

Student number: 40462364

Name: Imama Jawad

Course: ECS805

Date: 28/03/2024

By submitting the work, I declare that:

1. I have read and understood the University regulations relating to academic offences, including collusion and plagiarism:
<http://www.qub.ac.uk/directorates/AcademicStudentAffairs/AcademicAffairs/GeneralRegulations/Procedures/ProceduresforDealingwithAcademicOffences/>
2. The submission is my own original work and no part of it has been submitted for any other assignments, except as otherwise permitted.
3. All sources used, published or unpublished, have been acknowledged.
4. I give my consent for the work to be scanned using a plagiarism detection software.

Library Imports

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.decomposition import PCA, NMF, KernelPCA
from sklearn.manifold import TSNE
from sklearn.preprocessing import LabelEncoder, StandardScaler,
MinMaxScaler
from sklearn.model_selection import train_test_split, GridSearchCV
from sklearn.ensemble import RandomForestClassifier,
BaggingClassifier, RandomForestRegressor
from sklearn.svm import SVC
from sklearn.linear_model import LogisticRegression, BayesianRidge
from sklearn.metrics import accuracy_score, confusion_matrix,
classification_report
from xgboost import XGBClassifier
from sklearn.datasets import load_iris
from sklearn.experimental import enable_iterative_imputer
from sklearn.impute import IterativeImputer
from mpl_toolkits.mplot3d import Axes3D
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense, Dropout
from tensorflow.keras.optimizers import Adam
from collections import defaultdict
```

Install additional libraries (if not already installed)

```
!pip install sweetviz ydata-profiling
```

```
# Comprehensive EDA Report
```

```
from ydata_profiling import ProfileReport
```

```
Requirement already satisfied: sweetviz in  
/usr/local/lib/python3.10/dist-packages (2.3.1)  
Requirement already satisfied: pandas!=1.0.0,!=1.0.1,!=1.0.2,>=0.25.3  
in /usr/local/lib/python3.10/dist-packages (from sweetviz) (2.2.3)  
Requirement already satisfied: numpy>=1.16.0 in  
/usr/local/lib/python3.10/dist-packages (from sweetviz) (1.26.4)  
Requirement already satisfied: matplotlib>=3.1.3 in  
/usr/local/lib/python3.10/dist-packages (from sweetviz) (3.7.5)  
Requirement already satisfied: tqdm>=4.43.0 in  
/usr/local/lib/python3.10/dist-packages (from sweetviz) (4.67.1)  
Requirement already satisfied: scipy>=1.3.2 in  
/usr/local/lib/python3.10/dist-packages (from sweetviz) (1.13.1)  
Requirement already satisfied: jinja2>=2.11.1 in  
/usr/local/lib/python3.10/dist-packages (from sweetviz) (3.1.4)  
Requirement already satisfied: importlib-resources>=1.2.0 in  
/usr/local/lib/python3.10/dist-packages (from sweetviz) (5.13.0)  
Requirement already satisfied: MarkupSafe>=2.0 in  
/usr/local/lib/python3.10/dist-packages (from jinja2>=2.11.1-  
>sweetviz) (3.0.2)  
Requirement already satisfied: contourpy>=1.0.1 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (1.3.1)  
Requirement already satisfied: cycler>=0.10 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (0.12.1)  
Requirement already satisfied: fonttools>=4.22.0 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (4.55.3)  
Requirement already satisfied: kiwisolver>=1.0.1 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (1.4.7)  
Requirement already satisfied: packaging>=20.0 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (24.2)  
Requirement already satisfied: pillow>=6.2.0 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (11.0.0)  
Requirement already satisfied: pyparsing>=2.3.1 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (3.2.0)  
Requirement already satisfied: python-dateutil>=2.7 in  
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1.3-  
>sweetviz) (2.9.0.post0)  
Requirement already satisfied: mkl_fft in  
/usr/local/lib/python3.10/dist-packages (from numpy>=1.16.0->sweetviz)
```

(1.3.8)
Requirement already satisfied: mkl_random in
/usr/local/lib/python3.10/dist-packages (from numpy>=1.16.0->sweetviz)
(1.2.4)
Requirement already satisfied: mkl_umath in
/usr/local/lib/python3.10/dist-packages (from numpy>=1.16.0->sweetviz)
(0.1.1)
Requirement already satisfied: mkl in /usr/local/lib/python3.10/dist-
packages (from numpy>=1.16.0->sweetviz) (2025.0.1)
Requirement already satisfied: tbb4py in
/usr/local/lib/python3.10/dist-packages (from numpy>=1.16.0->sweetviz)
(2022.0.0)
Requirement already satisfied: mkl-service in
/usr/local/lib/python3.10/dist-packages (from numpy>=1.16.0->sweetviz)
(2.4.1)
Requirement already satisfied: pytz>=2020.1 in
/usr/local/lib/python3.10/dist-packages (from pandas!=1.0.0,! =1.0.1,!
=1.0.2,>=0.25.3->sweetviz) (2025.1)
Requirement already satisfied: tzdata>=2022.7 in
/usr/local/lib/python3.10/dist-packages (from pandas!=1.0.0,! =1.0.1,!
=1.0.2,>=0.25.3->sweetviz) (2025.1)
Requirement already satisfied: six>=1.5 in
/usr/local/lib/python3.10/dist-packages (from python-dateutil>=2.7-
>matplotlib>=3.1.3->sweetviz) (1.17.0)
Requirement already satisfied: intel-openmp>=2024 in
/usr/local/lib/python3.10/dist-packages (from mkl->numpy>=1.16.0-
>sweetviz) (2024.2.0)
Requirement already satisfied: tbb==2022.* in
/usr/local/lib/python3.10/dist-packages (from mkl->numpy>=1.16.0-
>sweetviz) (2022.0.0)
Requirement already satisfied: tcmlib==1.* in
/usr/local/lib/python3.10/dist-packages (from tbb==2022.*->mkl-
>numpy>=1.16.0->sweetviz) (1.2.0)
Requirement already satisfied: intel-cmplr-lib-rt in
/usr/local/lib/python3.10/dist-packages (from mkl_umath-
>numpy>=1.16.0->sweetviz) (2024.2.0)
Requirement already satisfied: intel-cmplr-lib-ur==2024.2.0 in
/usr/local/lib/python3.10/dist-packages (from intel-openmp>=2024->mkl-
>numpy>=1.16.0->sweetviz) (2024.2.0)
Requirement already satisfied: ydata-profiling in
/usr/local/lib/python3.10/dist-packages (4.12.2)
Requirement already satisfied: scipy<1.16,>=1.4.1 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(1.13.1)
Requirement already satisfied: pandas!=1.4.0,<3,>1.1 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (2.2.3)
Requirement already satisfied: matplotlib>=3.5 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (3.7.5)
Requirement already satisfied: pydantic>=2 in

/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(2.11.0a2)
Requirement already satisfied: PyYAML<6.1,>=5.0.0 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (6.0.2)
Requirement already satisfied: jinja2<3.2,>=2.11.1 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (3.1.4)
Requirement already satisfied: visions<0.8.0,>=0.7.5 in
/usr/local/lib/python3.10/dist-packages (from
visions[type_image_path]<0.8.0,>=0.7.5->ydata-profiling) (0.7.6)
Requirement already satisfied: numpy<2.2,>=1.16.0 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(1.26.4)
Requirement already satisfied: htmlmin==0.1.12 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(0.1.12)
Requirement already satisfied: phik<0.13,>=0.11.1 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(0.12.4)
Requirement already satisfied: requests<3,>=2.24.0 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(2.32.3)
Requirement already satisfied: tqdm<5,>=4.48.2 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(4.67.1)
Requirement already satisfied: seaborn<0.14,>=0.10.1 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(0.12.2)
Requirement already satisfied: multimethod<2,>=1.4 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (1.12)
Requirement already satisfied: statsmodels<1,>=0.13.2 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling)
(0.14.4)
Requirement already satisfied: typeguard<5,>=3 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (4.4.1)
Requirement already satisfied: imagehash==4.3.1 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (4.3.1)
Requirement already satisfied: wordcloud>=1.9.3 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (1.9.4)
Requirement already satisfied: dacite>=1.8 in
/usr/local/lib/python3.10/dist-packages (from ydata-profiling) (1.9.2)
Requirement already satisfied: PyWavelets in
/usr/local/lib/python3.10/dist-packages (from imagehash==4.3.1->ydata-
profiling) (1.8.0)
Requirement already satisfied: pillow in
/usr/local/lib/python3.10/dist-packages (from imagehash==4.3.1->ydata-
profiling) (11.0.0)
Requirement already satisfied: MarkupSafe>=2.0 in
/usr/local/lib/python3.10/dist-packages (from jinja2<3.2,>=2.11.1-
>ydata-profiling) (3.0.2)

Requirement already satisfied: contourpy>=1.0.1 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (1.3.1)

Requirement already satisfied: cycler>=0.10 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (0.12.1)

Requirement already satisfied: fonttools>=4.22.0 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (4.55.3)

Requirement already satisfied: kiwisolver>=1.0.1 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (1.4.7)

Requirement already satisfied: packaging>=20.0 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (24.2)

Requirement already satisfied: pyparsing>=2.3.1 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (3.2.0)

Requirement already satisfied: python-dateutil>=2.7 in
/usr/local/lib/python3.10/dist-packages (from matplotlib>=3.5->ydata-
profiling) (2.9.0.post0)

Requirement already satisfied: mkl_fft in
/usr/local/lib/python3.10/dist-packages (from numpy<2.2,>=1.16.0-
>ydata-profiling) (1.3.8)

Requirement already satisfied: mkl_random in
/usr/local/lib/python3.10/dist-packages (from numpy<2.2,>=1.16.0-
>ydata-profiling) (1.2.4)

Requirement already satisfied: mkl_umath in
/usr/local/lib/python3.10/dist-packages (from numpy<2.2,>=1.16.0-
>ydata-profiling) (0.1.1)

Requirement already satisfied: mkl in /usr/local/lib/python3.10/dist-
packages (from numpy<2.2,>=1.16.0->ydata-profiling) (2025.0.1)

Requirement already satisfied: tbb4py in
/usr/local/lib/python3.10/dist-packages (from numpy<2.2,>=1.16.0-
>ydata-profiling) (2022.0.0)

Requirement already satisfied: mkl-service in
/usr/local/lib/python3.10/dist-packages (from numpy<2.2,>=1.16.0-
>ydata-profiling) (2.4.1)

Requirement already satisfied: pytz>=2020.1 in
/usr/local/lib/python3.10/dist-packages (from pandas!=1.4.0,<3,>1.1-
>ydata-profiling) (2025.1)

Requirement already satisfied: tzdata>=2022.7 in
/usr/local/lib/python3.10/dist-packages (from pandas!=1.4.0,<3,>1.1-
>ydata-profiling) (2025.1)

Requirement already satisfied: joblib>=0.14.1 in
/usr/local/lib/python3.10/dist-packages (from phik<0.13,>=0.11.1-
>ydata-profiling) (1.4.2)

Requirement already satisfied: annotated-types>=0.6.0 in
/usr/local/lib/python3.10/dist-packages (from pydantic>=2->ydata-

```

profiling) (0.7.0)
Requirement already satisfied: pydantic-core==2.29.0 in
/usr/local/lib/python3.10/dist-packages (from pydantic>=2->ydata-
profiling) (2.29.0)
Requirement already satisfied: typing-extensions>=4.12.2 in
/usr/local/lib/python3.10/dist-packages (from pydantic>=2->ydata-
profiling) (4.12.2)
Requirement already satisfied: charset-normalizer<4,>=2 in
/usr/local/lib/python3.10/dist-packages (from requests<3,>=2.24.0-
>ydata-profiling) (3.4.1)
Requirement already satisfied: idna<4,>=2.5 in
/usr/local/lib/python3.10/dist-packages (from requests<3,>=2.24.0-
>ydata-profiling) (3.10)
Requirement already satisfied: urllib3<3,>=1.21.1 in
/usr/local/lib/python3.10/dist-packages (from requests<3,>=2.24.0-
>ydata-profiling) (2.3.0)
Requirement already satisfied: certifi>=2017.4.17 in
/usr/local/lib/python3.10/dist-packages (from requests<3,>=2.24.0-
>ydata-profiling) (2025.1.31)
Requirement already satisfied: patsy>=0.5.6 in
/usr/local/lib/python3.10/dist-packages (from statsmodels<1,>=0.13.2-
>ydata-profiling) (1.0.1)
Requirement already satisfied: attrs>=19.3.0 in
/usr/local/lib/python3.10/dist-packages (from visions<0.8.0,>=0.7.5-
>visions[type_image_path]<0.8.0,>=0.7.5->ydata-profiling) (25.1.0)
Requirement already satisfied: networkx>=2.4 in
/usr/local/lib/python3.10/dist-packages (from visions<0.8.0,>=0.7.5-
>visions[type_image_path]<0.8.0,>=0.7.5->ydata-profiling) (3.4.2)
Requirement already satisfied: six>=1.5 in
/usr/local/lib/python3.10/dist-packages (from python-dateutil>=2.7-
>matplotlib>=3.5->ydata-profiling) (1.17.0)
Requirement already satisfied: intel-openmp>=2024 in
/usr/local/lib/python3.10/dist-packages (from mkl->numpy<2.2,>=1.16.0-
>ydata-profiling) (2024.2.0)
Requirement already satisfied: tbb==2022.* in
/usr/local/lib/python3.10/dist-packages (from mkl->numpy<2.2,>=1.16.0-
>ydata-profiling) (2022.0.0)
Requirement already satisfied: tcmlib==1.* in
/usr/local/lib/python3.10/dist-packages (from tbb==2022.*->mkl-
>numpy<2.2,>=1.16.0->ydata-profiling) (1.2.0)
Requirement already satisfied: intel-cmplr-lib-rt in
/usr/local/lib/python3.10/dist-packages (from mkl_umath-
>numpy<2.2,>=1.16.0->ydata-profiling) (2024.2.0)
Requirement already satisfied: intel-cmplr-lib-ur==2024.2.0 in
/usr/local/lib/python3.10/dist-packages (from intel-openmp>=2024->mkl-
>numpy<2.2,>=1.16.0->ydata-profiling) (2024.2.0)

```

```

import os
for dirname, _, filenames in os.walk('/kaggle/input'):

```

```

for filename in filenames:
    print(os.path.join(dirname, filename))

/kaggle/input/paccanatlas/
PanCanAtlas_9126RNASeqSamplesWithImmuneSubtypes_440Genes_SampleIdsOrdered_SampleIdWithSubtypes_RR020718_RownamesGenesWithSignature.csv
/kaggle/input/paccanatlas/1-s2.0-S1074761318301213-mmc2.xlsx
/kaggle/input/ai-for-health2/Groups_7302_80Percent_from_Original.csv
/kaggle/input/ai-for-health2/EBPlusPlusAdjustPANCAN_IlluminaHiSeq_RNASeqV2.geneExp.tsv

```

Section 1- Data Preparation and Pre-processing

Objective: Prepare the RNA-seq and immune subtype data by cleaning, filtering, and normalizing the datasets.

```

# Immune Subtypes Dataset
file_pathcolumns =
"/kaggle/input/paccanatlas/PanCanAtlas_9126RNASeqSamplesWithImmuneSubtypes_440Genes_SampleIdsOrdered_SampleIdWithSubtypes_RR020718_RownamesGenesWithSignature.csv"

df=pd.read_csv(file_pathcolumns,index_col=0)
df_T = df.T
df_T.head()

```

	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5	ADCY7_S3
AIMP2_S5 \				
TCGA.02.0047.GBM.C4	745.567	4287.78	241.556	1067.640
406.736				
TCGA.02.0055.GBM.C4	1154.310	9475.54	6098.950	556.132
537.088				
TCGA.02.2483.GBM.C4	1498.680	2307.12	433.984	497.309
752.148				
TCGA.02.2485.GBM.C4	1320.000	2685.71	911.905	316.667
785.552				
TCGA.02.2486.GBM.C4	1404.270	2843.90	321.951	637.805
792.963				
	ALKBH7_S5	ALOX5AP_S3	AMPD3_S3	APITD1_S5
APOC1_S3 \				
TCGA.02.0047.GBM.C4	518.148	1326.410	326.992	184.308
1370.66				
TCGA.02.0055.GBM.C4	942.957	4211.350	361.598	319.535
3093.48				
TCGA.02.2483.GBM.C4	656.042	566.543	196.728	311.443
3504.38				
TCGA.02.2485.GBM.C4	953.809	307.143	80.000	260.462

```
2482.86
TCGA.02.2486.GBM.C4      815.244      5671.950      542.683      494.488
12512.80
```

```

... VTA1_S5  WDHD1_S5  WDR54_S5  WDR77_S5
WIPF1_S3 \
TCGA.02.0047.GBM.C4 ... 1230.52  107.5630   321.460   925.044
2154.950
TCGA.02.0055.GBM.C4 ... 1183.46  162.5790   585.844   931.184
1947.580
TCGA.02.2483.GBM.C4 ... 1514.30  267.2300   561.055  1370.770
1042.740
TCGA.02.2485.GBM.C4 ... 1577.62  309.5240   422.381   811.905
849.524
TCGA.02.2486.GBM.C4 ... 1484.76   45.7317   669.512   954.878
2454.880
```

```

WNT2B_S2  WNT8B_S2  WSB2_S5  ZWILCH_S5  ZYX_S2
TCGA.02.0047.GBM.C4  43.0253    0.0000  4291.47    260.211    3867.97
TCGA.02.0055.GBM.C4  44.8493    0.0000  2494.18    467.560   11481.40
TCGA.02.2483.GBM.C4  10.1319    0.8443  4128.76    682.242    5066.81
TCGA.02.2485.GBM.C4  18.5714    0.9524  3609.52    440.386    5090.95
TCGA.02.2486.GBM.C4   7.3171    0.6098  2747.56    166.689    7800.00
```

```
[5 rows x 440 columns]
```

```
print(df_T.shape[0], "Rows-", df_T.shape[1], "Columns" )
```

```
9126 Rows- 440 Columns
```

```
# Original Whole Tsv dataset unfiltered
```

```
file_geneexpressions =
"/kaggle/input/ai-for-health2/EBPlusPlusAdjustPANCAN_IlluminaHiSeq_RNA
SeqV2.geneExp.tsv"
```

```
df_geneExpressions=pd.read_csv(file_geneexpressions, sep="\t", index_col=0)
```

```
df_geneExpressions_T=df_geneExpressions.T
df_geneExpressions_T.head()
```

```

gene_id      ?|100130426  ?|100133144  ?|
100134869  ?|10357 \
TCGA-OR-A5J1-01A-11R-A29S-07      0.0      3.2661      3.9385
149.1350
```


TCGA-0R-A5J2-01A-11R-A29S-07	0.0	2.6815	8.9948	
81.0777				
TCGA-0R-A5J3-01A-11R-A29S-07	0.0	1.7301	6.5650	
86.4879				
TCGA-0R-A5J5-01A-11R-A29S-07	0.0	0.0000	1.5492	
53.9117				
TCGA-0R-A5J6-01A-31R-A29S-07	0.0	0.0000	4.4709	
66.9063				
gene_id	? 10431	? 136542	? 155060	? 26823 ?
280660 \				
TCGA-0R-A5J1-01A-11R-A29S-07	2034.10	0.0	274.255	1.4409
0.0				
TCGA-0R-A5J2-01A-11R-A29S-07	1304.93	0.0	199.302	0.0000
0.0				
TCGA-0R-A5J3-01A-11R-A29S-07	1054.66	0.0	348.393	0.5925
0.0				
TCGA-0R-A5J5-01A-11R-A29S-07	2350.89	0.0	439.194	0.7746
0.0				
TCGA-0R-A5J6-01A-31R-A29S-07	1257.99	0.0	149.215	0.0000
0.0				
gene_id	? 317712	...	ZWILCH 55055	ZWINT 11130
\				
TCGA-0R-A5J1-01A-11R-A29S-07	0.0	...	183.9580	146.9740
TCGA-0R-A5J2-01A-11R-A29S-07	0.0	...	264.1250	438.4640
TCGA-0R-A5J3-01A-11R-A29S-07	0.0	...	90.6532	190.1940
TCGA-0R-A5J5-01A-11R-A29S-07	0.0	...	260.2630	840.4340
TCGA-0R-A5J6-01A-31R-A29S-07	0.0	...	64.2685	63.1508
gene_id	ZXDA 7789	ZXDB 158586	ZXDC 79364	\
TCGA-0R-A5J1-01A-11R-A29S-07	20.6532	351.585	1050.910	
TCGA-0R-A5J2-01A-11R-A29S-07	57.1735	279.023	2979.870	
TCGA-0R-A5J3-01A-11R-A29S-07	39.6978	495.334	914.827	
TCGA-0R-A5J5-01A-11R-A29S-07	17.8156	206.042	890.782	
TCGA-0R-A5J6-01A-31R-A29S-07	12.8537	117.919	894.730	
gene_id	ZYG11A 440590	ZYG11B 79699	ZYX 7791	\
TCGA-0R-A5J1-01A-11R-A29S-07	0.4803	648.415	1841.020	
TCGA-0R-A5J2-01A-11R-A29S-07	31.4052	1166.020	3059.990	
TCGA-0R-A5J3-01A-11R-A29S-07	0.5925	806.399	2655.610	
TCGA-0R-A5J5-01A-11R-A29S-07	11.6189	553.834	2367.930	
TCGA-0R-A5J6-01A-31R-A29S-07	7.8240	795.812	708.071	
gene_id	ZZEF1 23140	ZZZ3 26009		

```

TCGA-OR-A5J1-01A-11R-A29S-07    1157.540    596.062
TCGA-OR-A5J2-01A-11R-A29S-07    1895.990    801.637
TCGA-OR-A5J3-01A-11R-A29S-07    1482.450    437.269
TCGA-OR-A5J5-01A-11R-A29S-07    1140.200    512.781
TCGA-OR-A5J6-01A-31R-A29S-07     796.371    475.587

[5 rows x 20531 columns]

print(df_geneExpressions_T.shape[0], "Rows, ", df_geneExpressions_T.shape
[1], "Columns")

11069 Rows, 20531 Columns

df_T.columns
Index(['ACTL6A_S5', 'ADAM9_S2', 'ADAMTS1_S5', 'ADCY7_S3', 'AIMP2_S5',
      'ALKBH7_S5', 'ALOX5AP_S3', 'AMPD3_S3', 'APITD1_S5', 'APOC1_S3',
      ...,
      'VTA1_S5', 'WDHD1_S5', 'WDR54_S5', 'WDR77_S5', 'WIPF1_S3',
      'WNT2B_S2',
      'WNT8B_S2', 'WSB2_S5', 'ZWILCH_S5', 'ZYX_S2'],
      dtype='object', length=440)

df_geneExpressions_T.columns
Index(['?|100130426', '?|100133144', '?|100134869', '?|10357', '?|
10431',
      '?|136542', '?|155060', '?|26823', '?|280660', '?|317712',
      ...,
      'ZWILCH|55055', 'ZWINT|11130', 'ZXDA|7789', 'ZXDB|158586',
      'ZXDC|79364',
      'ZYG11A|440590', 'ZYG11B|79699', 'ZYX|7791', 'ZZEF1|23140',
      'ZZZ3|26009'],
      dtype='object', name='gene_id', length=20531)

```

Extracting genotypes information relevant to immune subtype sampling

```

#Filter the columns found in df_T in the tsv to extract genes relevant
to immune subtype genes
#Extract prefixes before '|' from df_geneExpressions_T columns
prefixes_df1 = df_geneExpressions_T.columns.str.split('|').str[0]

# Extract prefixes before '_' from df_T columns
prefixes_df2 = df_T.columns.str.split('_').str[0]

# Find the common prefixes
common_prefixes = prefixes_df1.intersection(prefixes_df2)

# Filter df_geneExpressions_T to keep only matching columns

```

```
df_filtered = df_geneExpressions_T.loc[:,
prefixes_df1.isin(common_prefixes)]

# Rename df_filtered columns to match naming convention from df_T
rename_mapping = {col: df_T.columns[prefixes_df2 == col.split('|')
[0]].values[0] for col in df_filtered.columns}
df_filtered = df_filtered.rename(columns=rename_mapping)

df_filtered.head()
```

gene_id	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5
ADCY7_S3 \			
TCGA-0R-A5J1-01A-11R-A29S-07	829.011	542.267	377.0410
183.477			
TCGA-0R-A5J2-01A-11R-A29S-07	935.310	1612.130	1453.5000
281.036			
TCGA-0R-A5J3-01A-11R-A29S-07	756.036	1101.470	777.9590
117.316			
TCGA-0R-A5J5-01A-11R-A29S-07	512.006	431.449	48.7994
333.075			
TCGA-0R-A5J6-01A-31R-A29S-07	410.760	2296.900	310.7240
250.368			

gene_id	AIMP2_S5	ALKBH7_S5	ALOX5AP_S3
AMPD3_S3 \			
TCGA-0R-A5J1-01A-11R-A29S-07	338.823	974.544	83.5735
163.785			
TCGA-0R-A5J2-01A-11R-A29S-07	1001.020	812.508	9.2605
332.975			
TCGA-0R-A5J3-01A-11R-A29S-07	1081.980	1422.010	53.3254
347.208			
TCGA-0R-A5J5-01A-11R-A29S-07	198.474	1307.510	10.0697
115.414			
TCGA-0R-A5J6-01A-31R-A29S-07	477.403	1442.970	326.3720
939.997			

gene_id	APITD1_S5	APOC1_S3	...	VTA1_S5
WDHD1_S5 \				
TCGA-0R-A5J1-01A-11R-A29S-07	321.7190	1875.120	...	1124.880
64.3612				
TCGA-0R-A5J2-01A-11R-A29S-07	61.1998	237.552	...	1008.190
162.6630				
TCGA-0R-A5J3-01A-11R-A29S-07	210.2740	4049.180	...	537.994
183.6770				
TCGA-0R-A5J5-01A-11R-A29S-07	208.9080	646.785	...	455.461
192.8740				
TCGA-0R-A5J6-01A-31R-A29S-07	225.7780	79054.200	...	989.176
21.2366				

gene_id	WDR54_S5	WDR77_S5	WIPF1_S3	WNT2B_S2
---------	----------	----------	----------	----------

```

\
TCGA-0R-A5J1-01A-11R-A29S-07    411.6230    951.489    571.086    11.5274
TCGA-0R-A5J2-01A-11R-A29S-07     64.0182    805.663    584.217    10.8710
TCGA-0R-A5J3-01A-11R-A29S-07   1183.8200    865.649    158.791     0.5925
TCGA-0R-A5J5-01A-11R-A29S-07    509.6820    574.748    343.919    17.0411
TCGA-0R-A5J6-01A-31R-A29S-07    483.9700    582.888    393.435    11.1771

gene_id      WNT8B_S2  WSB2_S5  ZWILCH_S5  ZYX_S2
TCGA-0R-A5J1-01A-11R-A29S-07    0.0000  1731.51   183.9580  1841.020
TCGA-0R-A5J2-01A-11R-A29S-07    0.4026  5007.11   264.1250  3059.990
TCGA-0R-A5J3-01A-11R-A29S-07    3.5550  2508.67    90.6532  2655.610
TCGA-0R-A5J5-01A-11R-A29S-07    0.7746  1889.23   260.2630  2367.930
TCGA-0R-A5J6-01A-31R-A29S-07    0.0000  3739.87    64.2685   708.071

[5 rows x 440 columns]

print("Rows:", df_filtered.shape[0], "Columns:" ,df_filtered.shape[1])
Rows: 11069 Columns: 440

```

Extracting relevant subtypes for Samples C1 to C6

```

df_T.index

Index(['TCGA.02.0047.GBM.C4', 'TCGA.02.0055.GBM.C4',
      'TCGA.02.2483.GBM.C4',
      'TCGA.02.2485.GBM.C4', 'TCGA.02.2486.GBM.C4',
      'TCGA.04.1348.OV.C2',
      'TCGA.04.1357.OV.C2', 'TCGA.04.1362.OV.C4',
      'TCGA.04.1364.OV.C1',
      'TCGA.04.1365.OV.C2',
      ...,
      'TCGA.ZP.A9D2.LIHC.C2', 'TCGA.ZP.A9D4.LIHC.C4',
      'TCGA.ZQ.A9CR.STAD.C1',
      'TCGA.ZR.A9CJ.ESCA.C1', 'TCGA.ZS.A9CD.LIHC.C3',
      'TCGA.ZS.A9CE.LIHC.C4',
      'TCGA.ZS.A9CF.LIHC.C4', 'TCGA.ZS.A9CG.LIHC.C3',
      'TCGA.ZU.A8S4.CHOL.C1',
      'TCGA.ZX.AA5X.CESC.C2'],
      dtype='object', length=9126)

df_filtered.index

Index(['TCGA-0R-A5J1-01A-11R-A29S-07', 'TCGA-0R-A5J2-01A-11R-A29S-07',
      'TCGA-0R-A5J3-01A-11R-A29S-07', 'TCGA-0R-A5J5-01A-11R-A29S-07',
      'TCGA-0R-A5J6-01A-31R-A29S-07', 'TCGA-0R-A5J7-01A-11R-A29S-07',

```

```

        'TCGA-OR-A5J8-01A-11R-A29S-07', 'TCGA-OR-A5J9-01A-11R-A29S-07',
        'TCGA-OR-A5JA-01A-11R-A29S-07', 'TCGA-OR-A5JB-01A-11R-A29S-07',
        ...
        'TCGA-CG-4449-01A-01R-1157-13', 'TCGA-CG-4462-01A-01R-1157-13',
        'TCGA-CG-4465-01A-01R-1157-13', 'TCGA-CG-4466-01A-01R-1157-13',
        'TCGA-CG-4469-01A-01R-1157-13', 'TCGA-CG-4472-01A-01R-1157-13',
        'TCGA-CG-4474-01A-02R-1157-13', 'TCGA-CG-4475-01A-01R-1157-13',
        'TCGA-CG-4476-01A-01R-1157-13', 'TCGA-CG-4477-01A-01R-1157-
13'],
        dtype='object', length=11069)

```

```

# Extract first three segments from the index
df_filtered_index_trimmed =
df_filtered.index.str.split('-').str[:3].str.join('-')
df_T_index_trimmed = df_T.index.str.split('.').str[:3].str.join('-')

# Subset df_filtered based on matching first three segments
df_filtered_subset =
df_filtered.loc[df_filtered_index_trimmed.isin(df_T_index_trimmed)]

df_filtered_subset.head()

```

gene_id	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5
ADCY7_S3 \			
TCGA-OR-A5J1-01A-11R-A29S-07	829.011	542.267	377.0410
183.477			
TCGA-OR-A5J2-01A-11R-A29S-07	935.310	1612.130	1453.5000
281.036			
TCGA-OR-A5J3-01A-11R-A29S-07	756.036	1101.470	777.9590
117.316			
TCGA-OR-A5J5-01A-11R-A29S-07	512.006	431.449	48.7994
333.075			
TCGA-OR-A5J6-01A-31R-A29S-07	410.760	2296.900	310.7240
250.368			

gene_id	AIMP2_S5	ALKBH7_S5	ALOX5AP_S3
AMPD3_S3 \			
TCGA-OR-A5J1-01A-11R-A29S-07	338.823	974.544	83.5735
163.785			
TCGA-OR-A5J2-01A-11R-A29S-07	1001.020	812.508	9.2605
332.975			
TCGA-OR-A5J3-01A-11R-A29S-07	1081.980	1422.010	53.3254
347.208			
TCGA-OR-A5J5-01A-11R-A29S-07	198.474	1307.510	10.0697
115.414			
TCGA-OR-A5J6-01A-31R-A29S-07	477.403	1442.970	326.3720
939.997			

gene_id	APITD1_S5	APOC1_S3	...	VTA1_S5
WDHD1_S5 \				

TCGA-OR-A5J1-01A-11R-A29S-07	321.7190	1875.120	...	1124.880
64.3612				
TCGA-OR-A5J2-01A-11R-A29S-07	61.1998	237.552	...	1008.190
162.6630				
TCGA-OR-A5J3-01A-11R-A29S-07	210.2740	4049.180	...	537.994
183.6770				
TCGA-OR-A5J5-01A-11R-A29S-07	208.9080	646.785	...	455.461
192.8740				
TCGA-OR-A5J6-01A-31R-A29S-07	225.7780	79054.200	...	989.176
21.2366				

gene_id	WDR54_S5	WDR77_S5	WIPF1_S3	WNT2B_S2
TCGA-OR-A5J1-01A-11R-A29S-07	411.6230	951.489	571.086	11.5274
TCGA-OR-A5J2-01A-11R-A29S-07	64.0182	805.663	584.217	10.8710
TCGA-OR-A5J3-01A-11R-A29S-07	1183.8200	865.649	158.791	0.5925
TCGA-OR-A5J5-01A-11R-A29S-07	509.6820	574.748	343.919	17.0411
TCGA-OR-A5J6-01A-31R-A29S-07	483.9700	582.888	393.435	11.1771

gene_id	WNT8B_S2	WSB2_S5	ZWILCH_S5	ZYX_S2
TCGA-OR-A5J1-01A-11R-A29S-07	0.0000	1731.51	183.9580	1841.020
TCGA-OR-A5J2-01A-11R-A29S-07	0.4026	5007.11	264.1250	3059.990
TCGA-OR-A5J3-01A-11R-A29S-07	3.5550	2508.67	90.6532	2655.610
TCGA-OR-A5J5-01A-11R-A29S-07	0.7746	1889.23	260.2630	2367.930
TCGA-OR-A5J6-01A-31R-A29S-07	0.0000	3739.87	64.2685	708.071

[5 rows x 440 columns]

```
print(df_filtered_subset.shape[0],df_filtered_subset.shape[1])
```

9879 440

```
print(df_T.shape[0],df_T.shape[1])
```

9126 440

```
df_filtered_subset.index
```

```
Index(['TCGA-OR-A5J1-01A-11R-A29S-07', 'TCGA-OR-A5J2-01A-11R-A29S-07',
      'TCGA-OR-A5J3-01A-11R-A29S-07', 'TCGA-OR-A5J5-01A-11R-A29S-07',
      'TCGA-OR-A5J6-01A-31R-A29S-07', 'TCGA-OR-A5J7-01A-11R-A29S-07',
      'TCGA-OR-A5J8-01A-11R-A29S-07', 'TCGA-OR-A5J9-01A-11R-A29S-07',
      'TCGA-OR-A5JA-01A-11R-A29S-07', 'TCGA-OR-A5JB-01A-11R-A29S-07',
      ...,
      'TCGA-CG-4449-01A-01R-1157-13', 'TCGA-CG-4462-01A-01R-1157-13',
      'TCGA-CG-4465-01A-01R-1157-13', 'TCGA-CG-4466-01A-01R-1157-13',
```

```

        'TCGA-CG-4469-01A-01R-1157-13', 'TCGA-CG-4472-01A-01R-1157-13',
        'TCGA-CG-4474-01A-02R-1157-13', 'TCGA-CG-4475-01A-01R-1157-13',
        'TCGA-CG-4476-01A-01R-1157-13', 'TCGA-CG-4477-01A-01R-1157-
13'],
        dtype='object', length=9879)

```

Extracting Primary solid tumors based on 01 in 4th component in sample TCGA Barcode

The study is based only on primary solid tumors

```

# Extracting Primary solid tumors based on 01 in 4th component in
sample TCGA Barcode as the study is based only on primary solid tumors
# Assuming df_filtered_subset.index is a list of row names
df_filtered_subset_index = df_filtered_subset.index # Replace with
your DataFrame index

# Sample valid row names

# Function to check if row name is valid
def is_valid_row_name(name):
    parts = name.split('-')
    if len(parts) > 3: # Ensure the row name has enough parts
        number = parts[3][:2] # Extract the 4th component and take
the first two characters
        return number.isdigit() and int(number)==1
    return False

# Filter valid row names
valid_rows = [name for name in df_filtered_subset_index if
is_valid_row_name(name)]

# Check if row names in df_filtered_subset.index are in valid row
names
rows_in_valid_names = [row for row in df_filtered_subset_index if row
in valid_rows]

# Filter the DataFrame to include only the rows with valid row names
df_filtered_valid_rows = df_filtered_subset.loc[rows_in_valid_names]

# Show the filtered DataFrame
df_filtered_valid_rows.shape[0]

9130

```

Handling Duplicate TCGA Samples

General Rules for Selecting Representative Samples

When dealing with duplicate TCGA sample barcodes, we follow these selection rules:

1. **Primary tumor samples (01/TP) take priority** over other sample types.
2. **Metastatic samples (06/TM) are selected over additional metastatic (07/TAM)** when primary tumors are unavailable.
3. **If multiple primary tumor samples exist**, follow the BCR annotations to select the recommended sample.
4. **When BCR annotations are missing**, select the first available aliquot (01R) unless a specific second aliquot (02R) is chosen in BCR.

Specific Recommendations for Our Cases

1 TCGA-21-1076 (Both 01A Samples)

- Both are **Primary Tumor (01/TP)**
- **Selection:** TCGA-21-1076-01A-01R-0692-07
- **Rationale:** First aliquot (01R) is preferred when both are primary tumors.

2 TCGA-06-0156 (Both 01A Samples)

- Both are **Primary Tumor (01/TP)**
- **Selection:** TCGA-06-0156-01A-02R-1849-01
- **Rationale:** BCR annotation specifically selected second aliquot (02R).

3 TCGA-06-0211 (01A vs 01B)

- 01A: **Primary Tumor (01/TP)**
- 01B: **Additional New Primary (05/TAP)**
- **Selection:** TCGA-06-0211-01A-01R-1849-01
- **Rationale:** Primary tumor (01A) is preferred over "New Primary" (05/TAP).

4 TCGA-23-1023 (01A vs 01R)

- 01A: **Primary Tumor (01/TP)**
- 01R: **Human Tumor Original Cells (08/THOC)**
- **Selection:** TCGA-23-1023-01A-02R-1564-13
- **Rationale:** Primary tumor (01A) takes precedence over cell cultures.

