10DPI_SAC_3_3	11293000.0	17459000.0
10DPI_SAC_3_2	10115000.0	16184000.0
10DPI_SAC_3_1	9660300.0	20254000.0
10DPI_SAC_2_3	7141400.0	11286000.0
10DPI_SAC_2_2	11842000.0	14575000.0
10DPI_SAC_2_1	8280200.0	18280000.0

070404;A0A0R4JOR1;A0A0U1RPE8 A0A0R4JOZ1;P08003 \

index

4DPI_interface_2_4	4863000.0	20661000.0
4DPI_interface_2_2	5048700.0	20709000.0
4DPI_interface_2_1	4421600.0	17844000.0
4DPI_interface_1_2	1316400.0	22085000.0
4DPI_interface_1_1	2682100.0	24030000.0
4DPI_cortex_3_3	2543300.0	7914900.0

4DPI_cortex_3_2	2405800.0	10756000.0
4DPI_cortex_2_3	3551100.0	9577000.0
4DPI_cortex_2_2	3017800.0	14510000.0
4DPI_cortex_2_1	3218600.0	9714800.0
4DPI_cortex_1_2	4396200.0	5588800.0
4DPI_cortex_1_1	559210.0	12618000.0
4DPI_SAC_3_3	6073100.0	24566000.0
4DPI_SAC_3_2	5000700.0	20284000.0
4DPI_SAC_2_3	7744500.0	14750000.0
4DPI_SAC_2_2	3349900.0	14702000.0
4DPI_SAC_2_1	7488800.0	22398000.0
4DPI_SAC_1_3	4729600.0	22668000.0
4DPI_SAC_1_2	2086200.0	16305000.0
4DPI_SAC_1_1	3328200.0	20802000.0
10DPI_interface_5_3	7822400.0	28551000.0
10DPI_interface_4_3	5549700.0	25417000.0
10DPI_interface_4_2	3494600.0	20562000.0
10DPI_interface_4_1	3959500.0	14898000.0
10DPI_interface_3_2	10313000.0	21001000.0
10DPI_interface_3_1	5659300.0	17979000.0
10DPI_cortex_4_4	6568100.0	16769000.0
10DPI_cortex_4_3	3419900.0	11998000.0
10DPI_cortex_3_3	2486500.0	10442000.0
10DPI_cortex_3_2	3308800.0	5336900.0
10DPI_SAC_4_3	6796900.0	23906000.0
10DPI_SAC_4_2	7350200.0	19011000.0
10DPI_SAC_4_1	4744200.0	19508000.0
10DPI_SAC_3_3	12653000.0	20440000.0
10DPI_SAC_3_2	11506000.0	19529000.0
10DPI_SAC_3_1	9135600.0	22306000.0
10DPI_SAC_2_3	12044000.0	13540000.0
10DPI_SAC_2_2	11489000.0	15259000.0
10DPI_SAC_2_1	6878200.0	18782000.0

E9QL31;A0A0R4J104;P98078;E9PX84;Q3TRE6;F6TQN9;Q9DCE6;EOCXT5;EOCZ53;EOCYJ2 \

index

4DPI_interface_2_4	10187000.0
4DPI_interface_2_2	10975000.0
4DPI_interface_2_1	8142400.0
4DPI_interface_1_2	704720.0
4DPI_interface_1_1	4963900.0
4DPI_cortex_3_3	26064000.0
4DPI_cortex_3_2	19354000.0
4DPI_cortex_2_3	18882000.0
4DPI_cortex_2_2	14360000.0
4DPI_cortex_2_1	29494000.0
4DPI_cortex_1_2	8878300.0

4DPI_cortex_1_1	10476000.0
4DPI_SAC_3_3	7031500.0
4DPI_SAC_3_2	5700800.0
4DPI_SAC_2_3	5403700.0
4DPI_SAC_2_2	4984100.0
4DPI_SAC_2_1	7089200.0
4DPI_SAC_1_3	9872000.0
4DPI_SAC_1_2	7354500.0
4DPI_SAC_1_1	9433100.0
10DPI_interface_5_3	8816700.0
10DPI_interface_4_3	6936800.0
10DPI_interface_4_2	3421900.0
10DPI_interface_4_1	3713000.0
10DPI_interface_3_2	5232300.0
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10DPI_cortex_4_4	7816500.0
10DPI_cortex_4_3	14825000.0
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10DPI_SAC_4_3	5934200.0
10DPI_SAC_4_2	2999000.0
10DPI_SAC_4_1	4308500.0
10DPI_SAC_3_3	7713200.0
10DPI_SAC_3_2	9466900.0
10DPI_SAC_3_1	8250600.0
10DPI_SAC_2_3	6240900.0
10DPI_SAC_2_2	3403200.0
10DPI_SAC_2_1	8101900.0

AOA0R4J195;Q9CSU0;F6YB25;F7DAY5;Q8VDS4 \

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4DPI_interface_2_1	549190.0
4DPI_interface_1_2	1506600.0
4DPI_interface_1_1	1400400.0
4DPI_cortex_3_3	734590.0
4DPI_cortex_3_2	544140.0
4DPI_cortex_2_3	1263200.0
4DPI_cortex_2_2	911490.0
4DPI_cortex_2_1	1163700.0
4DPI_cortex_1_2	857580.0
4DPI_cortex_1_1	813620.0
4DPI_SAC_3_3	2025800.0
4DPI_SAC_3_2	1299600.0
4DPI_SAC_2_3	1140400.0
4DPI_SAC_2_2	1866900.0

4DPI_SAC_2_1	1723900.0
4DPI_SAC_1_3	1446200.0
4DPI_SAC_1_2	2133800.0
4DPI_SAC_1_1	1344000.0
10DPI_interface_5_3	1129300.0
10DPI_interface_4_3	1920700.0
10DPI_interface_4_2	1817400.0
10DPI_interface_4_1	1819300.0
10DPI_interface_3_2	1001900.0
10DPI_interface_3_1	2526900.0
10DPI_cortex_4_4	1237700.0
10DPI_cortex_4_3	2119000.0
10DPI_cortex_3_3	881230.0
10DPI_cortex_3_2	797640.0
10DPI_SAC_4_3	1421000.0
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10DPI_SAC_3_3	1626400.0
10DPI_SAC_3_2	1196300.0
10DPI_SAC_3_1	819090.0
10DPI_SAC_2_3	2583700.0
10DPI_SAC_2_2	1870900.0
10DPI_SAC_2_1	2521700.0

AOA1BOGQU8;P35980;AOA1BOGSS8;AOA1BOGSF7;AOA1BOGSA8;AOA1BOGS28;AOA1BOGRZ3 \

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4DPI_interface_2_1	20121000.0
4DPI_interface_1_2	36610000.0
4DPI_interface_1_1	23111000.0
4DPI_cortex_3_3	16177000.0
4DPI_cortex_3_2	17586000.0
4DPI_cortex_2_3	21878000.0
4DPI_cortex_2_2	16561000.0
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4DPI_cortex_1_1	20512000.0
4DPI_SAC_3_3	16860000.0
4DPI_SAC_3_2	22906000.0
4DPI_SAC_2_3	22223000.0
4DPI_SAC_2_2	26184000.0
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4DPI_SAC_1_3	18432000.0
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10DPI_interface_4_3	10966000.0
10DPI_interface_4_2	11930000.0
10DPI_interface_4_1	15307000.0
10DPI_interface_3_2	23524000.0
10DPI_interface_3_1	14105000.0
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10DPI_cortex_4_3	10131000.0
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10DPI_SAC_4_2	17678000.0
10DPI_SAC_4_1	14233000.0
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10DPI_SAC_3_1	11445000.0
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P06151;Q564E2;AOA1BOGSR9;AOA1BOGSX0;D3YZQ9;AOA1BOGSL7;AOA1B
 OGT41;AOA1BOGQX5;AOA1BOGS79;AOA1BOGRW9;AOA1BOGRC1;AOA1BOGRS2;D3YVR7;D3YZE4;AOA1B
 OGRE9;AOA1BOGSR2;P00342 \

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4DPI_cortex_2_2	63940000.0
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4DPI_SAC_1_1	59981000.0
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10DPI_interface_3_2	51215000.0
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10DPI_cortex_3_3	38540000.0
10DPI_cortex_3_2	19880000.0
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10DPI_SAC_4_2	92821000.0
10DPI_SAC_4_1	91489000.0
10DPI_SAC_3_3	47923000.0
10DPI_SAC_3_2	87510000.0
10DPI_SAC_3_1	39045000.0
10DPI_SAC_2_3	27205000.0
10DPI_SAC_2_2	50918000.0
10DPI_SAC_2_1	16776000.0

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4DPI_interface_2_4	...	1.031646e+07	-1.178469e+07	2.929428e-07
4DPI_interface_2_2	...	2.282268e+07	-5.412626e+04	2.929428e-07
4DPI_interface_2_1	...	-2.718176e+07	2.199637e+06	2.929428e-07
4DPI_interface_1_2	...	1.065881e+07	-7.706425e+05	2.929428e-07
4DPI_interface_1_1	...	-1.414153e+07	1.404734e+06	2.929428e-07
4DPI_cortex_3_3	...	-3.510812e+06	-1.258512e+07	2.929428e-07
4DPI_cortex_3_2	...	-1.934336e+07	6.848204e+06	2.929428e-07
4DPI_cortex_2_3	...	-2.514064e+07	9.140492e+06	2.929428e-07
4DPI_cortex_2_2	...	4.206068e+07	-1.488583e+07	2.929428e-07
4DPI_cortex_2_1	...	9.972975e+06	1.967906e+06	2.929428e-07
4DPI_cortex_1_2	...	-1.634307e+07	-2.895322e+06	2.929428e-07
4DPI_cortex_1_1	...	6.039433e+06	1.772419e+07	2.929428e-07
4DPI_SAC_3_3	...	1.320228e+07	4.387337e+04	2.929428e-07
4DPI_SAC_3_2	...	-3.905192e+06	2.317435e+07	2.929428e-07
4DPI_SAC_2_3	...	1.960526e+06	-8.147903e+05	2.929428e-07
4DPI_SAC_2_2	...	-5.308582e+06	3.966172e+06	2.929428e-07
4DPI_SAC_2_1	...	-2.123865e+06	-4.008947e+06	2.929428e-07
4DPI_SAC_1_3	...	9.412037e+06	-1.014204e+07	2.929428e-07
4DPI_SAC_1_2	...	-8.900910e+06	6.486090e+06	2.929428e-07
4DPI_SAC_1_1	...	3.946169e+06	-1.159411e+07	2.929428e-07
10DPI_interface_5_3	...	1.110906e+07	9.848782e+06	2.929428e-07
10DPI_interface_4_3	...	-1.415164e+07	-2.113884e+06	2.929428e-07
10DPI_interface_4_2	...	-1.030126e+07	-2.357560e+07	2.929428e-07
10DPI_interface_4_1	...	2.432251e+07	2.044187e+07	2.929428e-07
10DPI_interface_3_2	...	1.381270e+07	-3.013517e+07	2.929428e-07
10DPI_interface_3_1	...	-2.592833e+07	5.400819e+06	2.929428e-07
10DPI_cortex_4_4	...	-4.740456e+06	-7.302414e+06	2.929428e-07
10DPI_cortex_4_3	...	1.500560e+07	2.581753e+06	2.929428e-07
10DPI_cortex_3_3	...	-8.358536e+06	3.333076e+06	2.929428e-07

10DPI_cortex_3_2	...	4.094822e+06	-3.480078e+06	2.929428e-07
10DPI_SAC_4_3	...	6.421815e+06	1.283613e+07	2.929428e-07
10DPI_SAC_4_2	...	4.099798e+06	-4.146932e+06	2.929428e-07
10DPI_SAC_4_1	...	-1.157357e+07	-3.995583e+06	2.929428e-07
10DPI_SAC_3_3	...	-2.697074e+07	-3.500647e+06	2.929428e-07
10DPI_SAC_3_2	...	8.463252e+04	-1.427429e+07	2.929428e-07
10DPI_SAC_3_1	...	1.863299e+07	4.169473e+07	2.929428e-07
10DPI_SAC_2_3	...	-5.895312e+06	-4.218407e+06	2.929428e-07
10DPI_SAC_2_2	...	-3.574784e+06	3.894834e+06	2.929428e-07
10DPI_SAC_2_1	...	9.418379e+06	-6.709035e+06	2.929428e-07

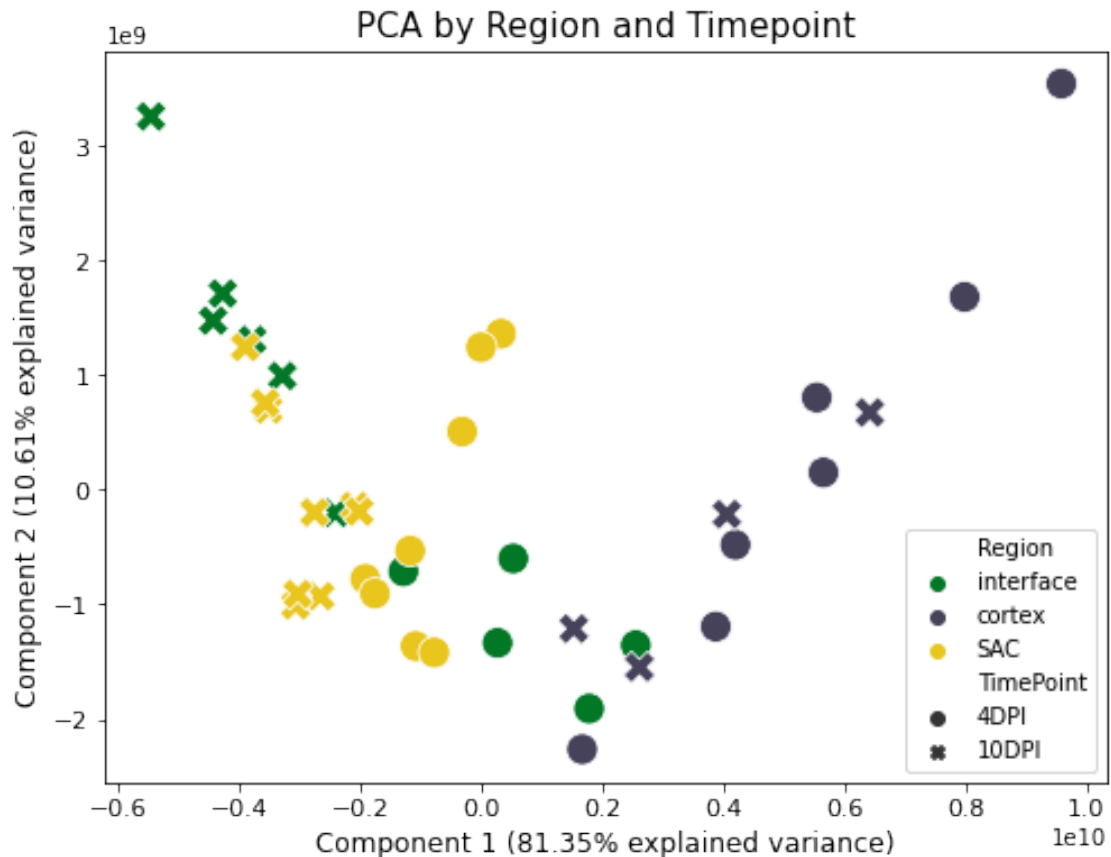
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4DPI_interface_2_2		1		4DPI	2	2	
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4DPI_interface_1_2		3		4DPI	1	2	
4DPI_interface_1_1		3		4DPI	1	1	
4DPI_cortex_3_3		3		4DPI	3	3	
4DPI_cortex_3_2		3		4DPI	3	2	
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4DPI_cortex_2_1		3		4DPI	2	1	
4DPI_cortex_1_2		2		4DPI	1	2	
4DPI_cortex_1_1		2		4DPI	1	1	
4DPI_SAC_3_3		1		4DPI	3	3	
4DPI_SAC_3_2		1		4DPI	3	2	
4DPI_SAC_2_3		1		4DPI	2	3	
4DPI_SAC_2_2		1		4DPI	2	2	
4DPI_SAC_2_1		1		4DPI	2	1	
4DPI_SAC_1_3		1		4DPI	1	3	
4DPI_SAC_1_2		1		4DPI	1	2	
4DPI_SAC_1_1		1		4DPI	1	1	
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10DPI_interface_4_3		0		10DPI	4	3	
10DPI_interface_4_2		0		10DPI	4	2	
10DPI_interface_4_1		0		10DPI	4	1	
10DPI_interface_3_2		1		10DPI	3	2	
10DPI_interface_3_1		0		10DPI	3	1	
10DPI_cortex_4_4		3		10DPI	4	4	
10DPI_cortex_4_3		3		10DPI	4	3	
10DPI_cortex_3_3		3		10DPI	3	3	
10DPI_cortex_3_2		2		10DPI	3	2	
10DPI_SAC_4_3		1		10DPI	4	3	
10DPI_SAC_4_2		1		10DPI	4	2	
10DPI_SAC_4_1		1		10DPI	4	1	
10DPI_SAC_3_3		0		10DPI	3	3	

10DPI_SAC_3_2	0	10DPI	3	2
10DPI_SAC_3_1	0	10DPI	3	1
10DPI_SAC_2_3	1	10DPI	2	3
10DPI_SAC_2_2	1	10DPI	2	2
10DPI_SAC_2_1	1	10DPI	2	1

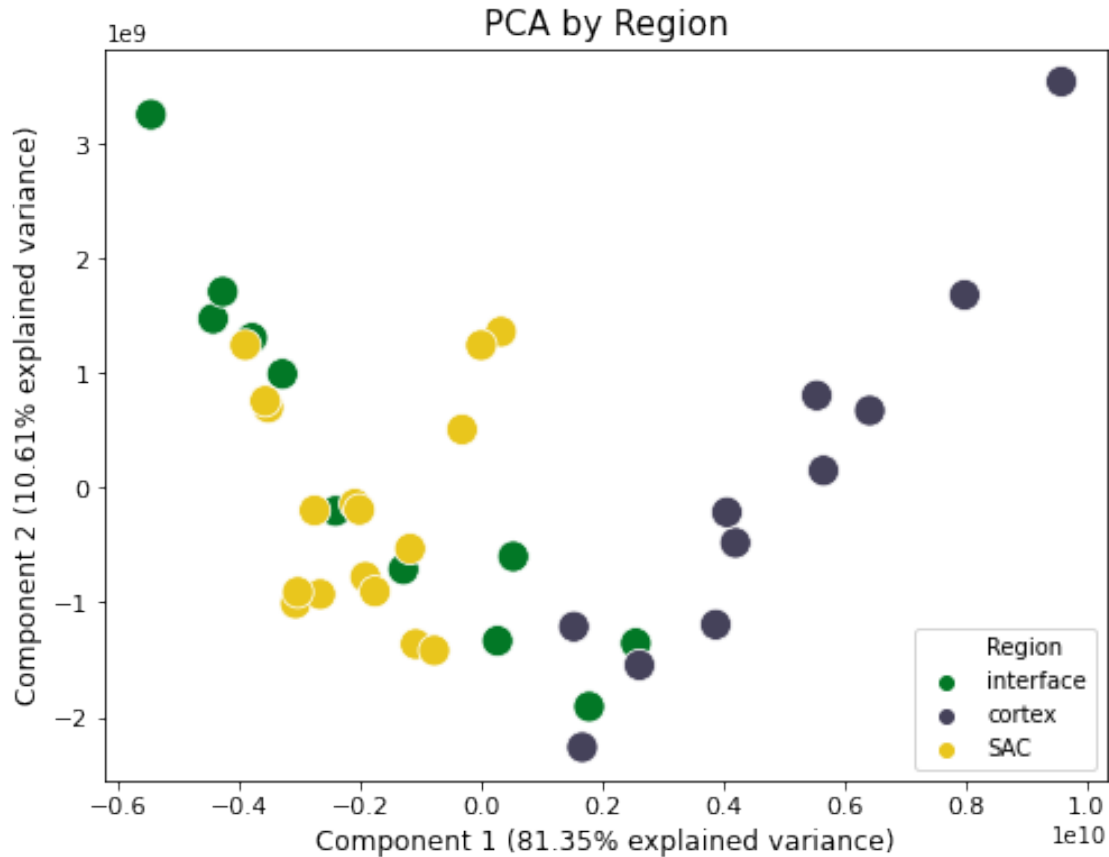
	Region	DPI_Region	BioRep_TechRep
index			
4DPI_interface_2_4	interface	4DPI_interface	2_4
4DPI_interface_2_2	interface	4DPI_interface	2_2
4DPI_interface_2_1	interface	4DPI_interface	2_1
4DPI_interface_1_2	interface	4DPI_interface	1_2
4DPI_interface_1_1	interface	4DPI_interface	1_1
4DPI_cortex_3_3	cortex	4DPI_cortex	3_3
4DPI_cortex_3_2	cortex	4DPI_cortex	3_2
4DPI_cortex_2_3	cortex	4DPI_cortex	2_3
4DPI_cortex_2_2	cortex	4DPI_cortex	2_2
4DPI_cortex_2_1	cortex	4DPI_cortex	2_1
4DPI_cortex_1_2	cortex	4DPI_cortex	1_2
4DPI_cortex_1_1	cortex	4DPI_cortex	1_1
4DPI_SAC_3_3	SAC	4DPI_SAC	3_3
4DPI_SAC_3_2	SAC	4DPI_SAC	3_2
4DPI_SAC_2_3	SAC	4DPI_SAC	2_3
4DPI_SAC_2_2	SAC	4DPI_SAC	2_2
4DPI_SAC_2_1	SAC	4DPI_SAC	2_1
4DPI_SAC_1_3	SAC	4DPI_SAC	1_3
4DPI_SAC_1_2	SAC	4DPI_SAC	1_2
4DPI_SAC_1_1	SAC	4DPI_SAC	1_1
10DPI_interface_5_3	interface	10DPI_interface	5_3
10DPI_interface_4_3	interface	10DPI_interface	4_3
10DPI_interface_4_2	interface	10DPI_interface	4_2
10DPI_interface_4_1	interface	10DPI_interface	4_1
10DPI_interface_3_2	interface	10DPI_interface	3_2
10DPI_interface_3_1	interface	10DPI_interface	3_1
10DPI_cortex_4_4	cortex	10DPI_cortex	4_4
10DPI_cortex_4_3	cortex	10DPI_cortex	4_3
10DPI_cortex_3_3	cortex	10DPI_cortex	3_3
10DPI_cortex_3_2	cortex	10DPI_cortex	3_2
10DPI_SAC_4_3	SAC	10DPI_SAC	4_3
10DPI_SAC_4_2	SAC	10DPI_SAC	4_2
10DPI_SAC_4_1	SAC	10DPI_SAC	4_1
10DPI_SAC_3_3	SAC	10DPI_SAC	3_3
10DPI_SAC_3_2	SAC	10DPI_SAC	3_2
10DPI_SAC_3_1	SAC	10DPI_SAC	3_1
10DPI_SAC_2_3	SAC	10DPI_SAC	2_3
10DPI_SAC_2_2	SAC	10DPI_SAC	2_2
10DPI_SAC_2_1	SAC	10DPI_SAC	2_1

[39 rows x 333 columns]

```
[82]: x_axis = df_segm_pca_kmeans['Component 1']
y_axis = df_segm_pca_kmeans['Component 2']
plt.figure(figsize=(8,6))
sns.scatterplot(x_axis, y_axis,
                hue=df_segm_pca_kmeans['Region'], palette=["#027826", "#45425A", "#E9C61D"],
                style=df_segm_pca_kmeans['TimePoint'], s=200, legend='full')
plt.title('PCA by Region and Timepoint', fontsize=15)
plt.tick_params(axis='both', which='major', labels=11)
plt.tick_params(axis='both', which='minor', labels=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
            variance)', fontsize=12)
plt.ylabel('Component 2 (' + str((pca_expl_var[1]*100).round(2)) + '% explained_
            variance)', fontsize=12)
plt.legend(loc='lower right')
plt.savefig('outs/' + date + '_combined_PCAwithRegionsandTimepoints.svg')
```



```
[83]: x_axis = df_segm_pca_kmeans['Component 1']
y_axis = df_segm_pca_kmeans['Component 2']
plt.figure(figsize=(8,6))
sns.scatterplot(x_axis, y_axis,
    →hue=df_segm_pca_kmeans['Region'],palette=["#027826","#45425A","#E9C61D"],s=200,
    →legend='full')
plt.title('PCA by Region', fontsize=15)
plt.tick_params(axis='both', which='major', labels=11)
plt.tick_params(axis='both', which='minor', labels=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
    →variance)', fontsize=12)
plt.ylabel('Component 2 (' + str((pca_expl_var[1]*100).round(2)) + '% explained_
    →variance)', fontsize=12)
plt.legend(loc='lower right')
plt.savefig('outs/' + date + '_combined_PCAbbyRegion.svg')
```

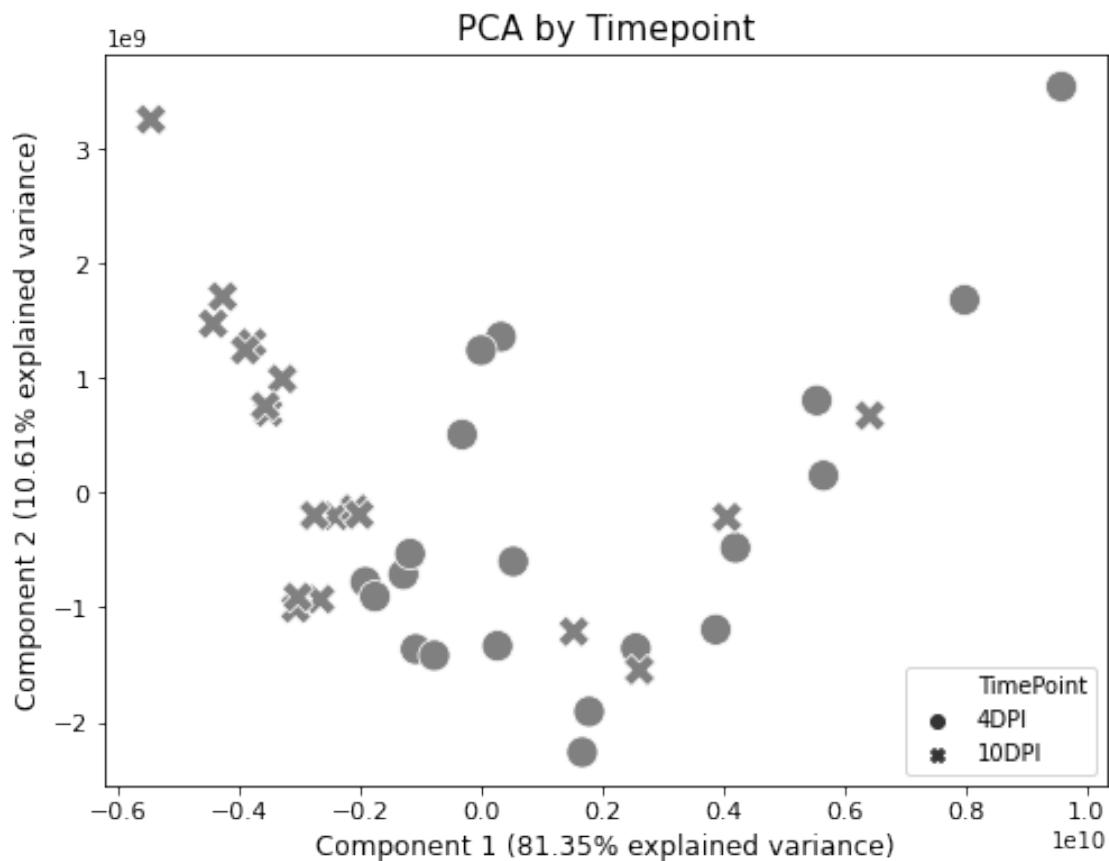


```
[84]: x_axis = df_segm_pca_kmeans['Component 1']
y_axis = df_segm_pca_kmeans['Component 2']
plt.figure(figsize=(8,6))
sns.scatterplot(x_axis,
```

```

        y_axis,
        color='grey',
        style=df_segm_pca_kmeans['TimePoint'],
        s=200,
        legend='full')
plt.title('PCA by Timepoint', fontsize=15)
plt.tick_params(axis='both', which='major', labels=11)
plt.tick_params(axis='both', which='minor', labels=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.ylabel('Component 2 (' + str((pca_expl_var[1]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.legend(loc='lower right')
plt.savefig('outs/' + date + '_combined_PCAByTimepoint.svg')

```



```

[174]: x_axis = df_segm_pca_kmeans['Component 1']
        y_axis = df_segm_pca_kmeans['Component 2']
        labels = df_segm_pca_kmeans['BioRep_TechRep']
        plt.figure(figsize=(8,6))
        sns.scatterplot(x_axis,

```

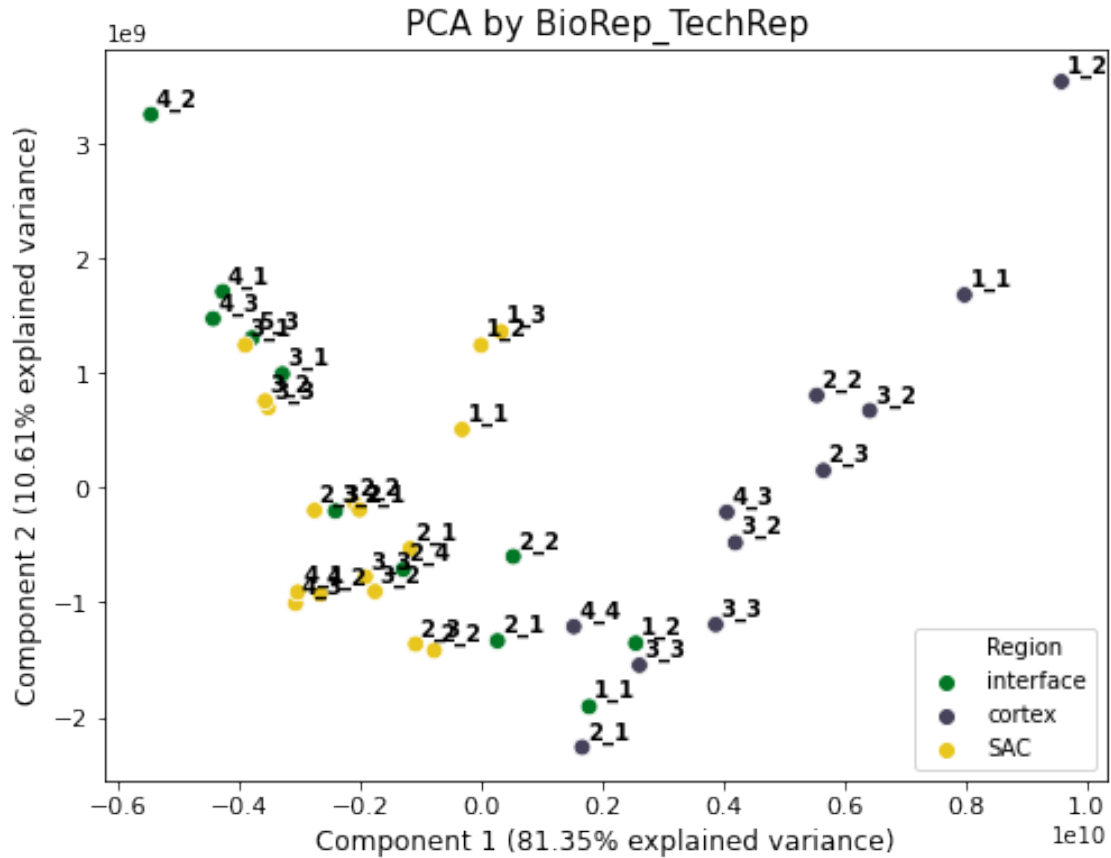
```

        y_axis,
        hue=df_segm_pca_kmeans['Region'],
        palette=["#027826", "#45425A", "#E9C61D"],
        s=60,
        legend='full')
plt.title('PCA by BioRep_TechRep', fontsize=15)
plt.tick_params(axis='both', which='major', labels=11)
plt.tick_params(axis='both', which='minor', labels=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.ylabel('Component 2 (' + str((pca_expl_var[1]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.legend(loc='lower right')

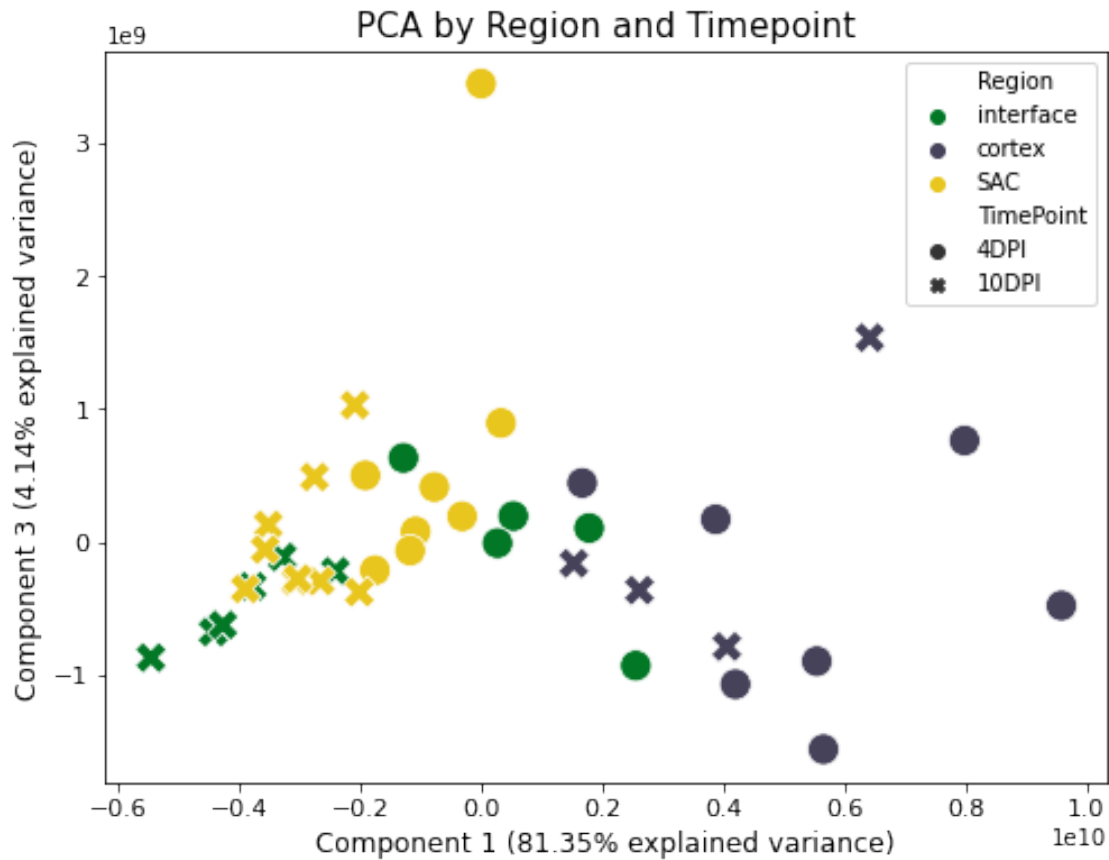
for line in range(0,df_segm_pca_kmeans.shape[0]):
    plt.annotate(xy = (x_axis[line], y_axis[line]),
        text = labels[line],
        horizontalalignment='left',
        verticalalignment='bottom',
        size='medium',
        color='black',
        weight='semibold',
        xytext=(2, 2),
        textcoords='offset points')

plt.savefig('outs/' + date + '_combined_PCAbyBioRep_TechRep.svg')

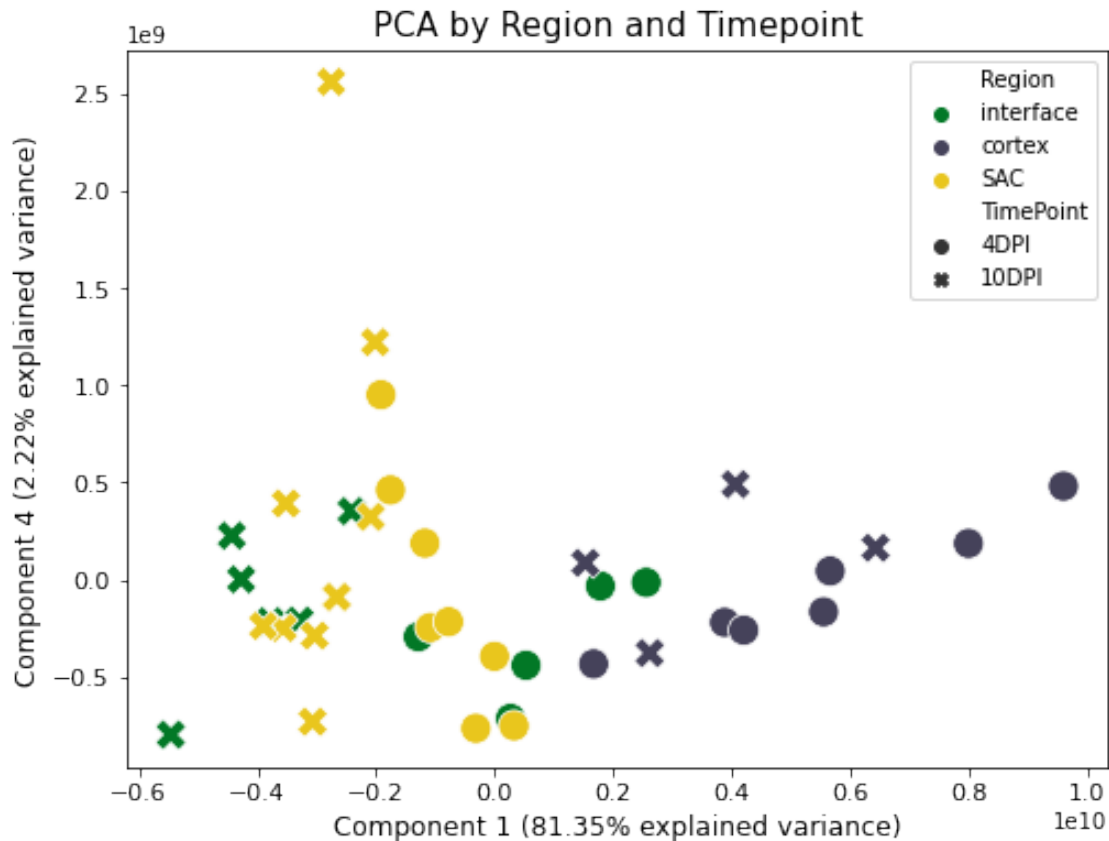
```