```
#Checking for duplicates , handling as menitonned in paper
# Get unique TCGA sample barcodes
unique_sample_barcodes = df_filtered_valid_rows.index.unique()

# Extract first three parts of each unique TCGA sample barcode
```



```

first_three_parts_set = {"-".join(sample_id.split("-")[:3]) for
sample_id in unique_sample_barcode}

print("Number of unique first-three-part TCGA sample barcodes:",
len(first_three_parts_set))

# Dictionary to store full TCGA sample barcodes for each first-three-
part key
sample_barcode_dict = defaultdict(list)

# Extract first three parts and store full TCGA sample barcodes
for sample_id in unique_sample_barcode:
    key = "-".join(sample_id.split("-")[:3]) # Extract first three
parts
    sample_barcode_dict[key].append(sample_id)

# Filter only the groups where the first three parts appear more than
once
duplicates = {key: val for key, val in sample_barcode_dict.items() if
len(val) > 1}

# Print results
print("Number of duplicated first-three-part TCGA sample barcodes:",
len(duplicates))
for key, sample_ids in duplicates.items():
    print(f"Duplicate Group ({key}):")
    print("\n".join(sample_ids)) # Print all full sample barcodes
within this duplicate group
    print("-" * 50)

```

```

Number of unique first-three-part TCGA sample barcodes: 9126
Number of duplicated first-three-part TCGA sample barcodes: 4
Duplicate Group (TCGA-21-1076):
TCGA-21-1076-01A-01R-0692-07
TCGA-21-1076-01A-02R-0692-07
-----
Duplicate Group (TCGA-06-0156):
TCGA-06-0156-01A-02R-1849-01
TCGA-06-0156-01A-03R-1849-01
-----
Duplicate Group (TCGA-06-0211):
TCGA-06-0211-01A-01R-1849-01
TCGA-06-0211-01B-01R-1849-01
-----
Duplicate Group (TCGA-23-1023):
TCGA-23-1023-01A-02R-1564-13
TCGA-23-1023-01R-01R-1564-13
-----

```

```

# List of TCGA sample IDs we want to remove
samples_to_remove = [
    "TCGA-21-1076-01A-02R",
    "TCGA-06-0156-01A-03R",
    "TCGA-06-0211-01B-01R",
    "TCGA-23-1023-01R-01R"
]

# Convert index to string to ensure proper matching
df_filtered_valid_rows.index =
df_filtered_valid_rows.index.astype(str)
# Use .str.contains() to remove indices that contain any of the
unwanted sample IDs
mask =
~df_filtered_valid_rows.index.str.contains("|".join(samples_to_remove)
, regex=True)

# Apply the mask to filter out the unwanted samples
df_filtered_valid_rows = df_filtered_valid_rows[mask]

# Display the number of samples after filtering
print(f"Final selected samples count: {len(df_filtered_valid_rows)}")

# Preview the final dataset
df_filtered_valid_rows.head()
print(df_filtered_valid_rows.shape[0])

Final selected samples count: 9126
9126

df_filtered_valid_rows = df_filtered_valid_rows.reset_index()

```

Joining Subtype Information

```

# Loading Table S1 from the research paper
file_pathcolumnsSample = "/kaggle/input/paccanatlans/1-s2.0-
S1074761318301213-mmc2.xlsx"

df_Sample=pd.read_excel(file_pathcolumnsSample)
df_Sample_T = df_Sample
df_Sample_T.head()

/usr/local/lib/python3.10/dist-packages/pandas/io/formats/
format.py:1458: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14

```

```

59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
58: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()

```

TCGA Participant Barcode	TCGA Study	Immune Subtype	TCGA Subtype \
0	TCGA-01-0639	OV	NaN
1	TCGA-02-0007	GBM	NaN
2	TCGA-02-0011	GBM	NaN
3	TCGA-02-0023	GBM	NaN
4	TCGA-02-0025	GBM	NaN

Leukocyte Fraction	Stromal Fraction	Intratumor Heterogeneity \
0	NaN	NaN
1	0.047660	NaN
2	0.062099	NaN
3	NaN	NaN
4	NaN	NaN

TIL Regional Fraction	Proliferation	Wound Healing	... T Cells
0	NaN	NaN	NaN ...
1	NaN	NaN	NaN ...
2	NaN	NaN	NaN ...
3	NaN	NaN	NaN ...
4	NaN	NaN	NaN ...

T Cells Follicular Helper T Cells gamma delta T Cells Regulatory Tregs \

0	NaN	NaN
NaN		
1	NaN	NaN
NaN		
2	NaN	NaN
NaN		
3	NaN	NaN
NaN		
4	NaN	NaN
NaN		

	Lymphocytes	Neutrophils.1	Eosinophils.1	Mast Cells	Dendritic Cells \
0	NaN	NaN	NaN	NaN	NaN
NaN					
1	NaN	NaN	NaN	NaN	NaN
NaN					
2	NaN	NaN	NaN	NaN	NaN
NaN					
3	NaN	NaN	NaN	NaN	NaN
NaN					
4	NaN	NaN	NaN	NaN	NaN
NaN					

	Macrophages
0	NaN
1	NaN
2	NaN
3	NaN
4	NaN

[5 rows x 64 columns]

```
print(df_Sample_T.shape[0], "Rows-", df_Sample_T.shape[1], "Columns" )
```

11080 Rows- 64 Columns

Viewing immune subtypes which are Not NA

```
df_Sample_NonNa = df_Sample_T[df_Sample_T["Immune Subtype"].notna()]
df_Sample_NonNa.head()
```

```
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/
format.py:1458: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
```

```

    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
58: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()

```

TCGA Participant Barcode	TCGA Study	Immune Subtype	TCGA Subtype \
9	TCGA-02-2485	GBM	C4
GBM_LGG.Classic-like			
13	TCGA-05-4410	LUAD	C3
LUAD.3			
14	TCGA-05-4424	LUAD	C2
LUAD.5			
15	TCGA-05-4425	LUAD	C6
LUAD.4			
17	TCGA-06-0156	GBM	C4
GBM_LGG.			

	Leukocyte Fraction	Stromal Fraction	Intratumor Heterogeneity \
9	0.049501	0.13	0.03
13	0.428618	0.76	0.39
14	0.326200	0.76	0.45
15	0.228394	0.39	0.08
17	NaN	NaN	NaN

TIL Regional Fraction	Proliferation	Wound Healing	... T Cells CD8 \
9	NaN	0.3223	0.150 ...
0.015523			
13	0.000000	-0.2923	-0.166 ...
0.109387			
14	4.438672	0.3447	0.020 ...
0.055662			
15	0.000000	-0.0918	-0.017 ...
0.052170			
17	NaN	0.4888	0.138 ...
0.096720			

T Cells Follicular Helper	T Cells gamma delta	T Cells Regulatory Tregs \
9	0.198057	0.054341

0.000000		
13	0.013077	0.000000
0.013376		
14	0.026872	0.000000
0.008113		
15	0.014315	0.000000
0.050522		
17	0.022861	0.000000
0.000000		

	Lymphocytes	Neutrophils.1	Eosinophils.1	Mast Cells	Dendritic Cells \
9	0.396934	0.000000	0.011272	0.094994	0.000000
13	0.726240	0.003574	0.000000	0.023075	0.000000
14	0.423194	0.012312	0.000000	0.033189	0.003214
15	0.325964	0.000000	0.000000	0.030454	0.033241
17	0.322847	0.079236	0.000000	0.082228	0.000142

	Macrophages
9	0.496801
13	0.247111
14	0.528092
15	0.610342
17	0.515548

[5 rows x 64 columns]

```
df_Sample_NonNa = df_Sample_NonNa.reset_index()
df_Sample_NonNa.set_index('TCGA Participant Barcode', inplace=True)
df_Sample_NonNa.drop(columns=['index', 'level_0'], inplace=True,
errors='ignore')
df_Sample_NonNa = df_Sample_NonNa.reset_index()

# Ensure row names are the index

# Extract the first three parts, replacing . with - for consistency
df_filtered_valid_rows['join_key'] =
df_filtered_valid_rows['index'].str.split('[-.]').str[:3].str.join('-')
df_Sample_NonNa['TCGA Participant Barcode'] = df_Sample_NonNa['TCGA
Participant Barcode'].str.split('[-.]').str[:3].str.join('-')

# Merge on the new join_key column
df_merged2 = df_Sample_NonNa.merge(df_filtered_valid_rows,
right_on='join_key', left_on='TCGA Participant Barcode',how='right')
```

```
# Drop the helper column if not needed
df_merged2.drop(columns=['join_key'], inplace=True)

df_merged2.head()

/usr/local/lib/python3.10/dist-packages/pandas/io/formats/
format.py:1458: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
58: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
```

TCGA Participant Barcode	TCGA Study	Immune Subtype	TCGA Subtype \
0	TCGA-OR-A5J1	ACC	C4
1	TCGA-OR-A5J2	ACC	C3
2	TCGA-OR-A5J3	ACC	C4
3	TCGA-OR-A5J5	ACC	C4
4	TCGA-OR-A5J6	ACC	C4
ACC.CIMP-high			
ACC.CIMP-low			
intermediate			
intermediate			
ACC.CIMP-low			

Leukocyte Fraction	Stromal Fraction	Intratumor Heterogeneity \
0	0.046374	0.10
1	0.057859	0.11
2	0.048460	0.07
3	0.016759	0.07
4	0.189302	0.31

TIL Regional Fraction	Proliferation	Wound Healing	...	VTA1_S5
-----------------------	---------------	---------------	-----	---------

\					
0	NaN	-1.2768	-0.0515	...	1124.880
1	NaN	-0.5351	-0.2730	...	1008.190
2	NaN	-1.0503	-0.1125	...	537.994
3	NaN	0.0606	0.0060	...	455.461
4	NaN	-2.2332	-0.4140	...	989.176

	WDHD1_S5	WDR54_S5	WDR77_S5	WIPF1_S3	WNT2B_S2	WNT8B_S2
WSB2_S5 \						
0	64.3612	411.6230	951.489	571.086	11.5274	0.0000
1	162.6630	64.0182	805.663	584.217	10.8710	0.4026
2	183.6770	1183.8200	865.649	158.791	0.5925	3.5550
3	192.8740	509.6820	574.748	343.919	17.0411	0.7746
4	21.2366	483.9700	582.888	393.435	11.1771	0.0000

	ZWILCH_S5	ZYX_S2
0	183.9580	1841.020
1	264.1250	3059.990
2	90.6532	2655.610
3	260.2630	2367.930
4	64.2685	708.071

[5 rows x 505 columns]

Set 'index_y' as the new index

Drop unnecessary columns

```
df_merged2.drop(columns=['index', 'level_0'], inplace=True,
errors='ignore')
```

df_merged2

```
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/
format.py:1458: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
```



```

    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
58: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()

```

	TCGA Participant Barcode	TCGA Study	Immune Subtype	\
0	TCGA-OR-A5J1	ACC	C4	
1	TCGA-OR-A5J2	ACC	C3	
2	TCGA-OR-A5J3	ACC	C4	
3	TCGA-OR-A5J5	ACC	C4	
4	TCGA-OR-A5J6	ACC	C4	
...	
9121	TCGA-CG-4472	STAD	C3	
9122	TCGA-CG-4474	STAD	C2	
9123	TCGA-CG-4475	STAD	C4	
9124	TCGA-CG-4476	STAD	C2	
9125	TCGA-CG-4477	STAD	C2	

	TCGA Subtype	Leukocyte Fraction	Stromal Fraction	\
0	ACC.CIMP-high	0.046374	0.10	
1	ACC.CIMP-low	0.057859	0.11	
2	ACC.CIMP-intermediate	0.048460	0.07	
3	ACC.CIMP-intermediate	0.016759	0.07	
4	ACC.CIMP-low	0.189302	0.31	
...	
9121	NaN	0.208206	NaN	
9122	GI.GS	0.292010	0.60	
9123	GI.CIN	0.174603	0.57	
9124	GI.CIN	0.280260	0.72	
9125	GI.CIN	0.301511	0.51	

	Intratumor Heterogeneity	TIL Regional Fraction	Proliferation	\
0	0.02	NaN	-1.2768	
1	0.16	NaN	-0.5351	
2	0.11	NaN	-1.0503	
3	0.15	NaN	0.0606	

4		0.06		NaN		-2.2332
...	
9121		NaN		9.149475		-0.2401
9122		0.32		13.349560		0.5126
9123		0.13		19.300401		0.5432
9124		0.25		7.341158		0.2374
9125		0.14		31.114986		0.8605
	Wound Healing	...	VTA1_S5	WDHD1_S5	WDR54_S5	
WDR77_S5 \						
0	-0.0515	...	1124.880000	64.361200	411.623000	951.489000
1	-0.2730	...	1008.190000	162.663000	64.018200	805.663000
2	-0.1125	...	537.994000	183.677000	1183.820000	865.649000
3	0.0060	...	455.461000	192.874000	509.682000	574.748000
4	-0.4140	...	989.176000	21.236600	483.970000	582.888000
...
...						
9121	-0.1060	...	1002.318114	203.007108	156.681161	642.255901
9122	0.1460	...	1596.401835	419.267633	71.266353	1004.123256
9123	0.1570	...	1071.598647	437.232142	369.559696	421.023012
9124	0.0310	...	1128.296079	346.089113	165.670587	1047.209793
9125	0.2880	...	1828.291508	411.704146	329.701859	1035.886535
	WIPF1_S3	WNT2B_S2	WNT8B_S2	WSB2_S5	ZWILCH_S5	ZYX_S2
0	571.086000	11.527400	0.000000	1731.510000	183.958000	1841.020000
1	584.217000	10.871000	0.402600	5007.110000	264.125000	3059.990000
2	158.791000	0.592500	3.555000	2508.670000	90.653200	2655.610000
3	343.919000	17.041100	0.774600	1889.230000	260.263000	2367.930000

```

4      393.435000    11.177100    0.000000    3739.870000    64.268500
708.071000
...      ...      ...      ...      ...      ...
...
9121  1964.794656    63.843908    2.351098    1716.985841    292.541116
3359.241568
9122  1782.515239    319.359797    1.061085    2914.005907    745.532095
4264.469081
9123  1400.004751    21.267203    0.283374    1535.171082    556.284407
3103.609391
9124  1945.712810    325.416409    0.574527    1652.487434    516.997610
3302.569055
9125  2546.771364    77.610289    15.219672    1446.443138    923.544322
2497.814797

```

```
[9126 rows x 504 columns]
```

```
df_merged2.shape[0]
```

```
9126
```

```
column_names_Table_S1 = df_merged2.columns[:64].tolist()
print(column_names_Table_S1)
```

```

['TCGA Participant Barcode', 'TCGA Study', 'Immune Subtype', 'TCGA
Subtype', 'Leukocyte Fraction', 'Stromal Fraction', 'Intratumor
Heterogeneity', 'TIL Regional Fraction', 'Proliferation', 'Wound
Healing', 'Macrophage Regulation', 'Lymphocyte Infiltration Signature
Score', 'IFN-gamma Response', 'TGF-beta Response', 'SNV Neoantigens',
'Indel Neoantigens', 'Silent Mutation Rate', 'Nonsilent Mutation
Rate', 'Number of Segments', 'Fraction Altered', 'Aneuploidy Score',
'Homologous Recombination Defects', 'BCR Evenness', 'BCR Shannon',
'BCR Richness', 'TCR Shannon', 'TCR Richness', 'TCR Evenness', 'CTA
Score', 'Th1 Cells', 'Th2 Cells', 'Th17 Cells', 'OS', 'OS Time',
'PFI', 'PFI Time', 'B Cells Memory', 'B Cells Naive', 'Dendritic Cells
Activated', 'Dendritic Cells Resting', 'Eosinophils', 'Macrophages
M0', 'Macrophages M1', 'Macrophages M2', 'Mast Cells Activated', 'Mast
Cells Resting', 'Monocytes', 'Neutrophils', 'NK Cells Activated', 'NK
Cells Resting', 'Plasma Cells', 'T Cells CD4 Memory Activated', 'T
Cells CD4 Memory Resting', 'T Cells CD4 Naive', 'T Cells CD8', 'T
Cells Follicular Helper', 'T Cells gamma delta', 'T Cells Regulatory
Tregs', 'Lymphocytes', 'Neutrophils.1', 'Eosinophils.1', 'Mast Cells',
'Dendritic Cells', 'Macrophages']

```

```
column_names_PanCan = df_merged2.columns[64:].tolist()
print(column_names_PanCan)
```

```

['ACTL6A_S5', 'ADAM9_S2', 'ADAMTS1_S5', 'ADCY7_S3', 'AIMP2_S5',
'ALKBH7_S5', 'ALOX5AP_S3', 'AMPD3_S3', 'APITD1_S5', 'APOC1_S3',
'APOE_S3', 'APOO_S5', 'ARHGAP1_S2', 'ARHGAP15_S3', 'ARHGDIA_S2',
'ARRB2_S3', 'B2M_S3', 'BCCIP_S5', 'BRCA2_S5', 'BRIP1_S5', 'BSG_S2',

```

'BTK_S3', 'C11orf24_S5', 'C12orf24_S5', 'C13orf1_S5', 'C13orf18_S3', 'C13orf27_S5', 'C16orf61_S5', 'C19orf48_S5', 'C1orf54_S3', 'C3AR1_S3', 'C3orf26_S5', 'CASP1_S3', 'CBX1_S5', 'CCBL2_S5', 'CCL2_S3', 'CCL5_S3', 'CCRL2_S3', 'CCT5_S5', 'CD14_S4', 'CD163_S3', 'CD19_S4', 'CD247_S4', 'CD33_S3', 'CD37_S4', 'CD3D_S4', 'CD3E_S4', 'CD3G_S4', 'CD44_S2', 'CD48_S3', 'CD52_S3', 'CD53_S3', 'CD59_S2', 'CD79A_S4', 'CD79B_S4', 'CD84_S3', 'CD86_S3', 'CD8A_S3', 'CD8B_S4', 'CD97_S3', 'CDCA4_S5', 'CDH6_S2', 'CDK2_S5', 'CECR1_S3', 'CELF2_S3', 'CENPJ_S5', 'CENPN_S5', 'CENPO_S5', 'CENPW_S5', 'CEP78_S5', 'CHEK1_S5', 'CKLF_S5', 'COL16A1_S2', 'COL1A2_S2', 'COL3A1_S2', 'COL6A1_S2', 'COL6A3_S2', 'COL8A1_S2', 'COPS6_S5', 'COQ2_S5', 'CORO1A_S3', 'CORO1C_S5', 'COTL1_S5', 'COX17_S5', 'CPEB4_S5', 'CPVL_S3', 'CSF1R_S3', 'CTNNA1_S2', 'CTNNB1_S2', 'CTSC_S3', 'CTSL1_S3', 'CTS5_S3', 'CXCL10_S1', 'CXCR4_S3', 'CYBB_S3', 'CYTIP_S3', 'DAPK1_S3', 'DBNDD1_S5', 'DCBLD2_S5', 'DCK_S5', 'DCLRE1B_S5', 'DDX58_S1', 'DDX60_S1', 'DHFR_S5', 'DLEU1_S5', 'DLEU2_S5', 'DOCK2_S3', 'DSP_S2', 'DUT_S5', 'DVL1_S2', 'DVL3_S2', 'DYNLT1_S5', 'EBNA1BP2_S5', 'EFNA5_S2', 'EIF2AK1_S5', 'EIF4EBP1_S5', 'EIF4G1_S5', 'EMP2_S5', 'ENO1_S5', 'EPHA2_S2', 'EPHB2_S2', 'EPHB3_S2', 'EPHB4_S2', 'ERLIN1_S5', 'EVI2A_S3', 'EVI2B_S3', 'EXOSC8_S5', 'EZR_S5', 'F3_S5', 'FAM167A_S5', 'FAM89B_S5', 'FARSA_S5', 'FARSB_S5', 'FBXO41_S5', 'FCER1G_S3', 'FCGR1A_S3', 'FCGR2A_S3', 'FCGR2B_S3', 'FCGR3A_S3', 'FGD1_S2', 'FGL2_S3', 'FGR_S3', 'FLI1_S3', 'FLNC_S5', 'FMNL1_S3', 'FN1_S2', 'FNBP1_S3', 'FPR3_S3', 'FYB_S3', 'GGH_S5', 'GIMAP4_S3', 'GLRX3_S5', 'GNG11_S5', 'GNPTAB_S3', 'GPLD1_S5', 'GPNMB_S3', 'GSTCD_S5', 'H2AFZ_S5', 'HAS2_S5', 'HAUS1_S5', 'HCK_S3', 'HCLS1_S3', 'HDC_S4', 'HERC5_S1', 'HERC6_S1', 'HLA-DMA_S3', 'HLA-DRB1_S3', 'HMGN2_S5', 'HMHA1_S3', 'HN1L_S5', 'HNRNPA2B1_S5', 'HNRNPR_S5', 'HSPB11_S5', 'HSPG2_S2', 'HYS1_S5', 'ICAM1_S2', 'ID2_S5', 'ID3_S5', 'IFI16_S3', 'IFI27_S1', 'IFI30_S3', 'IFI44_S1', 'IFI44L_S1', 'IFI6_S1', 'IFIH1_S1', 'IFIT1_S1', 'IFIT2_S1', 'IFIT3_S1', 'IFRD2_S5', 'IGF2R_S2', 'IGFBP2_S2', 'IGFBP3_S2', 'IGFBP4_S2', 'IGFBP5_S2', 'IGJ_S4', 'IKZF1_S3', 'IL10RA_S3', 'IL18_S3', 'IL7R_S3', 'IMP4_S5', 'IPO4_S5', 'ISG15_S1', 'ITGA3_S2', 'ITGA4_S2', 'ITGA5_S2', 'ITGA6_S5', 'ITGB2_S2', 'ITGB3_S2', 'ITGB5_S2', 'ITGB8_S2', 'JUNB_S5', 'JUP_S2', 'KIAA0090_S5', 'KRR1_S5', 'LAIR1_S3', 'LAMA4_S2', 'LAMB1_S2', 'LAPTM5_S3', 'LCK_S4', 'LCP1_S3', 'LCP2_S3', 'LHFPL2_S3', 'LILRB4_S3', 'LMNB2_S5', 'LOXL2_S5', 'LRMP_S3', 'LRP1_S2', 'LRRC17_S2', 'LRRC40_S5', 'LSM3_S5', 'LSM4_S5', 'LST1_S3', 'LTB_S4', 'LY86_S3', 'LYAR_S5', 'LYN_S3', 'LYZ_S3', 'MAGOHB_S5', 'MAP3K8_S5', 'MAPRE1_S5', 'MARCKSL1_S2', 'MARVELD2_S5', 'MCM3_S5', 'MCM7_S5', 'MCTS1_S5', 'MERTK_S3', 'MET_S5', 'MFSD11_S5', 'MKKS_S5', 'MLF1IP_S5', 'MMP11_S2', 'MMP1_S2', 'MMP14_S2', 'MMP17_S2', 'MMP19_S2', 'MMP2_S2', 'MMP3_S2', 'MNAT1_S5', 'MND4_S3', 'MPP1_S3', 'MRPL12_S5', 'MRPL37_S5', 'MRPS16_S5', 'MRPS28_S5', 'MRT04_S5', 'MS4A1_S4', 'MSN_S3', 'MT1F_S5', 'MT3_S5', 'MTA1_S2', 'MTHFD1_S5', 'MX1_S1', 'MYBL1_S5', 'MYBL2_S5', 'MYCBP_S5', 'MYL6_S5', 'MYO1F_S3', 'NCEH1_S5', 'NCF2_S3', 'NCKAP1L_S3', 'NCLN_S5', 'NEO1_S2', 'NID1_S2', 'NLN_S5', 'NME1_S5', 'NOP16_S5', 'NOTCH2_S2', 'NPC2_S3', 'NPL_S3', 'NRIP3_S5', 'NUDT1_S5',

```
'NUDT15_S5', 'NUP107_S5', 'NUP35_S5', 'NUP85_S5', 'NUP93_S5',
'NUPL1_S5', 'NUTF2_S5', 'OAS1_S1', 'OAS2_S1', 'OAS3_S1', 'OASL_S1',
'OSBPL3_S3', 'PA2G4_S5', 'PAICS_S5', 'PAK1_S2', 'PAK2_S2', 'PDAP1_S5',
'PDIA4_S5', 'PDLIM7_S5', 'PFKP_S5', 'PFN1_S5', 'PGM2_S5', 'PHF19_S5',
'PIK3CG_S3', 'PITPNC1_S5', 'PLAT_S2', 'PLAUR_S5', 'PLCG2_S3',
'PLEK_S3', 'PLG_S5', 'PLK4_S5', 'PLOD2_S5', 'PNN_S5', 'PN01_S5',
'POLE2_S5', 'POLR3K_S5', 'PPIH_S5', 'PSMA7_S5', 'PSMC3_S5',
'PSMD12_S5', 'PSMD14_S5', 'PSMD2_S5', 'PTPLB_S5', 'PTPRC_S3',
'PXN_S2', 'RAB3B_S5', 'RAC1_S2', 'RASSF2_S3', 'RASSF4_S3', 'RBM14_S5',
'RFC3_S5', 'RGS8_S5', 'RHOA_S2', 'RHOB_S2', 'RHOC_S5', 'RHOG_S2',
'RHQ_S2', 'RMND5B_S5', 'RNASE6_S3', 'RND3_S2', 'RNF138_S5',
'RNF41_S5', 'RPN1_S5', 'RPP40_S5', 'RSAD2_S1', 'RTP4_S1', 'RUNX3_S3',
'RUVBL1_S5', 'SAMD9_S1', 'SAMHD1_S3', 'SAMSN1_S3', 'SAR1A_S5',
'SAR1B_S5', 'SDC1_S5', 'SELL_S3', 'SELPLG_S3', 'SEMA3F_S2',
'SERPINE1_S2', 'SH2B3_S3', 'SH3BP5L_S5', 'SKA1_S5', 'SKA2_S5',
'SLC16A1_S5', 'SLC1A3_S3', 'SLC25A4_S5', 'SLC25A5_S5', 'SLC7A7_S3',
'SMC2_S5', 'SMO_S2', 'SMS_S5', 'SMURF2_S5', 'SNRPA1_S5', 'SNRPA_S5',
'SNRPB_S5', 'SNRPC_S5', 'SNRPD1_S5', 'SNRPE_S5', 'SNX17_S5',
'SP140_S3', 'SPAG17_S5', 'SPARC_S2', 'SRM_S5', 'STAT1_S1',
'STK17A_S5', 'STRA13_S5', 'SYK_S3', 'TAGLN_S5', 'TAP1_S1',
'TBXAS1_S3', 'TCEB1_S5', 'TCF7L2_S2', 'THBS1_S2', 'THBS2_S2',
'TIMP1_S2', 'TLR2_S3', 'TMEM130_S5', 'TNC_S2', 'TNFAIP3_S3',
'TNFRSF12A_S5', 'TNFRSF1A_S2', 'TNFRSF1B_S3', 'TNFSF13B_S3',
'TOMM40_S5', 'TPI1_S5', 'TPM1_S5', 'TPM2_S5', 'TPM3_S5', 'TPRKB_S5',
'TRA2B_S5', 'TUBA4A_S5', 'TUBG1_S5', 'UAP1_S5', 'UBE2J1_S5',
'UMPS_S5', 'UQCR10_S5', 'USPL1_S5', 'VCAN_S2', 'VDAC1_S5', 'VSIG4_S3',
'VTA1_S5', 'WDHD1_S5', 'WDR54_S5', 'WDR77_S5', 'WIPF1_S3', 'WNT2B_S2',
'WNT8B_S2', 'WSB2_S5', 'ZWILCH_S5', 'ZYX_S2']
```

More Preprocessing

```
# Remove display limits
pd.set_option('display.max_rows', None) # Show all rows
pd.set_option('display.max_columns', None) # Show all columns
pd.set_option('display.width', None) # Adjust width to fit content
pd.set_option('display.max_colwidth', None) # Show full column
content
#print(df_final.isna().sum())
```

```
df_merged2["Immune Subtype"].value_counts()
```

```
Immune Subtype
C2      2591
C1      2416
C3      2397
C4       1157
C5        385
```

```
C6      180
Name: count, dtype: int64
```

Balanced Representation of Immune Subtypes

I selected patient samples randomly while ensuring that each immune subtype was represented according to the required distribution: C1 (385 samples), C2 (414 samples), C3 (383 samples), C4 (462 samples), C5 (231 samples), and C6 (134 samples). This resulted in a total of 2009 samples in my dataset.

```
# Define the target sample sizes
target_samples = {
    "C1": 385,
    "C2": 414,
    "C3": 383,
    "C4": 462,
    "C5": 231,
    "C6": 134
}

# Perform stratified sampling
df_balanced = pd.concat([
    df_merged2[df_merged2["Immune Subtype"] ==
    subtype].sample(n=target_samples[subtype], random_state=42)
    for subtype in target_samples
])

# Verify the final distribution
print(df_balanced["Immune Subtype"].value_counts())

Immune Subtype
C4      462
C2      414
C1      385
C3      383
C5      231
C6      134
Name: count, dtype: int64

df_balanced.shape[0]

2009

# This Python 3 environment comes with many helpful analytics
libraries installed
# It is defined by the kaggle/python Docker image:
https://github.com/kaggle/docker-python
# For example, here's several helpful packages to load
```

```

import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
from mpl_toolkits.mplot3d import Axes3D
import matplotlib.pyplot as plt
from sklearn.decomposition import NMF
from sklearn.preprocessing import MinMaxScaler
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.svm import SVC
from sklearn.metrics import classification_report, confusion_matrix
df_balanced.head()

```

```

/usr/local/lib/python3.10/dist-packages/pandas/io/formats/
format.py:1458: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()

```

	TCGA Participant Barcode	TCGA Study	Immune Subtype	TCGA Subtype
1264	TCGA-WA-A7H4	HNSC	C1	NaN
548	TCGA-B5-A3F9	UCEC	C1	UCEC.CN_HIGH
6812	TCGA-49-6767	LUAD	C1	LUAD.3
7863	TCGA-AA-A03J	COAD	C1	GI.CIN
3228	TCGA-77-7337	LUSC	C1	NaN

	Leukocyte Fraction	Stromal Fraction	Intratumor Heterogeneity
1264	0.379482	0.82	0.40
548	0.219421	0.74	0.08
6812	0.348569	0.57	0.26
7863	0.241930	0.55	0.05
3228	0.297990	0.48	0.22

	TIL Regional Fraction	Proliferation	Wound Healing	\
1264	NaN	0.5746	0.1880	
548	8.503943	0.1406	0.1875	
6812	14.644431	0.6171	0.2375	
7863	8.225735	0.2587	0.1150	
3228	15.458627	0.9131	0.3410	
Score	Macrophage Regulation	Lymphocyte Infiltration	Signature	
1264	-0.1361		0.3799	
548	-0.1861		-0.1434	
6812	0.3779		0.6282	
7863	-0.3616		0.2081	
3228	0.2877		0.2484	
	IFN-gamma Response	TGF-beta Response	SNV Neoantigens	\
1264	0.0053	0.7164	27.0	
548	-0.1845	-0.1684	27.0	
6812	0.2599	0.2542	166.0	
7863	0.1503	0.1076	25.0	
3228	-0.1595	0.8806	176.0	
	Indel Neoantigens	Silent Mutation Rate	Nonsilent Mutation Rate	\
1264	0.0	0.511520	1.591394	
548	3.0	0.494144	1.095711	
6812	60.0	4.161290	11.800230	
7863	NaN	0.835700	2.328021	
3228	127.0	2.884412	8.474817	
	Number of Segments	Fraction Altered	Aneuploidy Score	\
1264	72.0	0.197422	10.0	
548	123.0	0.271444	12.0	
6812	172.0	0.581840	21.0	
7863	119.0	0.507269	18.0	
3228	169.0	0.578212	13.0	
	Homologous Recombination Defects	BCR Evenness	BCR Shannon	\
1264	34.0	0.922180	2.026238	
548	4.0	NaN	NaN	

6812	30.0	NaN	NaN
7863	10.0	NaN	NaN
3228	54.0	0.877917	3.549464

	BCR Richness	TCR Shannon	TCR Richness	TCR Evenness	CTA Score
\					
1264	9.0	0.636514	2.0	0.918296	5.581650
548	NaN	1.549826	5.0	0.962961	3.171152
6812	NaN	2.679544	17.0	0.945761	2.304206
7863	NaN	NaN	NaN	NaN	0.542844
3228	57.0	2.871476	18.0	0.993463	2.134452

	Th1 Cells	Th2 Cells	Th17 Cells	OS	OS Time	PFI	PFI
Time \							
1264	-1107.504119	335.699054	-633.233703	0.0	443.0	0.0	
443.0							
548	-706.967641	-589.271926	-1741.589978	0.0	4155.0	0.0	
4155.0							
6812	68.317547	282.227079	-2484.773402	0.0	677.0	0.0	
677.0							
7863	-800.007476	-581.504381	395.084039	0.0	1246.0	0.0	
1246.0							
3228	-159.628570	1137.349162	-1602.002392	1.0	3253.0	0.0	
3253.0							

	B Cells Memory	B Cells Naive	Dendritic Cells Activated	\
1264	0.000000	0.075893	0.000000	
548	0.039246	0.000000	0.002850	
6812	0.010711	0.014774	0.000000	
7863	0.015177	0.008338	0.000000	
3228	0.000000	0.049060	0.044275	

	Dendritic Cells Resting	Eosinophils	Macrophages M0
Macrophages M1 \			
1264	0.000000	0.000000	0.285709
0.026984			
548	0.000000	0.003538	0.076611
0.007250			
6812	0.000000	0.000000	0.252027
0.067449			
7863	0.018624	0.000000	0.034319
0.078631			
3228	0.002638	0.000000	0.128922
0.111607			

Macrophages M2		Mast Cells Activated		Mast Cells Resting	
Monocytes \					
1264	0.153365	0.000000	0.026059		
0.023109					
548	0.341072	0.000000	0.000000		
0.000000					
6812	0.164293	0.000000	0.033246		
0.000000					
7863	0.196650	0.042318	0.000000		
0.033560					
3228	0.218733	0.037657	0.000000		
0.011673					
Neutrophils		NK Cells Activated		NK Cells Resting	
\					
1264	0.003019	0.087552	0.008897	0.111249	
548	0.001251	0.064078	0.000000	0.035881	
6812	0.000000	0.111910	0.000000	0.000000	
7863	0.000000	0.093794	0.000000	0.026578	
3228	0.015157	0.000000	0.059039	0.095555	
T Cells CD4 Memory		Activated		T Cells CD4 Memory Resting	
\					
1264		0.000000	0.106067		
548		0.000000	0.070163		
6812		0.027125	0.043710		
7863		0.000000	0.000000		
3228		0.014349	0.107035		
T Cells CD4 Naive		T Cells CD8		T Cells Follicular Helper	
\					
1264	0.017419	0.027415	0.036103		
548	0.000000	0.169194	0.133036		
6812	0.000000	0.167443	0.060282		
7863	0.000000	0.277904	0.089188		
3228	0.000000	0.087035	0.016079		
T Cells gamma delta		T Cells Regulatory Tregs		Lymphocytes	
\					
1264	0.0	0.011161	0.481756		
548	0.0	0.055830	0.567428		
6812	0.0	0.047029	0.482986		
7863	0.0	0.084921	0.595899		
3228	0.0	0.001184	0.429337		
Neutrophils.1		Eosinophils.1		Mast Cells	
Macrophages				Dendritic Cells	
\					
1264	0.003019	0.000000	0.026059	0.000000	

0.489166				
548	0.001251	0.003538	0.000000	0.002850
0.424933				
6812	0.000000	0.000000	0.033246	0.000000
0.483769				
7863	0.000000	0.000000	0.042318	0.018624
0.343160				
3228	0.015157	0.000000	0.037657	0.046913
0.470935				

	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5	ADCY7_S3	AIMP2_S5
\					
1264	1414.260000	5574.520000	942.308000	946.314000	563.534000
548	1354.770000	1138.580000	1640.530000	1227.540000	482.911000
6812	1637.870000	5176.330000	161.538000	1004.730000	488.988000
7863	930.343078	1329.678264	216.256995	370.030352	1349.628189
3228	3289.310000	14204.300000	2384.660000	573.328000	1188.010000

	ALKBH7_S5	ALOX5AP_S3	AMPD3_S3	APITD1_S5	APOC1_S3	\
1264	521.635000	237.1790	587.340000	216.987000	138.622000	
548	872.698000	185.8710	353.850000	340.800000	1306.070000	
6812	299.408000	482.2480	276.331000	274.065000	1394.080000	
7863	1956.948165	131.8582	149.806773	125.083134	400.202368	
3228	317.341000	743.7580	308.820000	368.092000	126.118000	

	APOE_S3	AP00_S5	ARHGAP1_S2	ARHGAP15_S3	ARHGDIA_S2
\					
1264	1125.000000	110.577000	3838.940000	44.070500	8888.620000
548	4475.310000	620.729000	1648.980000	12.921500	6709.740000
6812	5431.360000	372.781000	1626.630000	71.597600	5318.340000
7863	2667.522991	252.673868	3276.934604	12.524268	15147.247057
3228	620.366000	315.296000	4495.270000	28.632300	8677.970000

	ARRB2_S3	B2M_S3	BCCIP_S5	BRCA2_S5
BRIP1_S5 \				
1264	523.237000	48758.800000	951.234000	224.359000
548	521.333000	66318.100000	1651.590000	70.571300
6812	695.858000	53988.100000	1667.410000	109.468000

7863	1306.517461	41306.987107	769.615704	146.312249	36.943463	
3228	564.124000	33636.100000	1540.930000	147.593000	209.970000	
	BSG_S2	BTK_S3	C11orf24_S5	C12orf24_S5	C13orf1_S5	\
1264	16931.90000	86.538500	1333.330000	66.50640	310.096000	
548	21927.30000	60.631700	754.417000	93.92940	273.836000	
6812	8614.20000	156.805000	1263.310000	68.04730	381.657000	
7863	19473.23631	28.087146	1035.470316	53.88891	492.691892	
3228	13157.20000	85.896900	1387.980000	84.19260	330.976000	
	C13orf18_S3	C13orf27_S5	C16orf61_S5	C19orf48_S5		
	C1orf54_S3	\				
1264	127.404000	193.910000	358.758000	976.763000	155.449000	
548	199.786000	180.404000	548.716000	1952.140000	203.265000	
6812	38.461500	166.864000	548.941000	863.314000	137.278000	
7863	2546.069144	117.261586	479.162611	4435.195669	66.573924	
3228	44.652700	250.533000	565.549000	1145.290000	81.806600	
	C3AR1_S3	C3orf26_S5	CASP1_S3	CBX1_S5	CCBL2_S5	\
1264	133.814000	787.660000	790.865000	1612.180000	716.090000	
548	154.561000	635.141000	120.766000	2532.120000	626.022000	
6812	370.414000	468.047000	199.408000	1936.090000	795.284000	
7863	130.132203	375.363702	463.236328	620.153334	424.498066	
3228	369.493000	1113.930000	672.177000	2124.930000	621.464000	
	CCL2_S3	CCL5_S3	CCRL2_S3	CCT5_S5		
	CD14_S4	\				
1264	351.763000	1123.400000	36.057700	7229.970000	1347.76000	
548	620.232000	500.957000	27.333900	6531.320000	1638.55000	
6812	642.604000	2890.530000	71.005900	10427.200000	1537.87000	
7863	122.172779	607.562311	370.657909	8111.215987	1392.85134	
3228	1044.060000	555.944000	62.377500	12827.600000	1825.99000	
	CD163_S3	CD19_S4	CD247_S4	CD33_S3	CD37_S4	
	CD3D_S4	\				
1264	320.513000	9.615400	101.763000	31.250000	169.8720	
	92.948700					
548	583.456000	5.466800	56.158800	67.092400	190.8410	
	93.432400					

6812	834.319000	1.775100	133.728000	71.005900	450.8880
200.592000					
7863	164.663887	8.810904	71.947677	30.009939	246.5297
109.859681					
3228	1267.660000	3.408600	63.400100	46.357100	182.0200
79.420500					
	CD3E_S4	CD3G_S4	CD44_S2	CD48_S3	CD52_S3
CD53_S3 \					
1264	187.500000	14.423100	18930.300000	94.551300	186.699000
338.14100					
548	194.320000	4.472800	2058.990000	157.543000	171.458000
465.17400					
6812	602.367000	56.804700	9745.560000	253.846000	629.586000
863.90500					
7863	189.354061	17.990105	3261.290314	96.453995	268.239955
315.95604					
3228	256.327000	41.244100	23292.700000	132.595000	219.514000
878.05700					
	CD59_S2	CD79A_S4	CD79B_S4	CD84_S3	CD86_S3
CD8A_S3 \					
1264	14369.400000	133.814000	56.089700	20.833300	233.173000
33.65380					
548	6228.660000	16.400400	18.388300	28.824900	104.863000
197.79800					
6812	6684.020000	37.869800	22.485200	62.721900	304.734000
652.07100					
7863	2817.520364	64.726949	33.807438	25.931826	77.030344
276.25988					
3228	13698.500000	174.862000	31.359200	117.597000	397.103000
208.60700					
	CD8B_S4	CD97_S3	CDCA4_S5	CDH6_S2	CDK2_S5 \
1264	24.038500	761.218000	1024.840000	84.13460	818.910000
548	229.605000	1288.170000	685.834000	69.57730	876.177000
6812	97.041400	1813.020000	786.982000	11.83430	1046.150000
7863	77.724818	6066.851717	480.374708	23.29196	787.869981
3228	102.940000	1295.950000	1040.310000	2406.48000	923.392000
	CECR1_S3	CELF2_S3	CENPJ_S5	CENPN_S5	CENPO_S5 \
1264	422.276000	397.436000	226.763000	702.941000	202.780000
548	1813.480000	328.007000	616.256000	295.157000	74.810500
6812	971.598000	270.414000	89.349100	584.787000	145.675000
7863	447.176864	700.887602	196.492764	630.422866	190.410746
3228	659.225000	306.775000	107.712000	1274.420000	436.735000
	CENPW_S5	CEP78_S5	CHEK1_S5	CKLF_S5	COL16A1_S2 \
1264	568.109000	243.590000	537.660000	213.14100	5395.030000
548	454.240000	284.273000	269.364000	227.61700	182.392000

6812	353.846000	223.077000	502.959000	897.04100	710.059000
7863	931.541342	89.391743	175.446418	465.70463	681.668235
3228	1029.400000	493.566000	662.633000	235.19400	862.378000

	COL1A2_S2	COL3A1_S2	COL6A1_S2	COL6A3_S2
COL8A1_S2 \				
1264	93153.000000	103386.000000	20055.300000	26830.900000
208.33300				
548	10381.900000	22155.900000	9287.580000	1272.270000
9.93960				
6812	21293.500000	21016.000000	18462.100000	8066.860000
395.85800				
7863	30497.210498	23821.008655	8179.105901	8199.892726
237.28691				
3228	105622.000000	155202.000000	6743.590000	34333.200000
896.12300				

	COPS6_S5	COQ2_S5	COR01A_S3	COR01C_S5
COTL1_S5 \				
1264	1607.370000	274.038000	713.141000	3767.630000
2705.130000				
548	2249.340000	250.478000	567.552000	1157.970000
3072.830000				
6812	8586.390000	269.822000	1411.830000	5149.700000
9766.860000				
7863	2323.238838	333.605287	3106.212752	3463.398929
2085.454571				
3228	1875.760000	253.941000	495.271000	8162.930000
3322.710000				

	COX17_S5	CPEB4_S5	CPVL_S3	CSF1R_S3	CTNNA1_S2
CTNNB1_S2 \					
1264	538.462000	811.699000	203.52600	724.359000	8176.2800
8176.970000					
548	778.769000	460.701000	230.10200	653.033000	7995.4300
6567.580000					
6812	839.645000	245.562000	393.49100	988.757000	6928.4000
13272.800000					
7863	591.715669	303.291589	222.88383	671.456392	8261.3316
9974.124921					
3228	399.489000	894.759000	299.95700	1158.930000	7950.5800
12085.200000					

	CTSC_S3	CTSL1_S3	CTSS_S3	CXCL10_S1
CXCR4_S3 \				
1264	5598.560000	1512.820000	835.737000	187.500000
298.077000				
548	3442.090000	2101.730000	1766.270000	308.625000
2493.850000				
6812	4711.830000	1643.790000	10181.700000	2600.000000

601.775000					
7863	2538.514223	658.990557	3005.643104	304.784386	
270.766523					
3228	11806.700000	2636.220000	2027.100000	568.215000	
1160.630000					
	CYBB_S3	CYTIP_S3	DAPK1_S3	DBNDD1_S5	DCBLD2_S5
DCK_S5 \					
1264	471.154000	129.80800	905.449000	197.115000	3229.170000
548.87800					
548	468.156000	163.50700	462.689000	824.988000	2031.160000
357.82600					
6812	1659.760000	208.87600	169.822000	862.130000	24820.100000
872.18900					
7863	350.122639	53.78859	304.985752	337.246512	503.510219
352.77352					
3228	1642.610000	252.91900	320.409000	167.703000	2947.080000
997.01800					
	DCLRE1B_S5	DDX58_S1	DDX60_S1	DHFR_S5	DLEU1_S5
DLEU2_S5 \					
1264	323.718000	770.833000	713.141000	225.561000	80.769200
69.487200					
548	238.551000	387.645000	578.983000	422.026000	201.471000
49.285600					
6812	264.497000	452.663000	763.905000	227.876000	92.112400
107.444000					
7863	142.552052	396.642255	705.860341	388.993479	134.812507
25.400413					
3228	276.438000	222.923000	731.487000	175.202000	170.703000
102.940000					
	DOCK2_S3	DSP_S2	DUT_S5	DVL1_S2	DVL3_S2
\					
1264	203.526000	44610.600000	1073.000000	2220.350000	2595.350000
548	100.390000	4294.410000	1422.170000	1487.960000	2026.190000
6812	321.302000	8887.570000	624.645000	1872.780000	2149.110000
7863	146.765031	8165.998316	1715.641893	3209.336764	2974.540775
3228	315.296000	52847.400000	577.541000	804.431000	4245.420000
	DYNLT1_S5	EBNA1BP2_S5	EFNA5_S2	EIF2AK1_S5	
EIF4EBP1_S5 \					
1264	1262.020000	2073.72000	518.429000	3613.550000	1009.620000
548	2670.780000	1220.58000	64.607500	5030.100000	1500.880000

6812	1210.650000	1886.98000	18.343200	4418.110000	1133.140000
7863	1160.197487	1246.37362	9.236864	5112.901411	1457.794599
3228	1212.100000	1114.96000	297.912000	5569.550000	3609.710000

	EIF4G1_S5	EMP2_S5	EN01_S5	EPHA2_S2	
EPHB2_S2 \					
1264	12786.100000	2140.220000	45858.200000	1617.790000	504.808000
548	6257.490000	1126.160000	28507.300000	631.663000	450.265000
6812	13284.600000	2677.510000	47065.700000	1069.230000	2076.330000
7863	10527.587171	1362.348884	23591.512322	3201.655891	4132.074258
3228	16915.900000	3410.310000	143228.000000	2470.900000	2208.440000

	EPHB3_S2	EPHB4_S2	ERLIN1_S5	EVI2A_S3	EVI2B_S3
EXOSC8_S5 \					
1264	1241.99000	2685.100000	955.12800	88.141000	157.051000
548	351.86200	1916.860000	1254.88000	81.504900	127.724000
6812	550.29600	13614.800000	2472.19000	200.592000	373.964000
7863	5667.72571	5248.529283	1297.53771	22.786344	81.030267
3228	2232.64000	1957.900000	2386.71000	232.467000	320.068000

	EZR_S5	F3_S5	FAM167A_S5	FAM89B_S5	FARSA_S5
\					
1264	11897.400000	8418.270000	90.544900	1355.770000	1358.970000
548	5497.600000	474.120000	20.873200	1135.600000	1468.080000
6812	17770.400000	1340.240000	69.822500	2065.680000	919.527000
7863	10746.871452	510.121969	39.215757	1077.730029	3382.821044
3228	9246.530000	1477.970000	29.314000	884.874000	1493.310000

	FARSB_S5	FBX041_S5	FCER1G_S3	FCGR1A_S3	FCGR2A_S3
FCGR2B_S3 \					
1264	1342.150000	324.519000	226.76300	27.24360	520.833000

120.994000					
548	1338.370000	51.189000	541.21200	121.24300	434.361000
99.396200					
6812	1501.780000	334.911000	937.27800	286.95300	1111.240000
200.592000					
7863	1201.770979	734.838076	215.44325	30.38253	477.218772
35.273426					
3228	1174.270000	85.215200	552.87600	85.53220	2778.360000
283.255000					

	FCGR3A_S3	FGD1_S2	FGL2_S3	FGR_S3	FLI1_S3
FLNC_S5 \					
1264	465.545000	204.327000	303.6860	165.064000	245.192000
5137.820000					
548	828.964000	830.952000	142.6330	87.965600	137.664000
91.444500					
6812	2390.530000	876.923000	492.3080	343.195000	191.124000
11415.400000					
7863	420.841219	253.368838	250.4135	157.269486	106.790429
994.628822					
3228	1908.480000	415.850000	512.3140	203.494000	254.282000
157.478000					

	FMNL1_S3	FN1_S2	FBNP1_S3	FPR3_S3	FYB_S3
\					
1264	574.103000	52455.900000	741.987000	255.609000	406.250000
548	138.658000	2513.230000	1015.830000	91.444500	315.086000
6812	1997.630000	54245.000000	1497.040000	469.822000	505.917000
7863	425.568343	7291.669478	639.341658	182.866036	84.920137
3228	345.292000	98427.600000	537.537000	563.443000	2971.280000

	GGH_S5	GIMAP4_S3	GLRX3_S5	GNG11_S5	GNPTAB_S3
GPLD1_S5 \					
1264	1058.490000	187.500000	1020.030000	451.122000	1433.490000
4.807700					
548	801.630000	278.806000	952.215000	611.783000	1293.140000
8.945700					
6812	420.710000	485.207000	2571.010000	155.030000	735.503000
8.875700					
7863	725.162161	182.790949	1104.235578	118.798676	664.235627
5.973108					
3228	1462.970000	341.542000	2100.380000	263.826000	2109.590000
9.203200					

	GPNMB_S3	GSTCD_S5	H2AFZ_S5	HAS2_S5	HAUS1_S5 \
1264	9549.680000	254.808000	5116.190000	198.638000	183.494000

548	2307.480000	318.565000	6503.990000	77.032000	531.273000
6812	6445.560000	238.462000	4285.210000	25.390500	150.296000
7863	656.968787	174.494972	4937.989857	53.483964	190.910932
3228	38693.500000	366.425000	5925.860000	753.104000	280.869000

	HCK_S3	HCLS1_S3	HDC_S4	HERC5_S1	HERC6_S1	HLA-DMA_S3 \
1264	326.122000	532.853000	66.506400	175.481000	601.763000	532.853000
548	266.879000	777.775000	1.490900	97.905200	428.397000	2299.030000
6812	675.740000	770.414000	1.775100	63.905300	179.882000	1369.230000
7863	343.856397	342.87048	51.860779	36.200517	333.053243	1201.901095
3228	450.959000	1998.13000	20.110800	247.806000	414.146000	1186.200000

	HLA-DRB1_S3	HMG2_S5	HMHA1_S3	HN1L_S5
	HNRNPA2B1_S5 \			
1264	2939.590000	5672.280000	722.756000	4440.640000
	20217.900000			
548	5338.540000	9221.480000	369.754000	4585.140000
	22632.000000			
6812	12539.900000	5878.700000	881.065000	2697.630000
	12819.500000			
7863	4294.439114	3542.639361	1596.510033	2639.516134
	9956.139522			
3228	4501.430000	6724.500000	230.081000	5817.460000
	10226.500000			

	HNRNPR_S5	HSPB11_S5	HSPG2_S2	HYLS1_S5
	ICAM1_S2 \			
1264	3596.150000	517.628000	12338.200000	217.949000
	1789.260000			
548	3508.680000	528.788000	4327.120000	124.742000
	685.834000			
6812	2786.390000	257.988000	5197.800000	156.805000
	7046.750000			
7863	1741.845332	139.147844	11231.842641	46.389359
	1215.882523			
3228	3903.880000	208.607000	13568.800000	298.935000
	3448.830000			

	ID2_S5	ID3_S5	IFI16_S3	IFI27_S1	IFI30_S3
	\				
1264	681.090000	822.917000	6826.120000	1596.150000	2146.550000
548	3584.230000	4047.910000	1736.450000	4601.050000	2636.480000

6812	600.592000	244.379000	2541.420000	5004.730000	5024.040000
7863	1846.198195	1449.654233	455.437316	8435.411981	4218.586672
3228	686.493000	932.595000	7451.210000	3560.630000	4289.150000
	IFI44_S1	IFI44L_S1	IFI6_S1	IFIH1_S1	IFIT1_S1
IFIT2_S1 \					
1264	956.731000	293.26900	2016.830000	541.667000	358.173000
173.077000					
548	437.840000	476.60500	5028.450000	457.719000	296.698000
255.945000					
6812	846.746000	2188.17000	748.521000	908.876000	218.935000
714.793000					
7863	126.583741	358.27975	3904.530045	365.052522	280.111308
218.129019					
3228	233.830000	89.30550	989.859000	632.297000	137.367000
158.500000					
	IFIT3_S1	IFRD2_S5	IGF2R_S2	IGFBP2_S2	IGFBP3_S2
\					
1264	522.436000	1132.650000	7806.090000	1262.820000	2625.000000
548	367.269000	967.299000	1889.020000	7163.480000	3840.670000
6812	1108.280000	1590.020000	5543.200000	101.183000	2001.180000
7863	746.590341	2946.979229	5759.380637	769.358703	2308.976639
3228	466.979000	821.133000	5435.710000	4615.590000	25963.400000
	IGFBP4_S2	IGFBP5_S2	IGJ_S4	IKZF1_S3	IL10RA_S3
\					
1264	7622.600000	4780.450000	6902.240000	229.167000	295.673000
548	25464.800000	2210.570000	294.213000	124.245000	228.611000
6812	5388.760000	5168.640000	202.367000	296.450000	513.610000
7863	6273.287026	3090.207707	879.747499	108.191389	310.704089
3228	9842.690000	20129.900000	3414.400000	206.562000	290.754000
	IL18_S3	IL7R_S3	IMP4_S5	IP04_S5	
ISG15_S1 \					
1264	874.199000	229.968000	741.186000	1429.780000	820.513000
548	60.134700	31.806800	695.276000	1465.530000	1994.880000

6812	1228.400000	37.869800	848.521000	1385.810000	1202.960000
7863	264.227307	30.011095	1463.603138	2068.202687	3030.829389
3228	539.923000	277.801000	795.228000	1700.700000	467.320000
ITGA3_S2 ITGA4_S2 ITGA5_S2 ITGA6_S5 ITGB2_S2					
\					
1264	15915.100000	282.340000	3802.080000	25782.900000	801.939000
548	918.421000	97.895300	583.456000	2266.230000	1002.910000
6812	15562.700000	183.462000	2215.380000	2964.500000	3699.830000
7863	3940.566087	123.141682	1132.763751	4288.080636	1357.033063
3228	6089.480000	358.493000	6613.380000	13324.200000	1438.310000
ITGB3_S2 ITGB5_S2 ITGB8_S2 JUNB_S5 JUP_S2					
\					
1264	217.949000	5173.880000	1781.250000	3051.280000	20550.500000
548	13.915500	2538.580000	906.493000	3195.090000	5095.050000
6812	91.716000	2510.060000	1984.020000	1322.490000	12150.900000
7863	50.598276	2787.829954	214.333668	5502.742214	19274.491393
3228	191.905000	2923.220000	975.884000	2970.600000	33214.500000
KIAA0090_S5 KRR1_S5 LAIR1_S3 LAMA4_S2 LAMB1_S2 \					
1264	2780.450000	1008.450000	183.49400	2374.200000	5812.500000
548	1177.350000	633.680000	313.09800	421.440000	7204.730000
6812	1236.090000	564.284000	499.40800	483.432000	4458.580000
7863	720.656034	352.075264	208.89802	783.651454	5772.950729
3228	4090.670000	1403.150000	293.82200	2424.540000	5426.160000
LAPTM5_S3 LCK_S4 LCP1_S3 LCP2_S3 LHFPL2_S3 \					
1264	1458.330000	125.801000	1877.400000	274.038000	1033.650000
548	2701.590000	130.706000	2575.350000	169.967000	733.544000
6812	6992.900000	328.994000	3215.380000	467.450000	971.598000
7863	1804.768877	147.246087	817.064039	158.550938	716.245256
3228	3877.970000	159.523000	3375.880000	522.539000	1385.600000
LILRB4_S3 LMNB2_S5 LOXL2_S5 LRMP_S3					
LRP1_S2 \					
1264	97.756400	3442.310000	1612.100000	52.083300	5866.990000

548	244.018000	2038.620000	444.301000	29.818900	2684.190000
6812	468.639000	2414.790000	1230.070000	21.301800	3021.890000
7863	202.107483	4983.516023	1275.832329	23.441881	11460.799714
3228	326.204000	3216.020000	2383.960000	326.885000	13292.900000

	LRRC17_S2	LRRC40_S5	LSM3_S5	LSM4_S5	LST1_S3
LTB_S4 \					
1264	420.67300	415.064000	673.878000	1681.090000	126.60300
	374.199000				
548	177.42200	448.774000	945.258000	3186.140000	89.45660
	162.016000				
6812	79.88170	540.237000	910.651000	1820.710000	170.41400
	1493.490000				
7863	27.08007	48.738581	469.580308	2784.870978	104.39482
	218.043389				
3228	117.25600	318.705000	682.403000	1516.150000	75.67110
	22.837700				

	LY86_S3	LYAR_S5	LYN_S3	LYZ_S3	
MAGOHB_S5 \					
1264	25.641000	749.199000	883.814000	13019.200000	275.641000
548	79.020000	386.651000	773.799000	834.431000	372.239000
6812	103.550000	1045.560000	2642.600000	6668.050000	364.497000
7863	38.308787	678.810049	562.685169	3068.651264	192.543608
3228	51.129100	860.332000	1834.510000	6712.910000	458.798000

	MAP3K8_S5	MAPRE1_S5	MARCKSL1_S2	MARVELD2_S5	MCM3_S5
\					
1264	154.647000	3822.120000	1948.720000	252.404000	4629.810000
548	268.370000	2547.030000	17631.900000	424.919000	3315.860000
6812	117.160000	3569.820000	7843.200000	400.592000	2574.560000
7863	80.427806	4168.603566	5833.127538	507.566001	3776.425366
3228	164.977000	7763.100000	1351.850000	402.897000	1764.290000

	MCM7_S5	MCTS1_S5	MERTK_S3	MET_S5	MFSD11_S5
\					

1264	2780.830000	1647.440000	106.570000	3673.080000	856.763000
548	2835.220000	1015.330000	442.313000	1659.920000	312.591000
6812	12199.600000	1617.160000	321.894000	40805.900000	667.325000
7863	7193.775818	891.683567	71.470187	2531.035123	445.909178
3228	3067.410000	774.435000	337.452000	4123.390000	753.769000
	MKKS_S5	MLF1IP_S5	MMP11_S2	MMP1_S2	
MMP14_S2 \					
1264	1009.620000	614.728000	8549.680000	33935.100000	
19246	0.000000				
548	1106.780000	681.395000	15438.200000	13.915500	
7597	3.500000				
6812	592.308000	610.467000	7279.290000	3834.910000	
29959	8.000000				
7863	433.255505	269.988979	6572.061363	1598.728465	
7607	7.90173				
3228	662.292000	379.163000	1861.780000	23431.800000	
12179	0.000000				
	MMP17_S2	MMP19_S2	MMP2_S2	MMP3_S2	MNAT1_S5 \
1264	199.51900	592.949000	11664.300000	5686.700000	314.103000
548	36.27960	793.181000	10317.300000	102.378000	483.562000
6812	90.53250	338.462000	4640.240000	13.017800	378.106000
7863	26.74886	308.075381	6768.362318	896.832598	90.714612
3228	66.46780	371.879000	14209.500000	1543.080000	366.425000
	MNDA_S3	MPP1_S3	MRPL12_S5	MRPL37_S5	MRPS16_S5 \
1264	84.134600	214.744000	1384.620000	2020.830000	1345.720000
548	57.152800	176.925000	1809.510000	1491.940000	2112.120000
6812	324.852000	820.710000	1376.920000	1696.450000	1979.640000
7863	36.692463	862.539391	3222.565362	3021.732273	2019.185734
3228	133.617000	245.079000	1209.710000	1497.060000	2782.770000
	MRPS28_S5	MRT04_S5	MS4A1_S4	MSN_S3	MT1F_S5 \
1264	588.878000	1703.530000	43.269200	11398.200000	167.468000
548	418.667000	1512.310000	1.987900	5326.140000	697.264000
6812	261.840000	1215.380000	11.834300	25971.600000	153.254000
7863	158.477752	976.047526	4.576277	2585.419706	459.608169
3228	461.283000	1683.510000	11.248400	17127.900000	172.135000
	MT3_S5	MTA1_S2	MTHFD1_S5	MX1_S1	MYBL1_S5 \
1264	3.205100	1064.900000	1481.530000	1167.470000	238.782000
548	147.106000	1970.030000	1389.490000	1137.590000	64.110500
6812	0.000000	1739.050000	2720.590000	1135.500000	431.361000
7863	91.901208	1555.890562	1733.931097	2361.615361	34.276176

3228	0.340900	998.722000	2440.870000	916.915000	78.738800
	MYBL2_S5	MYCBP_S5	MYL6_S5	MY01F_S3	
NCEH1_S5 \					
1264	1282.850000	720.353000	15859.000000	259.61500	411.859000
548	1157.970000	1166.910000	18998.100000	301.66700	694.282000
6812	1129.590000	715.414000	18542.000000	450.29600	1969.230000
7863	4624.135012	617.681299	14625.742112	438.45008	451.314543
3228	3666.640000	489.817000	13088.400000	201.44900	1193.350000
	NCF2_S3	NCKAP1L_S3	NCLN_S5	NE01_S2	
NID1_S2 \					
1264	375.801000	237.179000	2168.270000	1062.500000	1839.500000
548	82.001800	190.344000	2321.400000	2178.760000	1702.100000
6812	771.006000	445.562000	1911.240000	804.142000	1350.880000
7863	152.308393	256.994692	6476.962655	3104.024952	1136.524032
3228	1061.100000	351.087000	1691.690000	782.957000	5154.120000
	NLN_S5	NME1_S5	NOP16_S5	NOTCH2_S2	NPC2_S3 \
1264	435.096000	1711.670000	379.006000	2720.350000	2388.620000
548	422.434000	3232.230000	589.916000	2957.530000	6377.260000
6812	600.592000	2551.910000	322.485000	3546.150000	7479.290000
7863	374.185195	2190.579905	734.285336	712.373458	1552.987229
3228	511.973000	1949.150000	258.372000	3836.730000	6966.510000
	NPL_S3	NRIP3_S5	NUDT1_S5	NUDT15_S5	NUP107_S5
NUP35_S5 \					
1264	213.141000	96.955100	445.513000	854.199000	1161.860000
250.801000					
548	130.706000	16.400400	327.510000	688.771000	1000.420000
360.808000					
6812	120.118000	134.320000	288.757000	879.846000	1000.590000
251.479000					
7863	88.620145	14.386538	402.260555	600.915288	711.490968
148.455339					
3228	691.947000	247.124000	491.521000	854.865000	1263.570000
408.010000					
	NUP85_S5	NUP93_S5	NUPL1_S5	NUTF2_S5	
OAS1_S1 \					
1264	905.449000	1128.210000	1248.400000	1809.290000	560.096000

548	726.089000	1247.420000	756.405000	3353.130000	193.326000
6812	890.533000	1489.940000	1099.410000	3146.750000	1168.050000
7863	965.871197	1438.451723	1158.237576	3557.105678	1869.724499
3228	727.397000	1467.410000	1026.670000	2287.860000	477.887000

	OAS2_S1	OAS3_S1	OASL_S1	OSBPL3_S3	
PA2G4_S5 \					
1264	1220.350000	1335.740000	160.256000	1112.980000	2838.140000
548	385.657000	634.148000	114.306000	428.894000	4439.530000
6812	1243.200000	1981.070000	281.657000	815.385000	5613.610000
7863	935.073373	2756.552264	533.469981	596.266612	5580.006673
3228	1499.450000	1513.420000	82.147400	468.002000	4908.050000

	PAICS_S5	PAK1_S2	PAK2_S2	PDAP1_S5	
PDIA4_S5 \					
1264	3400.640000	1366.190000	3371.790000	2505.610000	5277.240000
548	4425.120000	2163.850000	2059.490000	3060.410000	12363.900000
6812	2313.610000	2145.560000	4628.990000	10816.600000	16620.700000
7863	3331.453327	1544.604524	1565.465184	3126.018611	6665.578363
3228	5149.380000	2339.330000	8009.890000	1954.840000	6801.870000

	PDLIM7_S5	PFKP_S5	PFN1_S5	PGM2_S5	PHF19_S5
\					
1264	3774.840000	1290.870000	18421.500000	1930.290000	897.43600
548	1167.900000	834.431000	14217.600000	537.733000	265.88500
6812	1638.460000	3407.100000	12803.000000	1115.980000	1609.47000
7863	1790.815205	3396.808005	21598.814199	543.057721	1775.78342
3228	838.858000	4072.260000	8874.990000	2447.720000	550.49000

	PIK3CG_S3	PITPNC1_S5	PLAT_S2	PLAUR_S5	PLCG2_S3
PLEK_S3 \					

1264	64.90380	311.699000	434.295000	758.013000	356.570000
229.167000					
548	7.95170	519.345000	550.158000	603.832000	254.951000
299.679000					
6812	47.33730	175.148000	2979.880000	2352.660000	207.101000
688.757000					
7863	23.68202	214.514288	659.910176	2324.934366	114.980755
161.426919					
3228	99.19050	411.760000	3454.960000	1480.700000	536.515000
561.057000					

	PLG_S5	PLK4_S5	PL0D2_S5	PNN_S5	PN01_S5
POLE2_S5 \					
1264	0.000000	506.410000	1907.850000	2253.210000	594.551000
126.60300					
548	23.855100	258.927000	2288.100000	2935.170000	393.112000
89.45660					
6812	0.000000	369.231000	1333.730000	1604.730000	665.681000
140.82800					
7863	3.938344	153.655922	439.280843	1478.458546	599.694356
67.87253					
3228	0.000000	387.218000	7612.440000	1761.910000	1266.980000
168.72600					

	POLR3K_S5	PPIH_S5	PSMA7_S5	PSMC3_S5	PSMD12_S5 \
1264	274.038000	492.78800	4498.400000	2745.990000	1370.990000
548	389.633000	743.98000	2975.920000	2598.220000	723.107000
6812	243.787000	580.47300	4809.470000	1921.890000	1149.110000
7863	574.739875	652.58254	12850.164704	1866.885196	501.157671
3228	508.223000	320.75000	6267.750000	2897.320000	2220.710000

	PSMD14_S5	PSMD2_S5	PTPLB_S5	PTPRC_S3	
PXN_S2 \					
1264	1602.560000	6587.340000	308.494000	516.026000	6759.01000
548	1436.770000	3072.830000	43.237300	339.438000	1946.43000
6812	1317.160000	5385.800000	243.195000	875.148000	7575.14000
7863	978.831589	4231.503959	75.527356	208.733265	7988.21454
3228	3146.480000	14749.000000	605.709000	1036.560000	4754.74000

	RAB3B_S5	RAC1_S2	RASSF2_S3	RASSF4_S3	RBM14_S5 \
1264	5.609000	6165.060000	365.385000	736.114000	1018.350000
548	1.987900	6536.290000	648.063000	825.982000	1355.860000
6812	26.627200	6388.170000	122.485000	1518.930000	1006.050000
7863	7.277026	7220.930029	122.210979	772.989068	2152.712306
3228	16.020500	11040.800000	252.578000	215.083000	1341.370000

	RFC3_S5	RGS8_S5	RHOA_S2	RHOB_S2	RHOC_S5
RHOG_S2 \					
1264	365.385000	0.0000	9258.810000	2931.09000	3935.10000
1266.830000					
548	442.810000	0.0000	9805.930000	11439.50000	3642.37000
898.541000					
6812	404.734000	0.0000	13426.000000	1077.51000	6773.37000
1168.050000					
7863	546.375419	NaN	9807.319352	3607.80426	5682.28049
1714.439988					
3228	546.740000	0.3409	10884.400000	2874.14000	3976.82000
885.215000					
	RH0Q_S2	RMND5B_S5	RNASE6_S3	RND3_S2	RNF138_S5 \
1264	1629.020000	351.763000	143.429000	1867.790000	368.590000
548	1649.260000	622.717000	124.742000	2087.320000	763.363000
6812	1291.900000	531.953000	175.148000	1360.360000	431.361000
7863	360.002602	937.189697	142.656267	438.776252	284.101016
3228	1348.960000	273.711000	116.574000	5216.870000	601.278000
	RNF41_S5	RPN1_S5	RPP40_S5	RSAD2_S1	RTP4_S1 \
1264	684.295000	6653.040000	213.141000	165.865000	92.147400
548	879.159000	11263.100000	134.682000	90.947500	193.326000
6812	828.402000	7985.800000	184.615000	308.876000	133.136000
7863	1102.745864	7524.043806	165.900124	630.334114	123.607153
3228	1178.700000	15401.800000	252.578000	294.844000	259.395000
	RUNX3_S3	RUVBL1_S5	SAMD9_S1	SAMHD1_S3	SAMSN1_S3 \
1264	657.853000	1214.70000	2354.970000	495.633000	111.378000
548	193.822000	1955.01000	236.066000	1195.850000	123.748000
6812	204.142000	1519.73000	305.325000	756.562000	193.491000
7863	1165.381881	1381.89423	380.608313	628.068511	74.518924
3228	1033.150000	2640.05000	1423.430000	1060.230000	184.406000
	SAR1A_S5	SAR1B_S5	SDC1_S5	SELL_S3	SELPLG_S3 \
1264	2307.690000	526.44200	16976.800000	138.622000	278.045000
548	2155.900000	437.84000	2425.760000	120.269000	216.187000
6812	1915.380000	364.49700	10459.200000	159.172000	585.799000
7863	1033.663388	291.35525	7002.780348	51.673906	422.963295
3228	3508.140000	580.82700	22700.000000	144.184000	383.468000
	SEMA3F_S2	SERPINE1_S2	SH2B3_S3	SH3BP5L_S5	
SKA1_S5 \					
1264	1383.810000	5875.00000	687.500000	1310.900000	93.750000
548	972.095000	117.28700	254.951000	594.389000	108.839000
6812	575.148000	1481.66000	1301.780000	553.254000	128.994000

7863	536.348425	237.63618	504.611923	983.091103	157.042894
3228	1699.530000	6642.01000	707.627000	593.779000	261.099000
SKA2_S5 SLC16A1_S5 SLC1A3_S3 SLC25A40_S5 SLC25A5_S5					
\					
1264	786.859000	3054.490000	477.564000	132.212000	11109.000000
548	842.880000	1665.380000	179.410000	59.140700	4494.690000
6812	793.491000	2778.700000	168.047000	346.154000	8710.650000
7863	216.737482	1117.373004	26.435782	120.723276	18811.796336
3228	831.018000	6389.430000	1772.130000	227.695000	10477.400000
SLC7A7_S3 SMC2_S5 SMO_S2 SMS_S5 SMURF2_S5 \					
1264	176.28200	1104.17000	475.160000	888.622000	483.974000
548	183.88300	605.82000	1728.000000	2884.480000	338.444000
6812	632.54400	1209.47000	931.953000	3064.500000	1871.010000
7863	363.96797	315.73208	326.766597	1884.414469	243.065824
3228	353.81300	2964.47000	509.587000	2040.730000	571.283000
SNRPA1_S5 SNRPA_S5 SNRPB_S5 SNRPC_S5					
SNRPD1_S5 \					
1264	424.680000	1115.380000	3641.030000	1903.040000	712.340000
548	595.880000	2843.230000	4689.010000	1951.640000	1430.310000
6812	614.201000	1108.280000	4876.920000	2188.760000	981.065000
7863	537.599672	2810.940332	4348.216621	2118.107643	686.240596
3228	855.901000	1157.900000	2653.260000	2204.690000	1225.050000
SNRPE_S5 SNX17_S5 SP140_S3 SPAG17_S5 SPARC_S2 \					
1264	600.962000	2597.760000	53.685900	41.666700	43480.000000
548	1544.620000	2713.520000	26.340000	289.740000	10374.000000
6812	1222.490000	2168.640000	97.041400	1.775100	12467.500000
7863	992.908169	2212.668745	21.068411	1.794282	12455.566615
3228	971.794000	2252.070000	33.063500	68.172100	53367.500000
SRM_S5 STAT1_S1 STK17A_S5 STRA13_S5					
SYK_S3 \					
1264	1790.060000	3429.490000	1818.110000	697.917000	607.372000
548	1435.280000	4222.850000	463.186000	896.553000	1712.600000

6812	2275.150000	8585.800000	2156.800000	507.692000	547.337000
7863	2229.710794	3373.295658	204.670983	944.119588	1408.615337
3228	2054.370000	7263.060000	1204.600000	260.418000	1351.850000
	TAGLN_S5	TAP1_S1	TBXAS1_S3	TCEB1_S5	TCF7L2_S2 \
1264	7978.370000	3900.640000	156.250000	1051.280000	436.699000
548	899.535000	4301.370000	378.202000	883.632000	704.222000
6812	1727.810000	7538.460000	170.414000	1136.090000	855.621000
7863	5412.55443	11272.08586	1303.232132	570.101003	977.233934
3228	5734.300000	3268.850000	170.089000	1231.530000	709.331000
	THBS1_S2	THBS2_S2	TIMP1_S2	TLR2_S3	TMEM130_S5
\					
1264	16249.200000	6189.100000	4869.390000	669.071000	9.61540
548	1140.570000	136.670000	4652.730000	343.911000	5.96380
6812	1164.500000	2704.730000	6509.470000	218.935000	27.81070
7863	2851.471087	1790.479229	8027.404873	107.980266	14.09625
3228	8540.260000	14896.000000	7015.590000	1279.930000	54.53770
	TNC_S2	TNFAIP3_S3	TNFRSF12A_S5	TNFRSF1A_S2	
TNFRSF1B_S3 \					
1264	51591.300000	1375.800000	3986.380000	3036.860000	
710.737000					
548	1885.550000	1716.570000	1070.000000	2660.840000	
1382.600000					
6812	18590.500000	947.929000	2894.080000	4164.500000	
1972.780000					
7863	2139.767469	1004.281681	2584.787646	3277.586534	
2469.421757					
3228	36879.800000	1954.840000	1190.290000	5124.500000	
652.067000					
	TNFSF13B_S3	TOMM40_S5	TPI1_S5	TPM1_S5	TPM2_S5
\					
1264	26.442300	1080.930000	6519.230000	7943.420000	10874.20000
548	18.885300	1375.150000	11693.000000	1983.870000	1485.48000
6812	99.408300	1551.480000	15510.100000	3089.760000	4177.51000
7863	28.323496	2911.791666	13301.708123	4638.258269	1984.92103
3228	72.262500	1441.840000	21580.200000	2700.970000	1286.41000

\					
	TPM3_S5	TPRKB_S5	TRA2B_S5	TUBA4A_S5	TUBG1_S5
1264	9447.920000	412.660000	3585.740000	7373.400000	1071.310000
548	7467.630000	722.113000	3100.170000	119.275000	984.022000
6812	8412.430000	335.503000	1897.630000	1947.930000	1028.400000
7863	8429.630182	211.853076	2050.314713	5324.263102	1245.895273
3228	10409.500000	435.620000	3686.750000	3058.880000	1813.040000
\					
	UAP1_S5	UBE2J1_S5	UMPS_S5	UQCR10_S5	USPL1_S5
1264	1302.880000	1580.93000	970.353000	1076.920000	534.455000
548	1015.830000	2069.93000	620.729000	1863.180000	506.921000
6812	999.408000	2207.10000	849.112000	1656.800000	478.106000
7863	576.843949	1253.38716	781.709909	1179.679495	286.464799
3228	1534.210000	2274.56000	1168.130000	743.076000	164.295000
\					
	VCAN_S2	VDAC1_S5	VSIG4_S3	VTA1_S5	WDHD1_S5
WDR54_S5					
1264	4942.310000	4340.540000	81.73080	1551.28000	425.481000
296.47400					
548	7417.440000	4943.470000	348.88100	1079.44000	187.362000
766.84100					
6812	2764.500000	4909.470000	621.89400	1128.40000	508.284000
151.47900					
7863	1249.379279	7151.557949	160.41858	716.27527	101.055152
140.38918					
3228	14117.100000	5529.780000	479.93200	1951.09000	611.163000
215.08300					
\					
	WDR77_S5	WIPF1_S3	WNT2B_S2	WNT8B_S2	WSB2_S5
ZWILCH_S5					
1264	906.250000	989.583000	28.044900	6.410300	1991.190000
483.061000					
548	4555.820000	750.441000	16.897300	1.490900	1495.420000
452.988000					
6812	1057.400000	1007.100000	5.325400	1.775100	2242.600000
325.953000					
7863	1725.383932	327.268858	31.568168	2.028401	2127.285705
266.422502					
3228	842.948000	1371.620000	249.169000	0.000000	2586.450000
498.921000					
\					
	ZYX_S2				
1264	5231.570000				