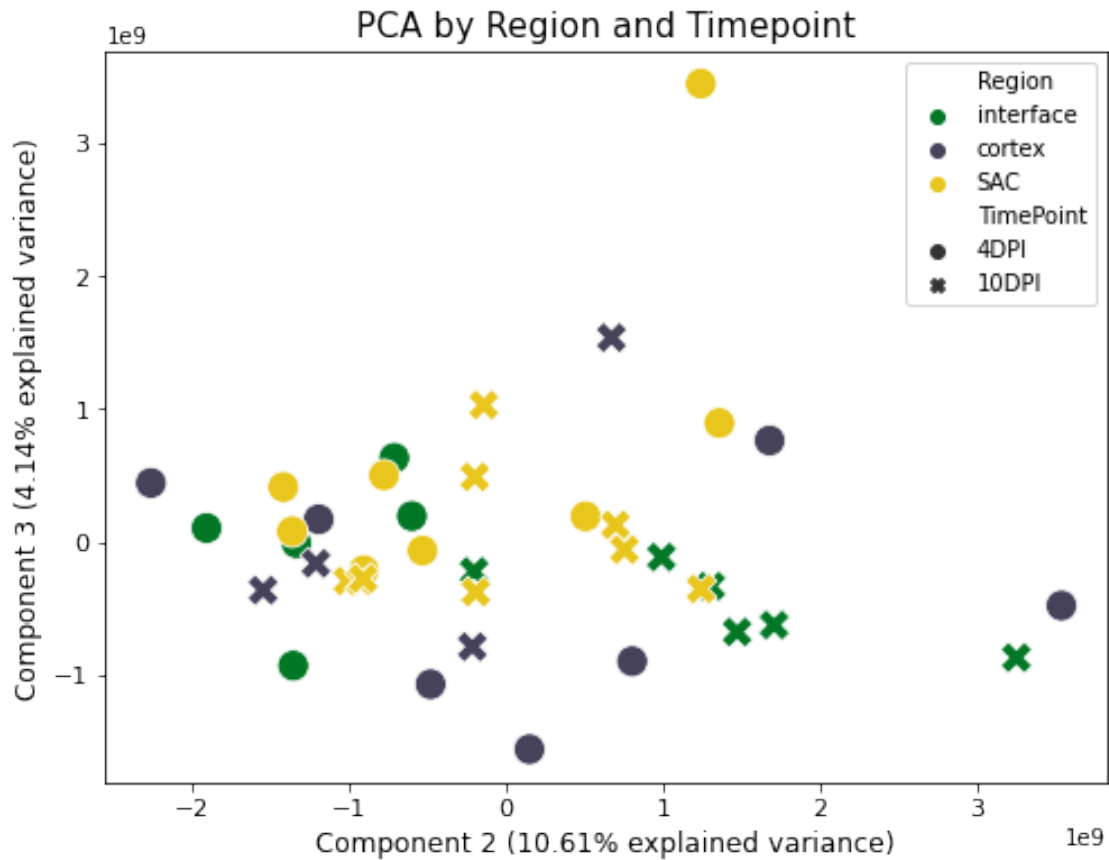
```
[175]: x_axis = df_segm_pca_kmeans['Component 1']
y_axis = df_segm_pca_kmeans['Component 3']
plt.figure(figsize=(8,6))
sns.scatterplot(x_axis,
                y_axis,
                hue=df_segm_pca_kmeans['Region'],
                palette=["#027826", "#45425A", "#E9C61D"],
                style=df_segm_pca_kmeans['TimePoint'],
                s=200,
                legend='full')
plt.title('PCA by Region and Timepoint', fontsize=15)
plt.tick_params(axis='both', which='major', labels=11)
plt.tick_params(axis='both', which='minor', labels=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.ylabel('Component 3 (' + str((pca_expl_var[2]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.legend(loc='upper right')
plt.savefig('outs/' + date + '_combined_PCAwithRegionsandTimepoints_PC13.svg')
```



```
[176]: x_axis = df_segmpca_kmeans['Component 1']
y_axis = df_segmpca_kmeans['Component 4']
plt.figure(figsize=(8,6))
sns.scatterplot(x_axis,
                y_axis,
                hue=df_segmpca_kmeans['Region'],
                palette=["#027826", "#45425A", "#E9C61D"],
                style=df_segmpca_kmeans['TimePoint'],
                s=200,
                legend='full')
plt.title('PCA by Region and Timepoint', fontsize=15)
plt.tick_params(axis='both', which='major', labelsize=11)
plt.tick_params(axis='both', which='minor', labelsize=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.ylabel('Component 4 (' + str((pca_expl_var[3]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.legend(loc='upper right')
plt.savefig('outs/' + date + '_combined_PCAwithRegionsandTimepoints_PC14.svg')
```



```
[177]: x_axis = df_segm_pca_kmeans['Component 2']
y_axis = df_segm_pca_kmeans['Component 3']
plt.figure(figsize=(8,6))
sns.scatterplot(x_axis, y_axis,
                hue=df_segm_pca_kmeans['Region'],
                palette=["#027826", "#45425A", "#E9C61D"],
                style=df_segm_pca_kmeans['TimePoint'],
                s=200,
                legend='full')
plt.title('PCA by Region and Timepoint', fontsize=15)
plt.tick_params(axis='both', which='major', labelsize=11)
plt.tick_params(axis='both', which='minor', labelsize=11)
plt.xlabel('Component 2 (' + str((pca_expl_var[1]*100).round(2)) + '% explained_↵
→variance)', fontsize=12)
plt.ylabel('Component 3 (' + str((pca_expl_var[2]*100).round(2)) + '% explained_↵
→variance)', fontsize=12)
plt.legend(loc='upper right')
plt.savefig('outs/' + date + '_combined_PCAwithRegionsandTimepoints_PC23.svg')
```

```
[178]: palette_kmeans = ["#dd7373", "#51a3a3", "#37d4fb", "#7b3e19"]
```

```
[179]: x_axis = df_segm_pca_kmeans['Component 1']
y_axis = df_segm_pca_kmeans['Component 2']
plt.figure(figsize=(8,6))

sns.scatterplot(x_axis, y_axis, hue=df_segm_pca_kmeans['Segment K-means_
→PCA'], palette=palette_kmeans, s=200, legend=None)

plt.title('k-Means Clusters in PCA Space', fontsize=15)
plt.tick_params(axis='both', which='major', labelsize=11)
plt.tick_params(axis='both', which='minor', labelsize=11)
plt.xlabel('Component 1 (' + str((pca_expl_var[0]*100).round(2)) + '% explained_
→variance)', fontsize=12)
plt.ylabel('Component 2 (' + str((pca_expl_var[1]*100).round(2)) + '% explained_
→variance)', fontsize=12)

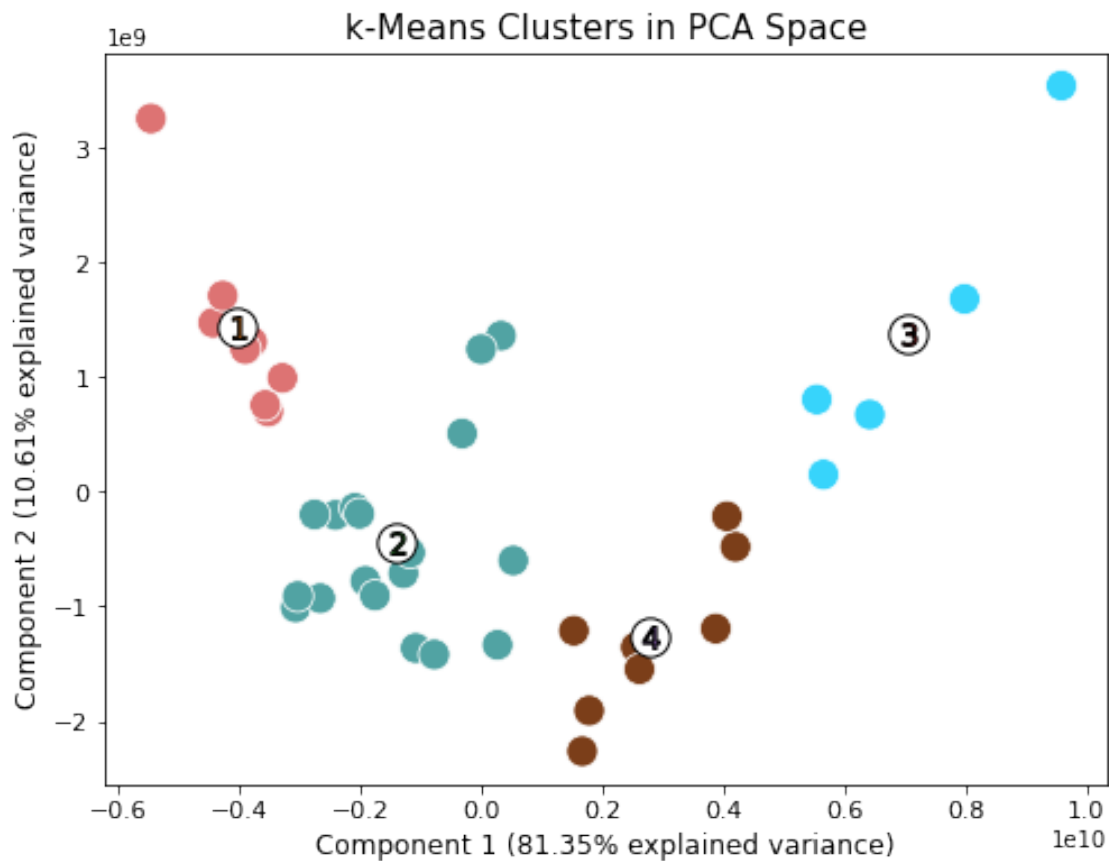
plt.scatter(cluster_centers[:, 0], cluster_centers[:, 1], marker='o',
            c="white", alpha=1, s=300, edgecolor='k')
```

```

for i, c in enumerate(cluster_centers):
    i = i+1
    plt.scatter(c[0], c[1], marker='o', alpha=1,
                s=100, edgecolor='k')

plt.savefig('outs/' + date + '_combined_kClustersinPCAspace.svg')

```



```

[180]: print(cluster_centers)
        print(type(cluster_centers))

```

```

[[-4.03019735e+09  1.42861860e+09 -3.62243311e+08 -1.35817336e+08
  1.06984274e+08  3.29454473e+07 -1.88805686e+07 -6.93786757e+07
 -3.30601347e+07  3.75564111e+06  2.05380309e+07  8.19662582e+06
  1.36503173e+07  2.08327123e+07  1.83816557e+07 -1.63459977e+07
  1.45807440e+07  1.69444361e+07  7.66743612e+06 -3.92161608e+06
 -8.04736787e+05  4.48151238e+06  3.27507424e+06  6.61627058e+06
  2.44923765e+06 -1.99081818e+06 -2.88351258e+06 -2.04113397e+06
 -1.47470614e+06 -4.03186553e+06 -3.00987622e+06  1.32354349e+06
 -1.13636306e+06  1.02620642e+06  5.05175036e+06  6.24579750e+05
 -2.90034849e+06  4.24022242e+06  2.92942842e-07]

```

```

[-1.40253631e+09 -4.50482097e+08  3.40041833e+08  6.28997019e+07
 -9.79711332e+07 -2.71040892e+07  2.27752926e+07  8.64032150e+07
  2.36974976e+06  9.84461085e+06 -7.81950527e+06 -4.57226219e+06
 -6.15663700e+06 -7.18565738e+06 -1.29394263e+07  1.33325889e+07
 -1.08417688e+07 -1.06147884e+07 -4.85661124e+06  3.16629298e+06
  1.29649164e+06 -1.78728383e+06  4.86709905e+05 -3.64280238e+06
 -1.35243966e+06  2.53569738e+05  4.50953066e+06  6.36056732e+05
  1.11333230e+06  2.42739822e+06  2.59040179e+06 -9.23552395e+05
  1.58763058e+05  1.05391964e+05 -1.96455976e+06  9.01510395e+04
  1.49715915e+06 -1.94459615e+06  2.92942842e-07]
[ 7.04060932e+09  1.36934796e+09 -1.26842256e+08  1.42314384e+08
  4.52560521e+07  1.64069463e+07  1.71095864e+07  1.88903315e+07
 -2.58743434e+07  6.08497911e+07  3.80342308e+07 -3.96927884e+07
 -8.62387196e+06 -7.87102830e+06  6.65039646e+06  1.33382000e+07
  1.99196938e+07 -1.28346415e+07 -2.25619600e+06 -2.22523794e+06
  2.80125176e+06 -2.31503553e+06  1.72611708e+07 -1.51500516e+07
 -7.97151832e+06 -1.46370109e+07 -2.86138679e+06  5.88732688e+06
  1.87291287e+06 -4.07563098e+06  3.69996399e+06  3.01382455e+06
 -3.31673591e+06 -5.40107936e+06  1.03092320e+05 -1.89425282e+06
  2.14224554e+06  1.12069078e+06  2.92942842e-07]
[ 2.78552322e+09 -1.27087635e+09 -3.23574402e+08 -9.46534829e+07
  8.51657433e+07  1.77844118e+07 -4.30573311e+07 -1.36835015e+08
  4.38996623e+07 -6.39371350e+07 -2.67155383e+07  2.68989569e+07
  5.59203597e+06  2.54409469e+05  6.57555582e+06 -2.19887024e+07
 -2.63657271e+06  1.49604887e+07  4.67006167e+06 -1.81176940e+06
 -3.86315175e+06  9.86773451e+05 -1.51584033e+07  1.10488170e+07
  5.57595053e+06  1.05684181e+07 -5.47456466e+06 -3.06957298e+06
 -2.20086208e+06  1.11748890e+06 -5.13100531e+06 -1.12919094e+06
  2.85210612e+06  2.11233626e+06 -6.95923611e+05  3.56488424e+05
 -1.80716306e+06 -5.65312807e+05  2.92942842e-07]]
<class 'numpy.ndarray'>

```

```
[181]: cluster_centers.shape
```

```
[181]: (4, 39)
```

```
[182]: components.shape
```

```
[182]: (39, 287)
```

```
[183]: centers = cluster_centers.dot(components)
centers.shape
```

```
[183]: (4, 287)
```

```
[184]: centers = pd.DataFrame(centers,
                             index=['Cluster 1', 'Cluster 2', 'Cluster 3', 'Cluster_
→4'],
                             columns = X.columns)
centers.to_csv('outs/' + date + '_combined_centers_annotated.csv')
```

```
[185]: centers_t = centers.T
       centers_t.shape
```

```
[185]: (287, 4)
```

```
[186]: centers
```

```
[186]: Protein IDs  AOA0J9YTY0;AOA0J9YUL3;Q8C1B7;AOA0J9YUV6;AOA0J9YVA6  \
Cluster 1          5.347031e+06
Cluster 2          5.557197e+05
Cluster 3         -4.514309e+06
Cluster 4         -3.775957e+06

Protein IDs  AOA0J9YUZ4;P63158;AOA0J9YUD8;D3YVC6;D3YZ18  AOA0R4J093;Q9DBP5  \
Cluster 1      6.832731e+06      6.331833e+05
Cluster 2      2.468034e+06      -8.206000e+05
Cluster 3     -6.670144e+06      2.444753e+06
Cluster 4     -8.216969e+06     -3.148042e+05

Protein IDs  AOA0R4J0I1;P07759;E9Q499;Q80X76  070404;AOA0R4J0R1;AOA0U1RPE8  \
Cluster 1      -1.365104e+07      2.010879e+06
Cluster 2       5.144795e+06      8.577997e+05
Cluster 3       1.707926e+07     -2.495012e+06
Cluster 4     -8.599288e+06     -2.381546e+06

Protein IDs  AOA0R4J0Z1;P08003  \
Cluster 1       3.832855e+06
Cluster 2       1.883494e+06
Cluster 3      -7.851255e+06
Cluster 4      -3.163682e+06

Protein IDs
E9QL31;AOA0R4J104;P98078;E9PX84;Q3TRE6;F6TQN9;Q9DCE6;E0CXT5;E0CZ53;E0CYJ2  \
Cluster 1      -2.891130e+06
Cluster 2     -2.992265e+06
Cluster 3       5.900767e+06
Cluster 4       5.935747e+06

Protein IDs  AOA0R4J195;Q9CSU0;F6YB25;F7DAY5;Q8VDS4  \
Cluster 1      114204.262820
Cluster 2      236712.735043
Cluster 3     -564013.487179
Cluster 4     -294299.487179

Protein IDs
AOA1BOGQU8;P35980;AOA1BOGSS8;AOA1BOGSF7;AOA1BOGSA8;AOA1BOGS28;AOA1BOGRZ3  \
Cluster 1      -5.603615e+06
Cluster 2       5.265530e+05
```

Cluster 3	1.781197e+06
Cluster 4	3.305622e+06

Protein IDs P06151;Q564E2;A0A1BOGSR9;A0A1BOGSX0;D3YZQ9;A0A1BOGSL7;A0A1BOGT41;A0A1BOGQX5;A0A1BOGS79;A0A1BOGRW9;A0A1BOGRC1;A0A1BOGRS2;D3YVR7;D3YZE4;A0A1BOGRE9;A0A1BOGSR2;P00342 \

Cluster 1	-1.792952e+06
Cluster 2	8.612312e+06
Cluster 3	-3.734277e+06
Cluster 4	-1.525083e+07

Protein IDs	...	Q9JKF1;A0A0U1RNG5;A0A0U1RPI2;A0A0U1RPU3	Q9JLJ2;Q3U367	\
Cluster 1	...	1.680077e+07	-6.899002e+06	
Cluster 2	...	5.252364e+06	-4.775488e+06	
Cluster 3	...	-1.551040e+07	1.191792e+07	
Cluster 4	...	-1.892459e+07	1.019515e+07	

Protein IDs	Q9QZQ8	Q9R1P4;A0A1BOGS70	Q9WU78	\
Cluster 1	8.566707e+06	3.967160e+06	4.199472e+06	
Cluster 2	4.460265e+05	2.049018e+06	9.130855e+04	
Cluster 3	-6.123298e+06	-5.780895e+06	-1.380523e+06	
Cluster 4	-5.743205e+06	-4.964390e+06	-3.542090e+06	

Protein IDs	Q9Z0P4;Q3UZP7;A0A1W2P879;A0A1W2P831	Q9Z1D1	Q9Z1Q5	\
Cluster 1	-882132.500000	4.513414e+05	2.564748e+06	
Cluster 2	-98701.666667	2.961831e+05	1.874555e+06	
Cluster 3	909390.000000	-1.005512e+06	-6.734190e+06	
Cluster 4	535842.500000	-4.893086e+05	-2.573627e+06	

Protein IDs	Q9Z2D6	Q9Z2I8;A0A0N4SWD1
Cluster 1	-1.676840e+06	-1.577885e+07
Cluster 2	-4.070078e+05	-8.384045e+06
Cluster 3	2.729420e+06	2.848534e+07
Cluster 4	8.867200e+05	1.683961e+07

[4 rows x 287 columns]

[187]: centers_t

[187]:	Cluster 1	\
Protein IDs		
A0A0J9YTY0;A0A0J9YUL3;Q8C1B7;A0A0J9YUV6;A0A0J9YVA6	5.347031e+06	
A0A0J9YUZ4;P63158;A0A0J9YUD8;D3YVC6;D3YZ18	6.832731e+06	
A0A0R4J093;Q9DBP5	6.331833e+05	
A0A0R4J0I1;P07759;E9Q499;Q80X76	-1.365104e+07	
O7O4O4;A0A0R4J0R1;A0A0U1RPE8	2.010879e+06	
...	...	
Q9Z0P4;Q3UZP7;A0A1W2P879;A0A1W2P831	-8.821325e+05	