```
548    3509.180000
6812   7905.920000
7863   6938.987139
3228   3666.980000
```

```
# Keep only the required columns
```

```
df_final = df_balanced.set_index("TCGA Participant Barcode")
[column_names_PanCan + ["Immune Subtype"]]
```

```
# Display the first few rows
```

```
df_final.head()
```

```
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/
format.py:1458: RuntimeWarning: invalid value encountered in greater
    has_large_values = (abs_vals > 1e6).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in less
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
/usr/local/lib/python3.10/dist-packages/pandas/io/formats/format.py:14
59: RuntimeWarning: invalid value encountered in greater
    has_small_values = ((abs_vals < 10 ** (-self.digits)) & (abs_vals >
0)).any()
```

	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5
ADCY7_S3 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1414.260000	5574.520000	942.308000
946.314000			
TCGA-B5-A3F9	1354.770000	1138.580000	1640.530000
1227.540000			
TCGA-49-6767	1637.870000	5176.330000	161.538000
1004.730000			
TCGA-AA-A03J	930.343078	1329.678264	216.256995
370.030352			
TCGA-77-7337	3289.310000	14204.300000	2384.660000
573.328000			

	AIMP2_S5	ALKBH7_S5	ALOX5AP_S3
AMPD3_S3 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	563.534000	521.635000	237.1790
587.340000			
TCGA-B5-A3F9	482.911000	872.698000	185.8710
353.850000			
TCGA-49-6767	488.988000	299.408000	482.2480
276.331000			
TCGA-AA-A03J	1349.628189	1956.948165	131.8582

149.806773			
TCGA-77-7337	1188.010000	317.341000	743.7580
308.820000			
	APITD1_S5	APOC1_S3	APOE_S3
AP00_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	216.987000	138.622000	1125.000000
110.577000			
TCGA-B5-A3F9	340.800000	1306.070000	4475.310000
620.729000			
TCGA-49-6767	274.065000	1394.080000	5431.360000
372.781000			
TCGA-AA-A03J	125.083134	400.202368	2667.522991
252.673868			
TCGA-77-7337	368.092000	126.118000	620.366000
315.296000			
	ARHGAP1_S2	ARHGAP15_S3	ARHGDIA_S2
ARRB2_S3 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	3838.940000	44.070500	8888.620000
523.237000			
TCGA-B5-A3F9	1648.980000	12.921500	6709.740000
521.333000			
TCGA-49-6767	1626.630000	71.597600	5318.340000
695.858000			
TCGA-AA-A03J	3276.934604	12.524268	15147.247057
1306.517461			
TCGA-77-7337	4495.270000	28.632300	8677.970000
564.124000			
	B2M_S3	BCCIP_S5	BRCA2_S5
BRIP1_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	48758.800000	951.234000	224.359000
169.071000			
TCGA-B5-A3F9	66318.100000	1651.590000	70.571300
51.189000			
TCGA-49-6767	53988.100000	1667.410000	109.468000
92.307700			
TCGA-AA-A03J	41306.987107	769.615704	146.312249
36.943463			
TCGA-77-7337	33636.100000	1540.930000	147.593000
209.970000			
	BSG_S2	BTK_S3	C11orf24_S5

C12orf24_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	16931.90000	86.538500	1333.330000
66.50640			
TCGA-B5-A3F9	21927.30000	60.631700	754.417000
93.92940			
TCGA-49-6767	8614.20000	156.805000	1263.310000
68.04730			
TCGA-AA-A03J	19473.23631	28.087146	1035.470316
53.88891			
TCGA-77-7337	13157.20000	85.896900	1387.980000
84.19260			

C13orf1_S5 C13orf18_S3 C13orf27_S5

C16orf61_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	310.096000	127.404000	193.910000
358.758000			
TCGA-B5-A3F9	273.836000	199.786000	180.404000
548.716000			
TCGA-49-6767	381.657000	38.461500	166.864000
548.941000			
TCGA-AA-A03J	492.691892	2546.069144	117.261586
479.162611			
TCGA-77-7337	330.976000	44.652700	250.533000
565.549000			

C19orf48_S5 C1orf54_S3 C3AR1_S3

C3orf26_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	976.763000	155.449000	133.814000
787.660000			
TCGA-B5-A3F9	1952.140000	203.265000	154.561000
635.141000			
TCGA-49-6767	863.314000	137.278000	370.414000
468.047000			
TCGA-AA-A03J	4435.195669	66.573924	130.132203
375.363702			
TCGA-77-7337	1145.290000	81.806600	369.493000
1113.930000			

CASP1_S3 CBX1_S5 CCBL2_S5

CCL2_S3 \

TCGA Participant Barcode

TCGA-WA-A7H4	790.865000	1612.180000	716.090000
351.763000			

TCGA-B5-A3F9	120.766000	2532.120000	626.022000	
620.232000				
TCGA-49-6767	199.408000	1936.090000	795.284000	
642.604000				
TCGA-AA-A03J	463.236328	620.153334	424.498066	
122.172779				
TCGA-77-7337	672.177000	2124.930000	621.464000	
1044.060000				
	CCL5_S3	CCRL2_S3	CCT5_S5	
CD14_S4 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	1123.400000	36.057700	7229.970000	
1347.76000				
TCGA-B5-A3F9	500.957000	27.333900	6531.320000	
1638.55000				
TCGA-49-6767	2890.530000	71.005900	10427.200000	
1537.87000				
TCGA-AA-A03J	607.562311	370.657909	8111.215987	
1392.85134				
TCGA-77-7337	555.944000	62.377500	12827.600000	
1825.99000				
	CD163_S3	CD19_S4	CD247_S4	CD33_S3
\				
TCGA Participant Barcode				
TCGA-WA-A7H4	320.513000	9.615400	101.763000	31.250000
TCGA-B5-A3F9	583.456000	5.466800	56.158800	67.092400
TCGA-49-6767	834.319000	1.775100	133.728000	71.005900
TCGA-AA-A03J	164.663887	8.810904	71.947677	30.009939
TCGA-77-7337	1267.660000	3.408600	63.400100	46.357100
	CD37_S4	CD3D_S4	CD3E_S4	CD3G_S4
\				
TCGA Participant Barcode				
TCGA-WA-A7H4	169.8720	92.948700	187.500000	14.423100
TCGA-B5-A3F9	190.8410	93.432400	194.320000	4.472800
TCGA-49-6767	450.8880	200.592000	602.367000	56.804700
TCGA-AA-A03J	246.5297	109.859681	189.354061	17.990105

TCGA-77-7337	182.0200	79.420500	256.327000	41.244100
CD53_S3 \	CD44_S2	CD48_S3	CD52_S3	
TCGA Participant Barcode				
TCGA-WA-A7H4	18930.300000	94.551300	186.699000	
338.14100				
TCGA-B5-A3F9	2058.990000	157.543000	171.458000	
465.17400				
TCGA-49-6767	9745.560000	253.846000	629.586000	
863.90500				
TCGA-AA-A03J	3261.290314	96.453995	268.239955	
315.95604				
TCGA-77-7337	23292.700000	132.595000	219.514000	
878.05700				
CD84_S3 \	CD59_S2	CD79A_S4	CD79B_S4	
TCGA Participant Barcode				
TCGA-WA-A7H4	14369.400000	133.814000	56.089700	
20.833300				
TCGA-B5-A3F9	6228.660000	16.400400	18.388300	
28.824900				
TCGA-49-6767	6684.020000	37.869800	22.485200	
62.721900				
TCGA-AA-A03J	2817.520364	64.726949	33.807438	
25.931826				
TCGA-77-7337	13698.500000	174.862000	31.359200	
117.597000				
CD97_S3 \	CD86_S3	CD8A_S3	CD8B_S4	
TCGA Participant Barcode				
TCGA-WA-A7H4	233.173000	33.65380	24.038500	
761.218000				
TCGA-B5-A3F9	104.863000	197.79800	229.605000	
1288.170000				
TCGA-49-6767	304.734000	652.07100	97.041400	
1813.020000				
TCGA-AA-A03J	77.030344	276.25988	77.724818	
6066.851717				
TCGA-77-7337	397.103000	208.60700	102.940000	
1295.950000				
CECR1_S3 \	CDCA4_S5	CDH6_S2	CDK2_S5	

TCGA Participant Barcode

TCGA-WA-A7H4	1024.840000	84.13460	818.910000
422.276000			
TCGA-B5-A3F9	685.834000	69.57730	876.177000
1813.480000			
TCGA-49-6767	786.982000	11.83430	1046.150000
971.598000			
TCGA-AA-A03J	480.374708	23.29196	787.869981
447.176864			
TCGA-77-7337	1040.310000	2406.48000	923.392000
659.225000			

CEL F2_S3 CENPJ_S5 CENPN_S5 CENPO_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	397.436000	226.763000	702.941000
202.780000			
TCGA-B5-A3F9	328.007000	616.256000	295.157000
74.810500			
TCGA-49-6767	270.414000	89.349100	584.787000
145.675000			
TCGA-AA-A03J	700.887602	196.492764	630.422866
190.410746			
TCGA-77-7337	306.775000	107.712000	1274.420000
436.735000			

CENPW_S5 CEP78_S5 CHEK1_S5 CKLF_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	568.109000	243.590000	537.660000
213.14100			
TCGA-B5-A3F9	454.240000	284.273000	269.364000
227.61700			
TCGA-49-6767	353.846000	223.077000	502.959000
897.04100			
TCGA-AA-A03J	931.541342	89.391743	175.446418
465.70463			
TCGA-77-7337	1029.400000	493.566000	662.633000
235.19400			

COL16A1_S2 COL1A2_S2 COL3A1_S2 \

TCGA Participant Barcode

TCGA-WA-A7H4	5395.030000	93153.000000	103386.000000
TCGA-B5-A3F9	182.392000	10381.900000	22155.900000
TCGA-49-6767	710.059000	21293.500000	21016.000000
TCGA-AA-A03J	681.668235	30497.210498	23821.008655
TCGA-77-7337	862.378000	105622.000000	155202.000000

	COL6A1_S2	COL6A3_S2	COL8A1_S2
COPS6_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	20055.300000	26830.900000	208.33300
1607.370000			
TCGA-B5-A3F9	9287.580000	1272.270000	9.93960
2249.340000			
TCGA-49-6767	18462.100000	8066.860000	395.85800
8586.390000			
TCGA-AA-A03J	8179.105901	8199.892726	237.28691
2323.238838			
TCGA-77-7337	6743.590000	34333.200000	896.12300
1875.760000			

	CQ2_S5	COR01A_S3	COR01C_S5
COTL1_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	274.038000	713.141000	3767.630000
2705.130000			
TCGA-B5-A3F9	250.478000	567.552000	1157.970000
3072.830000			
TCGA-49-6767	269.822000	1411.830000	5149.700000
9766.860000			
TCGA-AA-A03J	333.605287	3106.212752	3463.398929
2085.454571			
TCGA-77-7337	253.941000	495.271000	8162.930000
3322.710000			

	COX17_S5	CPEB4_S5	CPVL_S3
CSF1R_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	538.462000	811.699000	203.52600
724.359000			
TCGA-B5-A3F9	778.769000	460.701000	230.10200
653.033000			
TCGA-49-6767	839.645000	245.562000	393.49100
988.757000			
TCGA-AA-A03J	591.715669	303.291589	222.88383
671.456392			
TCGA-77-7337	399.489000	894.759000	299.95700
1158.930000			

	CTNNA1_S2	CTNNB1_S2	CTSC_S3
CTSL1_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4 1512.820000	8176.2800	8176.970000	5598.560000
TCGA-B5-A3F9 2101.730000	7995.4300	6567.580000	3442.090000
TCGA-49-6767 1643.790000	6928.4000	13272.800000	4711.830000
TCGA-AA-A03J 658.990557	8261.3316	9974.124921	2538.514223
TCGA-77-7337 2636.220000	7950.5800	12085.200000	11806.700000

CTSS_S3 CXCL10_S1 CXCR4_S3
CYBB_S3 \

TCGA Participant Barcode

TCGA-WA-A7H4 471.154000	835.737000	187.500000	298.077000
TCGA-B5-A3F9 468.156000	1766.270000	308.625000	2493.850000
TCGA-49-6767 1659.760000	10181.700000	2600.000000	601.775000
TCGA-AA-A03J 350.122639	3005.643104	304.784386	270.766523
TCGA-77-7337 1642.610000	2027.100000	568.215000	1160.630000

CYTIP_S3 DAPK1_S3 DBNDD1_S5
DCBLD2_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4 3229.170000	129.80800	905.449000	197.115000
TCGA-B5-A3F9 2031.160000	163.50700	462.689000	824.988000
TCGA-49-6767 24820.100000	208.87600	169.822000	862.130000
TCGA-AA-A03J 503.510219	53.78859	304.985752	337.246512
TCGA-77-7337 2947.080000	252.91900	320.409000	167.703000

DCK_S5 DCLRE1B_S5 DDX58_S1
DDX60_S1 \

TCGA Participant Barcode

TCGA-WA-A7H4 713.141000	548.87800	323.718000	770.833000
TCGA-B5-A3F9 578.983000	357.82600	238.551000	387.645000
TCGA-49-6767	872.18900	264.497000	452.663000

763.905000			
TCGA-AA-A03J	352.77352	142.552052	396.642255
705.860341			
TCGA-77-7337	997.01800	276.438000	222.923000
731.487000			
	DHFR_S5	DLEU1_S5	DLEU2_S5
DOCK2_S3 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	225.561000	80.769200	69.487200
203.526000			
TCGA-B5-A3F9	422.026000	201.471000	49.285600
100.390000			
TCGA-49-6767	227.876000	92.112400	107.444000
321.302000			
TCGA-AA-A03J	388.993479	134.812507	25.400413
146.765031			
TCGA-77-7337	175.202000	170.703000	102.940000
315.296000			
	DSP_S2	DUT_S5	DVL1_S2
DVL3_S2 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	44610.600000	1073.000000	2220.350000
2595.350000			
TCGA-B5-A3F9	4294.410000	1422.170000	1487.960000
2026.190000			
TCGA-49-6767	8887.570000	624.645000	1872.780000
2149.110000			
TCGA-AA-A03J	8165.998316	1715.641893	3209.336764
2974.540775			
TCGA-77-7337	52847.400000	577.541000	804.431000
4245.420000			
	DYNLT1_S5	EBNA1BP2_S5	EFNA5_S2
EIF2AK1_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1262.020000	2073.72000	518.429000
3613.550000			
TCGA-B5-A3F9	2670.780000	1220.58000	64.607500
5030.100000			
TCGA-49-6767	1210.650000	1886.98000	18.343200
4418.110000			
TCGA-AA-A03J	1160.197487	1246.37362	9.236864
5112.901411			
TCGA-77-7337	1212.100000	1114.96000	297.912000
5569.550000			

	EIF4EBP1_S5	EIF4G1_S5	EMP2_S5 \
TCGA Participant Barcode			
TCGA-WA-A7H4	1009.620000	12786.100000	2140.220000
TCGA-B5-A3F9	1500.880000	6257.490000	1126.160000
TCGA-49-6767	1133.140000	13284.600000	2677.510000
TCGA-AA-A03J	1457.794599	10527.587171	1362.348884
TCGA-77-7337	3609.710000	16915.900000	3410.310000
	EN01_S5	EPHA2_S2	EPHB2_S2
EPHB3_S2 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	45858.200000	1617.790000	504.808000
1241.99000			
TCGA-B5-A3F9	28507.300000	631.663000	450.265000
351.86200			
TCGA-49-6767	47065.700000	1069.230000	2076.330000
550.29600			
TCGA-AA-A03J	23591.512322	3201.655891	4132.074258
5667.72571			
TCGA-77-7337	143228.000000	2470.900000	2208.440000
2232.64000			
	EPHB4_S2	ERLIN1_S5	EVI2A_S3
EVI2B_S3 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	2685.100000	955.12800	88.141000
157.051000			
TCGA-B5-A3F9	1916.860000	1254.88000	81.504900
127.724000			
TCGA-49-6767	13614.800000	2472.19000	200.592000
373.964000			
TCGA-AA-A03J	5248.529283	1297.53771	22.786344
81.030267			
TCGA-77-7337	1957.900000	2386.71000	232.467000
320.068000			
	EXOSC8_S5	EZR_S5	F3_S5
FAM167A_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	427.72400	11897.400000	8418.270000
90.544900			
TCGA-B5-A3F9	481.60400	5497.600000	474.120000
20.873200			
TCGA-49-6767	585.28400	17770.400000	1340.240000
69.822500			
TCGA-AA-A03J	471.38021	10746.871452	510.121969

39.215757			
TCGA-77-7337	306.17100	9246.530000	1477.970000
29.314000			

	FAM89B_S5	FARSA_S5	FARSB_S5
FBX041_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	1355.770000	1358.970000	1342.150000
324.519000			
TCGA-B5-A3F9	1135.600000	1468.080000	1338.370000
51.189000			
TCGA-49-6767	2065.680000	919.527000	1501.780000
334.911000			
TCGA-AA-A03J	1077.730029	3382.821044	1201.770979
734.838076			
TCGA-77-7337	884.874000	1493.310000	1174.270000
85.215200			

	FCER1G_S3	FCGR1A_S3	FCGR2A_S3
FCGR2B_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	226.76300	27.24360	520.833000
120.994000			
TCGA-B5-A3F9	541.21200	121.24300	434.361000
99.396200			
TCGA-49-6767	937.27800	286.95300	1111.240000
200.592000			
TCGA-AA-A03J	215.44325	30.38253	477.218772
35.273426			
TCGA-77-7337	552.87600	85.53220	2778.360000
283.255000			

	FCGR3A_S3	FGD1_S2	FGL2_S3
FGR_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	465.545000	204.327000	303.6860
165.064000			
TCGA-B5-A3F9	828.964000	830.952000	142.6330
87.965600			
TCGA-49-6767	2390.530000	876.923000	492.3080
343.195000			
TCGA-AA-A03J	420.841219	253.368838	250.4135
157.269486			
TCGA-77-7337	1908.480000	415.850000	512.3140
203.494000			

	FLI1_S3	FLNC_S5	FMNL1_S3
--	---------	---------	----------

FN1_S2 \

TCGA Participant Barcode

TCGA-WA-A7H4	245.192000	5137.820000	574.103000
52455.900000			
TCGA-B5-A3F9	137.664000	91.444500	138.658000
2513.230000			
TCGA-49-6767	191.124000	11415.400000	1997.630000
54245.000000			
TCGA-AA-A03J	106.790429	994.628822	425.568343
7291.669478			
TCGA-77-7337	254.282000	157.478000	345.292000
98427.600000			

GGH_S5 \

TCGA Participant Barcode

FNBP1_S3 FPR3_S3 FYB_S3

TCGA-WA-A7H4	741.987000	255.609000	406.250000
1058.490000			
TCGA-B5-A3F9	1015.830000	91.444500	315.086000
801.630000			
TCGA-49-6767	1497.040000	469.822000	505.917000
420.710000			
TCGA-AA-A03J	639.341658	182.866036	84.920137
725.162161			
TCGA-77-7337	537.537000	563.443000	2971.280000
1462.970000			

GNPTAB_S3 \

TCGA Participant Barcode

GIMAP4_S3 GLRX3_S5 GNG11_S5

TCGA-WA-A7H4	187.500000	1020.030000	451.122000
1433.490000			
TCGA-B5-A3F9	278.806000	952.215000	611.783000
1293.140000			
TCGA-49-6767	485.207000	2571.010000	155.030000
735.503000			
TCGA-AA-A03J	182.790949	1104.235578	118.798676
664.235627			
TCGA-77-7337	341.542000	2100.380000	263.826000
2109.590000			

H2AFZ_S5 \

TCGA Participant Barcode

GPLD1_S5 GPNMB_S3 GSTCD_S5

TCGA-WA-A7H4	4.807700	9549.680000	254.808000
5116.190000			

TCGA-B5-A3F9 6503.990000	8.945700	2307.480000	318.565000	
TCGA-49-6767 4285.210000	8.875700	6445.560000	238.462000	
TCGA-AA-A03J 4937.989857	5.973108	656.968787	174.494972	
TCGA-77-7337 5925.860000	9.203200	38693.500000	366.425000	
	HAS2_S5	HAUS1_S5	HCK_S3	
HCLS1_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 532.85300	198.638000	183.494000	326.122000	
TCGA-B5-A3F9 777.77500	77.032000	531.273000	266.879000	
TCGA-49-6767 770.41400	25.390500	150.296000	675.740000	
TCGA-AA-A03J 342.87048	53.483964	190.910932	343.856397	
TCGA-77-7337 1998.13000	753.104000	280.869000	450.959000	
	HDC_S4	HERC5_S1	HERC6_S1	HLA-
DMA_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 532.853000	66.506400	175.481000	601.763000	
TCGA-B5-A3F9 2299.030000	1.490900	97.905200	428.397000	
TCGA-49-6767 1369.230000	1.775100	63.905300	179.882000	
TCGA-AA-A03J 1201.901095	51.860779	36.200517	333.053243	
TCGA-77-7337 1186.200000	20.110800	247.806000	414.146000	
	HLA-DRB1_S3	HMG2_S5	HMHA1_S3	
HN1L_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 4440.640000	2939.590000	5672.280000	722.756000	
TCGA-B5-A3F9 4585.140000	5338.540000	9221.480000	369.754000	
TCGA-49-6767 2697.630000	12539.900000	5878.700000	881.065000	
TCGA-AA-A03J	4294.439114	3542.639361	1596.510033	

2639.516134			
TCGA-77-7337	4501.430000	6724.500000	230.081000
5817.460000			

	HNRNPA2B1_S5	HNRNPR_S5	HSPB11_S5
HSPG2_S2 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	20217.900000	3596.150000	517.628000
12338.200000			
TCGA-B5-A3F9	22632.000000	3508.680000	528.788000
4327.120000			
TCGA-49-6767	12819.500000	2786.390000	257.988000
5197.800000			
TCGA-AA-A03J	9956.139522	1741.845332	139.147844
11231.842641			
TCGA-77-7337	10226.500000	3903.880000	208.607000
13568.800000			

	HYLS1_S5	ICAM1_S2	ID2_S5
ID3_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	217.949000	1789.260000	681.090000
822.917000			
TCGA-B5-A3F9	124.742000	685.834000	3584.230000
4047.910000			
TCGA-49-6767	156.805000	7046.750000	600.592000
244.379000			
TCGA-AA-A03J	46.389359	1215.882523	1846.198195
1449.654233			
TCGA-77-7337	298.935000	3448.830000	686.493000
932.595000			

	IFI16_S3	IFI27_S1	IFI30_S3
IFI44_S1 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	6826.120000	1596.150000	2146.550000
956.731000			
TCGA-B5-A3F9	1736.450000	4601.050000	2636.480000
437.840000			
TCGA-49-6767	2541.420000	5004.730000	5024.040000
846.746000			
TCGA-AA-A03J	455.437316	8435.411981	4218.586672
126.583741			
TCGA-77-7337	7451.210000	3560.630000	4289.150000
233.830000			

	IFI44L_S1	IFI6_S1	IFIH1_S1
--	-----------	---------	----------

IFIT1_S1 \

TCGA Participant Barcode

TCGA-WA-A7H4	293.26900	2016.830000	541.667000
358.173000			
TCGA-B5-A3F9	476.60500	5028.450000	457.719000
296.698000			
TCGA-49-6767	2188.17000	748.521000	908.876000
218.935000			
TCGA-AA-A03J	358.27975	3904.530045	365.052522
280.111308			
TCGA-77-7337	89.30550	989.859000	632.297000
137.367000			

IFIT2_S1

IFIT3_S1

IFRD2_S5

IGF2R_S2 \

TCGA Participant Barcode

TCGA-WA-A7H4	173.077000	522.436000	1132.650000
7806.090000			
TCGA-B5-A3F9	255.945000	367.269000	967.299000
1889.020000			
TCGA-49-6767	714.793000	1108.280000	1590.020000
5543.200000			
TCGA-AA-A03J	218.129019	746.590341	2946.979229
5759.380637			
TCGA-77-7337	158.500000	466.979000	821.133000
5435.710000			

IGFBP2_S2

IGFBP3_S2

IGFBP4_S2 \

TCGA Participant Barcode

TCGA-WA-A7H4	1262.820000	2625.000000	7622.600000
TCGA-B5-A3F9	7163.480000	3840.670000	25464.800000
TCGA-49-6767	101.183000	2001.180000	5388.760000
TCGA-AA-A03J	769.358703	2308.976639	6273.287026
TCGA-77-7337	4615.590000	25963.400000	9842.690000

IGFBP5_S2

IGJ_S4

IKZF1_S3

IL10RA_S3 \

TCGA Participant Barcode

TCGA-WA-A7H4	4780.450000	6902.240000	229.167000
295.673000			
TCGA-B5-A3F9	2210.570000	294.213000	124.245000
228.611000			
TCGA-49-6767	5168.640000	202.367000	296.450000
513.610000			
TCGA-AA-A03J	3090.207707	879.747499	108.191389
310.704089			
TCGA-77-7337	20129.900000	3414.400000	206.562000

290.754000

	IL18_S3	IL7R_S3	IMP4_S5
IP04_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	874.199000	229.968000	741.186000
1429.780000			
TCGA-B5-A3F9	60.134700	31.806800	695.276000
1465.530000			
TCGA-49-6767	1228.400000	37.869800	848.521000
1385.810000			
TCGA-AA-A03J	264.227307	30.011095	1463.603138
2068.202687			
TCGA-77-7337	539.923000	277.801000	795.228000
1700.700000			

	ISG15_S1	ITGA3_S2	ITGA4_S2
ITGA5_S2 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	820.513000	15915.100000	282.340000
3802.080000			
TCGA-B5-A3F9	1994.880000	918.421000	97.895300
583.456000			
TCGA-49-6767	1202.960000	15562.700000	183.462000
2215.380000			
TCGA-AA-A03J	3030.829389	3940.566087	123.141682
1132.763751			
TCGA-77-7337	467.320000	6089.480000	358.493000
6613.380000			

	ITGA6_S5	ITGB2_S2	ITGB3_S2
ITGB5_S2 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	25782.900000	801.939000	217.949000
5173.880000			
TCGA-B5-A3F9	2266.230000	1002.910000	13.915500
2538.580000			
TCGA-49-6767	2964.500000	3699.830000	91.716000
2510.060000			
TCGA-AA-A03J	4288.080636	1357.033063	50.598276
2787.829954			
TCGA-77-7337	13324.200000	1438.310000	191.905000
2923.220000			

	ITGB8_S2	JUNB_S5	JUP_S2
KIAA0090_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	1781.250000	3051.280000	20550.500000
2780.450000			
TCGA-B5-A3F9	906.493000	3195.090000	5095.050000
1177.350000			
TCGA-49-6767	1984.020000	1322.490000	12150.900000
1236.090000			
TCGA-AA-A03J	214.333668	5502.742214	19274.491393
720.656034			
TCGA-77-7337	975.884000	2970.600000	33214.500000
4090.670000			

KRR1_S5 LAIR1_S3 LAMA4_S2
 LAMB1_S2 \ TCGA Participant Barcode

TCGA-WA-A7H4	1008.450000	183.49400	2374.200000
5812.500000			
TCGA-B5-A3F9	633.680000	313.09800	421.440000
7204.730000			
TCGA-49-6767	564.284000	499.40800	483.432000
4458.580000			
TCGA-AA-A03J	352.075264	208.89802	783.651454
5772.950729			
TCGA-77-7337	1403.150000	293.82200	2424.540000
5426.160000			

LAPTM5_S3 LCK_S4 LCP1_S3
 LCP2_S3 \ TCGA Participant Barcode

TCGA-WA-A7H4	1458.330000	125.801000	1877.400000
274.038000			
TCGA-B5-A3F9	2701.590000	130.706000	2575.350000
169.967000			
TCGA-49-6767	6992.900000	328.994000	3215.380000
467.450000			
TCGA-AA-A03J	1804.768877	147.246087	817.064039
158.550938			
TCGA-77-7337	3877.970000	159.523000	3375.880000
522.539000			

LHFPL2_S3 LILRB4_S3 LMNB2_S5
 LOXL2_S5 \ TCGA Participant Barcode

TCGA-WA-A7H4	1033.650000	97.756400	3442.310000
1612.100000			
TCGA-B5-A3F9	733.544000	244.018000	2038.620000
444.301000			

TCGA-49-6767	971.598000	468.639000	2414.790000
1230.070000			
TCGA-AA-A03J	716.245256	202.107483	4983.516023
1275.832329			
TCGA-77-7337	1385.600000	326.204000	3216.020000
2383.960000			

	LRMP_S3	LRP1_S2	LRRC17_S2
--	---------	---------	-----------

LRRC40_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	52.083300	5866.990000	420.67300
415.064000			
TCGA-B5-A3F9	29.818900	2684.190000	177.42200
448.774000			
TCGA-49-6767	21.301800	3021.890000	79.88170
540.237000			
TCGA-AA-A03J	23.441881	11460.799714	27.08007
48.738581			
TCGA-77-7337	326.885000	13292.900000	117.25600
318.705000			

	LSM3_S5	LSM4_S5	LST1_S3
--	---------	---------	---------

LTB_S4 \

TCGA Participant Barcode

TCGA-WA-A7H4	673.878000	1681.090000	126.60300
374.199000			
TCGA-B5-A3F9	945.258000	3186.140000	89.45660
162.016000			
TCGA-49-6767	910.651000	1820.710000	170.41400
1493.490000			
TCGA-AA-A03J	469.580308	2784.870978	104.39482
218.043389			
TCGA-77-7337	682.403000	1516.150000	75.67110
22.837700			

	LY86_S3	LYAR_S5	LYN_S3
--	---------	---------	--------

LYZ_S3 \

TCGA Participant Barcode

TCGA-WA-A7H4	25.641000	749.199000	883.814000
13019.200000			
TCGA-B5-A3F9	79.020000	386.651000	773.799000
834.431000			
TCGA-49-6767	103.550000	1045.560000	2642.600000
6668.050000			
TCGA-AA-A03J	38.308787	678.810049	562.685169
3068.651264			
TCGA-77-7337	51.129100	860.332000	1834.510000

6712.910000

	MAGOHB_S5	MAP3K8_S5	MAPRE1_S5
MARCKSL1_S2 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	275.641000	154.647000	3822.120000
1948.720000			
TCGA-B5-A3F9	372.239000	268.370000	2547.030000
17631.900000			
TCGA-49-6767	364.497000	117.160000	3569.820000
7843.200000			
TCGA-AA-A03J	192.543608	80.427806	4168.603566
5833.127538			
TCGA-77-7337	458.798000	164.977000	7763.100000
1351.850000			

	MARVELD2_S5	MCM3_S5	MCM7_S5
MCTS1_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	252.404000	4629.810000	2780.830000
1647.440000			
TCGA-B5-A3F9	424.919000	3315.860000	2835.220000
1015.330000			
TCGA-49-6767	400.592000	2574.560000	12199.600000
1617.160000			
TCGA-AA-A03J	507.566001	3776.425366	7193.775818
891.683567			
TCGA-77-7337	402.897000	1764.290000	3067.410000
774.435000			

	MERTK_S3	MET_S5	MFSD11_S5
MKKS_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	106.570000	3673.080000	856.763000
1009.620000			
TCGA-B5-A3F9	442.313000	1659.920000	312.591000
1106.780000			
TCGA-49-6767	321.894000	40805.900000	667.325000
592.308000			
TCGA-AA-A03J	71.470187	2531.035123	445.909178
433.255505			
TCGA-77-7337	337.452000	4123.390000	753.769000
662.292000			

	MLF1IP_S5	MMP11_S2	MMP1_S2 \
TCGA Participant Barcode			
TCGA-WA-A7H4	614.728000	8549.680000	33935.100000

TCGA-B5-A3F9	681.395000	15438.200000	13.915500
TCGA-49-6767	610.467000	7279.290000	3834.910000
TCGA-AA-A03J	269.988979	6572.061363	1598.728465
TCGA-77-7337	379.163000	1861.780000	23431.800000

MMP2_S2 \

	MMP14_S2	MMP17_S2	MMP19_S2
--	----------	----------	----------

TCGA Participant Barcode

TCGA-WA-A7H4	19246.000000	199.51900	592.949000
11664.300000			
TCGA-B5-A3F9	7597.350000	36.27960	793.181000
10317.300000			
TCGA-49-6767	29959.800000	90.53250	338.462000
4640.240000			
TCGA-AA-A03J	7607.790173	26.74886	308.075381
6768.362318			
TCGA-77-7337	12179.000000	66.46780	371.879000
14209.500000			

MPP1_S3 \

	MMP3_S2	MNAT1_S5	MNDA_S3
--	---------	----------	---------

TCGA Participant Barcode

TCGA-WA-A7H4	5686.700000	314.103000	84.134600
214.744000			
TCGA-B5-A3F9	102.378000	483.562000	57.152800
176.925000			
TCGA-49-6767	13.017800	378.106000	324.852000
820.710000			
TCGA-AA-A03J	896.832598	90.714612	36.692463
862.539391			
TCGA-77-7337	1543.080000	366.425000	133.617000
245.079000			

MRPS28_S5 \

	MRPL12_S5	MRPL37_S5	MRPS16_S5
--	-----------	-----------	-----------

TCGA Participant Barcode

TCGA-WA-A7H4	1384.620000	2020.830000	1345.720000
588.878000			
TCGA-B5-A3F9	1809.510000	1491.940000	2112.120000
418.667000			
TCGA-49-6767	1376.920000	1696.450000	1979.640000
261.840000			
TCGA-AA-A03J	3222.565362	3021.732273	2019.185734
158.477752			
TCGA-77-7337	1209.710000	1497.060000	2782.770000
461.283000			

	MRT04_S5	MS4A1_S4	MSN_S3
MT1F_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1703.530000	43.269200	11398.200000
167.468000			
TCGA-B5-A3F9	1512.310000	1.987900	5326.140000
697.264000			
TCGA-49-6767	1215.380000	11.834300	25971.600000
153.254000			
TCGA-AA-A03J	976.047526	4.576277	2585.419706
459.608169			
TCGA-77-7337	1683.510000	11.248400	17127.900000
172.135000			
	MT3_S5	MTA1_S2	MTHFD1_S5
MX1_S1 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	3.205100	1064.900000	1481.530000
1167.470000			
TCGA-B5-A3F9	147.106000	1970.030000	1389.490000
1137.590000			
TCGA-49-6767	0.000000	1739.050000	2720.590000
1135.500000			
TCGA-AA-A03J	91.901208	1555.890562	1733.931097
2361.615361			
TCGA-77-7337	0.340900	998.722000	2440.870000
916.915000			
	MYBL1_S5	MYBL2_S5	MYCBP_S5
MYL6_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	238.782000	1282.850000	720.353000
15859.000000			
TCGA-B5-A3F9	64.110500	1157.970000	1166.910000
18998.100000			
TCGA-49-6767	431.361000	1129.590000	715.414000
18542.000000			
TCGA-AA-A03J	34.276176	4624.135012	617.681299
14625.742112			
TCGA-77-7337	78.738800	3666.640000	489.817000
13088.400000			
	MY01F_S3	NCEH1_S5	NCF2_S3
NCKAP1L_S3 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	259.61500	411.859000	375.801000

237.179000			
TCGA-B5-A3F9	301.66700	694.282000	82.001800
190.344000			
TCGA-49-6767	450.29600	1969.230000	771.006000
445.562000			
TCGA-AA-A03J	438.45008	451.314543	152.308393
256.994692			
TCGA-77-7337	201.44900	1193.350000	1061.100000
351.087000			

	NCLN_S5	NE01_S2	NID1_S2
--	---------	---------	---------

NLN_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	2168.270000	1062.500000	1839.500000
435.096000			
TCGA-B5-A3F9	2321.400000	2178.760000	1702.100000
422.434000			
TCGA-49-6767	1911.240000	804.142000	1350.880000
600.592000			
TCGA-AA-A03J	6476.962655	3104.024952	1136.524032
374.185195			
TCGA-77-7337	1691.690000	782.957000	5154.120000
511.973000			

	NME1_S5	NOP16_S5	NOTCH2_S2
--	---------	----------	-----------

NPC2_S3 \

TCGA Participant Barcode

TCGA-WA-A7H4	1711.670000	379.006000	2720.350000
2388.620000			
TCGA-B5-A3F9	3232.230000	589.916000	2957.530000
6377.260000			
TCGA-49-6767	2551.910000	322.485000	3546.150000
7479.290000			
TCGA-AA-A03J	2190.579905	734.285336	712.373458
1552.987229			
TCGA-77-7337	1949.150000	258.372000	3836.730000
6966.510000			

	NPL_S3	NRIP3_S5	NUDT1_S5
--	--------	----------	----------

NUDT15_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	213.141000	96.955100	445.513000
854.199000			
TCGA-B5-A3F9	130.706000	16.400400	327.510000
688.771000			
TCGA-49-6767	120.118000	134.320000	288.757000
879.846000			

TCGA-AA-A03J	88.620145	14.386538	402.260555
600.915288			
TCGA-77-7337	691.947000	247.124000	491.521000
854.865000			
	NUP107_S5	NUP35_S5	NUP85_S5
NUP93_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1161.860000	250.801000	905.449000
1128.210000			
TCGA-B5-A3F9	1000.420000	360.808000	726.089000
1247.420000			
TCGA-49-6767	1000.590000	251.479000	890.533000
1489.940000			
TCGA-AA-A03J	711.490968	148.455339	965.871197
1438.451723			
TCGA-77-7337	1263.570000	408.010000	727.397000
1467.410000			
	NUPL1_S5	NUTF2_S5	OAS1_S1
OAS2_S1 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1248.400000	1809.290000	560.096000
1220.350000			
TCGA-B5-A3F9	756.405000	3353.130000	193.326000
385.657000			
TCGA-49-6767	1099.410000	3146.750000	1168.050000
1243.200000			
TCGA-AA-A03J	1158.237576	3557.105678	1869.724499
935.073373			
TCGA-77-7337	1026.670000	2287.860000	477.887000
1499.450000			
	OAS3_S1	OASL_S1	OSBPL3_S3
PA2G4_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1335.740000	160.256000	1112.980000
2838.140000			
TCGA-B5-A3F9	634.148000	114.306000	428.894000
4439.530000			
TCGA-49-6767	1981.070000	281.657000	815.385000
5613.610000			
TCGA-AA-A03J	2756.552264	533.469981	596.266612
5580.006673			
TCGA-77-7337	1513.420000	82.147400	468.002000
4908.050000			

PDAP1_S5 \ TCGA Participant Barcode	PAICS_S5	PAK1_S2	PAK2_S2
TCGA-WA-A7H4 2505.610000	3400.640000	1366.190000	3371.790000
TCGA-B5-A3F9 3060.410000	4425.120000	2163.850000	2059.490000
TCGA-49-6767 10816.600000	2313.610000	2145.560000	4628.990000
TCGA-AA-A03J 3126.018611	3331.453327	1544.604524	1565.465184
TCGA-77-7337 1954.840000	5149.380000	2339.330000	8009.890000
TCGA Participant Barcode	PDIA4_S5	PDLIM7_S5	PFKP_S5 \
TCGA-WA-A7H4	5277.240000	3774.840000	1290.870000
TCGA-B5-A3F9	12363.900000	1167.900000	834.431000
TCGA-49-6767	16620.700000	1638.460000	3407.100000
TCGA-AA-A03J	6665.578363	1790.815205	3396.808005
TCGA-77-7337	6801.870000	838.858000	4072.260000
PIK3CG_S3 \	PFN1_S5	PGM2_S5	PHF19_S5
TCGA Participant Barcode			
TCGA-WA-A7H4 64.90380	18421.500000	1930.290000	897.43600
TCGA-B5-A3F9 7.95170	14217.600000	537.733000	265.88500
TCGA-49-6767 47.33730	12803.000000	1115.980000	1609.47000
TCGA-AA-A03J 23.68202	21598.814199	543.057721	1775.78342
TCGA-77-7337 99.19050	8874.990000	2447.720000	550.49000
PLCG2_S3 \	PITPNC1_S5	PLAT_S2	PLAUR_S5
TCGA Participant Barcode			
TCGA-WA-A7H4 356.570000	311.699000	434.295000	758.013000
TCGA-B5-A3F9 254.951000	519.345000	550.158000	603.832000
TCGA-49-6767 207.101000	175.148000	2979.880000	2352.660000
TCGA-AA-A03J 114.980755	214.514288	659.910176	2324.934366

TCGA-77-7337	411.760000	3454.960000	1480.700000
536.515000			

	PLEK_S3	PLG_S5	PLK4_S5
PL0D2_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	229.167000	0.000000	506.410000
1907.850000			
TCGA-B5-A3F9	299.679000	23.855100	258.927000
2288.100000			
TCGA-49-6767	688.757000	0.000000	369.231000
1333.730000			
TCGA-AA-A03J	161.426919	3.938344	153.655922
439.280843			
TCGA-77-7337	561.057000	0.000000	387.218000
7612.440000			

	PNN_S5	PN01_S5	POLE2_S5
P0LR3K_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	2253.210000	594.551000	126.60300
274.038000			
TCGA-B5-A3F9	2935.170000	393.112000	89.45660
389.633000			
TCGA-49-6767	1604.730000	665.681000	140.82800
243.787000			
TCGA-AA-A03J	1478.458546	599.694356	67.87253
574.739875			
TCGA-77-7337	1761.910000	1266.980000	168.72600
508.223000			

	PPIH_S5	PSMA7_S5	PSMC3_S5
PSMD12_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	492.78800	4498.400000	2745.990000
1370.990000			
TCGA-B5-A3F9	743.98000	2975.920000	2598.220000
723.107000			
TCGA-49-6767	580.47300	4809.470000	1921.890000
1149.110000			
TCGA-AA-A03J	652.58254	12850.164704	1866.885196
501.157671			
TCGA-77-7337	320.75000	6267.750000	2897.320000
2220.710000			

	PSMD14_S5	PSMD2_S5	PTPLB_S5
PTPRC_S3 \			

TCGA Participant Barcode

TCGA-WA-A7H4	1602.560000	6587.340000	308.494000
516.026000			
TCGA-B5-A3F9	1436.770000	3072.830000	43.237300
339.438000			
TCGA-49-6767	1317.160000	5385.800000	243.195000
875.148000			
TCGA-AA-A03J	978.831589	4231.503959	75.527356
208.733265			
TCGA-77-7337	3146.480000	14749.000000	605.709000
1036.560000			

RASSF2_S3 \

TCGA Participant Barcode

	PXN_S2	RAB3B_S5	RAC1_S2
TCGA-WA-A7H4	6759.01000	5.609000	6165.060000
365.385000			
TCGA-B5-A3F9	1946.43000	1.987900	6536.290000
648.063000			
TCGA-49-6767	7575.14000	26.627200	6388.170000
122.485000			
TCGA-AA-A03J	7988.21454	7.277026	7220.930029
122.210979			
TCGA-77-7337	4754.74000	16.020500	11040.800000
252.578000			

RGS8_S5 \

TCGA Participant Barcode

	RASSF4_S3	RBM14_S5	RFC3_S5
TCGA-WA-A7H4	736.114000	1018.350000	365.385000
0.0000			
TCGA-B5-A3F9	825.982000	1355.860000	442.810000
0.0000			
TCGA-49-6767	1518.930000	1006.050000	404.734000
0.0000			
TCGA-AA-A03J	772.989068	2152.712306	546.375419
NaN			
TCGA-77-7337	215.083000	1341.370000	546.740000
0.3409			

RHOG_S2 \

TCGA Participant Barcode

	RHOA_S2	RHOB_S2	RHOC_S5
TCGA-WA-A7H4	9258.810000	2931.09000	3935.10000
1266.830000			
TCGA-B5-A3F9	9805.930000	11439.50000	3642.37000

898.541000			
TCGA-49-6767	13426.000000	1077.51000	6773.37000
1168.050000			
TCGA-AA-A03J	9807.319352	3607.80426	5682.28049
1714.439988			
TCGA-77-7337	10884.400000	2874.14000	3976.82000
885.215000			

	RHOQ_S2	RMND5B_S5	RNASE6_S3
--	---------	-----------	-----------

RND3_S2 \

TCGA Participant Barcode

TCGA-WA-A7H4	1629.020000	351.763000	143.429000
1867.790000			
TCGA-B5-A3F9	1649.260000	622.717000	124.742000
2087.320000			
TCGA-49-6767	1291.900000	531.953000	175.148000
1360.360000			
TCGA-AA-A03J	360.002602	937.189697	142.656267
438.776252			
TCGA-77-7337	1348.960000	273.711000	116.574000
5216.870000			

	RNF138_S5	RNF41_S5	RPN1_S5
--	-----------	----------	---------

RPP40_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	368.590000	684.295000	6653.040000
213.141000			
TCGA-B5-A3F9	763.363000	879.159000	11263.100000
134.682000			
TCGA-49-6767	431.361000	828.402000	7985.800000
184.615000			
TCGA-AA-A03J	284.101016	1102.745864	7524.043806
165.900124			
TCGA-77-7337	601.278000	1178.700000	15401.800000
252.578000			

	RSAD2_S1	RTP4_S1	RUNX3_S3
--	----------	---------	----------

RUVBL1_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	165.865000	92.147400	657.853000
1214.70000			
TCGA-B5-A3F9	90.947500	193.326000	193.822000
1955.01000			
TCGA-49-6767	308.876000	133.136000	204.142000
1519.73000			
TCGA-AA-A03J	630.334114	123.607153	1165.381881
1381.89423			

TCGA-77-7337	294.844000	259.395000	1033.150000
2640.05000			

	SAMD9_S1	SAMHD1_S3	SAMSN1_S3
SAR1A_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	2354.970000	495.633000	111.378000
2307.690000			
TCGA-B5-A3F9	236.066000	1195.850000	123.748000
2155.900000			
TCGA-49-6767	305.325000	756.562000	193.491000
1915.380000			
TCGA-AA-A03J	380.608313	628.068511	74.518924
1033.663388			
TCGA-77-7337	1423.430000	1060.230000	184.406000
3508.140000			

	SAR1B_S5	SDC1_S5	SELL_S3
SELPLG_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	526.44200	16976.800000	138.622000
278.045000			
TCGA-B5-A3F9	437.84000	2425.760000	120.269000
216.187000			
TCGA-49-6767	364.49700	10459.200000	159.172000
585.799000			
TCGA-AA-A03J	291.35525	7002.780348	51.673906
422.963295			
TCGA-77-7337	580.82700	22700.000000	144.184000
383.468000			

	SEMA3F_S2	SERPINE1_S2	SH2B3_S3
SH3BP5L_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	1383.810000	5875.00000	687.500000
1310.900000			
TCGA-B5-A3F9	972.095000	117.28700	254.951000
594.389000			
TCGA-49-6767	575.148000	1481.66000	1301.780000
553.254000			
TCGA-AA-A03J	536.348425	237.63618	504.611923
983.091103			
TCGA-77-7337	1699.530000	6642.01000	707.627000
593.779000			

	SKA1_S5	SKA2_S5	SLC16A1_S5
SLC1A3_S3 \			

TCGA Participant Barcode

TCGA-WA-A7H4 477.564000	93.750000	786.859000	3054.490000
TCGA-B5-A3F9 179.410000	108.839000	842.880000	1665.380000
TCGA-49-6767 168.047000	128.994000	793.491000	2778.700000
TCGA-AA-A03J 26.435782	157.042894	216.737482	1117.373004
TCGA-77-7337 1772.130000	261.099000	831.018000	6389.430000

SMC2_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4 1104.17000	132.212000	11109.000000	176.28200
TCGA-B5-A3F9 605.82000	59.140700	4494.690000	183.88300
TCGA-49-6767 1209.47000	346.154000	8710.650000	632.54400
TCGA-AA-A03J 315.73208	120.723276	18811.796336	363.96797
TCGA-77-7337 2964.47000	227.695000	10477.400000	353.81300

SNRPA1_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4 424.680000	475.160000	888.622000	483.974000
TCGA-B5-A3F9 595.880000	1728.000000	2884.480000	338.444000
TCGA-49-6767 614.201000	931.953000	3064.500000	1871.010000
TCGA-AA-A03J 537.599672	326.766597	1884.414469	243.065824
TCGA-77-7337 855.901000	509.587000	2040.730000	571.283000

SNRPD1_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4 712.340000	1115.380000	3641.030000	1903.040000
TCGA-B5-A3F9	2843.230000	4689.010000	1951.640000

1430.310000			
TCGA-49-6767	1108.280000	4876.920000	2188.760000
981.065000			
TCGA-AA-A03J	2810.940332	4348.216621	2118.107643
686.240596			
TCGA-77-7337	1157.900000	2653.260000	2204.690000
1225.050000			

	SNRPE_S5	SNX17_S5	SP140_S3
--	----------	----------	----------

SPAG17_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	600.962000	2597.760000	53.685900
41.666700			
TCGA-B5-A3F9	1544.620000	2713.520000	26.340000
289.740000			
TCGA-49-6767	1222.490000	2168.640000	97.041400
1.775100			
TCGA-AA-A03J	992.908169	2212.668745	21.068411
1.794282			
TCGA-77-7337	971.794000	2252.070000	33.063500
68.172100			

	SPARC_S2	SRM_S5	STAT1_S1
--	----------	--------	----------

STK17A_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	43480.000000	1790.060000	3429.490000
1818.110000			
TCGA-B5-A3F9	10374.000000	1435.280000	4222.850000
463.186000			
TCGA-49-6767	12467.500000	2275.150000	8585.800000
2156.800000			
TCGA-AA-A03J	12455.566615	2229.710794	3373.295658
204.670983			
TCGA-77-7337	53367.500000	2054.370000	7263.060000
1204.600000			

	STRA13_S5	SYK_S3	TAGLN_S5
--	-----------	--------	----------

TAP1_S1 \

TCGA Participant Barcode

TCGA-WA-A7H4	697.917000	607.372000	7978.37000
3900.64000			
TCGA-B5-A3F9	896.553000	1712.600000	899.53500
4301.37000			
TCGA-49-6767	507.692000	547.337000	1727.81000
7538.46000			
TCGA-AA-A03J	944.119588	1408.615337	5412.55443
11272.08586			

TCGA-77-7337 3268.85000	260.418000	1351.850000	5734.30000
THBS1_S2 \	TBXAS1_S3	TCEB1_S5	TCF7L2_S2
TCGA Participant Barcode			
TCGA-WA-A7H4 16249.200000	156.250000	1051.280000	436.699000
TCGA-B5-A3F9 1140.570000	378.202000	883.632000	704.222000
TCGA-49-6767 1164.500000	170.414000	1136.090000	855.621000
TCGA-AA-A03J 2851.471087	1303.232132	570.101003	977.233934
TCGA-77-7337 8540.260000	170.089000	1231.530000	709.331000
TMEM130_S5 \	THBS2_S2	TIMP1_S2	TLR2_S3
TCGA Participant Barcode			
TCGA-WA-A7H4 9.61540	6189.100000	4869.390000	669.071000
TCGA-B5-A3F9 5.96380	136.670000	4652.730000	343.911000
TCGA-49-6767 27.81070	2704.730000	6509.470000	218.935000
TCGA-AA-A03J 14.09625	1790.479229	8027.404873	107.980266
TCGA-77-7337 54.53770	14896.000000	7015.590000	1279.930000
TCGA Participant Barcode	TNC_S2	TNFAIP3_S3	TNFRSF12A_S5 \
TCGA-WA-A7H4	51591.300000	1375.800000	3986.380000
TCGA-B5-A3F9	1885.550000	1716.570000	1070.000000
TCGA-49-6767	18590.500000	947.929000	2894.080000
TCGA-AA-A03J	2139.767469	1004.281681	2584.787646
TCGA-77-7337	36879.800000	1954.840000	1190.290000
TOMM40_S5 \	TNFRSF1A_S2	TNFRSF1B_S3	TNFSF13B_S3
TCGA Participant Barcode			
TCGA-WA-A7H4 1080.930000	3036.860000	710.737000	26.442300
TCGA-B5-A3F9 1375.150000	2660.840000	1382.600000	18.885300
TCGA-49-6767	4164.500000	1972.780000	99.408300

1551.480000			
TCGA-AA-A03J	3277.586534	2469.421757	28.323496
2911.791666			
TCGA-77-7337	5124.500000	652.067000	72.262500
1441.840000			
	TPI1_S5	TPM1_S5	TPM2_S5 \
TCGA Participant Barcode			
TCGA-WA-A7H4	6519.230000	7943.420000	10874.20000
TCGA-B5-A3F9	11693.000000	1983.870000	1485.48000
TCGA-49-6767	15510.100000	3089.760000	4177.51000
TCGA-AA-A03J	13301.708123	4638.258269	1984.92103
TCGA-77-7337	21580.200000	2700.970000	1286.41000
	TPM3_S5	TPRKB_S5	TRA2B_S5
TUBA4A_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	9447.920000	412.660000	3585.740000
7373.400000			
TCGA-B5-A3F9	7467.630000	722.113000	3100.170000
119.275000			
TCGA-49-6767	8412.430000	335.503000	1897.630000
1947.930000			
TCGA-AA-A03J	8429.630182	211.853076	2050.314713
5324.263102			
TCGA-77-7337	10409.500000	435.620000	3686.750000
3058.880000			
	TUBG1_S5	UAP1_S5	UBE2J1_S5
UMPS_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1071.310000	1302.880000	1580.93000
970.353000			
TCGA-B5-A3F9	984.022000	1015.830000	2069.93000
620.729000			
TCGA-49-6767	1028.400000	999.408000	2207.10000
849.112000			
TCGA-AA-A03J	1245.895273	576.843949	1253.38716
781.709909			
TCGA-77-7337	1813.040000	1534.210000	2274.56000
1168.130000			
	UQCR10_S5	USPL1_S5	VCAN_S2
VDAC1_S5 \			
TCGA Participant Barcode			
TCGA-WA-A7H4	1076.920000	534.455000	4942.310000
4340.540000			

TCGA-B5-A3F9	1863.180000	506.921000	7417.440000	
4943.470000				
TCGA-49-6767	1656.800000	478.106000	2764.500000	
4909.470000				
TCGA-AA-A03J	1179.679495	286.464799	1249.379279	
7151.557949				
TCGA-77-7337	743.076000	164.295000	14117.100000	
5529.780000				
	VSIG4_S3	VTA1_S5	WDHD1_S5	WDR54_S5
\				
TCGA Participant Barcode				
TCGA-WA-A7H4	81.73080	1551.28000	425.481000	296.47400
TCGA-B5-A3F9	348.88100	1079.44000	187.362000	766.84100
TCGA-49-6767	621.89400	1128.40000	508.284000	151.47900
TCGA-AA-A03J	160.41858	716.27527	101.055152	140.38918
TCGA-77-7337	479.93200	1951.09000	611.163000	215.08300
	WDR77_S5	WIPF1_S3	WNT2B_S2	
WNT8B_S2 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	906.250000	989.583000	28.044900	
6.410300				
TCGA-B5-A3F9	4555.820000	750.441000	16.897300	
1.490900				
TCGA-49-6767	1057.400000	1007.100000	5.325400	
1.775100				
TCGA-AA-A03J	1725.383932	327.268858	31.568168	
2.028401				
TCGA-77-7337	842.948000	1371.620000	249.169000	
0.000000				
	WSB2_S5	ZWILCH_S5	ZYX_S2	Immune
Subtype				
TCGA Participant Barcode				
TCGA-WA-A7H4	1991.190000	483.061000	5231.570000	
C1				
TCGA-B5-A3F9	1495.420000	452.988000	3509.180000	
C1				
TCGA-49-6767	2242.600000	325.953000	7905.920000	
C1				
TCGA-AA-A03J	2127.285705	266.422502	6938.987139	
C1				

TCGA-77-7337 C1	2586.450000	498.921000	3666.980000
--------------------	-------------	------------	-------------

Imputing Missing Information