Q9Z1D1	4.513414e+05
Q9Z1Q5	2.564748e+06
Q9Z2D6	-1.676840e+06
Q9Z2I8;AOA0N4SWD1	-1.577885e+07

Cluster 2 \

Protein IDs

AOA0J9YTY0;AOA0J9YUL3;Q8C1B7;AOA0J9YUV6;AOA0J9YVA6	5.557197e+05
AOA0J9YUZ4;P63158;AOA0J9YUD8;D3YVC6;D3YZ18	2.468034e+06
AOA0R4J093;Q9DBP5	-8.206000e+05
AOA0R4J0I1;P07759;E9Q499;Q80X76	5.144795e+06
O70404;AOA0R4J0R1;AOA0U1RPE8	8.577997e+05
...	...
Q9Z0P4;Q3UZP7;AOA1W2P879;AOA1W2P831	-9.870167e+04
Q9Z1D1	2.961831e+05
Q9Z1Q5	1.874555e+06
Q9Z2D6	-4.070078e+05
Q9Z2I8;AOA0N4SWD1	-8.384045e+06

Cluster 3

Cluster 4

Protein IDs

AOA0J9YTY0;AOA0J9YUL3;Q8C1B7;AOA0J9YUV6;AOA0J9YVA6	-4.514309e+06	-3.775957e+06
AOA0J9YUZ4;P63158;AOA0J9YUD8;D3YVC6;D3YZ18	-6.670144e+06	-8.216969e+06
AOA0R4J093;Q9DBP5	2.444753e+06	-3.148042e+05
AOA0R4J0I1;P07759;E9Q499;Q80X76	1.707926e+07	-8.599288e+06
O70404;AOA0R4J0R1;AOA0U1RPE8	-2.495012e+06	-2.381546e+06
...
Q9Z0P4;Q3UZP7;AOA1W2P879;AOA1W2P831	9.093900e+05	5.358425e+05
Q9Z1D1	-1.005512e+06	-4.893086e+05
Q9Z1Q5	-6.734190e+06	-2.573627e+06
Q9Z2D6	2.729420e+06	8.867200e+05
Q9Z2I8;AOA0N4SWD1	2.848534e+07	1.683961e+07

[287 rows x 4 columns]

```
[188]: #make temp df of pGroups_proc with only proteins present in all samples
temp = pGroups_proc.copy()
temp = temp.set_index(' Protein IDs')
temp = temp.loc[centers_t.index]

#plot only mouse markers; should be all proteins since this is the dataset with
→proteins present in all amples
#first pull only mouse index then pull centers_t based on that index

mouse_idx = temp.index[temp['Fasta headers'].str.contains('musculus',
→regex=False, na=False)].tolist()
centers_t_mouse = centers_t.loc[mouse_idx]
```

```
centers_t_mouse
centers_t_mouse.index = centers_t_mouse.index.str.split(';',0).str[0]
centers_t_mouse.to_csv('outs/' + date + '_mouse_centers_annotated.csv')
centers_t_mouse
```

```
[188]:
```

	Cluster 1	Cluster 2	Cluster 3	Cluster 4
Protein IDs				
AOA0J9YTY0	5.347031e+06	5.557197e+05	-4.514309e+06	-3.775957e+06
AOA0J9YUZ4	6.832731e+06	2.468034e+06	-6.670144e+06	-8.216969e+06
AOA0R4J093	6.331833e+05	-8.206000e+05	2.444753e+06	-3.148042e+05
AOA0R4J0I1	-1.365104e+07	5.144795e+06	1.707926e+07	-8.599288e+06
O70404	2.010879e+06	8.577997e+05	-2.495012e+06	-2.381546e+06
...
Q9Z0P4	-8.821325e+05	-9.870167e+04	9.093900e+05	5.358425e+05
Q9Z1D1	4.513414e+05	2.961831e+05	-1.005512e+06	-4.893086e+05
Q9Z1Q5	2.564748e+06	1.874555e+06	-6.734190e+06	-2.573627e+06
Q9Z2D6	-1.676840e+06	-4.070078e+05	2.729420e+06	8.867200e+05
Q9Z2I8	-1.577885e+07	-8.384045e+06	2.848534e+07	1.683961e+07

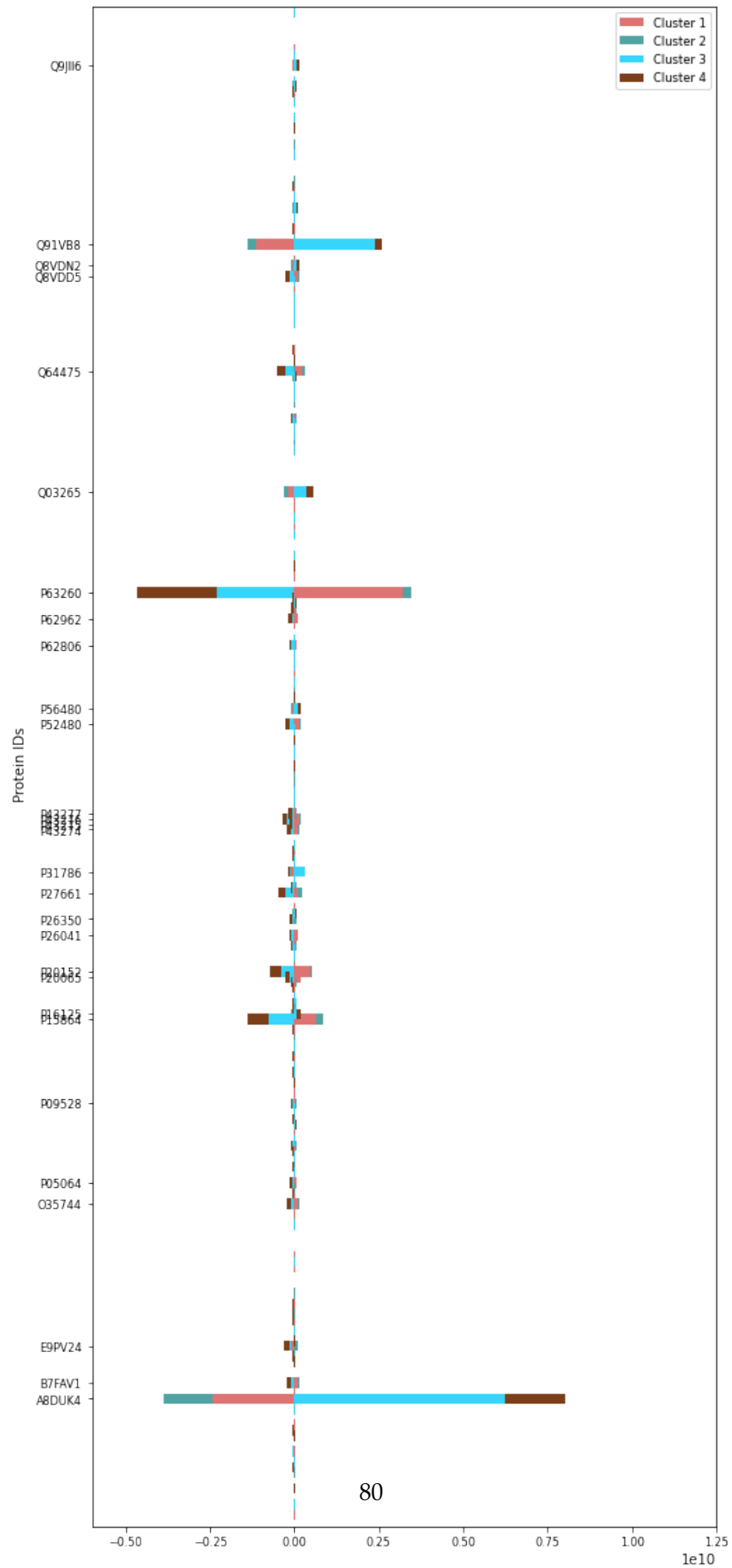
[287 rows x 4 columns]

```
[189]: #plot centroids with top 10% proteins labeled

summed = centers_t_mouse.abs().sum(axis=1)
sorted_mylist = sorted(((v, i) for i, v in enumerate(summed)), reverse=True)[0:
→29]
final = [lis[1] for lis in sorted_mylist]
finalID = centers_t_mouse.iloc[final].index

centers_t_mouse.to_csv('outs/' + date + '_finalID_mouse_all.csv')

centers_t_mouse.plot(kind="barh", stacked=True, width=2, figsize = (8,20),
→color=palette_kmeans)
plt.tick_params(axis='both', which='major', labelsize=8)
plt.yticks(final, finalID)
plt.xlim(-0.60e10, 1.25e10)
plt.legend(fontsize='small')
plt.savefig('outs/' + date + '_mouse_kmeansCentroids_all_top29.svg')
```



0.4 Gene Ontology Enrichment Analysis

0.4.1 Extract top 100 proteins driving the clustering and run gene ontology enrichment analysis using their LFQ intensity values after normalization

```
[190]: allProt = pd.read_csv('outs/' + date + '_finalID_mouse_all.csv', index_col = 0)
allProt.shape
```

```
[190]: (287, 4)
```

```
[191]: summed = allProt.abs().sum(axis=1)
sorted_mylist = sorted(((v, i) for i, v in enumerate(summed)), reverse=True)[0:
    →100]
final = [lis[1] for lis in sorted_mylist]
finalID = allProt.iloc[final].index
finalID
```

```
[191]: Index(['A8DUK4', 'P63260', 'Q91VB8', 'P15864', 'P20152', 'Q03265', 'Q64475',
    'P27661', 'P43276', 'P31786', 'P20065', 'P52480', 'Q8VDD5', 'E9PV24',
    'B7FAV1', 'P43274', 'O35744', 'P62962', 'P56480', 'P16125', 'P26041',
    'P43277', 'Q8VDN2', 'P62806', 'P05064', 'P43275', 'Q9JII6', 'P26350',
    'P09528', 'P27773', 'Q61233', 'Q99KI0', 'P20029', 'P63017', 'P07356',
    'P63038', 'P26039', 'Q9DCW4', 'Q64433', 'P18760', 'D3Z2H9', 'P26443',
    'O88569', 'P17742', 'AOA1D5RLD8', 'AOA1W2P768', 'P11499', 'Q99PT1',
    'P08249', 'P40124', 'E9Q7Q3', 'P40142', 'E9Q3W4', 'Q91Z25', 'Q6IRU2',
    'P13020', 'A2AL12', 'Q9Z2I8', 'E9Q616', 'Q61838', 'P63101', 'P70441',
    'P24270', 'P14733', 'P06745', 'P17182', 'Q9DCD0', 'P05213', 'P38647',
    'P26645', 'Q8BWT1', 'P45952', 'Q9JKF1', 'P99029', 'A3KGU5', 'P08113',
    'P10922', 'AOA1W2P6F6', 'O08749', 'P09103', 'Q61029', 'P17751',
    'P05202', 'Q921I1', 'Q62433', 'P99024', 'P14152', 'O35639',
    'AOA0R4J0I1', 'Q60932', 'P61979', 'Q9DCX2', 'AOA1L1SV25', 'F8WIP8',
    'P10126', 'P48036', 'Q8QZT1', 'E9PZF0', 'F8WIT2', 'P09405'],
    dtype='object', name=' Protein IDs')
```

```
[192]: #import lfq intensity data
lfq = pd.read_csv('outs/' + date + '_combined_lfq.csv', index_col=0)
lfq = lfq.transpose()
lfq = lfq.drop(['TimePoint', 'TechRep', 'BioRep', 'Region'],
    →'DPI_Region_BioRep', 'DPI_Region', 'BioRep_TechRep'], axis=0)
lfq
```

```
[192]: index                                4DPI_interface_2_4 \
G3UZW7;AOA023T778;Q9CQL1;P61327                                0
O70589;AOA067XG53;F6Y9I5                                        0
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...            0
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1                0
AOA075B5P3;AOA0A6YVP0;P01867                                0
```

...	...
Q9Z2X1; J3QMT0; J3QM80; J3QP45; J3QMV8; J3QNH2	5.246e+06
Q9Z2Z6	0
Q9Z315; A0A494B9E9	954220
S4R2J6; S4R165; S4R1X7; S4R2C1; S4R187; S4R2F5	0
S4R192	0
index	4DPI_interface_2_2 \
G3UZW7; A0A023T778; Q9CQL1; P61327	0
O70589; A0A067XG53; F6Y9I5	0
A0A140T8M9; A0A140T8M0; A0A0B4J1I0; A0A075B5N0; A0A...	0
A0A075B5M7; A0A0G2JDV4; A0A0B4J1J2; A0A0B4J1J1	0
A0A075B5P3; A0A0A6YVP0; P01867	0
...	...
Q9Z2X1; J3QMT0; J3QM80; J3QP45; J3QMV8; J3QNH2	3.0364e+06
Q9Z2Z6	668000
Q9Z315; A0A494B9E9	1.3774e+06
S4R2J6; S4R165; S4R1X7; S4R2C1; S4R187; S4R2F5	0
S4R192	0
index	4DPI_interface_2_1 \
G3UZW7; A0A023T778; Q9CQL1; P61327	0
O70589; A0A067XG53; F6Y9I5	0
A0A140T8M9; A0A140T8M0; A0A0B4J1I0; A0A075B5N0; A0A...	0
A0A075B5M7; A0A0G2JDV4; A0A0B4J1J2; A0A0B4J1J1	0
A0A075B5P3; A0A0A6YVP0; P01867	0
...	...
Q9Z2X1; J3QMT0; J3QM80; J3QP45; J3QMV8; J3QNH2	3.9714e+06
Q9Z2Z6	0
Q9Z315; A0A494B9E9	922440
S4R2J6; S4R165; S4R1X7; S4R2C1; S4R187; S4R2F5	0
S4R192	0
index	4DPI_interface_1_2 \
G3UZW7; A0A023T778; Q9CQL1; P61327	0
O70589; A0A067XG53; F6Y9I5	0
A0A140T8M9; A0A140T8M0; A0A0B4J1I0; A0A075B5N0; A0A...	0
A0A075B5M7; A0A0G2JDV4; A0A0B4J1J2; A0A0B4J1J1	0
A0A075B5P3; A0A0A6YVP0; P01867	0
...	...
Q9Z2X1; J3QMT0; J3QM80; J3QP45; J3QMV8; J3QNH2	9.2853e+06
Q9Z2Z6	1.4414e+06
Q9Z315; A0A494B9E9	0
S4R2J6; S4R165; S4R1X7; S4R2C1; S4R187; S4R2F5	0
S4R192	0
index	4DPI_interface_1_1 \

G3UZW7;AOA023T778;Q9CQL1;P61327	0
O70589;AOA067XG53;F6Y9I5	0
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0
AOA075B5P3;AOA0A6YVP0;P01867	0
...	...
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	6.0399e+06
Q9Z2Z6	1.1892e+06
Q9Z315;AOA494B9E9	1.1176e+06
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0
S4R192	0

index	4DPI_cortex_3_3 \
G3UZW7;AOA023T778;Q9CQL1;P61327	0
O70589;AOA067XG53;F6Y9I5	0
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0
AOA075B5P3;AOA0A6YVP0;P01867	0
...	...
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	2.6449e+06
Q9Z2Z6	1.3931e+06
Q9Z315;AOA494B9E9	638210
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0
S4R192	0

index	4DPI_cortex_3_2 \
G3UZW7;AOA023T778;Q9CQL1;P61327	0
O70589;AOA067XG53;F6Y9I5	0
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0
AOA075B5P3;AOA0A6YVP0;P01867	0
...	...
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	0
Q9Z2Z6	950880
Q9Z315;AOA494B9E9	661230
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0
S4R192	0

index	4DPI_cortex_2_3 \
G3UZW7;AOA023T778;Q9CQL1;P61327	0
O70589;AOA067XG53;F6Y9I5	0
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0
AOA075B5P3;AOA0A6YVP0;P01867	0
...	...
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	2.4288e+06
Q9Z2Z6	1.0941e+06

Q9Z315;A0A494B9E9	0	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	662690	
S4R192	0	
index	4DPI_cortex_2_2	\
G3UZW7;A0A023T778;Q9CQL1;P61327	0	
O70589;A0A067XG53;F6Y9I5	0	
A0A140T8M9;A0A140T8M0;A0A0B4J1I0;A0A075B5N0;A0A...	0	
A0A075B5M7;A0A0G2JDV4;A0A0B4J1J2;A0A0B4J1J1	0	
A0A075B5P3;A0A0A6YVP0;P01867	0	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	2.6804e+06	
Q9Z2Z6	1.1699e+06	
Q9Z315;A0A494B9E9	494860	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	4DPI_cortex_2_1	... \
G3UZW7;A0A023T778;Q9CQL1;P61327	0	...
O70589;A0A067XG53;F6Y9I5	497950	...
A0A140T8M9;A0A140T8M0;A0A0B4J1I0;A0A075B5N0;A0A...	0	...
A0A075B5M7;A0A0G2JDV4;A0A0B4J1J2;A0A0B4J1J1	0	...
A0A075B5P3;A0A0A6YVP0;P01867	0	...
...
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	2.0776e+06	...
Q9Z2Z6	1.1146e+06	...
Q9Z315;A0A494B9E9	674540	...
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	...
S4R192	0	...
index	10DPI_cortex_3_2	\
G3UZW7;A0A023T778;Q9CQL1;P61327	0	
O70589;A0A067XG53;F6Y9I5	434700	
A0A140T8M9;A0A140T8M0;A0A0B4J1I0;A0A075B5N0;A0A...	498950	
A0A075B5M7;A0A0G2JDV4;A0A0B4J1J2;A0A0B4J1J1	0	
A0A075B5P3;A0A0A6YVP0;P01867	0	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	2.1112e+06	
Q9Z2Z6	0	
Q9Z315;A0A494B9E9	556170	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	10DPI_SAC_4_3	\
G3UZW7;A0A023T778;Q9CQL1;P61327	0	
O70589;A0A067XG53;F6Y9I5	0	
A0A140T8M9;A0A140T8M0;A0A0B4J1I0;A0A075B5N0;A0A...	1.6741e+06	

AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	202930	
AOA075B5P3;AOA0A6YVP0;P01867	1.2076e+06	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	6.5735e+06	
Q9Z2Z6	338860	
Q9Z315;AOA494B9E9	1.1378e+06	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	10DPI_SAC_4_2	\
G3UZW7;AOA023T778;Q9CQL1;P61327	0	
O70589;AOA067XG53;F6Y9I5	0	
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	431080	
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	290690	
AOA075B5P3;AOA0A6YVP0;P01867	1.5261e+06	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	6.941e+06	
Q9Z2Z6	0	
Q9Z315;AOA494B9E9	1.3022e+06	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	10DPI_SAC_4_1	\
G3UZW7;AOA023T778;Q9CQL1;P61327	0	
O70589;AOA067XG53;F6Y9I5	0	
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	548100	
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0	
AOA075B5P3;AOA0A6YVP0;P01867	1.1656e+06	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	6.2944e+06	
Q9Z2Z6	0	
Q9Z315;AOA494B9E9	0	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	326950	
index	10DPI_SAC_3_3	\
G3UZW7;AOA023T778;Q9CQL1;P61327	0	
O70589;AOA067XG53;F6Y9I5	0	
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0	
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0	
AOA075B5P3;AOA0A6YVP0;P01867	0	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	4.6115e+06	
Q9Z2Z6	0	
Q9Z315;AOA494B9E9	1.7312e+06	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	423330	

index	10DPI_SAC_3_2	\
G3UZW7;AOA023T778;Q9CQL1;P61327	0	
O70589;AOA067XG53;F6Y9I5	0	
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0	
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0	
AOA075B5P3;AOA0A6YVP0;P01867	0	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	6.5767e+06	
Q9Z2Z6	0	
Q9Z315;AOA494B9E9	1.4447e+06	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	10DPI_SAC_3_1	\
G3UZW7;AOA023T778;Q9CQL1;P61327	0	
O70589;AOA067XG53;F6Y9I5	0	
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0	
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0	
AOA075B5P3;AOA0A6YVP0;P01867	724260	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	5.9742e+06	
Q9Z2Z6	0	
Q9Z315;AOA494B9E9	1.0595e+06	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	10DPI_SAC_2_3	\
G3UZW7;AOA023T778;Q9CQL1;P61327	0	
O70589;AOA067XG53;F6Y9I5	0	
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0	
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0	
AOA075B5P3;AOA0A6YVP0;P01867	0	
...	...	
Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	2.8952e+06	
Q9Z2Z6	0	
Q9Z315;AOA494B9E9	3.3433e+06	
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	
S4R192	0	
index	10DPI_SAC_2_2	10DPI_SAC_2_1
G3UZW7;AOA023T778;Q9CQL1;P61327	0	0
O70589;AOA067XG53;F6Y9I5	0	0
AOA140T8M9;AOA140T8M0;AOA0B4J1I0;AOA075B5N0;AOA...	0	0
AOA075B5M7;AOA0G2JDV4;AOA0B4J1J2;AOA0B4J1J1	0	0
AOA075B5P3;AOA0A6YVP0;P01867	0	0
...