```
df_final.isnull().sum().sum()

gene_cols = df_final.columns[:-1]
df_final[gene_cols] =
df_final[gene_cols].fillna(df_final[gene_cols].mean())

df_final.isnull().sum().sum()

0
```

Outputting the final csv

```
#Final 2009 Rows CSV * (440 Input Gene features + 1 Output Immune  
Subtype feature)
df_final.to_csv("Final_csv_2009_samples.csv")
```

Visualizing the balanced dataset

```
# Create the plot
plt.figure(figsize=(10, 6))

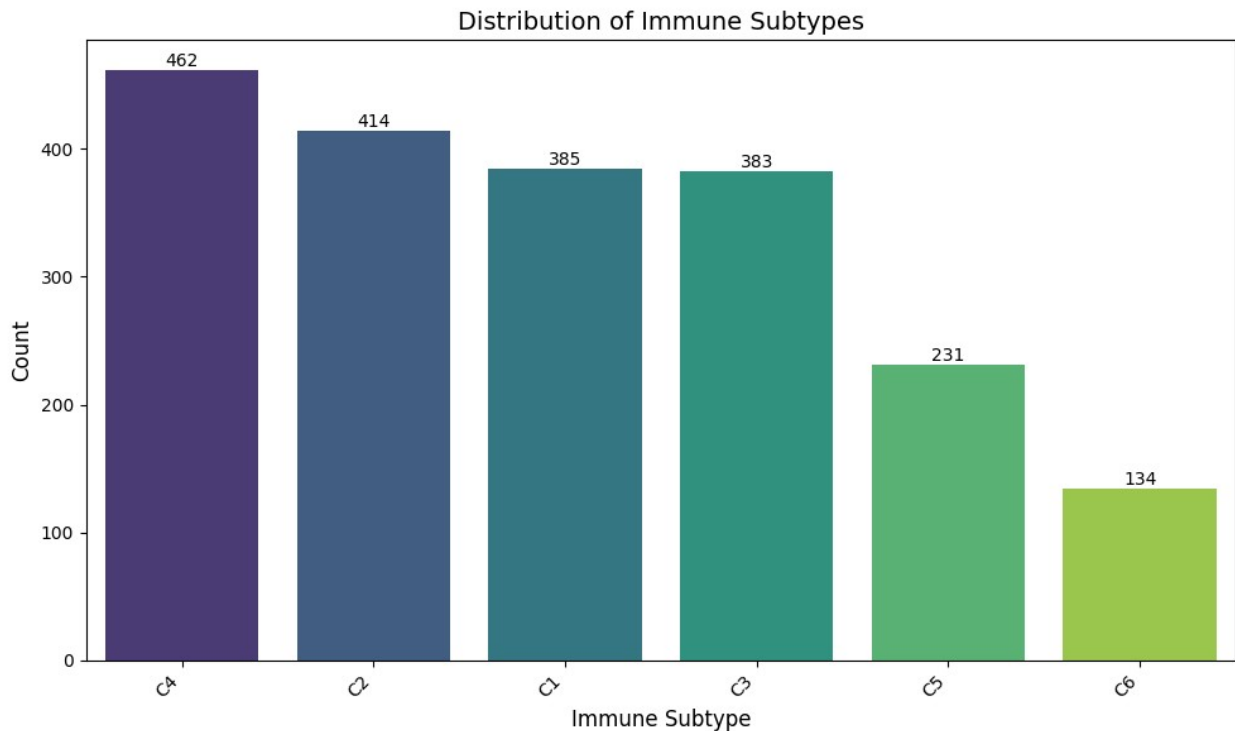
# Bar plot using Seaborn
ax = sns.countplot(data=df_final, x='Immune Subtype',
                    order=df_final['Immune  
Subtype'].value_counts().index,
                    palette='viridis')

# Add counts on top of bars
for p in ax.patches:
    ax.annotate(f'{p.get_height():.0f}',
                (p.get_x() + p.get_width() / 2., p.get_height()),
                ha='center', va='center',
                xytext=(0, 5),
                textcoords='offset points')

# Customize the plot
plt.title('Distribution of Immune Subtypes', fontsize=14)
plt.xlabel('Immune Subtype', fontsize=12)
plt.ylabel('Count', fontsize=12)
plt.xticks(rotation=45, ha='right') # Rotate x-labels for better  
readability

# Adjust layout to prevent label cutoff
```

```
plt.tight_layout()
plt.show()
```



```
# next steps, remove missingness , nmf - feature reduction,  
normalization, standardization, outliers etc  
# perform classification
```

```
# Group features by suffix (S1, S2, S3, etc.)
```

```
feature_groups = {}  
for suffix in ['S1', 'S2', 'S3', 'S4', 'S5']:  
    feature_groups[suffix] = [col for col in  
df_final_normalized.columns if col.endswith(f"_{suffix}")]
```

```
# Compute mean expression for each feature group per immune subtype
```

```
group_means = pd.DataFrame()  
for group, features in feature_groups.items():  
    if features:  
        group_means[group] =  
df_final_normalized[features].mean(axis=1)
```

```
# Add immune subtype labels
```

```
group_means['Immune Subtype'] = df_final_normalized['Immune Subtype']
```

```
# Plot each feature group across different immune subtypes
```

```
plt.figure(figsize=(12, 6))
```

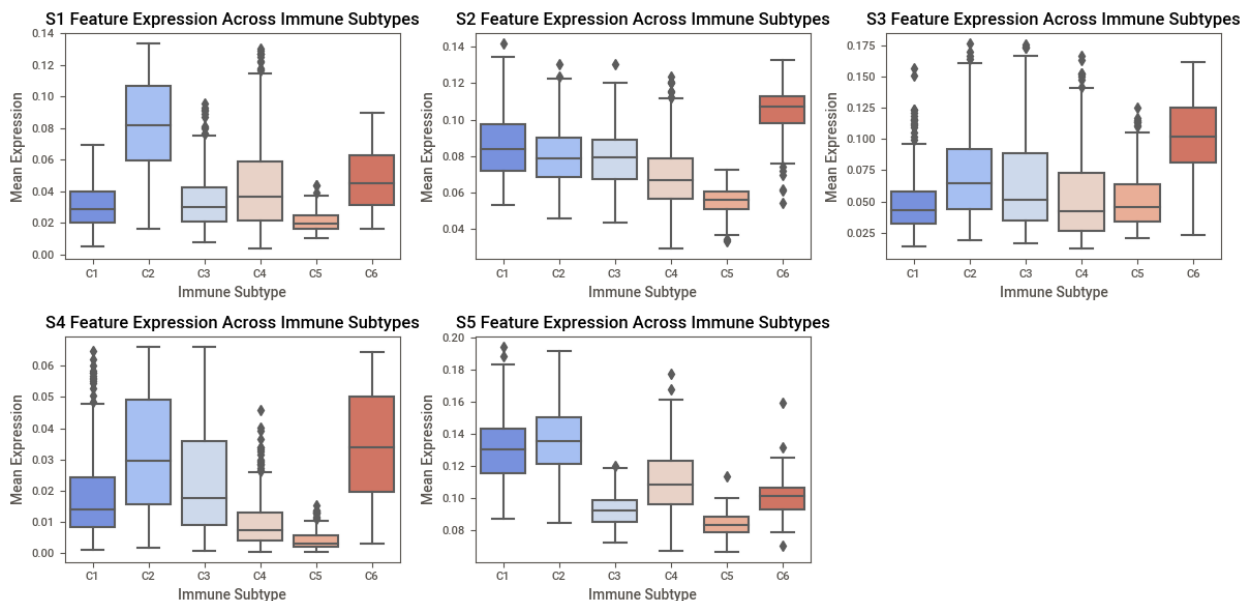


```

for i, group in enumerate(feature_groups.keys(), 1):
    plt.subplot(2, 3, i)
    sns.boxplot(x='Immune Subtype', y=group, data=group_means,
palette='coolwarm')
    plt.title(f"{group} Feature Expression Across Immune Subtypes")
    plt.xlabel("Immune Subtype")
    plt.ylabel("Mean Expression")

plt.tight_layout()
plt.show()

```



```

import numpy as np

# Convert categorical immune subtype labels to numerical
immune_subtype_numeric =
LabelEncoder().fit_transform(df_final_normalized['Immune Subtype'])

# Compute correlation for each feature group
correlations = {}
for suffix in ['S1', 'S2', 'S3', 'S4', 'S5']:
    feature_subset = [col for col in df_final_normalized.columns if
col.endswith(f"_{suffix}")]
    if feature_subset:
        mean_expression =
df_final_normalized[feature_subset].mean(axis=1)
        correlations[suffix] = np.corrcoef(mean_expression,
immune_subtype_numeric)[0, 1]

# Print correlation values
for group, corr in correlations.items():

```

```

    print(f"Correlation between {group} features and immune subtype:
{corr:.4f}")

Correlation between S1 features and immune subtype: -0.1655
Correlation between S2 features and immune subtype: -0.1757
Correlation between S3 features and immune subtype: 0.1368
Correlation between S4 features and immune subtype: -0.1786
Correlation between S5 features and immune subtype: -0.5616

# Count the number of features for each suffix
feature_counts = {suffix: len(features) for suffix, features in
feature_groups.items()}
print(feature_counts)

{'S1': 24, 'S2': 80, 'S3': 111, 'S4': 15, 'S5': 210}

```

Scaling and Normalization

```

from sklearn.preprocessing import StandardScaler, MinMaxScaler

# Select only numeric columns for preprocessing
numeric_cols = df_final.select_dtypes(include=['number']).columns

# Standardization (Z-score: mean=0, std=1)
scaler = StandardScaler()
df_final_standardized = df_final.copy()
df_final_standardized[numeric_cols] =
scaler.fit_transform(df_final[numeric_cols])