Q9Z2X1;J3QMT0;J3QM80;J3QP45;J3QMV8;J3QNH2	3.8473e+06	6.8296e+06
Q9Z2Z6	0	0
Q9Z315;AOA494B9E9	800330	2.3382e+06
S4R2J6;S4R165;S4R1X7;S4R2C1;S4R187;S4R2F5	0	0
S4R192	0	0

[3613 rows x 39 columns]

[193]: *#for ID's with multiple proteins per protein group, select first protein in
 ↳protein group for analysis*
 lfq.index = lfq.index.str.split(';',0).str[0]
 lfq

[193]:

index	4DPI_interface_2_4	4DPI_interface_2_2	4DPI_interface_2_1	\
G3UZW7	0	0	0	
O70589	0	0	0	
AOA140T8M9	0	0	0	
AOA075B5M7	0	0	0	
AOA075B5P3	0	0	0	
...	
Q9Z2X1	5.246e+06	3.0364e+06	3.9714e+06	
Q9Z2Z6	0	668000	0	
Q9Z315	954220	1.3774e+06	922440	
S4R2J6	0	0	0	
S4R192	0	0	0	

index	4DPI_interface_1_2	4DPI_interface_1_1	4DPI_cortex_3_3	\
G3UZW7	0	0	0	
O70589	0	0	0	
AOA140T8M9	0	0	0	
AOA075B5M7	0	0	0	
AOA075B5P3	0	0	0	
...	
Q9Z2X1	9.2853e+06	6.0399e+06	2.6449e+06	
Q9Z2Z6	1.4414e+06	1.1892e+06	1.3931e+06	
Q9Z315	0	1.1176e+06	638210	
S4R2J6	0	0	0	
S4R192	0	0	0	

index	4DPI_cortex_3_2	4DPI_cortex_2_3	4DPI_cortex_2_2	4DPI_cortex_2_1	\
G3UZW7	0	0	0	0	
O70589	0	0	0	497950	
AOA140T8M9	0	0	0	0	
AOA075B5M7	0	0	0	0	
AOA075B5P3	0	0	0	0	
...	
Q9Z2X1	0	2.4288e+06	2.6804e+06	2.0776e+06	
Q9Z2Z6	950880	1.0941e+06	1.1699e+06	1.1146e+06	

Q9Z315	661230	0	494860	674540
S4R2J6	0	662690	0	0
S4R192	0	0	0	0

index	...	10DPI_cortex_3_2	10DPI_SAC_4_3	10DPI_SAC_4_2	10DPI_SAC_4_1	\
G3UZW7	...	0	0	0	0	
O70589	...	434700	0	0	0	
AOA140T8M9	...	498950	1.6741e+06	431080	548100	
AOA075B5M7	...	0	202930	290690	0	
AOA075B5P3	...	0	1.2076e+06	1.5261e+06	1.1656e+06	
...	
Q9Z2X1	...	2.1112e+06	6.5735e+06	6.941e+06	6.2944e+06	
Q9Z2Z6	...	0	338860	0	0	
Q9Z315	...	556170	1.1378e+06	1.3022e+06	0	
S4R2J6	...	0	0	0	0	
S4R192	...	0	0	0	326950	

index	10DPI_SAC_3_3	10DPI_SAC_3_2	10DPI_SAC_3_1	10DPI_SAC_2_3	\
G3UZW7	0	0	0	0	
O70589	0	0	0	0	
AOA140T8M9	0	0	0	0	
AOA075B5M7	0	0	0	0	
AOA075B5P3	0	0	724260	0	
...	
Q9Z2X1	4.6115e+06	6.5767e+06	5.9742e+06	2.8952e+06	
Q9Z2Z6	0	0	0	0	
Q9Z315	1.7312e+06	1.4447e+06	1.0595e+06	3.3433e+06	
S4R2J6	0	0	0	0	
S4R192	423330	0	0	0	

index	10DPI_SAC_2_2	10DPI_SAC_2_1
G3UZW7	0	0
O70589	0	0
AOA140T8M9	0	0
AOA075B5M7	0	0
AOA075B5P3	0	0
...
Q9Z2X1	3.8473e+06	6.8296e+06
Q9Z2Z6	0	0
Q9Z315	800330	2.3382e+06
S4R2J6	0	0
S4R192	0	0

[3613 rows x 39 columns]

```
[194]: #select top 100 proteins based on highest k-means centroid values
df = lfq.loc[finalID]
```


df

```
[194]: index      4DPI_interface_2_4 4DPI_interface_2_2 4DPI_interface_2_1 \
      Protein IDs
A8DUK4      1.7442e+09      3.4139e+09      2.9423e+09
P63260      4.1307e+09      3.378e+09      2.9545e+09
Q91VB8      1.7297e+09      1.9171e+09      1.4639e+09
P15864      9.1421e+08      6.0707e+08      3.477e+08
P20152      4.5815e+08      4.2805e+08      3.6485e+08
...
P48036      5.4002e+07      4.5385e+07      6.3113e+07
Q8QZT1      1.1984e+07      1.7382e+07      1.5672e+07
E9PZF0      3.8056e+07      5.7943e+07      3.3903e+07
F8WIT2      2.2663e+07      1.4908e+07      3.6708e+07
P09405      6.0954e+07      4.9944e+07      5.7182e+07

index      4DPI_interface_1_2 4DPI_interface_1_1 4DPI_cortex_3_3 \
      Protein IDs
A8DUK4      5.1737e+09      3.8777e+09      5.782e+09
P63260      1.771e+09      1.6077e+09      1.2741e+09
Q91VB8      1.3123e+09      1.8905e+09      2.7786e+09
P15864      4.7936e+08      5.7903e+08      2.5711e+08
P20152      3.079e+08      3.2536e+08      5.2257e+07
...
P48036      5.3553e+07      5.4752e+07      1.362e+07
Q8QZT1      8.09e+06      1.1016e+07      5.4671e+07
E9PZF0      2.0192e+07      3.1333e+07      6.8917e+07
F8WIT2      1.3592e+06      4.1288e+06      1.5675e+06
P09405      3.2236e+07      4.5901e+07      1.6173e+07

index      4DPI_cortex_3_2 4DPI_cortex_2_3 4DPI_cortex_2_2 4DPI_cortex_2_1 \
      Protein IDs
A8DUK4      6.8564e+09      8.5305e+09      8.4491e+09      3.4106e+09
P63260      1.8186e+09      1.5937e+09      2.1845e+09      1.5055e+09
Q91VB8      1.9416e+09      2.0437e+09      2.8438e+09      2.0836e+09
P15864      2.659e+08      3.1331e+08      2.7457e+08      3.3855e+08
P20152      7.7997e+07      1.2761e+08      1.2123e+08      3.4905e+07
...
P48036      2.8752e+07      4.4965e+07      4.3174e+07      2.4549e+07
Q8QZT1      3.6034e+07      3.6496e+07      4.7283e+07      5.1152e+07
E9PZF0      6.386e+07      3.55e+07      5.2153e+07      5.4563e+07
F8WIT2      4.3106e+06      1.0905e+07      1.2193e+07      2.284e+06
P09405      1.7219e+07      3.9117e+07      3.0935e+07      7.6649e+06

index      ... 10DPI_cortex_3_2 10DPI_SAC_4_3 10DPI_SAC_4_2 10DPI_SAC_4_1 \
      Protein IDs ...
A8DUK4      ...      8.122e+09      5.3066e+08      9.5197e+08      6.2291e+08
```

P63260	...	1.3877e+09	4.8895e+09	4.6028e+09	4.8811e+09
Q91VB8	...	5.3114e+09	3.0269e+08	4.4618e+08	3.8388e+08
P15864	...	3.7537e+08	8.0293e+08	1.3161e+09	1.2945e+09
P20152	...	5.8747e+07	7.1322e+08	8.472e+08	6.4955e+08
...
P48036	...	2.8541e+07	7.5051e+07	7.6192e+07	6.2647e+07
Q8QZT1	...	2.434e+07	6.1046e+06	3.5123e+06	7.2743e+06
E9PZF0	...	4.9142e+07	4.5855e+07	3.8078e+07	3.7067e+07
F8WIT2	...	1.9329e+06	2.8045e+07	2.4982e+07	2.3535e+07
P09405	...	2.367e+07	3.7192e+07	4.091e+07	4.1109e+07

index	10DPI_SAC_3_3	10DPI_SAC_3_2	10DPI_SAC_3_1	10DPI_SAC_2_3	\
Protein IDs					
A8DUK4	8.5632e+08	8.6326e+08	9.2512e+08	1.0979e+09	
P63260	6.279e+09	6.4949e+09	7.117e+09	4.7165e+09	
Q91VB8	1.0176e+09	8.2537e+08	6.0648e+08	1.3203e+09	
P15864	2.0244e+09	1.426e+09	1.6338e+09	3.6126e+09	
P20152	9.1203e+08	8.0969e+08	7.6861e+08	6.5963e+08	
...	
P48036	6.6472e+07	9.4073e+07	7.7432e+07	5.382e+07	
Q8QZT1	5.6125e+06	385710	9.5847e+06	6.7139e+06	
E9PZF0	2.3838e+07	1.8537e+07	2.1536e+07	1.8987e+07	
F8WIT2	2.866e+07	4.5345e+07	4.2387e+07	1.0955e+07	
P09405	4.5695e+07	3.9475e+07	5.1029e+07	3.6213e+07	

index	10DPI_SAC_2_2	10DPI_SAC_2_1
Protein IDs		
A8DUK4	1.2606e+09	1.8661e+09
P63260	4.8012e+09	4.7378e+09
Q91VB8	2.0021e+09	8.6326e+08
P15864	1.3788e+09	2.4812e+09
P20152	9.2283e+08	6.4163e+08
...
P48036	4.0377e+07	5.0462e+07
Q8QZT1	1.0046e+07	2.324e+06
E9PZF0	2.9525e+07	2.8315e+07
F8WIT2	1.5971e+07	4.3004e+06
P09405	5.2582e+07	4.8543e+07

[100 rows x 39 columns]

```
[195]: #transpose and change datatype to int64
df2 = df.transpose()
df2 = df2.astype('int64')
```

```
[196]: #standardize values by removing the mean and scaling to unit variance
df3 = StandardScaler().fit_transform(df2)
df3 = pd.DataFrame(df3, columns=df2.columns, index=df2.index)
```

df3

[196]:	Protein IDs	A8DUK4	P63260	Q91VB8	P15864	P20152	\
	index						
	4DPI_interface_2_4	-0.540090	0.021508	-0.022581	-0.236924	-0.124103	
	4DPI_interface_2_2	0.019054	-0.337894	0.113558	-0.642223	-0.213255	
	4DPI_interface_2_1	-0.138874	-0.540108	-0.215675	-0.984486	-0.400444	
	4DPI_interface_1_2	0.608371	-1.105209	-0.325806	-0.810748	-0.569122	
	4DPI_interface_1_1	0.174370	-1.183182	0.094234	-0.679225	-0.517408	
	4DPI_cortex_3_3	0.812077	-1.342470	0.739405	-1.104028	-1.326299	
	4DPI_cortex_3_2	1.171869	-1.082481	0.131356	-1.092429	-1.250061	
	4DPI_cortex_2_3	1.732487	-1.189867	0.205528	-1.029867	-1.103115	
	4DPI_cortex_2_2	1.705228	-0.907770	0.786771	-1.080988	-1.122011	
	4DPI_cortex_2_1	0.017949	-1.231981	0.234514	-0.996560	-1.377694	
	4DPI_cortex_1_2	3.162245	-0.851188	2.487419	-0.892101	-1.028298	
	4DPI_cortex_1_1	2.284197	-1.246783	2.586436	-1.133441	-1.165106	
	4DPI_SAC_3_3	-0.653982	0.011098	-0.271394	1.321765	0.137459	
	4DPI_SAC_3_2	-0.555428	0.006037	-0.709916	0.867429	0.003198	
	4DPI_SAC_2_3	-0.512329	-0.250133	-0.485555	-0.434520	-0.619947	
	4DPI_SAC_2_2	-0.480516	-0.370315	-0.212042	-0.525994	-0.592846	
	4DPI_SAC_2_1	-0.371781	0.039127	-0.427148	0.498867	0.228388	
	4DPI_SAC_1_3	0.162783	0.504720	0.867335	-0.745891	-0.339045	
	4DPI_SAC_1_2	-0.281163	0.401154	2.496209	-0.308182	-0.292751	
	4DPI_SAC_1_1	-0.043099	0.343665	0.096850	-0.764444	-0.312506	
	10DPI_interface_5_3	-0.777652	1.419721	-0.753751	0.824938	1.692848	
	10DPI_interface_4_3	-0.866100	1.606369	-1.090698	1.305138	1.805398	
	10DPI_interface_4_2	-0.873578	2.688393	-1.119343	0.698258	1.568746	
	10DPI_interface_4_1	-0.800096	1.695420	-0.998278	1.009681	1.820800	
	10DPI_interface_3_2	-0.632316	0.429326	-0.690294	0.874819	1.296107	
	10DPI_interface_3_1	-0.719082	1.182173	-0.586279	0.609317	1.189421	
	10DPI_cortex_4_4	0.249718	-0.833378	-0.002022	-0.349340	-0.885863	
	10DPI_cortex_4_3	1.169458	-1.045285	0.320527	-0.289708	-0.986684	
	10DPI_cortex_3_3	0.492771	-1.152337	0.040476	-1.045689	-1.388084	
	10DPI_cortex_3_2	1.595690	-1.288228	2.579389	-0.947973	-1.307077	
	10DPI_SAC_4_3	-0.946477	0.383822	-1.059250	-0.383768	0.631378	
	10DPI_SAC_4_2	-0.805390	0.246927	-0.955010	0.293407	1.028207	
	10DPI_SAC_4_1	-0.915585	0.379811	-1.000268	0.264904	0.442797	
	10DPI_SAC_3_3	-0.837421	1.047284	-0.539895	1.228074	1.220225	
	10DPI_SAC_3_2	-0.835097	1.150373	-0.679543	0.438430	0.917108	
	10DPI_SAC_3_1	-0.814382	1.447415	-0.838558	0.712641	0.795435	
	10DPI_SAC_2_3	-0.756522	0.301217	-0.319995	3.323850	0.472652	
	10DPI_SAC_2_2	-0.702037	0.341660	0.175307	0.376145	1.252213	
	10DPI_SAC_2_1	-0.499269	0.311387	-0.652017	1.830863	0.419339	
	Protein IDs	Q03265	Q64475	P27661	P43276	P31786	...
	index						...
	4DPI_interface_2_4	-0.225207	0.487461	0.304723	-0.028911	0.134433	...

4DPI_interface_2_2	0.089789	-0.433295	-0.113263	-0.522751	-0.504536	...
4DPI_interface_2_1	-0.369529	-0.155440	-0.818871	-1.047645	-1.302654	...
4DPI_interface_1_2	-0.241213	0.125845	-0.007414	0.002511	-1.015394	...
4DPI_interface_1_1	0.168928	-1.043352	-0.340969	-0.393190	-0.514812	...
4DPI_cortex_3_3	1.809980	-1.645795	-1.310597	-1.351087	0.551226	...
4DPI_cortex_3_2	1.360204	-1.052557	-1.656457	-1.324128	0.710533	...
4DPI_cortex_2_3	2.563874	-1.384593	-0.422396	-1.128870	2.363262	...
4DPI_cortex_2_2	1.560213	-1.012349	-1.323757	-0.949139	1.326104	...
4DPI_cortex_2_1	2.644484	-1.569062	-1.415339	-1.385625	-0.472758	...
4DPI_cortex_1_2	0.906860	-0.699643	-0.763131	-0.753384	1.490523	...
4DPI_cortex_1_1	0.987024	-1.281794	-0.206395	-1.246260	0.739623	...
4DPI_SAC_3_3	-0.525400	1.972168	2.332589	0.926510	-0.777197	...
4DPI_SAC_3_2	-0.485897	0.522564	0.658175	0.951647	-0.857984	...
4DPI_SAC_2_3	-0.125960	0.362906	-0.387693	0.358201	1.684032	...
4DPI_SAC_2_2	-0.046063	1.084758	0.075848	0.243042	1.603192	...
4DPI_SAC_2_1	-0.187086	-0.117659	0.580671	0.241366	-0.301172	...
4DPI_SAC_1_3	-0.217939	-0.801061	1.767597	-0.877708	0.019972	...
4DPI_SAC_1_2	-0.500477	-0.780309	1.001093	-0.349095	0.899931	...
4DPI_SAC_1_1	-0.382281	-0.266356	0.663553	-0.823244	0.018918	...
10DPI_interface_5_3	-0.901580	1.013422	-0.382157	0.127778	-0.526985	...
10DPI_interface_4_3	-0.827514	1.870624	0.372136	1.542423	-0.595809	...
10DPI_interface_4_2	-0.759250	1.245463	0.559159	0.560974	-0.991206	...
10DPI_interface_4_1	-0.808740	0.584194	0.445623	1.316398	-0.703630	...
10DPI_interface_3_2	-0.899163	1.029865	0.112986	0.569039	-0.266707	...
10DPI_interface_3_1	-0.726168	0.444367	0.102009	0.726513	-0.385806	...
10DPI_cortex_4_4	0.463414	-0.368067	0.051774	-0.178477	-0.821938	...
10DPI_cortex_4_3	0.495828	-0.910387	0.338825	-0.099504	0.317561	...
10DPI_cortex_3_3	0.805517	-1.033478	-1.298924	-1.414690	-0.097914	...
10DPI_cortex_3_2	1.836598	-1.234055	-1.243058	-1.222343	2.401152	...
10DPI_SAC_4_3	-0.887629	-0.012392	-0.042908	-0.335688	-1.490924	...
10DPI_SAC_4_2	-0.928505	-0.264724	0.373718	0.088606	-1.390982	...
10DPI_SAC_4_1	-0.804905	0.156136	-0.493826	-0.204923	-1.472079	...
10DPI_SAC_3_3	-0.772893	0.738915	0.963448	1.472510	-0.798119	...
10DPI_SAC_3_2	-0.810746	0.665529	0.529107	0.355426	-0.462956	...
10DPI_SAC_3_1	-0.729155	1.164043	0.115897	0.340239	-0.021449	...
10DPI_SAC_2_3	-0.941055	2.246173	2.216491	2.602791	0.750162	...
10DPI_SAC_2_2	-0.734416	0.680466	1.143132	0.986525	0.257537	...
10DPI_SAC_2_1	-0.853945	-0.328529	-2.481400	2.224163	0.504851	...

Protein IDs	P61979	Q9DCX2	A0A1L1SV25	F8WIP8	P10126	\
index						
4DPI_interface_2_4	0.573323	-0.334925	0.018748	0.107914	1.212955	
4DPI_interface_2_2	-0.566791	0.321537	1.474015	0.968526	-0.854949	
4DPI_interface_2_1	-0.193876	0.310361	0.307097	-0.023676	-0.441144	
4DPI_interface_1_2	-1.254921	-0.812839	0.199176	1.497482	1.173070	
4DPI_interface_1_1	-0.503401	0.292111	1.774868	0.328190	-1.647993	
4DPI_cortex_3_3	-1.516123	2.398939	0.720059	-0.229218	-0.556935	

4DPI_cortex_3_2	-0.977258	1.324569	0.731844	0.454642	0.938331
4DPI_cortex_2_3	-0.687040	2.349517	1.431181	2.344207	-0.239975
4DPI_cortex_2_2	-0.905276	2.138293	2.043236	3.298078	-0.636912
4DPI_cortex_2_1	-0.815326	0.759878	-0.269183	-0.294759	-1.810814
4DPI_cortex_1_2	0.065149	1.148796	-0.013677	2.515370	0.541643
4DPI_cortex_1_1	0.772643	1.394336	2.132254	0.542312	0.812029
4DPI_SAC_3_3	0.751048	-0.350119	-0.709544	0.106389	-0.141509
4DPI_SAC_3_2	-0.764715	-0.412290	-0.747891	0.087704	-1.318569
4DPI_SAC_2_3	-0.978095	0.036965	-1.984384	0.284609	-0.330962
4DPI_SAC_2_2	0.503350	-0.281836	-1.038753	-0.296284	0.070711
4DPI_SAC_2_1	0.015152	-0.096370	0.178716	0.518885	-0.455685
4DPI_SAC_1_3	-0.344091	-0.479700	0.296030	0.670741	0.446418
4DPI_SAC_1_2	-0.576277	-0.367495	-0.237596	-0.092830	0.151312
4DPI_SAC_1_1	0.586827	-0.592533	-0.200805	0.315036	1.908031
10DPI_interface_5_3	0.013980	-0.907178	-0.050588	-0.081766	2.083774
10DPI_interface_4_3	0.920569	-1.111181	0.140489	-1.168785	0.730058
10DPI_interface_4_2	1.032170	-0.882729	0.063615	-1.169293	0.702471
10DPI_interface_4_1	0.202920	-0.728393	-0.880639	-1.214748	-0.650622
10DPI_interface_3_2	0.118885	-0.491401	-1.009320	-0.610496	0.617134
10DPI_interface_3_1	-0.119885	-0.893251	-0.182499	-0.552520	1.172655
10DPI_cortex_4_4	-1.469306	0.694825	1.663895	-0.550092	0.606789
10DPI_cortex_4_3	0.562498	-0.194255	0.216405	-0.609141	-1.383216
10DPI_cortex_3_3	-2.290354	1.912836	0.919092	-1.265154	-1.059899
10DPI_cortex_3_2	-1.737204	0.966998	-0.815013	-1.216605	-1.192724
10DPI_SAC_4_3	1.407205	-0.724979	0.791728	-0.444244	-1.382800
10DPI_SAC_4_2	2.253920	-0.875848	-0.514998	-0.913915	-0.916771
10DPI_SAC_4_1	2.103817	-0.870530	-0.130691	-0.379494	-0.759516
10DPI_SAC_3_3	1.343202	-0.625610	-1.035522	-0.322308	0.767741
10DPI_SAC_3_2	0.129320	-1.125903	-0.712475	-0.495334	1.261565
10DPI_SAC_3_1	0.764775	-0.839165	-0.457866	-0.619811	1.450602
10DPI_SAC_2_3	0.705905	-0.551101	-1.993058	-0.614956	-0.233784
10DPI_SAC_2_2	0.323393	-0.616799	-0.699494	-0.551729	0.151312
10DPI_SAC_2_1	0.549887	-0.883532	-1.418453	-0.322929	-0.783821

Protein IDs	P48036	Q8QZT1	E9PZF0	F8WIT2	P09405
index					
4DPI_interface_2_4	-0.024608	-0.362371	-0.030764	0.333282	1.597443
4DPI_interface_2_2	-0.503046	0.048106	1.395860	-0.255785	0.741501
4DPI_interface_2_1	0.481258	-0.081927	-0.328686	1.400135	1.304200
4DPI_interface_1_2	-0.049537	-0.658481	-1.312265	-1.284947	-0.635158
4DPI_interface_1_1	0.017034	-0.435981	-0.513049	-1.074569	0.427190
4DPI_cortex_3_3	-2.266719	2.883655	2.183097	-1.269124	-1.883931
4DPI_cortex_3_2	-1.426552	1.466451	1.820325	-1.060759	-1.802613
4DPI_cortex_2_3	-0.526365	1.501583	-0.214122	-0.559851	-0.100214
4DPI_cortex_2_2	-0.625806	2.321853	0.980506	-0.462015	-0.736300
4DPI_cortex_2_1	-1.659913	2.616062	1.153391	-1.214699	-2.545370
4DPI_cortex_1_2	-1.030787	0.707471	1.691200	-0.514503	-0.741976

4DPI_cortex_1_1	-0.595879	0.885410	2.194431	-0.125894	-0.369901
4DPI_SAC_3_3	-1.048333	-0.082839	-0.089157	-0.533417	1.397490
4DPI_SAC_3_2	-0.713698	-0.405868	-0.592748	-0.343139	0.263853
4DPI_SAC_2_3	0.411022	-0.084284	0.319239	-0.942338	-1.200577
4DPI_SAC_2_2	-0.711533	0.207415	-0.308169	-0.712401	0.020676
4DPI_SAC_2_1	-0.634745	-0.365261	0.010556	-0.392664	1.233921
4DPI_SAC_1_3	-0.324874	0.654773	0.692699	0.203771	0.307000
4DPI_SAC_1_2	-0.207055	-0.169680	-0.464053	-0.223654	-0.654438
4DPI_SAC_1_1	-0.554571	0.140422	0.212710	0.059371	0.638882
10DPI_interface_5_3	0.992343	-0.835014	-0.679406	2.108762	1.231277
10DPI_interface_4_3	0.235904	-0.576910	-0.186289	1.100246	1.105335
10DPI_interface_4_2	0.341730	-0.686738	-0.772089	0.692647	0.991598
10DPI_interface_4_1	0.772141	-0.715809	-1.494046	0.663326	1.152602
10DPI_interface_3_2	1.074627	-0.620855	-1.031775	1.966034	-0.021150
10DPI_interface_3_1	1.524249	-0.677940	-1.059968	1.545065	0.638104
10DPI_cortex_4_4	1.000782	-0.770248	-0.080477	-0.645693	-0.976057
10DPI_cortex_4_3	2.065094	0.627398	0.954609	-0.752614	-1.150666
10DPI_cortex_3_3	-0.488110	0.434782	0.854680	-0.998138	-1.193425
10DPI_cortex_3_2	-1.438267	0.577210	0.764507	-1.241369	-1.301098
10DPI_SAC_4_3	1.144086	-0.809456	0.528709	0.742096	-0.249867
10DPI_SAC_4_2	1.207437	-1.006581	-0.029186	0.509432	0.039178
10DPI_SAC_4_1	0.455384	-0.720509	-0.101711	0.399519	0.054649
10DPI_SAC_3_3	0.667758	-0.846876	-1.050714	0.788812	0.411175
10DPI_SAC_3_2	2.200236	-1.244334	-1.430989	2.056198	-0.072382
10DPI_SAC_3_1	1.276285	-0.544820	-1.215851	1.831509	0.825852
10DPI_SAC_2_3	-0.034713	-0.763123	-1.398708	-0.556053	-0.325977
10DPI_SAC_2_2	-0.781103	-0.509742	-0.642748	-0.175040	0.946585
10DPI_SAC_2_1	-0.221158	-1.096942	-0.729549	-1.061534	0.632585

[39 rows x 100 columns]

```
[197]: #load in kmeans results
kmeans = pd.read_csv('outs/' + date + '_combined_df_segm_pca_kmeans.csv',
    ↪index_col=0)
kmeans
```

```
[197]: AOA0J9YTY0;AOA0J9YUL3;Q8C1B7;AOA0J9YUV6;AOA0J9YVA6 \
index
4DPI_interface_2_4      8884400.0
4DPI_interface_2_2      4805100.0
4DPI_interface_2_1      4679400.0
4DPI_interface_1_2      2752000.0
4DPI_interface_1_1      6446300.0
4DPI_cortex_3_3         3966100.0
4DPI_cortex_3_2         3174000.0
4DPI_cortex_2_3         3505600.0
4DPI_cortex_2_2         3623400.0
4DPI_cortex_2_1         5408800.0
```

4DPI_cortex_1_2	3500500.0
4DPI_cortex_1_1	4027400.0
4DPI_SAC_3_3	10933000.0
4DPI_SAC_3_2	9909200.0
4DPI_SAC_2_3	7513200.0
4DPI_SAC_2_2	5452600.0
4DPI_SAC_2_1	11385000.0
4DPI_SAC_1_3	8059300.0
4DPI_SAC_1_2	5600200.0
4DPI_SAC_1_1	6698000.0
10DPI_interface_5_3	14004000.0
10DPI_interface_4_3	11083000.0
10DPI_interface_4_2	10662000.0
10DPI_interface_4_1	10168000.0
10DPI_interface_3_2	10654000.0
10DPI_interface_3_1	13143000.0
10DPI_cortex_4_4	4613200.0
10DPI_cortex_4_3	7330400.0
10DPI_cortex_3_3	2525300.0
10DPI_cortex_3_2	4286400.0
10DPI_SAC_4_3	9362000.0
10DPI_SAC_4_2	10750000.0
10DPI_SAC_4_1	10526000.0
10DPI_SAC_3_3	22060000.0
10DPI_SAC_3_2	16201000.0
10DPI_SAC_3_1	11879000.0
10DPI_SAC_2_3	10519000.0
10DPI_SAC_2_2	11053000.0
10DPI_SAC_2_1	12673000.0

AOA0J9YUZ4;P63158;AOA0J9YUD8;D3YVC6;D3YZ18 \

index

4DPI_interface_2_4	13795000.0
4DPI_interface_2_2	16043000.0
4DPI_interface_2_1	14321000.0
4DPI_interface_1_2	2559000.0
4DPI_interface_1_1	18740000.0
4DPI_cortex_3_3	10538000.0
4DPI_cortex_3_2	5800300.0
4DPI_cortex_2_3	7019000.0
4DPI_cortex_2_2	9811200.0
4DPI_cortex_2_1	7080900.0
4DPI_cortex_1_2	13352000.0
4DPI_cortex_1_1	15997000.0
4DPI_SAC_3_3	24605000.0
4DPI_SAC_3_2	18164000.0
4DPI_SAC_2_3	11153000.0

4DPI_SAC_2_2	7476200.0
4DPI_SAC_2_1	24354000.0
4DPI_SAC_1_3	15659000.0
4DPI_SAC_1_2	18231000.0
4DPI_SAC_1_1	17122000.0
10DPI_interface_5_3	19714000.0
10DPI_interface_4_3	41455000.0
10DPI_interface_4_2	18782000.0
10DPI_interface_4_1	26015000.0
10DPI_interface_3_2	28269000.0
10DPI_interface_3_1	18962000.0
10DPI_cortex_4_4	6571500.0
10DPI_cortex_4_3	12236000.0
10DPI_cortex_3_3	2883700.0
10DPI_cortex_3_2	3060800.0
10DPI_SAC_4_3	19296000.0
10DPI_SAC_4_2	26852000.0
10DPI_SAC_4_1	17329000.0
10DPI_SAC_3_3	20678000.0
10DPI_SAC_3_2	18474000.0
10DPI_SAC_3_1	22727000.0
10DPI_SAC_2_3	21473000.0
10DPI_SAC_2_2	33776000.0
10DPI_SAC_2_1	13833000.0

AOAOR4J093;Q9DBP5 AOAOR4J0I1;P07759;E9Q499;Q80X76 \

index

4DPI_interface_2_4	5749000.0	73676000.0
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4DPI_interface_2_1	9228800.0	105190000.0
4DPI_interface_1_2	6292300.0	22129000.0
4DPI_interface_1_1	18973000.0	38062000.0
4DPI_cortex_3_3	14700000.0	32753000.0
4DPI_cortex_3_2	12382000.0	24879000.0
4DPI_cortex_2_3	15484000.0	63258000.0
4DPI_cortex_2_2	11188000.0	47764000.0
4DPI_cortex_2_1	7899100.0	44351000.0
4DPI_cortex_1_2	15600000.0	60586000.0
4DPI_cortex_1_1	14481000.0	84161000.0
4DPI_SAC_3_3	10172000.0	33907000.0
4DPI_SAC_3_2	11552000.0	28193000.0
4DPI_SAC_2_3	13766000.0	56852000.0
4DPI_SAC_2_2	9574300.0	37292000.0
4DPI_SAC_2_1	12936000.0	60285000.0
4DPI_SAC_1_3	6236600.0	79677000.0
4DPI_SAC_1_2	4541100.0	63271000.0
4DPI_SAC_1_1	5351400.0	53311000.0

10DPI_interface_5_3	12167000.0	16442000.0
10DPI_interface_4_3	11927000.0	20993000.0
10DPI_interface_4_2	12228000.0	37156000.0
10DPI_interface_4_1	9103100.0	37811000.0
10DPI_interface_3_2	12785000.0	17304000.0
10DPI_interface_3_1	13679000.0	23913000.0
10DPI_cortex_4_4	15905000.0	21923000.0
10DPI_cortex_4_3	3591100.0	28778000.0
10DPI_cortex_3_3	2846000.0	17751000.0
10DPI_cortex_3_2	8662600.0	16765000.0
10DPI_SAC_4_3	12920000.0	26820000.0
10DPI_SAC_4_2	12945000.0	18110000.0
10DPI_SAC_4_1	11294000.0	17675000.0
10DPI_SAC_3_3	11293000.0	17459000.0
10DPI_SAC_3_2	10115000.0	16184000.0
10DPI_SAC_3_1	9660300.0	20254000.0
10DPI_SAC_2_3	7141400.0	11286000.0
10DPI_SAC_2_2	11842000.0	14575000.0
10DPI_SAC_2_1	8280200.0	18280000.0

070404;AOA0R4JOR1;AOA0U1RPE8 AOA0R4JOZ1;P08003 \

index

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4DPI_interface_2_2	5048700.0	20709000.0
4DPI_interface_2_1	4421600.0	17844000.0
4DPI_interface_1_2	1316400.0	22085000.0
4DPI_interface_1_1	2682100.0	24030000.0
4DPI_cortex_3_3	2543300.0	7914900.0
4DPI_cortex_3_2	2405800.0	10756000.0
4DPI_cortex_2_3	3551100.0	9577000.0
4DPI_cortex_2_2	3017800.0	14510000.0
4DPI_cortex_2_1	3218600.0	9714800.0
4DPI_cortex_1_2	4396200.0	5588800.0
4DPI_cortex_1_1	559210.0	12618000.0
4DPI_SAC_3_3	6073100.0	24566000.0
4DPI_SAC_3_2	5000700.0	20284000.0
4DPI_SAC_2_3	7744500.0	14750000.0
4DPI_SAC_2_2	3349900.0	14702000.0
4DPI_SAC_2_1	7488800.0	22398000.0
4DPI_SAC_1_3	4729600.0	22668000.0
4DPI_SAC_1_2	2086200.0	16305000.0
4DPI_SAC_1_1	3328200.0	20802000.0
10DPI_interface_5_3	7822400.0	28551000.0
10DPI_interface_4_3	5549700.0	25417000.0
10DPI_interface_4_2	3494600.0	20562000.0
10DPI_interface_4_1	3959500.0	14898000.0
10DPI_interface_3_2	10313000.0	21001000.0

10DPI_interface_3_1	5659300.0	17979000.0
10DPI_cortex_4_4	6568100.0	16769000.0
10DPI_cortex_4_3	3419900.0	11998000.0
10DPI_cortex_3_3	2486500.0	10442000.0
10DPI_cortex_3_2	3308800.0	5336900.0
10DPI_SAC_4_3	6796900.0	23906000.0
10DPI_SAC_4_2	7350200.0	19011000.0
10DPI_SAC_4_1	4744200.0	19508000.0
10DPI_SAC_3_3	12653000.0	20440000.0
10DPI_SAC_3_2	11506000.0	19529000.0
10DPI_SAC_3_1	9135600.0	22306000.0
10DPI_SAC_2_3	12044000.0	13540000.0
10DPI_SAC_2_2	11489000.0	15259000.0
10DPI_SAC_2_1	6878200.0	18782000.0

E9QL31;A0A0R4J104;P98078;E9PX84;Q3TRE6;F6TQN9;Q9DCE6;E0CXT5;E0CZ53;E0CYJ2 \

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4DPI_interface_1_1	4963900.0
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4DPI_cortex_3_2	19354000.0
4DPI_cortex_2_3	18882000.0
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4DPI_SAC_3_2	5700800.0
4DPI_SAC_2_3	5403700.0
4DPI_SAC_2_2	4984100.0
4DPI_SAC_2_1	7089200.0
4DPI_SAC_1_3	9872000.0
4DPI_SAC_1_2	7354500.0
4DPI_SAC_1_1	9433100.0
10DPI_interface_5_3	8816700.0
10DPI_interface_4_3	6936800.0
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10DPI_interface_4_1	3713000.0
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10DPI_cortex_3_3	22599000.0
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10DPI_SAC_4_3	5934200.0
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10DPI_SAC_4_1	4308500.0
10DPI_SAC_3_3	7713200.0
10DPI_SAC_3_2	9466900.0
10DPI_SAC_3_1	8250600.0
10DPI_SAC_2_3	6240900.0
10DPI_SAC_2_2	3403200.0
10DPI_SAC_2_1	8101900.0

AOA0R4J195;Q9CSU0;F6YB25;F7DAY5;Q8VDS4 \

index

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4DPI_interface_2_2	959390.0
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4DPI_interface_1_2	1506600.0
4DPI_interface_1_1	1400400.0
4DPI_cortex_3_3	734590.0
4DPI_cortex_3_2	544140.0
4DPI_cortex_2_3	1263200.0
4DPI_cortex_2_2	911490.0
4DPI_cortex_2_1	1163700.0
4DPI_cortex_1_2	857580.0
4DPI_cortex_1_1	813620.0
4DPI_SAC_3_3	2025800.0
4DPI_SAC_3_2	1299600.0
4DPI_SAC_2_3	1140400.0
4DPI_SAC_2_2	1866900.0
4DPI_SAC_2_1	1723900.0
4DPI_SAC_1_3	1446200.0
4DPI_SAC_1_2	2133800.0
4DPI_SAC_1_1	1344000.0
10DPI_interface_5_3	1129300.0
10DPI_interface_4_3	1920700.0
10DPI_interface_4_2	1817400.0
10DPI_interface_4_1	1819300.0
10DPI_interface_3_2	1001900.0
10DPI_interface_3_1	2526900.0
10DPI_cortex_4_4	1237700.0
10DPI_cortex_4_3	2119000.0
10DPI_cortex_3_3	881230.0
10DPI_cortex_3_2	797640.0
10DPI_SAC_4_3	1421000.0
10DPI_SAC_4_2	3697900.0
10DPI_SAC_4_1	2148600.0
10DPI_SAC_3_3	1626400.0
10DPI_SAC_3_2	1196300.0