# Normalization (Min-Max Scaling: range 0-1)
min_max_scaler = MinMaxScaler()
df_final_normalized = df_final.copy()
df_final_normalized[numeric_cols] =
min_max_scaler.fit_transform(df_final[numeric_cols])

df_final_normalized.head()

```

	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5	ADCY7_S3
AIMP2_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.129424	0.169197	0.022398	0.234976
0.180090				
TCGA-B5-A3F9	0.123828	0.033567	0.039275	0.306215
0.153894				
TCGA-49-6767	0.150459	0.157022	0.003525	0.249774
0.155868				
TCGA-AA-A03J	0.083904	0.039410	0.004848	0.088996
0.435516				

TCGA-77-7337 0.383001	0.305805	0.433055	0.057262	0.140494
	ALKBH7_S5	ALOX5AP_S3	AMPD3_S3	
APITD1_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.036046	0.025048	0.083360	0.183705
TCGA-B5-A3F9	0.062357	0.019597	0.049824	0.299588
TCGA-49-6767	0.019391	0.051084	0.038690	0.237127
TCGA-AA-A03J	0.143618	0.013858	0.020518	0.097687
TCGA-77-7337	0.020735	0.078867	0.043357	0.325132
	APOC1_S3	APOE_S3	AP00_S5	ARHGAP1_S2 \
TCGA Participant Barcode				
TCGA-WA-A7H4	0.000159	0.001151	0.023024	0.199662
TCGA-B5-A3F9	0.001520	0.004717	0.137673	0.067752
TCGA-49-6767	0.001623	0.005735	0.081950	0.066406
TCGA-AA-A03J	0.000464	0.002793	0.054958	0.165810
TCGA-77-7337	0.000145	0.000614	0.069031	0.239196
	ARHGAP15_S3	ARHGDIA_S2	ARRB2_S3	B2M_S3
\				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.034113	0.193792	0.062812	0.149856
TCGA-B5-A3F9	0.010200	0.143697	0.062524	0.207232
TCGA-49-6767	0.055246	0.111708	0.088977	0.166943
TCGA-AA-A03J	0.009896	0.337684	0.181536	0.125507
TCGA-77-7337	0.022262	0.188949	0.069010	0.100442
	BCCIP_S5	BRCA2_S5	BRIP1_S5	BSG_S2
BTK_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.137327	0.135782	0.095438	0.044576
0.034921				
TCGA-B5-A3F9	0.263794	0.042710	0.028896	0.060415
0.024206				
TCGA-49-6767	0.266651	0.066250	0.052106	0.018203
0.063984				

TCGA-AA-A03J	0.104531	0.088548	0.020854	0.052634
0.010746				
TCGA-77-7337	0.243811	0.089324	0.118525	0.032608
0.034656				

	C11orf24_S5	C12orf24_S5	C13orf1_S5
C13orf18_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	0.152497	0.022158	0.155023
0.016833			
TCGA-B5-A3F9	0.077849	0.031294	0.136670
0.026442			
TCGA-49-6767	0.143469	0.022671	0.191243
0.005027			
TCGA-AA-A03J	0.114090	0.017954	0.247442
0.337895			
TCGA-77-7337	0.159544	0.028050	0.165591
0.005849			

	C13orf27_S5	C16orf61_S5	C19orf48_S5
C1orf54_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	0.076541	0.045475	0.019964
0.091298			
TCGA-B5-A3F9	0.070951	0.076200	0.041087
0.120955			
TCGA-49-6767	0.065347	0.076236	0.017507
0.080028			
TCGA-AA-A03J	0.044817	0.064950	0.094862
0.036174			
TCGA-77-7337	0.099976	0.078923	0.023613
0.045622			

	C3AR1_S3	C3orf26_S5	CASP1_S3	CBX1_S5
CCBL2_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.033466	0.252371	0.130334	0.104727
0.357476				
TCGA-B5-A3F9	0.038793	0.198402	0.018940	0.171738
0.306802				
TCGA-49-6767	0.094222	0.139277	0.032013	0.128322
0.402032				
TCGA-AA-A03J	0.032520	0.106481	0.075871	0.032466
0.193422				
TCGA-77-7337	0.093986	0.367820	0.110604	0.142077
0.304238				

CD14_S4 \ TCGA Participant Barcode	CCL2_S3	CCL5_S3	CCRL2_S3	CCT5_S5
TCGA-WA-A7H4 0.018676	0.017832	0.039156	0.006166	0.139557
TCGA-B5-A3F9 0.022774	0.031532	0.017448	0.004626	0.123400
TCGA-49-6767 0.021355	0.032674	0.100785	0.012335	0.213498
TCGA-AA-A03J 0.019312	0.006116	0.021166	0.065235	0.159937
TCGA-77-7337 0.025416	0.053161	0.019366	0.010812	0.269010
CD37_S4 \ TCGA Participant Barcode	CD163_S3	CD19_S4	CD247_S4	CD33_S3
TCGA-WA-A7H4 0.015664	0.009689	0.003777	0.063614	0.033109
TCGA-B5-A3F9 0.017639	0.017677	0.002194	0.035106	0.071084
TCGA-49-6767 0.042144	0.025298	0.000785	0.083596	0.075230
TCGA-AA-A03J 0.022887	0.004955	0.003470	0.044976	0.031795
TCGA-77-7337 0.016808	0.038462	0.001408	0.039633	0.049115
CD48_S3 \ TCGA Participant Barcode	CD3D_S4	CD3E_S4	CD3G_S4	CD44_S2
TCGA-WA-A7H4 0.014924	0.040169	0.023715	0.016702	0.075255
TCGA-B5-A3F9 0.024867	0.040378	0.024578	0.005270	0.008014
TCGA-49-6767 0.040068	0.086689	0.076188	0.065392	0.038649
TCGA-AA-A03J 0.015225	0.047478	0.023950	0.020800	0.012806
TCGA-77-7337 0.020929	0.034323	0.032421	0.047515	0.092642
CD79B_S4 \ TCGA Participant Barcode	CD52_S3	CD53_S3	CD59_S2	CD79A_S4
TCGA-WA-A7H4	0.018208	0.038376	0.113421	0.012376

0.009356				
TCGA-B5-A3F9	0.016722	0.053371	0.043971	0.001537
0.003067				
TCGA-49-6767	0.061403	0.100435	0.047856	0.003519
0.003751				
TCGA-AA-A03J	0.026161	0.035758	0.014871	0.005998
0.005639				
TCGA-77-7337	0.021409	0.102105	0.107698	0.016165
0.005231				

	CD84_S3	CD86_S3	CD8A_S3	CD8B_S4
CD97_S3 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.003355	0.110400	0.004560	0.002090
0.030530				
TCGA-B5-A3F9	0.004642	0.049360	0.026804	0.019961
0.052208				
TCGA-49-6767	0.010102	0.144443	0.088363	0.008436
0.073799				
TCGA-AA-A03J	0.004176	0.036119	0.037436	0.006757
0.248789				
TCGA-77-7337	0.018940	0.188385	0.028269	0.008949
0.052528				

	CDCA4_S5	CDH6_S2	CDK2_S5	CECR1_S3
CELF2_S3 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.239033	0.004352	0.029561	0.022926
0.054613				
TCGA-B5-A3F9	0.158576	0.003591	0.031723	0.101669
0.044890				
TCGA-49-6767	0.182581	0.000571	0.038141	0.054018
0.036824				
TCGA-AA-A03J	0.109814	0.001170	0.028389	0.024335
0.097108				
TCGA-77-7337	0.242704	0.125795	0.033506	0.036337
0.041916				

	CENPJ_S5	CENPN_S5	CENPO_S5	CENPW_S5
CEP78_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.077392	0.058463	0.176886	0.036657
0.158284				
TCGA-B5-A3F9	0.215607	0.022141	0.065257	0.029258
0.187278				
TCGA-49-6767	0.028629	0.047939	0.127073	0.022736
0.143665				

TCGA-AA-A03J 0.048390	0.066650	0.052004	0.166096	0.060270
TCGA-77-7337 0.336437	0.035145	0.109367	0.380965	0.066628
TCGA Participant Barcode	CHEK1_S5	CKLF_S5	COL16A1_S2	COL1A2_S2 \
TCGA-WA-A7H4	0.341859	0.055799	0.179046	0.049270
TCGA-B5-A3F9	0.168456	0.060539	0.006053	0.005468
TCGA-49-6767	0.319432	0.279732	0.023565	0.011242
TCGA-AA-A03J	0.107756	0.138498	0.022623	0.016113
TCGA-77-7337	0.422631	0.063020	0.028620	0.055869
TCGA Participant Barcode	COL3A1_S2	COL6A1_S2	COL6A3_S2	
TCGA-WA-A7H4	0.053382	0.090780	0.168243	0.020451
TCGA-B5-A3F9	0.011437	0.041633	0.007959	0.000976
TCGA-49-6767	0.010848	0.083508	0.050570	0.038859
TCGA-AA-A03J	0.012296	0.036574	0.051404	0.023293
TCGA-77-7337	0.080138	0.030022	0.215291	0.087967
TCGA Participant Barcode	COPS6_S5	CQ2_S5	COR01A_S3	COR01C_S5
TCGA-WA-A7H4 0.089469	0.073187	0.151087	0.039494	0.153347
TCGA-B5-A3F9 0.102082	0.111060	0.135605	0.031174	0.040879
TCGA-49-6767 0.331710	0.484917	0.148317	0.079423	0.212910
TCGA-AA-A03J 0.068212	0.115420	0.190231	0.176253	0.140236
TCGA-77-7337 0.110654	0.089021	0.137881	0.027044	0.342771
TCGA Participant Barcode	COX17_S5	CPEB4_S5	CPVL_S3	CSF1R_S3
TCGA-WA-A7H4 0.332514	0.077845	0.066910	0.008639	0.035377
TCGA-B5-A3F9	0.119659	0.037612	0.009777	0.031854

0.324371				
TCGA-49-6767	0.130252	0.019654	0.016773	0.048437
0.276330				
TCGA-AA-A03J	0.087111	0.024473	0.009468	0.032764
0.336343				
TCGA-77-7337	0.053663	0.073843	0.012768	0.056842
0.322352				
	CTNNB1_S2	CTSC_S3	CTSL1_S3	CTSS_S3
CXCL10_S1 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.163481	0.095917	0.023556	0.018769
0.003976				
TCGA-B5-A3F9	0.128645	0.058378	0.034386	0.040411
0.006545				
TCGA-49-6767	0.273783	0.080481	0.025965	0.236131
0.055138				
TCGA-AA-A03J	0.202381	0.042648	0.007856	0.069236
0.006464				
TCGA-77-7337	0.248077	0.203987	0.044214	0.046477
0.012050				
	CXCR4_S3	CYBB_S3	CYTIP_S3	DAPK1_S3
DBNDD1_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.009499	0.018678	0.030886	0.082398
0.015780				
TCGA-B5-A3F9	0.083557	0.018557	0.038931	0.041218
0.066342				
TCGA-49-6767	0.019742	0.066786	0.049761	0.013979
0.069333				
TCGA-AA-A03J	0.008578	0.013780	0.012738	0.026551
0.027064				
TCGA-77-7337	0.038591	0.066092	0.060276	0.027985
0.013411				
	DCBLD2_S5	DCK_S5	DCLRE1B_S5	DDX58_S1
DDX60_S1 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.110837	0.160330	0.287794	0.053464
0.072989				
TCGA-B5-A3F9	0.069558	0.103752	0.207635	0.026454
0.059039				
TCGA-49-6767	0.854769	0.256074	0.232055	0.031037
0.078268				
TCGA-AA-A03J	0.016922	0.102256	0.117280	0.027088
0.072232				

TCGA-77-7337 0.074897	0.101117	0.293041	0.243294	0.014842
	DHFR_S5	DLEU1_S5	DLEU2_S5	DOCK2_S3
DSP_S2 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 0.184312	0.085747	0.097426	0.145469	0.044880
TCGA-B5-A3F9 0.017743	0.165736	0.245137	0.103178	0.021801
TCGA-49-6767 0.036720	0.086690	0.111307	0.224930	0.071234
TCGA-AA-A03J 0.033738	0.152287	0.163563	0.053175	0.032179
TCGA-77-7337 0.218342	0.065244	0.207484	0.215501	0.069890
	DUT_S5	DVL1_S2	DVL3_S2	DYNLT1_S5 \
TCGA Participant Barcode				
TCGA-WA-A7H4	0.202159	0.284554	0.069952	0.117494
TCGA-B5-A3F9	0.286724	0.189871	0.051480	0.268827
TCGA-49-6767	0.093573	0.239620	0.055469	0.111976
TCGA-AA-A03J	0.357800	0.412409	0.082258	0.106556
TCGA-77-7337	0.082165	0.101505	0.123503	0.112132
	EBNA1BP2_S5	EFNA5_S2	EIF2AK1_S5	
EIF4EBP1_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 0.045329	0.254913	0.100997	0.180324	
TCGA-B5-A3F9 0.067787	0.138149	0.012586	0.275381	
TCGA-49-6767 0.050975	0.229355	0.003574	0.234314	
TCGA-AA-A03J 0.065817	0.141679	0.001799	0.280938	
TCGA-77-7337 0.164195	0.123693	0.058037	0.311581	
	EIF4G1_S5	EMP2_S5	EN01_S5	EPHA2_S2
EPHB2_S2 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 0.030938	0.224737	0.073334	0.176186	0.100160
TCGA-B5-A3F9 0.027591	0.099652	0.037854	0.106735	0.038517
TCGA-49-6767	0.234288	0.092133	0.181019	0.065869

0.127376				
TCGA-AA-A03J	0.181465	0.046118	0.087059	0.199168
0.253530				
TCGA-77-7337	0.303863	0.117772	0.565930	0.153488
0.135483				
	EPHB3_S2	EPHB4_S2	ERLIN1_S5	EVI2A_S3
EVI2B_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.032859	0.146180	0.144039	0.018113
0.023936				
TCGA-B5-A3F9	0.009309	0.103594	0.204526	0.016738
0.019382				
TCGA-49-6767	0.014559	0.752040	0.450168	0.041415
0.057620				
TCGA-AA-A03J	0.149952	0.288277	0.213134	0.004571
0.012131				
TCGA-77-7337	0.059069	0.105869	0.432919	0.048020
0.049250				
	EXOSC8_S5	EZR_S5	F3_S5	FAM167A_S5 \
TCGA Participant Barcode				
TCGA-WA-A7H4	0.161982	0.171173	0.210631	0.010754
TCGA-B5-A3F9	0.184557	0.076877	0.011839	0.002479
TCGA-49-6767	0.227998	0.257706	0.033512	0.008293
TCGA-AA-A03J	0.180274	0.154220	0.012739	0.004658
TCGA-77-7337	0.111053	0.132114	0.036959	0.003482
	FAM89B_S5	FARSA_S5	FARSB_S5	FBX041_S5
FCER1G_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.204180	0.102053	0.329130	0.047208
0.018273				
TCGA-B5-A3F9	0.169258	0.111014	0.328010	0.007366
0.044371				
TCGA-49-6767	0.316784	0.065963	0.376409	0.048723
0.077242				
TCGA-AA-A03J	0.160079	0.268263	0.287552	0.107019
0.017333				
TCGA-77-7337	0.129489	0.113086	0.279407	0.012326
0.045339				
	FCGR1A_S3	FCGR2A_S3	FCGR2B_S3	
FCGR3A_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.006308	0.030941	0.012875	0.013803

TCGA-B5-A3F9	0.028072	0.025621	0.010577	0.024701
TCGA-49-6767	0.066439	0.067265	0.021345	0.071530
TCGA-AA-A03J	0.007035	0.028258	0.003753	0.012462
TCGA-77-7337	0.019803	0.169831	0.030141	0.057074
FLNC_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.020552	0.006781	0.032155	0.083692
0.068794				
TCGA-B5-A3F9	0.088063	0.003095	0.017032	0.046122
0.001224				
TCGA-49-6767	0.093016	0.011098	0.067096	0.064801
0.152849				
TCGA-AA-A03J	0.025836	0.005562	0.030626	0.035335
0.013318				
TCGA-77-7337	0.043341	0.011555	0.039693	0.086868
0.002109				
FYB_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.101362	0.027169	0.046967	0.056917
0.053084				
TCGA-B5-A3F9	0.022260	0.001267	0.065727	0.020283
0.041086				
TCGA-49-6767	0.359959	0.028097	0.098693	0.104719
0.066201				
TCGA-AA-A03J	0.074379	0.003745	0.039936	0.040684
0.010794				
TCGA-77-7337	0.059797	0.051011	0.032962	0.125610
0.390670				
GNPTAB_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.027823	0.025146	0.160717	0.010046
0.057354				
TCGA-B5-A3F9	0.021042	0.039299	0.149249	0.013832
0.051172				
TCGA-49-6767	0.010985	0.071292	0.422999	0.003070
0.026608				
TCGA-AA-A03J	0.019023	0.024416	0.174956	0.002216

0.023469				
TCGA-77-7337	0.038501	0.049023	0.343412	0.005633
0.087136				
	GPLD1_S5	GPNMB_S3	GSTCD_S5	H2AFZ_S5
HAS2_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.000408	0.053609	0.109785	0.152008
0.037437				
TCGA-B5-A3F9	0.000736	0.012891	0.137458	0.199343
0.014518				
TCGA-49-6767	0.000730	0.036157	0.102690	0.123665
0.004785				
TCGA-AA-A03J	0.000500	0.003611	0.074926	0.145930
0.010080				
TCGA-77-7337	0.000756	0.217466	0.158231	0.179624
0.141935				
	HAUS1_S5	HCK_S3	HCLS1_S3	HDC_S4
HERC5_S1 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.020346	0.047622	0.057652	0.001211
0.025456				
TCGA-B5-A3F9	0.067098	0.038908	0.086070	0.000030
0.014123				
TCGA-49-6767	0.015883	0.099041	0.085216	0.000035
0.009157				
TCGA-AA-A03J	0.021343	0.050230	0.035609	0.000945
0.005109				
TCGA-77-7337	0.033436	0.065982	0.227667	0.000368
0.036022				
	HERC6_S1	HLA-DMA_S3	HLA-DRB1_S3	HMG2_S5
\				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.075657	0.022540	0.047645	0.136413
TCGA-B5-A3F9	0.053794	0.103293	0.086854	0.244664
TCGA-49-6767	0.022453	0.060781	0.204553	0.142709
TCGA-AA-A03J	0.041770	0.053130	0.069789	0.071459
TCGA-77-7337	0.051997	0.052412	0.073172	0.168506
	HMHA1_S3	HN1L_S5	HNRNPA2B1_S5	HNRNPR_S5
\				

TCGA Participant Barcode

TCGA-WA-A7H4	0.071628	0.271469	0.475610	0.399541
TCGA-B5-A3F9	0.034324	0.280583	0.548851	0.388752
TCGA-49-6767	0.088358	0.161530	0.251151	0.299662
TCGA-AA-A03J	0.163964	0.157864	0.164280	0.170824
TCGA-77-7337	0.019563	0.358311	0.172482	0.437498

ID2_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	0.218957	0.188063	0.109928	0.034119
0.037530				
TCGA-B5-A3F9	0.224201	0.064808	0.062105	0.012642
0.209001				
TCGA-49-6767	0.096953	0.078204	0.078556	0.136449
0.032776				
TCGA-AA-A03J	0.041110	0.171041	0.021903	0.022959
0.106346				
TCGA-77-7337	0.073749	0.206996	0.151481	0.066420
0.037849				

ID3_S5 IFI16_S3 IFI27_S1 IFI30_S3

IFI44_S1 \

TCGA Participant Barcode

TCGA-WA-A7H4	0.030801	0.174878	0.010585	0.038855
0.092889				
TCGA-B5-A3F9	0.159889	0.044124	0.030673	0.047912
0.042425				
TCGA-49-6767	0.007644	0.064804	0.033371	0.092049
0.082192				
TCGA-AA-A03J	0.055888	0.011214	0.056305	0.077159
0.012154				
TCGA-77-7337	0.035191	0.190937	0.023718	0.078464
0.022584				

IFI44L_S1 IFI6_S1 IFIH1_S1 IFIT1_S1

IFIT2_S1 \

TCGA Participant Barcode

TCGA-WA-A7H4	0.008884	0.016181	0.052465	0.012104
0.013984				
TCGA-B5-A3F9	0.014465	0.041187	0.044148	0.009991

0.020951				
TCGA-49-6767	0.066570	0.005650	0.088845	0.007317
0.059526				
TCGA-AA-A03J	0.010863	0.031855	0.034967	0.009420
0.017772				
TCGA-77-7337	0.002674	0.007654	0.061444	0.004512
0.012758				

	IFIT3_S1	IFRD2_S5	IGF2R_S2	IGFBP2_S2
IGFBP3_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.032078	0.174951	0.417760	0.008048
0.011803				
TCGA-B5-A3F9	0.022286	0.148126	0.090358	0.046128
0.017299				
TCGA-49-6767	0.069046	0.249150	0.292550	0.000552
0.008982				
TCGA-AA-A03J	0.046223	0.469290	0.304512	0.004864
0.010374				
TCGA-77-7337	0.028578	0.124414	0.286602	0.029685
0.117323				

	IGFBP4_S2	IGFBP5_S2	IGJ_S4	IKZF1_S3
IL10RA_S3 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.076637	0.006510	0.077137	0.030634
0.043147				
TCGA-B5-A3F9	0.259275	0.002993	0.003288	0.016425
0.033162				
TCGA-49-6767	0.053771	0.007041	0.002262	0.039746
0.075596				
TCGA-AA-A03J	0.062825	0.004197	0.009832	0.014251
0.045385				
TCGA-77-7337	0.099363	0.027515	0.038158	0.027573
0.042415				

	IL18_S3	IL7R_S3	IMP4_S5	IP04_S5
ISG15_S1 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.083119	0.079098	0.139524	0.207161
0.014778				
TCGA-B5-A3F9	0.005650	0.010949	0.130361	0.212707
0.037194				
TCGA-49-6767	0.116826	0.013034	0.160946	0.200339
0.022078				
TCGA-AA-A03J	0.025072	0.010332	0.283705	0.306210
0.056967				

TCGA-77-7337 0.008037	0.051308	0.095548	0.150310	0.249193
ITGB2_S2 \	ITGA3_S2	ITGA4_S2	ITGA5_S2	ITGA6_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.021288	0.276907	0.053032	0.086581	0.373463
TCGA-B5-A3F9 0.026752	0.015699	0.018388	0.012094	0.032617
TCGA-49-6767 0.100076	0.270769	0.034460	0.049861	0.042737
TCGA-AA-A03J 0.036380	0.068338	0.023130	0.024807	0.061921
TCGA-77-7337 0.038590	0.105767	0.067336	0.151642	0.192889
JUP_S2 \	ITGB3_S2	ITGB5_S2	ITGB8_S2	JUNB_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.081848	0.017033	0.202443	0.024935	0.028278
TCGA-B5-A3F9 0.020165	0.001087	0.097532	0.012690	0.029653
TCGA-49-6767 0.048325	0.007168	0.096397	0.027774	0.011750
TCGA-AA-A03J 0.076755	0.003954	0.107455	0.003000	0.051715
TCGA-77-7337 0.132390	0.014997	0.112845	0.013661	0.027507
LAMB1_S2 \	KIAA0090_S5	KRR1_S5	LAIR1_S3	LAMA4_S2
TCGA Participant Barcode				
TCGA-WA-A7H4 0.134072	0.442496	0.266041	0.031268	0.063886
TCGA-B5-A3F9 0.166282	0.161326	0.163491	0.054181	0.010380
TCGA-49-6767 0.102749	0.171628	0.144502	0.087119	0.012079
TCGA-AA-A03J 0.133157	0.081225	0.086434	0.035759	0.020305
TCGA-77-7337 0.125134	0.672297	0.374044	0.050773	0.065265
LHFPL2_S3 \	LAPTM5_S3	LCK_S4	LCP1_S3	LCP2_S3

TCGA Participant Barcode

TCGA-WA-A7H4 0.132281	0.029875	0.023022	0.057339	0.062984
TCGA-B5-A3F9 0.092902	0.057293	0.023920	0.078932	0.038561
TCGA-49-6767 0.124139	0.151932	0.060208	0.098733	0.108373
TCGA-AA-A03J 0.090632	0.037515	0.026947	0.024536	0.035882
TCGA-77-7337 0.178464	0.083237	0.029194	0.103698	0.121301

	LILRB4_S3	LMNB2_S5	LOXL2_S5	LRMP_S3
LRP1_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.021220	0.008157	0.250072	0.030669	0.004127
TCGA-B5-A3F9 0.009356	0.020459	0.144390	0.008254	0.002365
TCGA-49-6767 0.010615	0.039352	0.172712	0.023336	0.001690
TCGA-AA-A03J 0.042072	0.016934	0.366108	0.024214	0.001860
TCGA-77-7337 0.048902	0.027372	0.233035	0.045484	0.025887

	LRRC17_S2	LRRC40_S5	LSM3_S5	LSM4_S5
LST1_S3 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.045123	0.060840	0.178373	0.109577	0.100458
TCGA-B5-A3F9 0.031602	0.025620	0.193141	0.163247	0.198695
TCGA-49-6767 0.061070	0.011497	0.233212	0.156403	0.109571
TCGA-AA-A03J 0.037040	0.003852	0.017883	0.069174	0.172504
TCGA-77-7337 0.026585	0.016909	0.136157	0.111263	0.089692

	LTB_S4	LY86_S3	LYAR_S5	LYN_S3
LYZ_S3 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.031317	0.042660	0.009661	0.210344	0.114421
TCGA-B5-A3F9	0.018470	0.030685	0.104753	0.099920

0.001993				
TCGA-49-6767	0.170264	0.040346	0.296658	0.346249
0.016032				
TCGA-AA-A03J	0.024858	0.014650	0.189843	0.072093
0.007370				
TCGA-77-7337	0.002604	0.019700	0.242711	0.239733
0.016140				
	MAGOHB_S5	MAP3K8_S5	MAPRE1_S5	MARCKSL1_S2
\				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.075697	0.035770	0.145664	0.040757
TCGA-B5-A3F9	0.107498	0.062074	0.090899	0.385772
TCGA-49-6767	0.104949	0.027099	0.134828	0.170430
TCGA-AA-A03J	0.048341	0.018603	0.160546	0.126210
TCGA-77-7337	0.135993	0.038159	0.314931	0.027627
	MARVELD2_S5	MCM3_S5	MCM7_S5	MCTS1_S5
MERTK_S3 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.113350	0.216977	0.074077	0.221840
0.010786				
TCGA-B5-A3F9	0.190823	0.152350	0.075562	0.121013
0.046835				
TCGA-49-6767	0.179898	0.115889	0.331314	0.217010
0.033905				
TCGA-AA-A03J	0.227938	0.175003	0.194599	0.101290
0.007017				
TCGA-77-7337	0.180933	0.076036	0.081903	0.082588
0.035576				
	MET_S5	MFSD11_S5	MKKS_S5	MLF1IP_S5
MMP11_S2 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.028999	0.368101	0.106318	0.169909
0.080923				
TCGA-B5-A3F9	0.013101	0.097814	0.118294	0.188447
0.146142				
TCGA-49-6767	0.322232	0.274009	0.054881	0.168724
0.068895				
TCGA-AA-A03J	0.019980	0.164033	0.035276	0.074050
0.062199				

TCGA-77-7337 0.017603	0.032555	0.316945	0.063507	0.104407
MMP2_S2 \	MMP1_S2	MMP14_S2	MMP17_S2	MMP19_S2
TCGA Participant Barcode				
TCGA-WA-A7H4 0.076244	0.194250	0.138782	0.028384	0.021184
TCGA-B5-A3F9 0.067427	0.000080	0.054637	0.005161	0.028822
TCGA-49-6767 0.030269	0.021952	0.216173	0.012879	0.011475
TCGA-AA-A03J 0.044198	0.009151	0.054713	0.003805	0.010316
TCGA-77-7337 0.092904	0.134127	0.087733	0.009456	0.012750
MRPL12_S5 \	MMP3_S2	MNAT1_S5	MNDA_S3	MPP1_S3
TCGA Participant Barcode				
TCGA-WA-A7H4 0.053989	0.100101	0.297820	0.034854	0.028984
TCGA-B5-A3F9 0.071177	0.001807	0.472321	0.023676	0.023693
TCGA-49-6767 0.053678	0.000234	0.363728	0.134574	0.113771
TCGA-AA-A03J 0.128341	0.015790	0.067786	0.015200	0.119624
TCGA-77-7337 0.046913	0.027166	0.351699	0.055352	0.033229
MS4A1_S4 \	MRPL37_S5	MRPS16_S5	MRPS28_S5	MRT04_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.001867	0.215832	0.081801	0.152298	0.296662
TCGA-B5-A3F9 0.000091	0.149825	0.159533	0.102605	0.261444
TCGA-49-6767 0.000514	0.175349	0.146096	0.056819	0.206756
TCGA-AA-A03J 0.000202	0.340748	0.150107	0.026642	0.162677
TCGA-77-7337 0.000489	0.150464	0.227553	0.115047	0.292974
MTHFD1_S5 \	MSN_S3	MT1F_S5	MT3_S5	MTA1_S2

TCGA Participant Barcode

TCGA-WA-A7H4 0.053998	0.225881	0.001859	0.000017	0.108317
TCGA-B5-A3F9 0.049750	0.102073	0.007738	0.000714	0.205529
TCGA-49-6767 0.111187	0.523028	0.001701	0.000002	0.180721
TCGA-AA-A03J 0.065647	0.046191	0.005101	0.000447	0.161050
TCGA-77-7337 0.098276	0.342708	0.001910	0.000003	0.101209

	MX1_S1	MYBL1_S5	MYBL2_S5	MYCBP_S5
MYL6_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.092348	0.023341	0.044450	0.036125	0.094650
TCGA-B5-A3F9 0.113473	0.022720	0.011306	0.032607	0.157460
TCGA-49-6767 0.110403	0.022676	0.080992	0.031808	0.093955
TCGA-AA-A03J 0.084048	0.048149	0.005645	0.130239	0.080209
TCGA-77-7337 0.073702	0.018135	0.014082	0.103269	0.062224

	MY01F_S3	NCEH1_S5	NCF2_S3	NCKAP1L_S3
NCLN_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.124847	0.060112	0.033299	0.088336	0.041589
TCGA-B5-A3F9 0.134272	0.070195	0.056313	0.018637	0.033273
TCGA-49-6767 0.109027	0.105832	0.160207	0.182093	0.078590
TCGA-AA-A03J 0.390046	0.102991	0.036514	0.035316	0.045108
TCGA-77-7337 0.095514	0.046165	0.096981	0.250913	0.061815

	NE01_S2	NID1_S2	NLN_S5	NME1_S5
NOP16_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.164285	0.103046	0.011977	0.239173	0.064625
TCGA-B5-A3F9	0.214586	0.011057	0.232128	0.127969

0.271940				
TCGA-49-6767	0.077230	0.008704	0.331265	0.099628
0.135435				
TCGA-AA-A03J	0.307042	0.007268	0.205279	0.084575
0.345630				
TCGA-77-7337	0.075113	0.034183	0.281952	0.074518
0.102710				

	NOTCH2_S2	NPC2_S3	NPL_S3	NRIP3_S5
NUDT1_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.111047	0.015887	0.049165	0.003531
0.080380				
TCGA-B5-A3F9	0.120854	0.047381	0.029214	0.000597
0.057976				
TCGA-49-6767	0.145194	0.056083	0.026651	0.004892
0.050619				
TCGA-AA-A03J	0.028016	0.009288	0.019027	0.000524
0.072168				
TCGA-77-7337	0.157209	0.052034	0.165051	0.009000
0.089115				

	NUDT15_S5	NUP107_S5	NUP35_S5	NUP85_S5
NUP93_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.158879	0.083663	0.173388	0.174390
0.121559				
TCGA-B5-A3F9	0.124530	0.071152	0.272161	0.137736
0.136196				
TCGA-49-6767	0.164204	0.071165	0.173997	0.171342
0.165973				
TCGA-AA-A03J	0.106287	0.048762	0.081494	0.186738
0.159651				
TCGA-77-7337	0.159017	0.091544	0.314542	0.138004
0.163207				

	NUPL1_S5	NUTF2_S5	OAS1_S1	OAS2_S1
OAS3_S1 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.210659	0.143198	0.035967	0.045707
0.045250				
TCGA-B5-A3F9	0.120356	0.307269	0.012002	0.013871
0.020847				
TCGA-49-6767	0.183313	0.285336	0.075691	0.046578
0.067696				
TCGA-AA-A03J	0.194111	0.328946	0.121539	0.034826
0.094669				

TCGA-77-7337 0.051430	0.169962	0.194058	0.030596	0.056352
PAK1_S2 \	OASL_S1	OSBPL3_S3	PA2G4_S5	PAICS_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.072260	0.018917	0.153589	0.079970	0.166344
TCGA-B5-A3F9 0.117644	0.013454	0.058412	0.160077	0.224232
TCGA-49-6767 0.116603	0.033349	0.112185	0.218809	0.104921
TCGA-AA-A03J 0.082411	0.063285	0.081699	0.217128	0.162434
TCGA-77-7337 0.127628	0.009631	0.063854	0.183514	0.265156
PFKP_S5 \	PAK2_S2	PDAP1_S5	PDIA4_S5	PDLIM7_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.049983	0.241153	0.168676	0.135598	0.250676
TCGA-B5-A3F9 0.032116	0.126378	0.210840	0.335843	0.075176
TCGA-49-6767 0.132821	0.351109	0.800304	0.456125	0.106854
TCGA-AA-A03J 0.132418	0.083170	0.215826	0.174828	0.117111
TCGA-77-7337 0.158858	0.646806	0.126817	0.178679	0.053024
PITPNC1_S5 \	PFN1_S5	PGM2_S5	PHF19_S5	PIK3CG_S3
TCGA Participant Barcode				
TCGA-WA-A7H4 0.076846	0.255600	0.210470	0.191977	0.024237
TCGA-B5-A3F9 0.131413	0.191584	0.054117	0.053123	0.002969
TCGA-49-6767 0.040962	0.170042	0.119041	0.348527	0.017677
TCGA-AA-A03J 0.051307	0.303984	0.054715	0.385093	0.008843
TCGA-77-7337 0.103141	0.110226	0.268565	0.115697	0.037040
PLG_S5 \	PLAT_S2	PLAUR_S5	PLCG2_S3	PLEK_S3

TCGA Participant Barcode

TCGA-WA-A7H4 0.000005	0.001419	0.039017	0.006507	0.023230
TCGA-B5-A3F9 0.000152	0.001808	0.030971	0.004564	0.030490
TCGA-49-6767 0.000005	0.009950	0.122240	0.003650	0.070551
TCGA-AA-A03J 0.000029	0.002175	0.120793	0.001889	0.016255
TCGA-77-7337 0.000005	0.011542	0.076734	0.009946	0.057403

	PLK4_S5	PL0D2_S5	PNN_S5	PN01_S5
POLE2_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.136847	0.272897	0.015910	0.147280	0.161602
TCGA-B5-A3F9 0.096535	0.139299	0.019097	0.197264	0.096395
TCGA-49-6767 0.152285	0.198844	0.011098	0.099750	0.184627
TCGA-AA-A03J 0.073111	0.082470	0.003602	0.090495	0.163267
TCGA-77-7337 0.182561	0.208554	0.063720	0.111271	0.379271

	POLR3K_S5	PPIH_S5	PSMA7_S5	PSMC3_S5
PSMD12_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.139899	0.089785	0.144393	0.125138	0.167270
TCGA-B5-A3F9 0.062331	0.137554	0.232093	0.073134	0.156777
TCGA-49-6767 0.113334	0.077284	0.175007	0.135764	0.108752
TCGA-AA-A03J 0.035758	0.214049	0.200183	0.410415	0.104846
TCGA-77-7337 0.241631	0.186561	0.084329	0.185575	0.178016

	PSMD14_S5	PSMD2_S5	PTPLB_S5	PTPRC_S3
PXN_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4 0.135201	0.268905	0.157737	0.084729	0.024735
TCGA-B5-A3F9	0.238535	0.063664	0.011875	0.016258

0.030791				
TCGA-49-6767	0.216624	0.125575	0.066794	0.041973
0.152908				
TCGA-AA-A03J	0.154648	0.094678	0.020744	0.009985
0.161869				
TCGA-77-7337	0.551728	0.376199	0.166360	0.049721
0.091718				

	RAB3B_S5	RAC1_S2	RASSF2_S3	RASSF4_S3
RBM14_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.001371	0.153141	0.006755	0.025563
0.282242				
TCGA-B5-A3F9	0.000577	0.167645	0.012288	0.028804
0.387320				
TCGA-49-6767	0.005981	0.161858	0.002001	0.053798
0.278413				
TCGA-AA-A03J	0.001737	0.194394	0.001996	0.026893
0.635406				
TCGA-77-7337	0.003655	0.343638	0.004547	0.006769
0.382809				

	RFC3_S5	RGS8_S5	RHOA_S2	RHOB_S2
RHOC_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.095780	0.000000	0.254724	0.040654
0.129998				
TCGA-B5-A3F9	0.117169	0.000000	0.276097	0.170179
0.119639				
TCGA-49-6767	0.106650	0.000000	0.417517	0.012437
0.230437				
TCGA-AA-A03J	0.145781	0.014786	0.276152	0.050956
0.191826				
TCGA-77-7337	0.145882	0.002612	0.318228	0.039787
0.131475				

	RHOG_S2	RHOQ_S2	RMND5B_S5	RNASE6_S3
RND3_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.174512	0.107443	0.097324	0.034370
0.133599				
TCGA-B5-A3F9	0.117589	0.108968	0.181132	0.029793
0.149408				
TCGA-49-6767	0.159244	0.082045	0.153058	0.042139
0.097058				
TCGA-AA-A03J	0.243694	0.011838	0.278400	0.034181
0.030692				

TCGA-77-7337 0.374773	0.115530	0.086344	0.073182	0.027792
RSAD2_S1 \	RNF138_S5	RNF41_S5	RPN1_S5	RPP40_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.011339	0.087150	0.089728	0.169853	0.203581
TCGA-B5-A3F9 0.006195	0.197635	0.124705	0.339800	0.126313
TCGA-49-6767 0.021158	0.104717	0.115594	0.218984	0.175488
TCGA-AA-A03J 0.043230	0.063504	0.164837	0.201962	0.157057
TCGA-77-7337 0.020195	0.152272	0.178470	0.492370	0.242419
SAMHD1_S3 \	RTP4_S1	RUNX3_S3	RUVBL1_S5	SAMD9_S1
TCGA Participant Barcode				
TCGA-WA-A7H4 0.052966	0.022262	0.040771	0.176109	0.080709
TCGA-B5-A3F9 0.130442	0.047336	0.011941	0.304404	0.007888
TCGA-49-6767 0.081837	0.032420	0.012582	0.228971	0.010268
TCGA-AA-A03J 0.067619	0.030058	0.072303	0.205084	0.012855
TCGA-77-7337 0.115436	0.063709	0.064088	0.423121	0.048694
SELL_S3 \	SAMSN1_S3	SAR1A_S5	SAR1B_S5	SDC1_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.004675	0.047820	0.206790	0.100884	0.098025
TCGA-B5-A3F9 0.004051	0.053131	0.188092	0.083722	0.013992
TCGA-49-6767 0.005374	0.083075	0.158464	0.069516	0.060385
TCGA-AA-A03J 0.001718	0.031994	0.049852	0.055348	0.040424
TCGA-77-7337 0.004864	0.079174	0.354664	0.111418	0.131077
\	SELPLG_S3	SEMA3F_S2	SERPINE1_S2	SH2B3_S3

TCGA Participant Barcode

TCGA-WA-A7H4	0.037912	0.107456	0.047018	0.071774
TCGA-B5-A3F9	0.029117	0.075353	0.000925	0.022738
TCGA-49-6767	0.081667	0.044402	0.011848	0.141412
TCGA-AA-A03J	0.058516	0.041377	0.001889	0.051041
TCGA-77-7337	0.052901	0.132073	0.053159	0.074056

SH3BP5L_S5 SKA1_S5 SKA2_S5

SLC16A1_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	0.397680	0.023298	0.150416	0.187154
TCGA-B5-A3F9	0.161268	0.027082	0.163312	0.101799
TCGA-49-6767	0.147696	0.032137	0.151942	0.170208
TCGA-AA-A03J	0.289520	0.039170	0.019166	0.068127
TCGA-77-7337	0.161067	0.065265	0.160582	0.392069

SLC1A3_S3 SLC25A40_S5 SLC25A5_S5

SLC7A7_S3 \

TCGA Participant Barcode

TCGA-WA-A7H4	0.004853	0.113213	0.066678
0.008365			
TCGA-B5-A3F9	0.001806	0.050642	0.025590
0.008739			
TCGA-49-6767	0.001690	0.296410	0.051780
0.030765			
TCGA-AA-A03J	0.000242	0.103375	0.114527
0.017580			
TCGA-77-7337	0.018085	0.194974	0.062755
0.017081			

SMC2_S5 SM0_S2 SMS_S5 SMURF2_S5

SNRPA1_S5 \

TCGA Participant Barcode

TCGA-WA-A7H4	0.111563	0.061107	0.010822	0.117046
0.098420				
TCGA-B5-A3F9	0.059632	0.223859	0.042159	0.080097
0.150227				

TCGA-49-6767 0.155771	0.122536	0.120447	0.044985	0.469200
TCGA-AA-A03J 0.132591	0.029403	0.041830	0.026457	0.055882
TCGA-77-7337 0.228911	0.305418	0.065579	0.028911	0.139213
SNRPE_S5 \	SNRPA_S5	SNRPB_S5	SNRPC_S5	SNRPD1_S5
TCGA Participant Barcode				
TCGA-WA-A7H4 0.033202	0.129965	0.110974	0.189107	0.082965
TCGA-B5-A3F9 0.106333	0.357502	0.147041	0.194742	0.211044
TCGA-49-6767 0.081369	0.129030	0.153509	0.222232	0.130903
TCGA-AA-A03J 0.063577	0.353250	0.135313	0.214041	0.078309
TCGA-77-7337 0.061940	0.135565	0.076978	0.224079	0.174427
SRM_S5 \	SNX17_S5	SP140_S3	SPAG17_S5	SPARC_S2
TCGA Participant Barcode				
TCGA-WA-A7H4 0.102953	0.217200	0.025957	0.016643	0.045811
TCGA-B5-A3F9 0.081692	0.230067	0.012544	0.115356	0.010303
TCGA-49-6767 0.132022	0.169501	0.047222	0.000769	0.012549
TCGA-AA-A03J 0.129299	0.174395	0.009958	0.000777	0.012536
TCGA-77-7337 0.118792	0.178775	0.015842	0.027190	0.056416
TAGLN_S5 \	STAT1_S1	STK17A_S5	STRA13_S5	SYK_S3
TCGA Participant Barcode				
TCGA-WA-A7H4 0.038161	0.044481	0.241405	0.076512	0.059110
TCGA-B5-A3F9 0.003984	0.055906	0.059119	0.099121	0.168666
TCGA-49-6767 0.007983	0.118732	0.286971	0.054861	0.053159
TCGA-AA-A03J 0.025773	0.043672	0.024340	0.104535	0.138534
TCGA-77-7337	0.099685	0.158866	0.026717	0.132907

0.027327

	TAP1_S1	TBXAS1_S3	TCEB1_S5	TCF7L2_S2
THBS1_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.078096	0.035709	0.130174	0.099239
0.178477				
TCGA-B5-A3F9	0.086456	0.087330	0.103268	0.162975
0.012453				
TCGA-49-6767	0.153990	0.039003	0.143786	0.199046
0.012716				
TCGA-AA-A03J	0.231884	0.302471	0.052949	0.228019
0.031254				
TCGA-77-7337	0.064915	0.038927	0.159103	0.164192
0.093766				

	THBS2_S2	TIMP1_S2	TLR2_S3	TMEM130_S5
TNC_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.065712	0.028905	0.148018	0.000175
0.422519				
TCGA-B5-A3F9	0.001430	0.027609	0.075820	0.000108
0.015413				
TCGA-49-6767	0.028705	0.038713	0.048070	0.000505
0.152232				
TCGA-AA-A03J	0.018995	0.047791	0.023434	0.000256
0.017496				
TCGA-77-7337	0.158186	0.041740	0.283652	0.000990
0.302027				

	TNFAIP3_S3	TNFRSF12A_S5	TNFRSF1A_S2
TNFRSF1B_S3 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	0.080005	0.168056	0.252215
0.044209			
TCGA-B5-A3F9	0.100122	0.044841	0.219536
0.089376			
TCGA-49-6767	0.054746	0.121907	0.350215
0.129051			
TCGA-AA-A03J	0.058073	0.108840	0.273136
0.162439			
TCGA-77-7337	0.114188	0.049923	0.433645
0.040264			

	TNFSF13B_S3	TOMM40_S5	TPI1_S5
TPM1_S5 \			
TCGA Participant Barcode			

TCGA-WA-A7H4	0.007273	0.030447	0.055560	0.085022
TCGA-B5-A3F9	0.005195	0.039365	0.121625	0.019840
TCGA-49-6767	0.027343	0.044709	0.170366	0.031936
TCGA-AA-A03J	0.007791	0.085939	0.142167	0.048872
TCGA-77-7337	0.019876	0.041386	0.247876	0.027684

TUBA4A_S5 \

	TPM2_S5	TPM3_S5	TPRKB_S5	TRA2B_S5
--	---------	---------	----------	----------

TCGA Participant Barcode

TCGA-WA-A7H4	0.076884	0.248836	0.224479	0.370910
0.153193				
TCGA-B5-A3F9	0.010281	0.191961	0.432516	0.307541
0.002220				
TCGA-49-6767	0.029378	0.219096	0.172608	0.150605
0.040278				
TCGA-AA-A03J	0.013824	0.219590	0.089482	0.170531
0.110547				
TCGA-77-7337	0.008869	0.276453	0.239914	0.384092
0.063399				

TUBG1_S5 UAP1_S5 UBE2J1_S5 UMPS_S5

UQCR10_S5 \

	TUBG1_S5	UAP1_S5	UBE2J1_S5	UMPS_S5
--	----------	---------	-----------	---------

TCGA Participant Barcode

TCGA-WA-A7H4	0.249644	0.107880	0.182237	0.252513
0.037382				
TCGA-B5-A3F9	0.227805	0.080434	0.249096	0.145911
0.072831				
TCGA-49-6767	0.238908	0.078863	0.267850	0.215546
0.063526				
TCGA-AA-A03J	0.293324	0.038459	0.137453	0.194995
0.042015				
TCGA-77-7337	0.435219	0.129999	0.277074	0.312816
0.022331				

USPL1_S5 VCAN_S2 VDAC1_S5 VSIG4_S3

VTA1_S5 \

	USPL1_S5	VCAN_S2	VDAC1_S5	VSIG4_S3
--	----------	---------	----------	----------

TCGA Participant Barcode

TCGA-WA-A7H4	0.201070	0.072346	0.195607	0.003054
0.269151				
TCGA-B5-A3F9	0.190156	0.108592	0.229437	0.013031
0.175206				

TCGA-49-6767 0.184954	0.178736	0.040453	0.227530	0.023227
TCGA-AA-A03J 0.102899	0.102779	0.018265	0.353334	0.005992
TCGA-77-7337 0.348754	0.054357	0.206705	0.262335	0.017925
	WDHD1_S5	WDR54_S5	WDR77_S5	WIPF1_S3
WNT2B_S2 \				
TCGA Participant Barcode				
TCGA-WA-A7H4 0.010692	0.079282	0.072920	0.102072	0.089859
TCGA-B5-A3F9 0.006442	0.034451	0.188849	0.633403	0.067037
TCGA-49-6767 0.002030	0.094871	0.037184	0.124078	0.091530
TCGA-AA-A03J 0.012035	0.018202	0.034451	0.221328	0.026653
TCGA-77-7337 0.094993	0.114240	0.052860	0.092856	0.126317
	WNT8B_S2	WSB2_S5	ZWILCH_S5	ZYX_S2 \
TCGA Participant Barcode				
TCGA-WA-A7H4	0.009518	0.124660	0.154684	0.191966
TCGA-B5-A3F9	0.002612	0.089291	0.144194	0.126821
TCGA-49-6767	0.003011	0.142595	0.099884	0.293115
TCGA-AA-A03J	0.003367	0.134369	0.079119	0.256544
TCGA-77-7337	0.000520	0.167126	0.160216	0.132789
	Immune Subtype			
TCGA Participant Barcode				
TCGA-WA-A7H4	C1			
TCGA-B5-A3F9	C1			
TCGA-49-6767	C1			
TCGA-AA-A03J	C1			
TCGA-77-7337	C1			