10DPI_SAC_3_1	819090.0
10DPI_SAC_2_3	2583700.0
10DPI_SAC_2_2	1870900.0
10DPI_SAC_2_1	2521700.0

AOA1BOGQU8;P35980;AOA1BOGSS8;AOA1BOGSF7;AOA1BOGSA8;AOA1BOGS28;AOA1BOGRZ3 \

index

4DPI_interface_2_4	11981000.0
4DPI_interface_2_2	22897000.0
4DPI_interface_2_1	20121000.0
4DPI_interface_1_2	36610000.0
4DPI_interface_1_1	23111000.0
4DPI_cortex_3_3	16177000.0
4DPI_cortex_3_2	17586000.0
4DPI_cortex_2_3	21878000.0
4DPI_cortex_2_2	16561000.0
4DPI_cortex_2_1	23720000.0
4DPI_cortex_1_2	22475000.0
4DPI_cortex_1_1	20512000.0
4DPI_SAC_3_3	16860000.0
4DPI_SAC_3_2	22906000.0
4DPI_SAC_2_3	22223000.0
4DPI_SAC_2_2	26184000.0
4DPI_SAC_2_1	24626000.0
4DPI_SAC_1_3	18432000.0
4DPI_SAC_1_2	15517000.0
4DPI_SAC_1_1	15370000.0
10DPI_interface_5_3	11978000.0
10DPI_interface_4_3	10966000.0
10DPI_interface_4_2	11930000.0
10DPI_interface_4_1	15307000.0
10DPI_interface_3_2	23524000.0
10DPI_interface_3_1	14105000.0
10DPI_cortex_4_4	19951000.0
10DPI_cortex_4_3	10131000.0
10DPI_cortex_3_3	20335000.0
10DPI_cortex_3_2	15715000.0
10DPI_SAC_4_3	16438000.0
10DPI_SAC_4_2	17678000.0
10DPI_SAC_4_1	14233000.0
10DPI_SAC_3_3	13098000.0
10DPI_SAC_3_2	7518100.0
10DPI_SAC_3_1	11445000.0
10DPI_SAC_2_3	13901000.0
10DPI_SAC_2_2	12134000.0
10DPI_SAC_2_1	12099000.0

P06151;Q564E2;AOA1BOGSR9;AOA1BOGSX0;D3YZQ9;AOA1BOGSL7;AOA1B
 OGT41;AOA1BOGQX5;AOA1BOGS79;AOA1BOGRW9;AOA1BOGRC1;AOA1BOGRS2;D3YVR7;D3YZE4;AOA1B
 OGRE9;AOA1BOGSR2;P00342 \

index

4DPI_interface_2_4	48765000.0
4DPI_interface_2_2	84281000.0
4DPI_interface_2_1	62405000.0
4DPI_interface_1_2	61114000.0
4DPI_interface_1_1	36589000.0
4DPI_cortex_3_3	33949000.0
4DPI_cortex_3_2	29980000.0
4DPI_cortex_2_3	62446000.0
4DPI_cortex_2_2	63940000.0
4DPI_cortex_2_1	25525000.0
4DPI_cortex_1_2	45436000.0
4DPI_cortex_1_1	61932000.0
4DPI_SAC_3_3	58681000.0
4DPI_SAC_3_2	56833000.0
4DPI_SAC_2_3	55551000.0
4DPI_SAC_2_2	50517000.0
4DPI_SAC_2_1	84878000.0
4DPI_SAC_1_3	81791000.0
4DPI_SAC_1_2	81333000.0
4DPI_SAC_1_1	59981000.0
10DPI_interface_5_3	38969000.0
10DPI_interface_4_3	61117000.0
10DPI_interface_4_2	41654000.0
10DPI_interface_4_1	71086000.0
10DPI_interface_3_2	51215000.0
10DPI_interface_3_1	34041000.0
10DPI_cortex_4_4	51373000.0
10DPI_cortex_4_3	36612000.0
10DPI_cortex_3_3	38540000.0
10DPI_cortex_3_2	19880000.0
10DPI_SAC_4_3	79881000.0
10DPI_SAC_4_2	92821000.0
10DPI_SAC_4_1	91489000.0
10DPI_SAC_3_3	47923000.0
10DPI_SAC_3_2	87510000.0
10DPI_SAC_3_1	39045000.0
10DPI_SAC_2_3	27205000.0
10DPI_SAC_2_2	50918000.0
10DPI_SAC_2_1	16776000.0

... 36 37 38 \

index

4DPI_interface_2_4 ... 1.031646e+07 -1.178469e+07 2.929428e-07

4DPI_interface_2_2	...	2.282268e+07	-5.412626e+04	2.929428e-07
4DPI_interface_2_1	...	-2.718176e+07	2.199637e+06	2.929428e-07
4DPI_interface_1_2	...	1.065881e+07	-7.706425e+05	2.929428e-07
4DPI_interface_1_1	...	-1.414153e+07	1.404734e+06	2.929428e-07
4DPI_cortex_3_3	...	-3.510812e+06	-1.258512e+07	2.929428e-07
4DPI_cortex_3_2	...	-1.934336e+07	6.848204e+06	2.929428e-07
4DPI_cortex_2_3	...	-2.514064e+07	9.140492e+06	2.929428e-07
4DPI_cortex_2_2	...	4.206068e+07	-1.488583e+07	2.929428e-07
4DPI_cortex_2_1	...	9.972975e+06	1.967906e+06	2.929428e-07
4DPI_cortex_1_2	...	-1.634307e+07	-2.895322e+06	2.929428e-07
4DPI_cortex_1_1	...	6.039433e+06	1.772419e+07	2.929428e-07
4DPI_SAC_3_3	...	1.320228e+07	4.387337e+04	2.929428e-07
4DPI_SAC_3_2	...	-3.905192e+06	2.317435e+07	2.929428e-07
4DPI_SAC_2_3	...	1.960526e+06	-8.147903e+05	2.929428e-07
4DPI_SAC_2_2	...	-5.308582e+06	3.966172e+06	2.929428e-07
4DPI_SAC_2_1	...	-2.123865e+06	-4.008947e+06	2.929428e-07
4DPI_SAC_1_3	...	9.412037e+06	-1.014204e+07	2.929428e-07
4DPI_SAC_1_2	...	-8.900910e+06	6.486090e+06	2.929428e-07
4DPI_SAC_1_1	...	3.946169e+06	-1.159411e+07	2.929428e-07
10DPI_interface_5_3	...	1.110906e+07	9.848782e+06	2.929428e-07
10DPI_interface_4_3	...	-1.415164e+07	-2.113884e+06	2.929428e-07
10DPI_interface_4_2	...	-1.030126e+07	-2.357560e+07	2.929428e-07
10DPI_interface_4_1	...	2.432251e+07	2.044187e+07	2.929428e-07
10DPI_interface_3_2	...	1.381270e+07	-3.013517e+07	2.929428e-07
10DPI_interface_3_1	...	-2.592833e+07	5.400819e+06	2.929428e-07
10DPI_cortex_4_4	...	-4.740456e+06	-7.302414e+06	2.929428e-07
10DPI_cortex_4_3	...	1.500560e+07	2.581753e+06	2.929428e-07
10DPI_cortex_3_3	...	-8.358536e+06	3.333076e+06	2.929428e-07
10DPI_cortex_3_2	...	4.094822e+06	-3.480078e+06	2.929428e-07
10DPI_SAC_4_3	...	6.421815e+06	1.283613e+07	2.929428e-07
10DPI_SAC_4_2	...	4.099798e+06	-4.146932e+06	2.929428e-07
10DPI_SAC_4_1	...	-1.157357e+07	-3.995583e+06	2.929428e-07
10DPI_SAC_3_3	...	-2.697074e+07	-3.500647e+06	2.929428e-07
10DPI_SAC_3_2	...	8.463252e+04	-1.427429e+07	2.929428e-07
10DPI_SAC_3_1	...	1.863299e+07	4.169473e+07	2.929428e-07
10DPI_SAC_2_3	...	-5.895312e+06	-4.218407e+06	2.929428e-07
10DPI_SAC_2_2	...	-3.574784e+06	3.894834e+06	2.929428e-07
10DPI_SAC_2_1	...	9.418379e+06	-6.709035e+06	2.929428e-07

	Segment	K-means	PCA	TimePoint	BioRep	TechRep	\
index							
4DPI_interface_2_4		1		4DPI	2	4	
4DPI_interface_2_2		1		4DPI	2	2	
4DPI_interface_2_1		1		4DPI	2	1	
4DPI_interface_1_2		3		4DPI	1	2	
4DPI_interface_1_1		3		4DPI	1	1	
4DPI_cortex_3_3		3		4DPI	3	3	

4DPI_cortex_3_2	3	4DPI	3	2
4DPI_cortex_2_3	2	4DPI	2	3
4DPI_cortex_2_2	2	4DPI	2	2
4DPI_cortex_2_1	3	4DPI	2	1
4DPI_cortex_1_2	2	4DPI	1	2
4DPI_cortex_1_1	2	4DPI	1	1
4DPI_SAC_3_3	1	4DPI	3	3
4DPI_SAC_3_2	1	4DPI	3	2
4DPI_SAC_2_3	1	4DPI	2	3
4DPI_SAC_2_2	1	4DPI	2	2
4DPI_SAC_2_1	1	4DPI	2	1
4DPI_SAC_1_3	1	4DPI	1	3
4DPI_SAC_1_2	1	4DPI	1	2
4DPI_SAC_1_1	1	4DPI	1	1
10DPI_interface_5_3	0	10DPI	5	3
10DPI_interface_4_3	0	10DPI	4	3
10DPI_interface_4_2	0	10DPI	4	2
10DPI_interface_4_1	0	10DPI	4	1
10DPI_interface_3_2	1	10DPI	3	2
10DPI_interface_3_1	0	10DPI	3	1
10DPI_cortex_4_4	3	10DPI	4	4
10DPI_cortex_4_3	3	10DPI	4	3
10DPI_cortex_3_3	3	10DPI	3	3
10DPI_cortex_3_2	2	10DPI	3	2
10DPI_SAC_4_3	1	10DPI	4	3
10DPI_SAC_4_2	1	10DPI	4	2
10DPI_SAC_4_1	1	10DPI	4	1
10DPI_SAC_3_3	0	10DPI	3	3
10DPI_SAC_3_2	0	10DPI	3	2
10DPI_SAC_3_1	0	10DPI	3	1
10DPI_SAC_2_3	1	10DPI	2	3
10DPI_SAC_2_2	1	10DPI	2	2
10DPI_SAC_2_1	1	10DPI	2	1

	Region	DPI_Region	BioRep_TechRep
index			
4DPI_interface_2_4	interface	4DPI_interface	2_4
4DPI_interface_2_2	interface	4DPI_interface	2_2
4DPI_interface_2_1	interface	4DPI_interface	2_1
4DPI_interface_1_2	interface	4DPI_interface	1_2
4DPI_interface_1_1	interface	4DPI_interface	1_1
4DPI_cortex_3_3	cortex	4DPI_cortex	3_3
4DPI_cortex_3_2	cortex	4DPI_cortex	3_2
4DPI_cortex_2_3	cortex	4DPI_cortex	2_3
4DPI_cortex_2_2	cortex	4DPI_cortex	2_2
4DPI_cortex_2_1	cortex	4DPI_cortex	2_1
4DPI_cortex_1_2	cortex	4DPI_cortex	1_2

4DPI_cortex_1_1	cortex	4DPI_cortex	1_1
4DPI_SAC_3_3	SAC	4DPI_SAC	3_3
4DPI_SAC_3_2	SAC	4DPI_SAC	3_2
4DPI_SAC_2_3	SAC	4DPI_SAC	2_3
4DPI_SAC_2_2	SAC	4DPI_SAC	2_2
4DPI_SAC_2_1	SAC	4DPI_SAC	2_1
4DPI_SAC_1_3	SAC	4DPI_SAC	1_3
4DPI_SAC_1_2	SAC	4DPI_SAC	1_2
4DPI_SAC_1_1	SAC	4DPI_SAC	1_1
10DPI_interface_5_3	interface	10DPI_interface	5_3
10DPI_interface_4_3	interface	10DPI_interface	4_3
10DPI_interface_4_2	interface	10DPI_interface	4_2
10DPI_interface_4_1	interface	10DPI_interface	4_1
10DPI_interface_3_2	interface	10DPI_interface	3_2
10DPI_interface_3_1	interface	10DPI_interface	3_1
10DPI_cortex_4_4	cortex	10DPI_cortex	4_4
10DPI_cortex_4_3	cortex	10DPI_cortex	4_3
10DPI_cortex_3_3	cortex	10DPI_cortex	3_3
10DPI_cortex_3_2	cortex	10DPI_cortex	3_2
10DPI_SAC_4_3	SAC	10DPI_SAC	4_3
10DPI_SAC_4_2	SAC	10DPI_SAC	4_2
10DPI_SAC_4_1	SAC	10DPI_SAC	4_1
10DPI_SAC_3_3	SAC	10DPI_SAC	3_3
10DPI_SAC_3_2	SAC	10DPI_SAC	3_2
10DPI_SAC_3_1	SAC	10DPI_SAC	3_1
10DPI_SAC_2_3	SAC	10DPI_SAC	2_3
10DPI_SAC_2_2	SAC	10DPI_SAC	2_2
10DPI_SAC_2_1	SAC	10DPI_SAC	2_1

[39 rows x 333 columns]

```
[198]: #append k cluster ID to each sample
df3['k'] = kmeans['Segment K-means PCA']
df3
```

```
[198]: Protein IDs      A8DUK4      P63260      Q91VB8      P15864      P20152  \
index
4DPI_interface_2_4 -0.540090  0.021508 -0.022581 -0.236924 -0.124103
4DPI_interface_2_2  0.019054 -0.337894  0.113558 -0.642223 -0.213255
4DPI_interface_2_1 -0.138874 -0.540108 -0.215675 -0.984486 -0.400444
4DPI_interface_1_2  0.608371 -1.105209 -0.325806 -0.810748 -0.569122
4DPI_interface_1_1  0.174370 -1.183182  0.094234 -0.679225 -0.517408
4DPI_cortex_3_3     0.812077 -1.342470  0.739405 -1.104028 -1.326299
4DPI_cortex_3_2     1.171869 -1.082481  0.131356 -1.092429 -1.250061
4DPI_cortex_2_3     1.732487 -1.189867  0.205528 -1.029867 -1.103115
4DPI_cortex_2_2     1.705228 -0.907770  0.786771 -1.080988 -1.122011
4DPI_cortex_2_1     0.017949 -1.231981  0.234514 -0.996560 -1.377694
4DPI_cortex_1_2     3.162245 -0.851188  2.487419 -0.892101 -1.028298
```


4DPI_cortex_1_1	2.284197	-1.246783	2.586436	-1.133441	-1.165106
4DPI_SAC_3_3	-0.653982	0.011098	-0.271394	1.321765	0.137459
4DPI_SAC_3_2	-0.555428	0.006037	-0.709916	0.867429	0.003198
4DPI_SAC_2_3	-0.512329	-0.250133	-0.485555	-0.434520	-0.619947
4DPI_SAC_2_2	-0.480516	-0.370315	-0.212042	-0.525994	-0.592846
4DPI_SAC_2_1	-0.371781	0.039127	-0.427148	0.498867	0.228388
4DPI_SAC_1_3	0.162783	0.504720	0.867335	-0.745891	-0.339045
4DPI_SAC_1_2	-0.281163	0.401154	2.496209	-0.308182	-0.292751
4DPI_SAC_1_1	-0.043099	0.343665	0.096850	-0.764444	-0.312506
10DPI_interface_5_3	-0.777652	1.419721	-0.753751	0.824938	1.692848
10DPI_interface_4_3	-0.866100	1.606369	-1.090698	1.305138	1.805398
10DPI_interface_4_2	-0.873578	2.688393	-1.119343	0.698258	1.568746
10DPI_interface_4_1	-0.800096	1.695420	-0.998278	1.009681	1.820800
10DPI_interface_3_2	-0.632316	0.429326	-0.690294	0.874819	1.296107
10DPI_interface_3_1	-0.719082	1.182173	-0.586279	0.609317	1.189421
10DPI_cortex_4_4	0.249718	-0.833378	-0.002022	-0.349340	-0.885863
10DPI_cortex_4_3	1.169458	-1.045285	0.320527	-0.289708	-0.986684
10DPI_cortex_3_3	0.492771	-1.152337	0.040476	-1.045689	-1.388084
10DPI_cortex_3_2	1.595690	-1.288228	2.579389	-0.947973	-1.307077
10DPI_SAC_4_3	-0.946477	0.383822	-1.059250	-0.383768	0.631378
10DPI_SAC_4_2	-0.805390	0.246927	-0.955010	0.293407	1.028207
10DPI_SAC_4_1	-0.915585	0.379811	-1.000268	0.264904	0.442797
10DPI_SAC_3_3	-0.837421	1.047284	-0.539895	1.228074	1.220225
10DPI_SAC_3_2	-0.835097	1.150373	-0.679543	0.438430	0.917108
10DPI_SAC_3_1	-0.814382	1.447415	-0.838558	0.712641	0.795435
10DPI_SAC_2_3	-0.756522	0.301217	-0.319995	3.323850	0.472652
10DPI_SAC_2_2	-0.702037	0.341660	0.175307	0.376145	1.252213
10DPI_SAC_2_1	-0.499269	0.311387	-0.652017	1.830863	0.419339

Protein IDs	Q03265	Q64475	P27661	P43276	P31786	...	\
index						...	
4DPI_interface_2_4	-0.225207	0.487461	0.304723	-0.028911	0.134433	...	
4DPI_interface_2_2	0.089789	-0.433295	-0.113263	-0.522751	-0.504536	...	
4DPI_interface_2_1	-0.369529	-0.155440	-0.818871	-1.047645	-1.302654	...	
4DPI_interface_1_2	-0.241213	0.125845	-0.007414	0.002511	-1.015394	...	
4DPI_interface_1_1	0.168928	-1.043352	-0.340969	-0.393190	-0.514812	...	
4DPI_cortex_3_3	1.809980	-1.645795	-1.310597	-1.351087	0.551226	...	
4DPI_cortex_3_2	1.360204	-1.052557	-1.656457	-1.324128	0.710533	...	
4DPI_cortex_2_3	2.563874	-1.384593	-0.422396	-1.128870	2.363262	...	
4DPI_cortex_2_2	1.560213	-1.012349	-1.323757	-0.949139	1.326104	...	
4DPI_cortex_2_1	2.644484	-1.569062	-1.415339	-1.385625	-0.472758	...	
4DPI_cortex_1_2	0.906860	-0.699643	-0.763131	-0.753384	1.490523	...	
4DPI_cortex_1_1	0.987024	-1.281794	-0.206395	-1.246260	0.739623	...	
4DPI_SAC_3_3	-0.525400	1.972168	2.332589	0.926510	-0.777197	...	
4DPI_SAC_3_2	-0.485897	0.522564	0.658175	0.951647	-0.857984	...	
4DPI_SAC_2_3	-0.125960	0.362906	-0.387693	0.358201	1.684032	...	
4DPI_SAC_2_2	-0.046063	1.084758	0.075848	0.243042	1.603192	...	