```
df_final_normalized.to_csv("df_final_normalized.csv")
```

Outliers Handling

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.decomposition import NMF
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score
```

```

# Visualize Outliers
def visualize_outliers(df):
    plt.figure(figsize=(15, 5))
    sns.boxplot(data=df)
    plt.xticks(rotation=90)
    plt.title("Boxplot to detect outliers")
    plt.show()

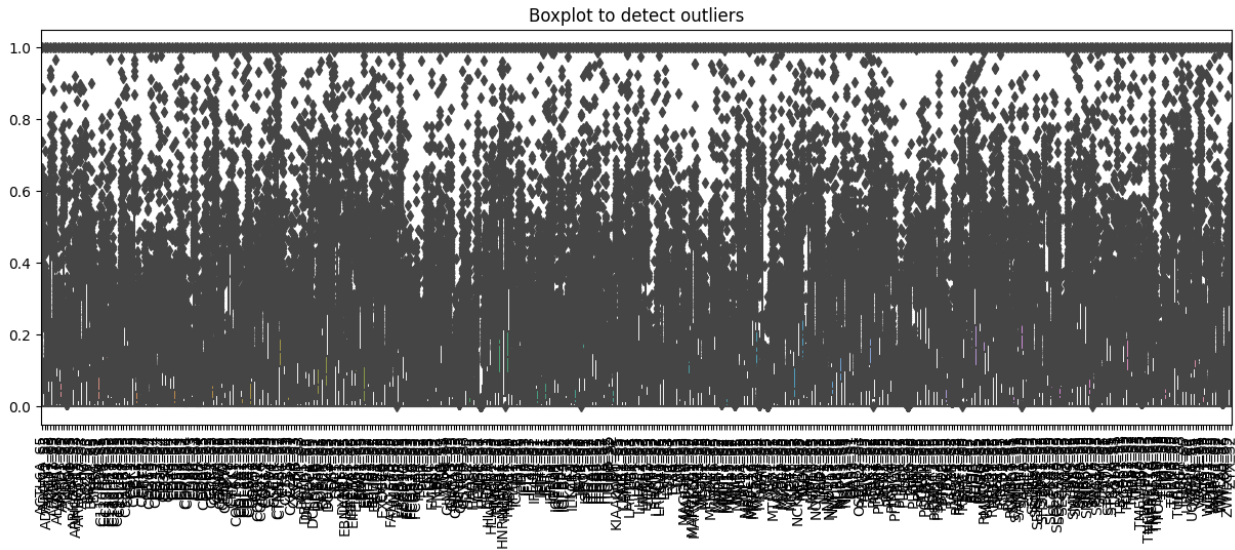
# Outlier Treatment using IQR Method
# Outlier Treatment using IQR Method (Handle Non-Numeric Columns)
def remove_outliers_iqr(df):
    numeric_df = df.select_dtypes(include=[np.number]) # Keep only
    numeric columns
    Q1 = numeric_df.quantile(0.25)
    Q3 = numeric_df.quantile(0.75)
    IQR = Q3 - Q1
    lower_bound = Q1 - 1.5 * IQR
    upper_bound = Q3 + 1.5 * IQR
    numeric_df = numeric_df.clip(lower=lower_bound, upper=upper_bound,
axis=1)

    # Merge back non-numeric columns
    df[numeric_df.columns] = numeric_df
    return df

# Load Data (Assuming df_final_normalized is pre-loaded)
# df_final_normalized = pd.read_csv("your_data.csv") # Uncomment if
loading from file

# Visualize and Remove Outliers
visualize_outliers(df_final_normalized)
df_final_normalized = remove_outliers_iqr(df_final_normalized)

```



Feature Reduction- Performing NMF

```
# Section 3: Perform NMF (excluding output variable)
X = df_final_normalized.drop(columns=['Immune Subtype']) # Exclude
target variable
nmf = NMF(n_components=50, random_state=42) # Choose appropriate
components
X_nmf = nmf.fit_transform(X)

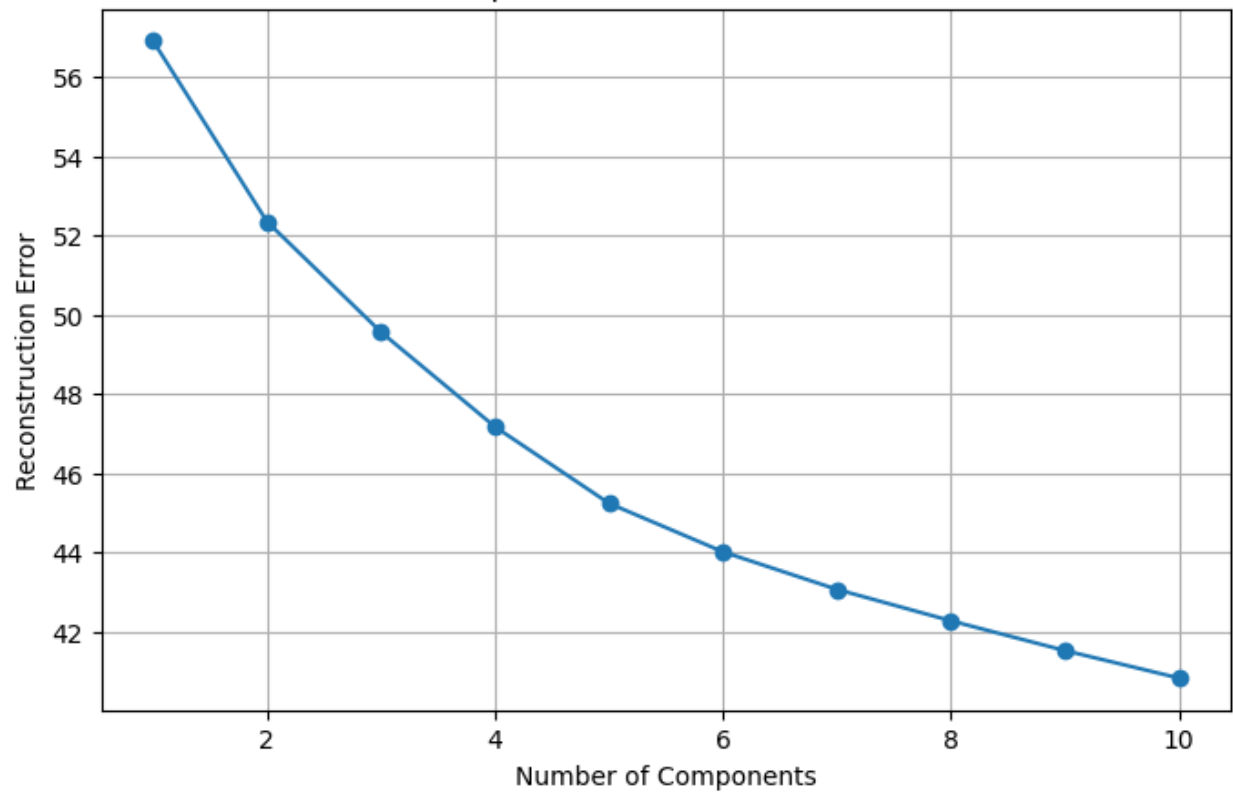
# Explained Reconstruction Error Plot
reconstruction_errors = []
components_range = range(1, 11)
for n in components_range:
    nmf_test = NMF(n_components=n, random_state=42)
    nmf_test.fit(X)
    reconstruction_errors.append(nmf_test.reconstruction_err_)

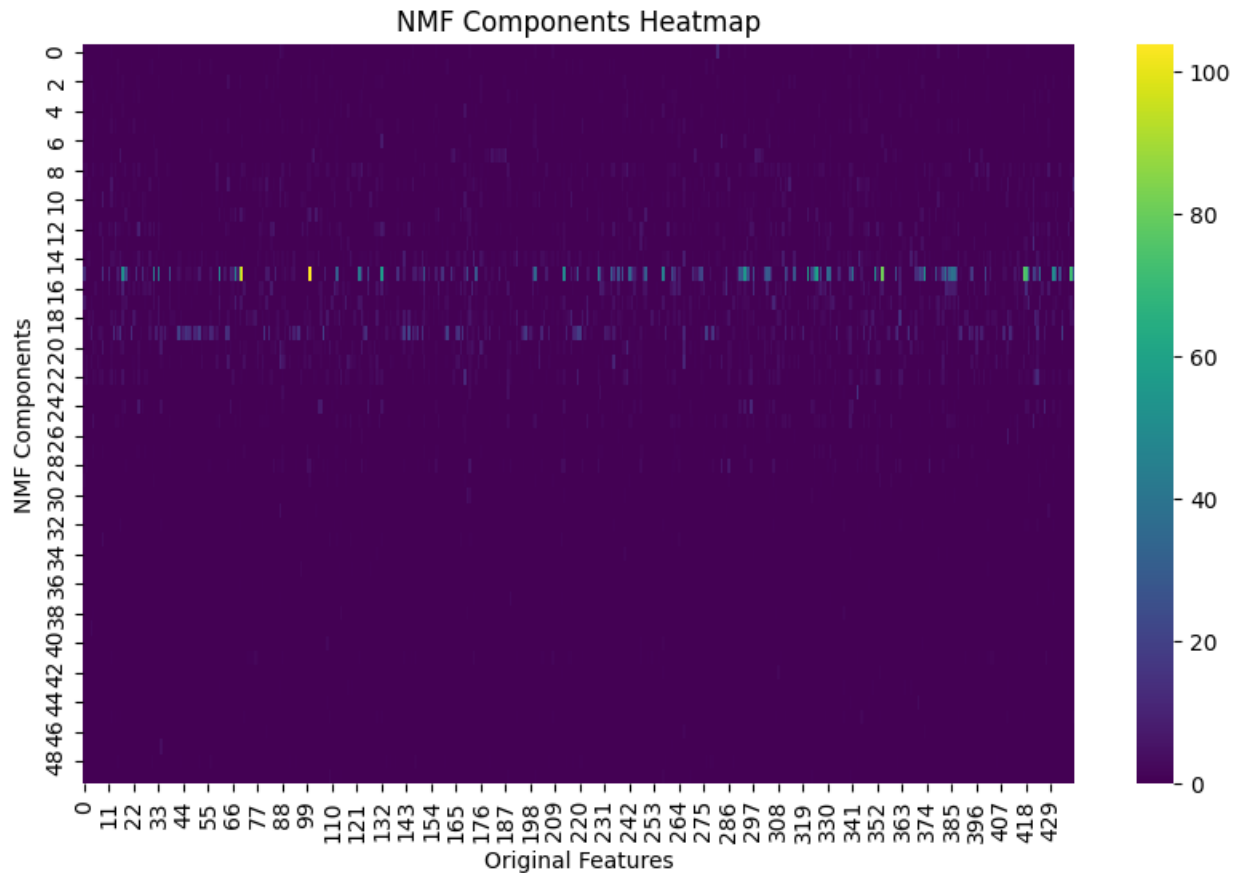
plt.figure(figsize=(8, 5))
plt.plot(components_range, reconstruction_errors, marker='o')
plt.xlabel('Number of Components')
plt.ylabel('Reconstruction Error')
plt.title('NMF Components vs. Reconstruction Error')
plt.grid()
plt.show()

# Visualizing NMF Components
plt.figure(figsize=(10, 6))
sns.heatmap(nmf.components_, cmap='viridis', annot=False)
plt.xlabel("Original Features")
plt.ylabel("NMF Components")
plt.title("NMF Components Heatmap")
plt.show()
```

[illegible]

NMF Components vs. Reconstruction Error





-knn,svc, xgb,nn different hyperparamters, cross val score, f1, recall -pros/cons for everything

- upto 6 minutes -pdf file will have comments as well its just a py version of it

Feature Correlation Plots

```
# Encode target variable
le = LabelEncoder()
y = le.fit_transform(df_final_normalized['Immune Subtype']) # Encode categorical labels

# Convert y to DataFrame
y_df = pd.DataFrame(y, columns=['Immune Subtype'])

# Drop target from X
X = df_final_normalized.drop(columns=['Immune Subtype'])

# Reset indices and concatenate
X = X.reset_index(drop=True)
y_df = y_df.reset_index(drop=True)
```

```

df = pd.concat([X, y_df], axis=1)

# Compute correlation matrix
corr_matrix = df.corr()

# 1. First look specifically at Immune Subtype correlations
print("Top correlations with Immune Subtype:")
print(corr_matrix["Immune Subtype"].sort_values(ascending=False).head(10))
print("\nBottom correlations with Immune Subtype:")
print(corr_matrix["Immune Subtype"].sort_values().head(10))

# 2. Better visualization focused on Immune Subtype
plt.figure(figsize=(8, 12))
immune_corrs = corr_matrix["Immune Subtype"].sort_values(ascending=False)
immune_corrs = immune_corrs[immune_corrs.index != "Immune Subtype"] # Remove self-correlation
sns.barplot(x=immune_corrs.values, y=immune_corrs.index, palette="coolwarm")
plt.title("Feature Correlations with Immune Subtype")
plt.xlabel("Correlation Coefficient")
plt.tight_layout()
plt.show()

# 3. Filtered heatmap version (adjusted threshold)
plt.figure(figsize=(15, 12))
threshold = 0.3 # More reasonable biological threshold

# Get top N features correlated with Immune Subtype
top_features = corr_matrix["Immune Subtype"].abs().sort_values(ascending=False).index[1:21] # Top 20 excluding self
filtered_corr = corr_matrix.loc[top_features, top_features]

sns.heatmap(filtered_corr, annot=True, cmap='coolwarm',
            fmt='.2f', vmin=-1, vmax=1, center=0,
            annot_kws={"size": 8})
plt.title("Top 20 Features Correlated with Immune Subtype")
plt.xticks(rotation=45, ha='right')
plt.tight_layout()
plt.show()

```

Top correlations with Immune Subtype:

Immune Subtype	1.000000
RASSF4_S3	0.471393
RASSF2_S3	0.467887
CPEB4_S5	0.456713
SLC1A3_S3	0.427795
CELF2_S3	0.415925

EVI2A_S3	0.410678
APOE_S3	0.408811
CSF1R_S3	0.407040
FAM167A_S5	0.404026

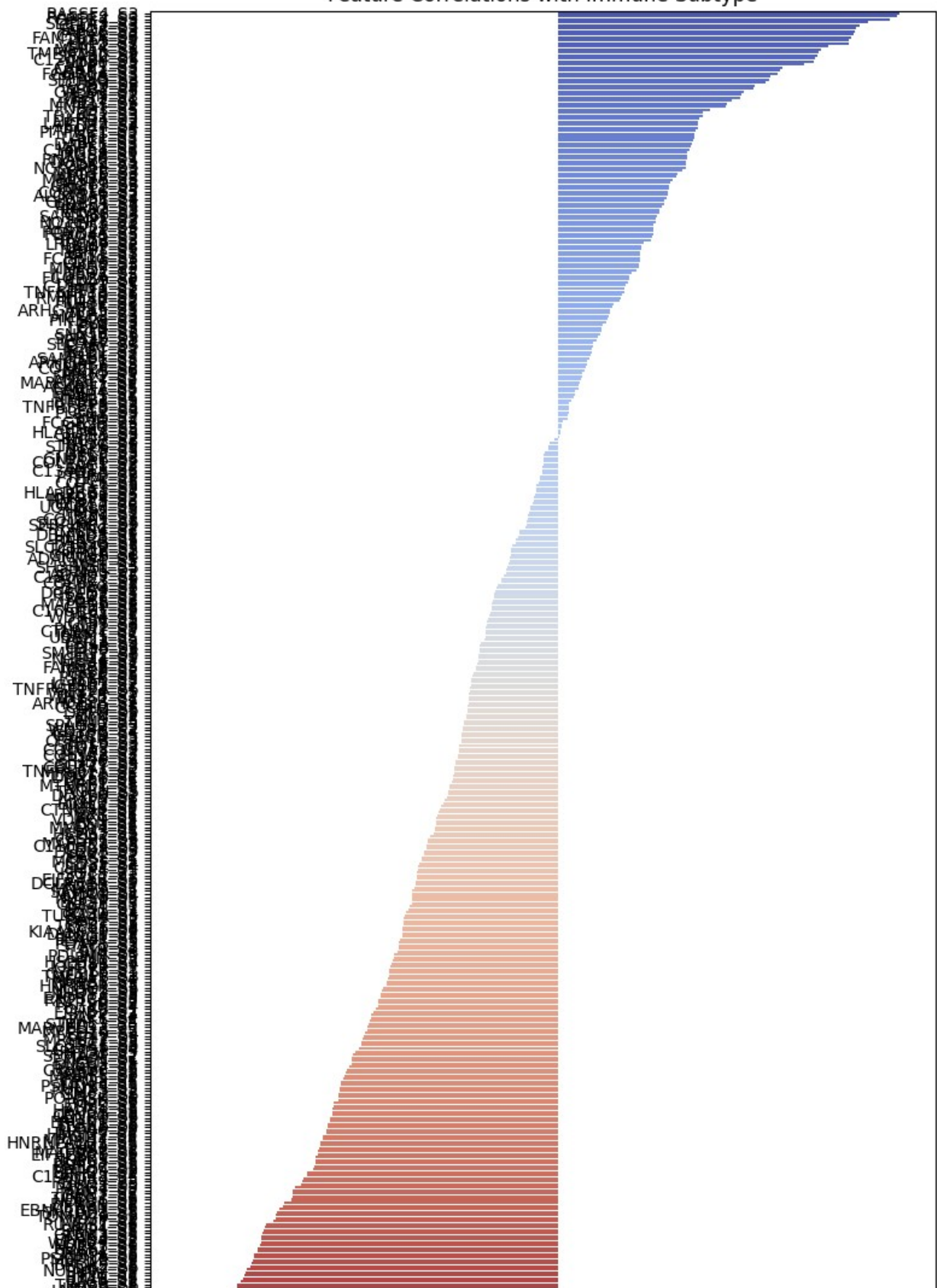
Name: Immune Subtype, dtype: float64

Bottom correlations with Immune Subtype:

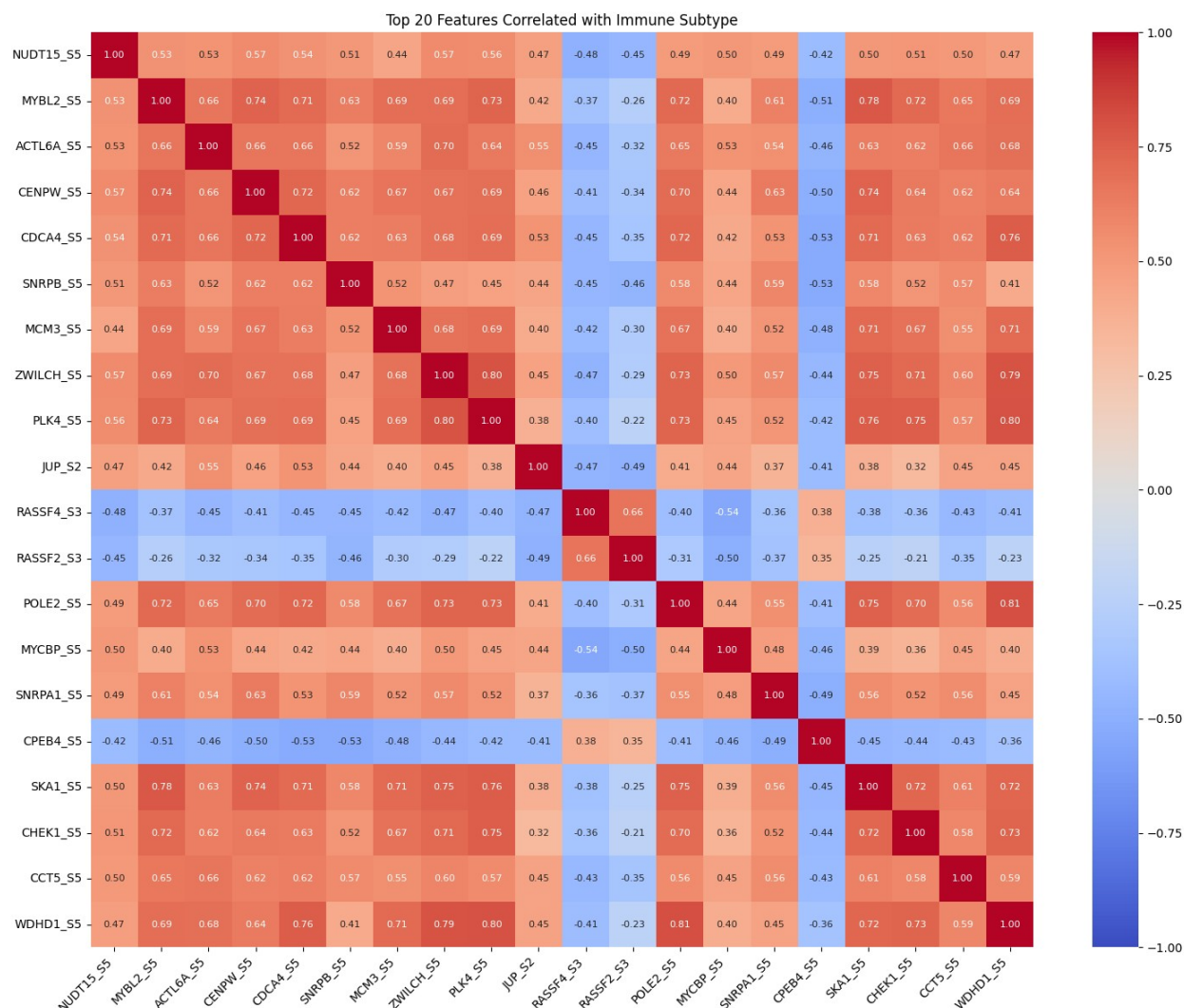
NUDT15_S5	-0.514211
MYBL2_S5	-0.513117
ACTL6A_S5	-0.505404
CENPW_S5	-0.496627
CDCA4_S5	-0.491714
SNRPB_S5	-0.487020
MCM3_S5	-0.479625
ZWILCH_S5	-0.479406
PLK4_S5	-0.476722
JUP_S2	-0.476225

Name: Immune Subtype, dtype: float64

Feature Correlations with Immune Subtype



```
/usr/local/lib/python3.10/dist-packages/matplotlib/colors.py:721:
RuntimeWarning: invalid value encountered in less
xa[xa < 0] = -1
```



```
X_selected = df_final_normalized[top_features]
X_selected.head()
```

	NUDT15_S5	MYBL2_S5	ACTL6A_S5	CENPW_S5
CDCA4_S5 \				
TCGA Participant Barcode				
TCGA-WA-A7H4	0.158879	0.036125	0.129424	0.036657
0.239033				
TCGA-B5-A3F9	0.124530	0.032607	0.123828	0.029258
0.158576				
TCGA-49-6767	0.164204	0.031808	0.150459	0.022736

0.182581				
TCGA-AA-A03J	0.106287	0.108375	0.083904	0.049675
0.109814				
TCGA-77-7337	0.159017	0.103269	0.251188	0.049675
0.242704				

	SNRPB_S5	MCM3_S5	ZWILCH_S5	PLK4_S5
JUP_S2 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.110974	0.216977	0.154684	0.272897
0.081848				
TCGA-B5-A3F9	0.147041	0.152350	0.144194	0.139299
0.020165				
TCGA-49-6767	0.153509	0.115889	0.099884	0.198844
0.048325				
TCGA-AA-A03J	0.135313	0.175003	0.079119	0.082470
0.076755				
TCGA-77-7337	0.076978	0.076036	0.160216	0.208554
0.102031				

	RASSF4_S3	RASSF2_S3	POLE2_S5	MYCBP_S5
SNRPA1_S5 \				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.025563	0.006755	0.136847	0.094650
0.098420				
TCGA-B5-A3F9	0.028804	0.012288	0.096535	0.157460
0.150227				
TCGA-49-6767	0.053798	0.002001	0.152285	0.093955
0.155771				
TCGA-AA-A03J	0.026893	0.001996	0.073111	0.080209
0.132591				
TCGA-77-7337	0.006769	0.004547	0.182561	0.062224
0.228911				