4DPI_SAC_2_1	-0.187086	-0.117659	0.580671	0.241366	-0.301172	...
4DPI_SAC_1_3	-0.217939	-0.801061	1.767597	-0.877708	0.019972	...
4DPI_SAC_1_2	-0.500477	-0.780309	1.001093	-0.349095	0.899931	...
4DPI_SAC_1_1	-0.382281	-0.266356	0.663553	-0.823244	0.018918	...
10DPI_interface_5_3	-0.901580	1.013422	-0.382157	0.127778	-0.526985	...
10DPI_interface_4_3	-0.827514	1.870624	0.372136	1.542423	-0.595809	...
10DPI_interface_4_2	-0.759250	1.245463	0.559159	0.560974	-0.991206	...
10DPI_interface_4_1	-0.808740	0.584194	0.445623	1.316398	-0.703630	...
10DPI_interface_3_2	-0.899163	1.029865	0.112986	0.569039	-0.266707	...
10DPI_interface_3_1	-0.726168	0.444367	0.102009	0.726513	-0.385806	...
10DPI_cortex_4_4	0.463414	-0.368067	0.051774	-0.178477	-0.821938	...
10DPI_cortex_4_3	0.495828	-0.910387	0.338825	-0.099504	0.317561	...
10DPI_cortex_3_3	0.805517	-1.033478	-1.298924	-1.414690	-0.097914	...
10DPI_cortex_3_2	1.836598	-1.234055	-1.243058	-1.222343	2.401152	...
10DPI_SAC_4_3	-0.887629	-0.012392	-0.042908	-0.335688	-1.490924	...
10DPI_SAC_4_2	-0.928505	-0.264724	0.373718	0.088606	-1.390982	...
10DPI_SAC_4_1	-0.804905	0.156136	-0.493826	-0.204923	-1.472079	...
10DPI_SAC_3_3	-0.772893	0.738915	0.963448	1.472510	-0.798119	...
10DPI_SAC_3_2	-0.810746	0.665529	0.529107	0.355426	-0.462956	...
10DPI_SAC_3_1	-0.729155	1.164043	0.115897	0.340239	-0.021449	...
10DPI_SAC_2_3	-0.941055	2.246173	2.216491	2.602791	0.750162	...
10DPI_SAC_2_2	-0.734416	0.680466	1.143132	0.986525	0.257537	...
10DPI_SAC_2_1	-0.853945	-0.328529	-2.481400	2.224163	0.504851	...

Protein IDs index	Q9DCX2	A0A1L1SV25	F8WIP8	P10126	P48036	\
4DPI_interface_2_4	-0.334925	0.018748	0.107914	1.212955	-0.024608	
4DPI_interface_2_2	0.321537	1.474015	0.968526	-0.854949	-0.503046	
4DPI_interface_2_1	0.310361	0.307097	-0.023676	-0.441144	0.481258	
4DPI_interface_1_2	-0.812839	0.199176	1.497482	1.173070	-0.049537	
4DPI_interface_1_1	0.292111	1.774868	0.328190	-1.647993	0.017034	
4DPI_cortex_3_3	2.398939	0.720059	-0.229218	-0.556935	-2.266719	
4DPI_cortex_3_2	1.324569	0.731844	0.454642	0.938331	-1.426552	
4DPI_cortex_2_3	2.349517	1.431181	2.344207	-0.239975	-0.526365	
4DPI_cortex_2_2	2.138293	2.043236	3.298078	-0.636912	-0.625806	
4DPI_cortex_2_1	0.759878	-0.269183	-0.294759	-1.810814	-1.659913	
4DPI_cortex_1_2	1.148796	-0.013677	2.515370	0.541643	-1.030787	
4DPI_cortex_1_1	1.394336	2.132254	0.542312	0.812029	-0.595879	
4DPI_SAC_3_3	-0.350119	-0.709544	0.106389	-0.141509	-1.048333	
4DPI_SAC_3_2	-0.412290	-0.747891	0.087704	-1.318569	-0.713698	
4DPI_SAC_2_3	0.036965	-1.984384	0.284609	-0.330962	0.411022	
4DPI_SAC_2_2	-0.281836	-1.038753	-0.296284	0.070711	-0.711533	
4DPI_SAC_2_1	-0.096370	0.178716	0.518885	-0.455685	-0.634745	
4DPI_SAC_1_3	-0.479700	0.296030	0.670741	0.446418	-0.324874	
4DPI_SAC_1_2	-0.367495	-0.237596	-0.092830	0.151312	-0.207055	
4DPI_SAC_1_1	-0.592533	-0.200805	0.315036	1.908031	-0.554571	
10DPI_interface_5_3	-0.907178	-0.050588	-0.081766	2.083774	0.992343	

10DPI_interface_4_3	-1.111181	0.140489	-1.168785	0.730058	0.235904
10DPI_interface_4_2	-0.882729	0.063615	-1.169293	0.702471	0.341730
10DPI_interface_4_1	-0.728393	-0.880639	-1.214748	-0.650622	0.772141
10DPI_interface_3_2	-0.491401	-1.009320	-0.610496	0.617134	1.074627
10DPI_interface_3_1	-0.893251	-0.182499	-0.552520	1.172655	1.524249
10DPI_cortex_4_4	0.694825	1.663895	-0.550092	0.606789	1.000782
10DPI_cortex_4_3	-0.194255	0.216405	-0.609141	-1.383216	2.065094
10DPI_cortex_3_3	1.912836	0.919092	-1.265154	-1.059899	-0.488110
10DPI_cortex_3_2	0.966998	-0.815013	-1.216605	-1.192724	-1.438267
10DPI_SAC_4_3	-0.724979	0.791728	-0.444244	-1.382800	1.144086
10DPI_SAC_4_2	-0.875848	-0.514998	-0.913915	-0.916771	1.207437
10DPI_SAC_4_1	-0.870530	-0.130691	-0.379494	-0.759516	0.455384
10DPI_SAC_3_3	-0.625610	-1.035522	-0.322308	0.767741	0.667758
10DPI_SAC_3_2	-1.125903	-0.712475	-0.495334	1.261565	2.200236
10DPI_SAC_3_1	-0.839165	-0.457866	-0.619811	1.450602	1.276285
10DPI_SAC_2_3	-0.551101	-1.993058	-0.614956	-0.233784	-0.034713
10DPI_SAC_2_2	-0.616799	-0.699494	-0.551729	0.151312	-0.781103
10DPI_SAC_2_1	-0.883532	-1.418453	-0.322929	-0.783821	-0.221158

Protein IDs	Q8QZT1	E9PZF0	F8WIT2	P09405	k
index					
4DPI_interface_2_4	-0.362371	-0.030764	0.333282	1.597443	1
4DPI_interface_2_2	0.048106	1.395860	-0.255785	0.741501	1
4DPI_interface_2_1	-0.081927	-0.328686	1.400135	1.304200	1
4DPI_interface_1_2	-0.658481	-1.312265	-1.284947	-0.635158	3
4DPI_interface_1_1	-0.435981	-0.513049	-1.074569	0.427190	3
4DPI_cortex_3_3	2.883655	2.183097	-1.269124	-1.883931	3
4DPI_cortex_3_2	1.466451	1.820325	-1.060759	-1.802613	3
4DPI_cortex_2_3	1.501583	-0.214122	-0.559851	-0.100214	2
4DPI_cortex_2_2	2.321853	0.980506	-0.462015	-0.736300	2
4DPI_cortex_2_1	2.616062	1.153391	-1.214699	-2.545370	3
4DPI_cortex_1_2	0.707471	1.691200	-0.514503	-0.741976	2
4DPI_cortex_1_1	0.885410	2.194431	-0.125894	-0.369901	2
4DPI_SAC_3_3	-0.082839	-0.089157	-0.533417	1.397490	1
4DPI_SAC_3_2	-0.405868	-0.592748	-0.343139	0.263853	1
4DPI_SAC_2_3	-0.084284	0.319239	-0.942338	-1.200577	1
4DPI_SAC_2_2	0.207415	-0.308169	-0.712401	0.020676	1
4DPI_SAC_2_1	-0.365261	0.010556	-0.392664	1.233921	1
4DPI_SAC_1_3	0.654773	0.692699	0.203771	0.307000	1
4DPI_SAC_1_2	-0.169680	-0.464053	-0.223654	-0.654438	1
4DPI_SAC_1_1	0.140422	0.212710	0.059371	0.638882	1
10DPI_interface_5_3	-0.835014	-0.679406	2.108762	1.231277	0
10DPI_interface_4_3	-0.576910	-0.186289	1.100246	1.105335	0
10DPI_interface_4_2	-0.686738	-0.772089	0.692647	0.991598	0
10DPI_interface_4_1	-0.715809	-1.494046	0.663326	1.152602	0
10DPI_interface_3_2	-0.620855	-1.031775	1.966034	-0.021150	1
10DPI_interface_3_1	-0.677940	-1.059968	1.545065	0.638104	0

10DPI_cortex_4_4	-0.770248	-0.080477	-0.645693	-0.976057	3
10DPI_cortex_4_3	0.627398	0.954609	-0.752614	-1.150666	3
10DPI_cortex_3_3	0.434782	0.854680	-0.998138	-1.193425	3
10DPI_cortex_3_2	0.577210	0.764507	-1.241369	-1.301098	2
10DPI_SAC_4_3	-0.809456	0.528709	0.742096	-0.249867	1
10DPI_SAC_4_2	-1.006581	-0.029186	0.509432	0.039178	1
10DPI_SAC_4_1	-0.720509	-0.101711	0.399519	0.054649	1
10DPI_SAC_3_3	-0.846876	-1.050714	0.788812	0.411175	0
10DPI_SAC_3_2	-1.244334	-1.430989	2.056198	-0.072382	0
10DPI_SAC_3_1	-0.544820	-1.215851	1.831509	0.825852	0
10DPI_SAC_2_3	-0.763123	-1.398708	-0.556053	-0.325977	1
10DPI_SAC_2_2	-0.509742	-0.642748	-0.175040	0.946585	1
10DPI_SAC_2_1	-1.096942	-0.729549	-1.061534	0.632585	1

[39 rows x 101 columns]

```
[199]: #change k to categorical value
df3['k'] = df3['k'].astype('category')
df3.dtypes
```

```
[199]: Protein IDs
A8DUK4      float64
P63260      float64
Q91VB8      float64
P15864      float64
P20152      float64
...
Q8QZT1      float64
E9PZF0      float64
F8WIT2      float64
P09405      float64
k           category
Length: 101, dtype: object
```

```
[200]: #calculate mean for each cluster
df4 = df3.groupby(['k']).mean()
df4
```

```
[200]: Protein IDs      A8DUK4      P63260      Q91VB8      P15864      P20152      Q03265 \
k
0          -0.815426      1.529644     -0.825793      0.853310      1.376248     -0.792006
1          -0.480723      0.123501     -0.181771      0.256979      0.167602     -0.501426
2           2.095969     -1.096767      1.729109     -1.016874     -1.145122      1.570914
3           0.587073     -1.122040      0.154086     -0.795966     -1.037652      0.938393

Protein IDs      Q64475      P27661      P43276      P31786      ...      P61979      Q9DCX2 \
k
0           0.965820      0.338153      0.805282     -0.560745      ...      0.535881     -0.889176
1           0.299041      0.382923      0.277885     -0.138400      ...      0.359382     -0.403366
```

```

2          -1.122487 -0.791747 -1.059999  1.664133  ... -0.498346  1.599588
3          -0.937107 -0.704888 -0.768024 -0.167937  ... -1.033024  0.797008

```

```

Protein IDs  A0A1L1SV25    F8WIP8    P10126    P48036    Q8QZT1    E9PZF0  \
k
0          -0.389436 -0.703070  0.939780  1.001331 -0.766055 -0.986169
1          -0.423258 -0.066153 -0.170091 -0.054757 -0.334929 -0.143749
2           0.955596  1.496673 -0.143188 -0.843421  1.198705  1.083304
3           0.744519 -0.083506 -0.467583 -0.350990  0.770455  0.632539

```

```

Protein IDs    F8WIT2    P09405
k
0           1.348321  0.785445
1           0.023201  0.373664
2          -0.580727 -0.649898
3          -1.037568 -1.220004

```

[4 rows x 100 columns]

```

[201]: #select proteins with values greater than 0 for each cluster; these will be
        ↪used for GO analysis
df5 = df4[df4>0]
df5

```

```

[201]: Protein IDs    A8DUK4    P63260    Q91VB8    P15864    P20152    Q03265  \
k
0           NaN  1.529644      NaN  0.853310  1.376248      NaN
1           NaN  0.123501      NaN  0.256979  0.167602      NaN
2          2.095969      NaN  1.729109      NaN      NaN  1.570914
3          0.587073      NaN  0.154086      NaN      NaN  0.938393

```

```

Protein IDs    Q64475    P27661    P43276    P31786  ...    P61979    Q9DCX2  \
k          ...
0          0.965820  0.338153  0.805282      NaN  ...  0.535881      NaN
1          0.299041  0.382923  0.277885      NaN  ...  0.359382      NaN
2           NaN      NaN      NaN  1.664133  ...      NaN  1.599588
3           NaN      NaN      NaN      NaN  ...      NaN  0.797008

```

```

Protein IDs  A0A1L1SV25    F8WIP8    P10126    P48036    Q8QZT1    E9PZF0  \
k
0           NaN      NaN  0.93978  1.001331      NaN      NaN
1           NaN      NaN      NaN      NaN      NaN      NaN
2          0.955596  1.496673      NaN      NaN  1.198705  1.083304
3          0.744519      NaN      NaN      NaN  0.770455  0.632539

```

```

Protein IDs    F8WIT2    P09405
k
0           1.348321  0.785445

```

1	0.023201	0.373664
2	NaN	NaN
3	NaN	NaN

[4 rows x 100 columns]

[202]: *#make list of proteins with values greater than 0 for each cluster*

```
proteinIDs = []

for i in range(0,4):
    proteinList = df5.iloc[i][-df5.iloc[i].isnull()].index
    proteinIDs.append(proteinList)

proteinIDs
```

[202]: [Index(['P63260', 'P15864', 'P20152', 'Q64475', 'P27661', 'P43276', 'P20065',
'P52480', 'Q8VDD5', 'B7FAV1', 'P43274', 'O35744', 'P62962', 'P26041',
'P43277', 'P62806', 'P05064', 'P43275', 'P09528', 'P27773', 'Q61233',
'P20029', 'P63017', 'P07356', 'P26039', 'P18760', 'D3Z2H9', 'O88569',
'P17742', 'AOA1D5RLD8', 'AOA1W2P768', 'P11499', 'Q99PT1', 'P40124',
'E9Q7Q3', 'P40142', 'E9Q3W4', 'Q91Z25', 'Q6IRU2', 'P13020', 'A2AL12',
'P63101', 'P14733', 'P06745', 'P17182', 'Q9DCD0', 'P05213', 'P26645',
'Q9JKF1', 'P08113', 'AOA1W2P6F6', 'P09103', 'Q61029', 'P99024',
'O35639', 'P61979', 'P10126', 'P48036', 'F8WIT2', 'P09405'],
dtype='object', name=' Protein IDs'),
Index(['P63260', 'P15864', 'P20152', 'Q64475', 'P27661', 'P43276', 'P20065',
'P52480', 'Q8VDD5', 'E9PV24', 'B7FAV1', 'P43274', 'O35744', 'P62962',
'P26041', 'P43277', 'P62806', 'P05064', 'P43275', 'P26350', 'P09528',
'P27773', 'Q61233', 'P20029', 'P63017', 'P07356', 'P26039', 'P18760',
'D3Z2H9', 'O88569', 'P17742', 'AOA1D5RLD8', 'P11499', 'Q99PT1',
'P40124', 'E9Q7Q3', 'P40142', 'E9Q3W4', 'Q91Z25', 'Q6IRU2', 'P13020',
'A2AL12', 'E9Q616', 'P63101', 'P14733', 'P06745', 'P17182', 'Q9DCD0',
'P05213', 'P26645', 'Q9JKF1', 'P08113', 'AOA1W2P6F6', 'P09103',
'Q61029', 'Q921I1', 'P99024', 'O35639', 'AOA0R4JOI1', 'P61979',
'F8WIT2', 'P09405'],
dtype='object', name=' Protein IDs'),
Index(['A8DUK4', 'Q91VB8', 'Q03265', 'P31786', 'P56480', 'P16125', 'Q8VDN2',
'Q9JII6', 'Q99KI0', 'P63038', 'Q9DCW4', 'Q64433', 'P26443', 'P17742',
'P08249', 'Q9Z2I8', 'Q61838', 'P70441', 'P24270', 'P05213', 'P38647',
'Q8BWT1', 'P45952', 'P99029', 'A3KGU5', 'O08749', 'P17751', 'P05202',
'Q921I1', 'Q62433', 'P14152', 'AOA0R4JOI1', 'Q60932', 'Q9DCX2',
'AOA1L1SV25', 'F8WIP8', 'Q8QZT1', 'E9PZF0'],
dtype='object', name=' Protein IDs'),
Index(['A8DUK4', 'Q91VB8', 'Q03265', 'P56480', 'P16125', 'Q8VDN2', 'Q9JII6',
'Q99KI0', 'P63038', 'Q9DCW4', 'Q64433', 'P26443', 'P08249', 'Q9Z2I8',
'Q61838', 'P70441', 'P24270', 'P38647', 'Q8BWT1', 'P45952', 'A3KGU5',
'P10922', 'O08749', 'P17751', 'P05202', 'Q921I1', 'Q62433', 'P14152',
'Q60932', 'Q9DCX2', 'AOA1L1SV25', 'Q8QZT1', 'E9PZF0'],

```
dtype='object', name=' Protein IDs')]
```

```
[203]: #save protein list as df
proteinIDs = pd.DataFrame(proteinIDs)
proteinIDs.to_csv('outs/' + date + '_topProteins_100.csv')
```

0.4.2 Run proteins from line above in PANTHER and generate gene ontology classifications, then proceed with code below to combine GO information per cluster into one dataframe

```
[205]: #combine individual GO results from PANTHER into a single df

path = 'outs/'

fileNames = ['/*proteinClass_100_noImputation.txt']
geneOntologyTerm = ['ProteinClass']

for file, go in zip(fileNames, geneOntologyTerm):

    files = glob.glob(path + file)

    li = []
    clusterID = []

    for f in files:
        c = f.split('/')[1].split('_')[1]
        clusterID.append(c)

    for filename, c in zip(files, clusterID):
        df = pd.read_csv(filename, index_col = None, header = None, sep = '\t')
        df['cluster'] = str(c)
        li.append(df)

    finalDf = pd.concat(li, axis=0, ignore_index=True).sort_values('cluster')
    finalDf.columns = ['idx', 'Category name (Accession)', '# genes', 'Percent of_
→gene hit against total # genes', 'Percent of gene hit against total #_
→Protein Class hits', 'cluster']

    finalDf.to_csv('outs/' + date + '_panther_combined_gene_ontology_' +_
→str(go) + '_100_noImputation.csv')
```

```
[206]: #extract and save gene counts in a single df

geneOntologyTerm = ['ProteinClass']

for go in geneOntologyTerm:
    df = pd.read_csv('outs/panther_combined_gene_ontology_' + str(go) +_
→'_100_noImputation.csv', index_col = 0)
```

```

df = df[['Category name (Accession)', '# genes', 'cluster']]
df = df.pivot(index = 'Category name (Accession)', columns = 'cluster')
df['# genes'].to_csv('outs/' + date + '_numberGenes_' + str(go) +
→ '_100_noImputation.csv')

```

0.5 Extract LFQ intensity data for top proteins based on highest k-means centroid values

```

[208]: summed = allProt.abs().sum(axis=1)
sorted_mylist = sorted(((v, i) for i, v in enumerate(summed)), reverse=True)[0:
→ 29]
final = [lis[1] for lis in sorted_mylist]
finalID = allProt.iloc[final].index
finalID

#get LFQ intensity data for top proteins based on highest k-means centroid
→ values
df = lfq.loc[finalID]
df.to_csv('outs/' + date + '_topProteinLFQ.csv')

```

```
[ ]:
```