	CPEB4_S5	SKA1_S5	CHEK1_S5	CCT5_S5
WDHD1_S5				
TCGA Participant Barcode				

TCGA-WA-A7H4	0.066910	0.023298	0.341859	0.139557
0.079282				
TCGA-B5-A3F9	0.037612	0.027082	0.168456	0.123400
0.034451				
TCGA-49-6767	0.019654	0.032137	0.319432	0.213498
0.094871				
TCGA-AA-A03J	0.024473	0.039170	0.107756	0.159937
0.018202				
TCGA-77-7337	0.073843	0.065265	0.422631	0.222876
0.114240				

```
df.head()
```

	ACTL6A_S5	ADAM9_S2	ADAMTS1_S5	ADCY7_S3	AIMP2_S5	ALKBH7_S5
ALOX5AP_S3 \						
0	0.129424	0.169197	0.022398	0.234976	0.180090	0.036046
	0.025048					
1	0.123828	0.033567	0.039275	0.306215	0.153894	0.062357
	0.019597					
2	0.150459	0.157022	0.003525	0.249774	0.155868	0.019391
	0.051084					
3	0.083904	0.039410	0.004848	0.088996	0.413569	0.143618
	0.013858					
4	0.251188	0.217491	0.057262	0.140494	0.383001	0.020735
	0.078867					

	AMPD3_S3	APITD1_S5	APOC1_S3	AP0E_S3	AP00_S5	ARHGAP1_S2
ARHGAP15_S3 \						
0	0.083360	0.183705	0.000159	0.001151	0.023024	0.199662
	0.034113					
1	0.049824	0.299588	0.001520	0.004717	0.137673	0.067752
	0.010200					
2	0.038690	0.237127	0.001623	0.005735	0.081950	0.066406
	0.055246					
3	0.020518	0.097687	0.000464	0.002793	0.054958	0.165810
	0.009896					
4	0.043357	0.325132	0.000145	0.000614	0.069031	0.239196
	0.022262					

	ARHGDIA_S2	ARRB2_S3	B2M_S3	BCCIP_S5	BRCA2_S5	BRIP1_S5
BSG_S2 \						
0	0.193792	0.062812	0.149856	0.137327	0.135782	0.095438
	0.044576					
1	0.143697	0.062524	0.207232	0.263794	0.042710	0.028896
	0.060415					
2	0.111708	0.088977	0.166943	0.266651	0.066250	0.052106
	0.018203					
3	0.310192	0.181536	0.125507	0.104531	0.088548	0.020854
	0.052634					
4	0.188949	0.069010	0.100442	0.243811	0.089324	0.118525
	0.032608					

	BTK_S3	C11orf24_S5	C12orf24_S5	C13orf1_S5	C13orf18_S3
C13orf27_S5 \					
0	0.034921	0.152497	0.022158	0.155023	0.016833
	0.076541				
1	0.024206	0.077849	0.031294	0.136670	0.026442
	0.070951				
2	0.063984	0.143469	0.022671	0.191243	0.005027
	0.065347				
3	0.010746	0.114090	0.017954	0.247442	0.043303

0.044817
4 0.034656 0.159544 0.028050 0.165591 0.005849
0.099976

	C16orf61_S5	C19orf48_S5	C1orf54_S3	C3AR1_S3	C3orf26_S5
CASP1_S3 \					
0	0.045475	0.019964	0.091298	0.033466	0.252371
0.130334					
1	0.076200	0.041087	0.120955	0.038793	0.198402
0.018940					
2	0.076236	0.017507	0.080028	0.094222	0.139277
0.032013					
3	0.064950	0.076582	0.036174	0.032520	0.106481
0.075871					
4	0.078923	0.023613	0.045622	0.093986	0.300077
0.110604					

	CBX1_S5	CCBL2_S5	CCL2_S3	CCL5_S3	CCRL2_S3	CCT5_S5
CD14_S4 \						
0	0.104727	0.357476	0.017832	0.039156	0.006166	0.139557
0.018676						
1	0.171738	0.306802	0.031532	0.017448	0.004626	0.123400
0.022774						
2	0.128322	0.402032	0.032674	0.069883	0.012335	0.213498
0.021355						
3	0.032466	0.193422	0.006116	0.021166	0.044520	0.159937
0.019312						
4	0.142077	0.304238	0.053161	0.019366	0.010812	0.222876
0.025416						

	CD163_S3	CD19_S4	CD247_S4	CD33_S3	CD37_S4	CD3D_S4
CD3E_S4 \						
0	0.009689	0.003777	0.063614	0.033109	0.015664	0.040169
0.023715						
1	0.017677	0.002194	0.035106	0.071084	0.017639	0.040378
0.024578						
2	0.025298	0.000785	0.083596	0.075230	0.042144	0.086689
0.076188						
3	0.004955	0.003470	0.044976	0.031795	0.022887	0.047478
0.023950						
4	0.038462	0.001408	0.039633	0.049115	0.016808	0.034323
0.032421						

	CD3G_S4	CD44_S2	CD48_S3	CD52_S3	CD53_S3	CD59_S2
CD79A_S4 \						
0	0.016702	0.075255	0.014924	0.018208	0.038376	0.113421
0.012376						
1	0.005270	0.008014	0.024867	0.016722	0.053371	0.043971
0.001537						
2	0.065392	0.038649	0.040068	0.061403	0.100435	0.047856

0.003519					
3 0.020800	0.012806	0.015225	0.026161	0.035758	0.014871
0.005998					
4 0.047515	0.092642	0.020929	0.021409	0.102105	0.107698
0.016165					
CD79B_S4	CD84_S3	CD86_S3	CD8A_S3	CD8B_S4	CD97_S3
CDCA4_S5 \					
0 0.009356	0.003355	0.110400	0.004560	0.002090	0.030530
0.239033					
1 0.003067	0.004642	0.049360	0.026804	0.016076	0.052208
0.158576					
2 0.003751	0.010102	0.144443	0.077024	0.008436	0.073799
0.182581					
3 0.005639	0.004176	0.036119	0.037436	0.006757	0.201381
0.109814					
4 0.005231	0.018940	0.188385	0.028269	0.008949	0.052528
0.242704					
CDH6_S2	CDK2_S5	CECR1_S3	CELF2_S3	CENPJ_S5	CENPN_S5
CENPO_S5 \					
0 0.004352	0.029561	0.022926	0.054613	0.077392	0.058463
0.176886					
1 0.003591	0.031723	0.101669	0.044890	0.183160	0.022141
0.065257					
2 0.000571	0.038141	0.054018	0.036824	0.028629	0.047939
0.127073					
3 0.001170	0.028389	0.024335	0.097108	0.066650	0.052004
0.166096					
4 0.028240	0.033506	0.036337	0.041916	0.035145	0.106980
0.339721					
CENPW_S5	CEP78_S5	CHEK1_S5	CKLF_S5	COL16A1_S2	COL1A2_S2
COL3A1_S2 \					
0 0.036657	0.158284	0.341859	0.055799	0.130675	0.049270
0.053382					
1 0.029258	0.187278	0.168456	0.060539	0.006053	0.005468
0.011437					
2 0.022736	0.143665	0.319432	0.279732	0.023565	0.011242
0.010848					
3 0.049675	0.048390	0.107756	0.138498	0.022623	0.016113
0.012296					
4 0.049675	0.336437	0.422631	0.063020	0.028620	0.055869
0.059424					
COL6A1_S2	COL6A3_S2	COL8A1_S2	COPS6_S5	CQ2_S5	COR01A_S3
COR01C_S5 \					
0 0.090780	0.168243	0.020451	0.073187	0.151087	0.039494
0.153347					
1 0.041633	0.007959	0.000976	0.111060	0.135605	0.031174

0.040879
2 0.083508 0.050570 0.038859 0.235384 0.148317 0.079423
0.212910
3 0.036574 0.051404 0.023293 0.115420 0.190231 0.173819
0.140236
4 0.030022 0.203530 0.087967 0.089021 0.137881 0.027044
0.337473

COTL1_S5 COX17_S5 CPEB4_S5 CPVL_S3 CSF1R_S3 CTNNA1_S2
CTNNB1_S2 \

0 0.089469 0.077845 0.066910 0.008639 0.035377 0.332514
0.163481
1 0.102082 0.119659 0.037612 0.009777 0.031854 0.324371
0.128645
2 0.274675 0.130252 0.019654 0.016773 0.048437 0.276330
0.273783
3 0.068212 0.087111 0.024473 0.009468 0.032764 0.336343
0.202381
4 0.110654 0.053663 0.073843 0.012768 0.056842 0.322352
0.248077

CTSC_S3 CTSL1_S3 CTSS_S3 CXCL10_S1 CXCR4_S3 CYBB_S3
CYTIP_S3 \

0 0.095917 0.023556 0.018769 0.003976 0.009499 0.018678
0.030886
1 0.058378 0.034386 0.040411 0.006545 0.083557 0.018557
0.038931
2 0.080481 0.025965 0.171797 0.032839 0.019742 0.066786
0.049761
3 0.042648 0.007856 0.069236 0.006464 0.008578 0.013780
0.012738
4 0.191799 0.044214 0.046477 0.012050 0.038591 0.066092
0.060276

DAPK1_S3 DBNDD1_S5 DCBLD2_S5 DCK_S5 DCLRE1B_S5 DDX58_S1
DDX60_S1 \

0 0.082398 0.015780 0.107223 0.160330 0.287794 0.053464
0.072989
1 0.041218 0.066342 0.069558 0.103752 0.207635 0.026454
0.059039
2 0.013979 0.069333 0.107223 0.256074 0.232055 0.031037
0.078268
3 0.026551 0.027064 0.016922 0.102256 0.117280 0.027088
0.072232
4 0.027985 0.013411 0.101117 0.293041 0.243294 0.014842
0.074897

DHFR_S5 DLEU1_S5 DLEU2_S5 DOCK2_S3 DSP_S2 DUT_S5
DVL1_S2 \

0 0.085747 0.097426 0.145469 0.044880 0.118441 0.202159

0.284554						
1	0.165736	0.245137	0.103178	0.021801	0.017743	0.286724
0.189871						
2	0.086690	0.111307	0.224930	0.071234	0.036720	0.093573
0.239620						
3	0.152287	0.163563	0.053175	0.032179	0.033738	0.357800
0.412409						
4	0.065244	0.207484	0.215501	0.069890	0.118441	0.082165
0.101505						
	DVL3_S2	DYNLT1_S5	EBNA1BP2_S5	EFNA5_S2	EIF2AK1_S5	EIF4EBP1_S5
\						
0	0.069952	0.117494	0.254913	0.100997	0.180324	0.045329
1	0.051480	0.268827	0.138149	0.012586	0.275381	0.067787
2	0.055469	0.111976	0.229355	0.003574	0.234314	0.050975
3	0.082258	0.106556	0.141679	0.001799	0.280938	0.065817
4	0.123503	0.112132	0.123693	0.058037	0.311581	0.112676
	EIF4G1_S5	EMP2_S5	EN01_S5	EPHA2_S2	EPHB2_S2	EPHB3_S2
EPHB4_S2	\					
0	0.224737	0.073334	0.176186	0.100160	0.030938	0.032859
0.146180						
1	0.099652	0.037854	0.106735	0.038517	0.027591	0.009309
0.103594						
2	0.234288	0.092133	0.181019	0.065869	0.119018	0.014559
0.315124						
3	0.181465	0.046118	0.087059	0.199168	0.119018	0.072677
0.288277						
4	0.303863	0.117772	0.311886	0.153488	0.119018	0.059069
0.105869						
	ERLIN1_S5	EVI2A_S3	EVI2B_S3	EXOSC8_S5	EZR_S5	F3_S5
FAM167A_S5	\					
0	0.144039	0.018113	0.023936	0.161982	0.171173	0.084577
0.010754						
1	0.204526	0.016738	0.019382	0.184557	0.076877	0.011839
0.002479						
2	0.411496	0.041415	0.057620	0.227998	0.257706	0.033512
0.008293						
3	0.213134	0.004571	0.012131	0.180274	0.154220	0.012739
0.004658						
4	0.411496	0.048020	0.049250	0.111053	0.132114	0.036959
0.003482						
	FAM89B_S5	FARSA_S5	FARSB_S5	FBX041_S5	FCER1G_S3	FCGR1A_S3

FCGR2A_S3 \						
0	0.204180	0.102053	0.329130	0.047208	0.018273	0.006308
0.030941						
1	0.169258	0.111014	0.328010	0.007366	0.044371	0.028072
0.025621						
2	0.316784	0.065963	0.376409	0.048723	0.077242	0.066439
0.067265						
3	0.160079	0.230402	0.287552	0.107019	0.017333	0.007035
0.028258						
4	0.129489	0.113086	0.279407	0.012326	0.045339	0.019803
0.161422						

FCGR2B_S3	FCGR3A_S3	FGD1_S2	FGL2_S3	FGR_S3	FLI1_S3
FLNC_S5 \					
0	0.012875	0.013803	0.020552	0.006781	0.032155
0.031307					
1	0.010577	0.024701	0.088063	0.003095	0.017032
0.001224					
2	0.021345	0.071530	0.093016	0.011098	0.067096
0.031307					
3	0.003753	0.012462	0.025836	0.005562	0.030626
0.013318					
4	0.030141	0.057074	0.043341	0.011555	0.039693
0.002109					

FMNL1_S3	FN1_S2	FNBP1_S3	FPR3_S3	FYB_S3	GGH_S5
GIMAP4_S3 \					
0	0.101362	0.027169	0.046967	0.056917	0.053084
0.025146					
1	0.022260	0.001267	0.065727	0.020283	0.041086
0.039299					
2	0.272864	0.028097	0.098693	0.104719	0.066201
0.071292					
3	0.074379	0.003745	0.039936	0.040684	0.010794
0.024416					
4	0.059797	0.051011	0.032962	0.125610	0.181051
0.049023					

GLRX3_S5	GNG11_S5	GNPTAB_S3	GPLD1_S5	GPNMB_S3	GSTCD_S5
H2AFZ_S5 \					
0	0.160717	0.010046	0.057354	0.000408	0.053609
0.152008					
1	0.149249	0.013832	0.051172	0.000736	0.012891
0.199343					
2	0.331305	0.003070	0.026608	0.000730	0.036157
0.123665					
3	0.174956	0.002216	0.023469	0.000500	0.003611
0.145930					
4	0.331305	0.005633	0.087136	0.000756	0.101994
0.179624					

	HAS2_S5	HAUS1_S5	HCK_S3	HCLS1_S3	HDC_S4	HERC5_S1
HERC6_S1 \						
0	0.037437	0.020346	0.047622	0.057652	0.001211	0.025456
	0.075657					
1	0.014518	0.067098	0.038908	0.086070	0.000030	0.014123
	0.053794					
2	0.004785	0.015883	0.099041	0.085216	0.000035	0.009157
	0.022453					
3	0.010080	0.021343	0.050230	0.035609	0.000945	0.005109
	0.041770					
4	0.061951	0.033436	0.065982	0.227667	0.000368	0.036022
	0.051997					

	HLA-DMA_S3	HLA-DRB1_S3	HMG2_S5	HMHA1_S3	HN1L_S5	HNRNPA2B1_S5
\						
0	0.022540	0.047645	0.136413	0.071628	0.271469	0.475610
1	0.103293	0.086854	0.244664	0.034324	0.280583	0.548851
2	0.060781	0.204553	0.142709	0.088358	0.161530	0.251151
3	0.053130	0.069789	0.071459	0.163964	0.157864	0.164280
4	0.052412	0.073172	0.168506	0.019563	0.358311	0.172482

	HNRNPR_S5	HSPB11_S5	HSPG2_S2	HYLS1_S5	ICAM1_S2	ID2_S5
ID3_S5 \						
0	0.399541	0.218957	0.188063	0.109928	0.034119	0.037530
	0.030801					
1	0.388752	0.224201	0.064808	0.062105	0.012642	0.209001
	0.159889					
2	0.299662	0.096953	0.078204	0.078556	0.118333	0.032776
	0.007644					
3	0.170824	0.041110	0.171041	0.021903	0.022959	0.106346
	0.055888					
4	0.437498	0.073749	0.206996	0.151481	0.066420	0.037849
	0.035191					

	IFI16_S3	IFI27_S1	IFI30_S3	IFI44_S1	IFI44L_S1	IFI6_S1
IFIH1_S1 \						
0	0.174878	0.010585	0.038855	0.092889	0.008884	0.016181
	0.052465					
1	0.044124	0.030673	0.047912	0.042425	0.014465	0.041187
	0.044148					
2	0.064804	0.033371	0.092049	0.082192	0.066570	0.005650
	0.088845					
3	0.011214	0.056305	0.077159	0.012154	0.010863	0.031855
	0.034967					

4	0.190937	0.023718	0.078464	0.022584	0.002674	0.007654
0.061444						

	IFIT1_S1	IFIT2_S1	IFIT3_S1	IFRD2_S5	IGF2R_S2	IGFBP2_S2
IGFBP3_S2 \						
0	0.012104	0.013984	0.032078	0.174951	0.417760	0.008048
0.011803						
1	0.009991	0.020951	0.022286	0.148126	0.090358	0.046128
0.017299						
2	0.007317	0.059526	0.069046	0.249150	0.292550	0.000552
0.008982						
3	0.009420	0.017772	0.046223	0.406637	0.304512	0.004864
0.010374						
4	0.004512	0.012758	0.028578	0.124414	0.286602	0.029685
0.073992						

	IGFBP4_S2	IGFBP5_S2	IGJ_S4	IKZF1_S3	IL10RA_S3	IL18_S3
IL7R_S3 \						
0	0.076637	0.006510	0.055104	0.030634	0.043147	0.083119
0.079098						
1	0.259275	0.002993	0.003288	0.016425	0.033162	0.005650
0.010949						
2	0.053771	0.007041	0.002262	0.039746	0.075596	0.109420
0.013034						
3	0.062825	0.004197	0.009832	0.014251	0.045385	0.025072
0.010332						
4	0.099363	0.027515	0.038158	0.027573	0.042415	0.051308
0.094333						

	IMP4_S5	IP04_S5	ISG15_S1	ITGA3_S2	ITGA4_S2	ITGA5_S2
ITGA6_S5 \						
0	0.139524	0.207161	0.014778	0.251271	0.053032	0.086581
0.139174						
1	0.130361	0.212707	0.037194	0.015699	0.018388	0.012094
0.032617						
2	0.160946	0.200339	0.022078	0.251271	0.034460	0.049861
0.042737						
3	0.283705	0.306210	0.056967	0.068338	0.023130	0.024807
0.061921						
4	0.150310	0.249193	0.008037	0.105767	0.067336	0.151642
0.139174						

	ITGB2_S2	ITGB3_S2	ITGB5_S2	ITGB8_S2	JUNB_S5	JUP_S2
KIAA0090_S5 \						
0	0.021288	0.017033	0.202443	0.024935	0.028278	0.081848
0.409317						
1	0.026752	0.001087	0.097532	0.012690	0.029653	0.020165
0.161326						
2	0.100076	0.007168	0.096397	0.027774	0.011750	0.048325
0.171628						

3	0.036380	0.003954	0.107455	0.003000	0.051715	0.076755
0.081225						
4	0.038590	0.014997	0.112845	0.013661	0.027507	0.102031
0.409317						

	KRR1_S5	LAIR1_S3	LAMA4_S2	LAMB1_S2	LAPTM5_S3	LCK_S4
LCP1_S3 \						
0	0.266041	0.031268	0.063886	0.134072	0.029875	0.023022
0.057339						
1	0.163491	0.054181	0.010380	0.166282	0.057293	0.023920
0.078932						
2	0.144502	0.087119	0.012079	0.102749	0.151932	0.060208
0.098733						
3	0.086434	0.035759	0.020305	0.133157	0.037515	0.026947
0.024536						
4	0.355570	0.050773	0.065265	0.125134	0.083237	0.029194
0.103698						

	LCP2_S3	LHFPL2_S3	LILRB4_S3	LMNB2_S5	LOXL2_S5	LRMP_S3
LRP1_S2 \						
0	0.062984	0.132281	0.008157	0.250072	0.030669	0.004127
0.021220						
1	0.038561	0.092902	0.020459	0.144390	0.008254	0.002365
0.009356						
2	0.108373	0.124139	0.039352	0.172712	0.023336	0.001690
0.010615						
3	0.035882	0.090632	0.016934	0.366108	0.024214	0.001860
0.042072						
4	0.121301	0.178464	0.027372	0.233035	0.045484	0.025887
0.048902						

	LRRC17_S2	LRRC40_S5	LSM3_S5	LSM4_S5	LST1_S3	LTB_S4
LY86_S3 \						
0	0.060840	0.178373	0.109577	0.100458	0.045123	0.042660
0.009661						
1	0.025620	0.193141	0.163247	0.198695	0.031602	0.018470
0.030685						
2	0.011497	0.233212	0.156403	0.109571	0.061070	0.057014
0.040346						
3	0.003852	0.017883	0.069174	0.172504	0.037040	0.024858
0.014650						
4	0.016909	0.136157	0.111263	0.089692	0.026585	0.002604
0.019700						

	LYAR_S5	LYN_S3	LYZ_S3	MAGOHB_S5	MAP3K8_S5	MAPRE1_S5
MARCKSL1_S2 \						
0	0.210344	0.114421	0.031317	0.075697	0.035770	0.145664
0.040757						
1	0.104753	0.099920	0.001993	0.107498	0.062074	0.090899
0.385772						

2	0.229836	0.346249	0.016032	0.104949	0.027099	0.134828
	0.170430					
3	0.189843	0.072093	0.007370	0.048341	0.018603	0.160546
	0.126210					
4	0.229836	0.239733	0.016140	0.135993	0.038159	0.246152
	0.027627					

	MARVELD2_S5	MCM3_S5	MCM7_S5	MCTS1_S5	MERTK_S3	MET_S5
MFSD11_S5 \						
0	0.113350	0.216977	0.074077	0.217205	0.010786	0.028999
	0.368101					
1	0.190823	0.152350	0.075562	0.121013	0.046835	0.013101
	0.097814					
2	0.179898	0.115889	0.199362	0.217010	0.033905	0.072149
	0.274009					
3	0.227938	0.175003	0.194599	0.101290	0.007017	0.019980
	0.164033					
4	0.180933	0.076036	0.081903	0.082588	0.035576	0.032555
	0.316945					

	MKKS_S5	MLF1IP_S5	MMP11_S2	MMP1_S2	MMP14_S2	MMP17_S2
MMP19_S2 \						
0	0.106318	0.169909	0.073249	0.007719	0.138782	0.028384
	0.021184					
1	0.118294	0.188447	0.073249	0.000080	0.054637	0.005161
	0.028822					
2	0.054881	0.168724	0.068895	0.007719	0.169203	0.012879
	0.011475					
3	0.035276	0.074050	0.062199	0.007719	0.054713	0.003805
	0.010316					
4	0.063507	0.104407	0.017603	0.007719	0.087733	0.009456
	0.012750					

	MMP2_S2	MMP3_S2	MNAT1_S5	MNDA_S3	MPP1_S3	MRPL12_S5
MRPL37_S5 \						
0	0.076244	0.004697	0.297820	0.034854	0.028984	0.053989
	0.215832					
1	0.067427	0.001807	0.460132	0.023676	0.023693	0.071177
	0.149825					
2	0.030269	0.000234	0.363728	0.134574	0.113771	0.053678
	0.175349					
3	0.044198	0.004697	0.067786	0.015200	0.119624	0.128341
	0.340748					
4	0.092904	0.004697	0.351699	0.055352	0.033229	0.046913
	0.150464					

	MRPS16_S5	MRPS28_S5	MRT04_S5	MS4A1_S4	MSN_S3	MT1F_S5
MT3_S5 \						
0	0.081801	0.152298	0.296662	0.001867	0.225881	0.001859
	0.000017					

1	0.159533	0.102605	0.261444	0.000091	0.102073	0.007738
0.000714						
2	0.146096	0.056819	0.206756	0.000514	0.523028	0.001701
0.000002						
3	0.150107	0.026642	0.162677	0.000202	0.046191	0.005101
0.000447						
4	0.227553	0.115047	0.292974	0.000489	0.342708	0.001910
0.000003						

	MTA1_S2	MTHFD1_S5	MX1_S1	MYBL1_S5	MYBL2_S5	MYCBP_S5
MYL6_S5 \						
0	0.108317	0.053998	0.023341	0.044450	0.036125	0.094650
0.092348						
1	0.205529	0.049750	0.022720	0.011306	0.032607	0.157460
0.113473						
2	0.180721	0.111187	0.022676	0.059719	0.031808	0.093955
0.110403						
3	0.161050	0.065647	0.048149	0.005645	0.108375	0.080209
0.084048						
4	0.101209	0.098276	0.018135	0.014082	0.103269	0.062224
0.073702						

	MYO1F_S3	NCEH1_S5	NCF2_S3	NCKAP1L_S3	NCLN_S5	NEO1_S2
NID1_S2 \						
0	0.060112	0.033299	0.088336	0.041589	0.124847	0.103046
0.011977						
1	0.070195	0.056313	0.018637	0.033273	0.134272	0.214586
0.011057						
2	0.105832	0.160207	0.182093	0.078590	0.109027	0.077230
0.008704						
3	0.102991	0.036514	0.035316	0.045108	0.249041	0.307042
0.007268						
4	0.046165	0.096981	0.237740	0.061815	0.095514	0.075113
0.034183						

	NLN_S5	NME1_S5	NOP16_S5	NOTCH2_S2	NPC2_S3	NPL_S3
NRIP3_S5 \						
0	0.239173	0.064625	0.164285	0.111047	0.015887	0.049165
0.003531						
1	0.232128	0.127969	0.271940	0.120854	0.047381	0.029214
0.000597						
2	0.331265	0.099628	0.135435	0.145194	0.056083	0.026651
0.004892						
3	0.205279	0.084575	0.345630	0.028016	0.009288	0.019027
0.000524						
4	0.281952	0.074518	0.102710	0.157209	0.052034	0.165051
0.009000						

	NUDT1_S5	NUDT15_S5	NUP107_S5	NUP35_S5	NUP85_S5	NUP93_S5
NUPL1_S5 \						

0	0.080380	0.158879	0.083663	0.173388	0.174390	0.121559
0.210659						
1	0.057976	0.124530	0.071152	0.272161	0.137736	0.136196
0.120356						
2	0.050619	0.164204	0.071165	0.173997	0.171342	0.165973
0.183313						
3	0.072168	0.106287	0.048762	0.081494	0.186738	0.159651
0.194111						
4	0.089115	0.159017	0.091544	0.314542	0.138004	0.163207
0.169962						

	NUTF2_S5	OAS1_S1	OAS2_S1	OAS3_S1	OASL_S1	OSBPL3_S3
PA2G4_S5 \						
0	0.143198	0.035967	0.045707	0.045250	0.018917	0.153589
0.079970						
1	0.307269	0.012002	0.013871	0.020847	0.013454	0.058412
0.160077						
2	0.285336	0.075691	0.046578	0.067696	0.033349	0.112185
0.218809						
3	0.328946	0.121539	0.034826	0.094669	0.063285	0.081699
0.217128						
4	0.194058	0.030596	0.056352	0.051430	0.009631	0.063854
0.183514						

	PAICS_S5	PAK1_S2	PAK2_S2	PDAP1_S5	PDIA4_S5	PDLIM7_S5
PFKP_S5 \						
0	0.166344	0.072260	0.241153	0.168676	0.135598	0.207377
0.049983						
1	0.224232	0.117644	0.126378	0.210840	0.335843	0.075176
0.032116						
2	0.104921	0.116603	0.351109	0.331346	0.456125	0.106854
0.132821						
3	0.162434	0.082411	0.083170	0.215826	0.174828	0.117111
0.132418						
4	0.265156	0.127628	0.389034	0.126817	0.178679	0.053024
0.158858						

	PFN1_S5	PGM2_S5	PHF19_S5	PIK3CG_S3	PITPNC1_S5	PLAT_S2
PLAUR_S5 \						
0	0.255600	0.176835	0.191977	0.024237	0.076846	0.001419
0.039017						
1	0.191584	0.054117	0.053123	0.002969	0.131413	0.001808
0.030971						
2	0.170042	0.119041	0.348527	0.017677	0.040962	0.009950
0.122240						
3	0.303984	0.054715	0.385093	0.008843	0.051307	0.002175
0.120793						
4	0.110226	0.176835	0.115697	0.037040	0.103141	0.011542
0.076734						

	PLCG2_S3	PLEK_S3	PLG_S5	PLK4_S5	PLOD2_S5	PNN_S5
PN01_S5 \						
0	0.006507	0.023230	0.000005	0.272897	0.015910	0.147280
	0.161602					
1	0.004564	0.030490	0.000023	0.139299	0.019097	0.197264
	0.096395					
2	0.003650	0.070551	0.000005	0.198844	0.011098	0.099750
	0.184627					
3	0.001889	0.016255	0.000023	0.082470	0.003602	0.090495
	0.163267					
4	0.009946	0.057403	0.000005	0.208554	0.048598	0.111271
	0.322468					

	POLE2_S5	POLR3K_S5	PPIH_S5	PSMA7_S5	PSMC3_S5	PSMD12_S5
PSMD14_S5 \						
0	0.136847	0.089785	0.144393	0.125138	0.167270	0.139899
	0.268905					
1	0.096535	0.137554	0.232093	0.073134	0.156777	0.062331
	0.238535					
2	0.152285	0.077284	0.175007	0.135764	0.108752	0.113334
	0.216624					
3	0.073111	0.214049	0.200183	0.229206	0.104846	0.035758
	0.154648					
4	0.182561	0.186561	0.084329	0.185575	0.178016	0.194763
	0.482800					

	PSMD2_S5	PTPLB_S5	PTPRC_S3	PXN_S2	RAB3B_S5	RAC1_S2
RASSF2_S3 \						
0	0.157737	0.084729	0.024735	0.135201	0.001371	0.153141
	0.006755					
1	0.063664	0.011875	0.016258	0.030791	0.000577	0.167645
	0.012288					
2	0.125575	0.066794	0.041973	0.139315	0.005981	0.161858
	0.002001					
3	0.094678	0.020744	0.009985	0.139315	0.001737	0.194394
	0.001996					
4	0.223142	0.147774	0.049721	0.091718	0.003655	0.343638
	0.004547					

	RASSF4_S3	RBM14_S5	RFC3_S5	RGS8_S5	RHOA_S2	RHOB_S2
RHOC_S5 \						
0	0.025563	0.282242	0.095780	0.000000	0.254724	0.040654
	0.129998					
1	0.028804	0.387320	0.117169	0.000000	0.276097	0.170179
	0.119639					
2	0.053798	0.278413	0.106650	0.000000	0.417517	0.012437
	0.230437					
3	0.026893	0.587550	0.145781	0.014786	0.276152	0.050956
	0.191826					
4	0.006769	0.382809	0.145882	0.002612	0.318228	0.039787

0.131475

	RHOG_S2	RHOQ_S2	RMND5B_S5	RNASE6_S3	RND3_S2	RNF138_S5
RNF41_S5 \						
0	0.174512	0.107443	0.097324	0.034370	0.133599	0.087150
	0.089728					
1	0.117589	0.108968	0.181132	0.029793	0.149408	0.197635
	0.124705					
2	0.159244	0.082045	0.153058	0.042139	0.097058	0.104717
	0.115594					
3	0.243694	0.011838	0.278400	0.034181	0.030692	0.063504
	0.164837					
4	0.115530	0.086344	0.073182	0.027792	0.198128	0.152272
	0.178470					

	RPN1_S5	RPP40_S5	RSAD2_S1	RTP4_S1	RUNX3_S3	RUUBL1_S5
SAMD9_S1 \						
0	0.169853	0.203581	0.011339	0.022262	0.040771	0.176109
	0.065363					
1	0.339800	0.126313	0.006195	0.047336	0.011941	0.304404
	0.007888					
2	0.218984	0.175488	0.021158	0.032420	0.012582	0.228971
	0.010268					
3	0.201962	0.157057	0.043230	0.030058	0.069775	0.205084
	0.012855					
4	0.482965	0.242419	0.020195	0.063709	0.064088	0.419433
	0.048694					

	SAMHD1_S3	SAMSN1_S3	SAR1A_S5	SAR1B_S5	SDC1_S5	SELL_S3
SELPLG_S3 \						
0	0.052966	0.047820	0.206790	0.100884	0.098025	0.004675
	0.037912					
1	0.130442	0.053131	0.188092	0.083722	0.013992	0.004051
	0.029117					
2	0.081837	0.083075	0.158464	0.069516	0.060385	0.005374
	0.081667					
3	0.067619	0.031994	0.049852	0.055348	0.040424	0.001718
	0.058516					
4	0.115436	0.079174	0.354664	0.111418	0.131077	0.004864
	0.052901					

	SEMA3F_S2	SERPINE1_S2	SH2B3_S3	SH3BP5L_S5	SKA1_S5	SKA2_S5	\
0	0.107456	0.047018	0.071774	0.397680	0.023298	0.150416	
1	0.075353	0.000925	0.022738	0.161268	0.027082	0.163312	
2	0.044402	0.011848	0.141412	0.147696	0.032137	0.151942	
3	0.041377	0.001889	0.051041	0.289520	0.039170	0.019166	
4	0.132073	0.053159	0.074056	0.161067	0.065265	0.160582	

	SLC16A1_S5	SLC1A3_S3	SLC25A40_S5	SLC25A5_S5	SLC7A7_S3	SMC2_S5	\
--	------------	-----------	-------------	------------	-----------	---------	---

0	0.187154	0.004853	0.113213	0.066678	0.008365	0.111563
1	0.101799	0.001806	0.050642	0.025590	0.008739	0.059632
2	0.170208	0.001690	0.296410	0.051780	0.030765	0.122536
3	0.068127	0.000242	0.103375	0.096260	0.017580	0.029403
4	0.342128	0.018085	0.194974	0.062755	0.017081	0.181299

	SMO_S2	SMS_S5	SMURF2_S5	SNRPA1_S5	SNRPA_S5	SNRPB_S5
SNRPC_S5 \						
0	0.061107	0.010822	0.117046	0.098420	0.129965	0.110974
	0.189107					
1	0.223859	0.042159	0.080097	0.150227	0.357502	0.147041
	0.194742					
2	0.120447	0.044985	0.240120	0.155771	0.129030	0.153509
	0.222232					
3	0.041830	0.026457	0.055882	0.132591	0.353250	0.135313
	0.214041					
4	0.065579	0.028911	0.139213	0.228911	0.135565	0.076978
	0.224079					

	SNRPD1_S5	SNRPE_S5	SNX17_S5	SP140_S3	SPAG17_S5	SPARC_S2
SRM_S5 \						
0	0.082965	0.033202	0.217200	0.025957	0.016643	0.045811
	0.102953					
1	0.211044	0.106333	0.230067	0.012544	0.066766	0.010303
	0.081692					
2	0.130903	0.081369	0.169501	0.047222	0.000769	0.012549
	0.132022					
3	0.078309	0.063577	0.174395	0.009958	0.000777	0.012536
	0.129299					
4	0.174427	0.061940	0.178775	0.015842	0.027190	0.056416
	0.118792					

	STAT1_S1	STK17A_S5	STRA13_S5	SYK_S3	TAGLN_S5	TAP1_S1
TBXAS1_S3 \						
0	0.044481	0.229143	0.076512	0.059110	0.038161	0.078096
	0.035709					
1	0.055906	0.059119	0.099121	0.168666	0.003984	0.086456
	0.087330					
2	0.118732	0.229143	0.054861	0.053159	0.007983	0.153990
	0.039003					
3	0.043672	0.024340	0.104535	0.138534	0.025773	0.173724
	0.236709					
4	0.099685	0.158866	0.026717	0.132907	0.027327	0.064915
	0.038927					

	TCEB1_S5	TCF7L2_S2	THBS1_S2	THBS2_S2	TIMP1_S2	TLR2_S3
TMEM130_S5 \						
0	0.130174	0.099239	0.178477	0.065712	0.028905	0.148018
	0.000175					
1	0.103268	0.162975	0.012453	0.001430	0.027609	0.075820
	0.000108					
2	0.143786	0.199046	0.012716	0.028705	0.038713	0.048070
	0.000505					
3	0.052949	0.228019	0.031254	0.018995	0.047791	0.023434
	0.000256					
4	0.159103	0.164192	0.093766	0.106684	0.041740	0.263277
	0.000990					

	TNC_S2	TNFAIP3_S3	TNFRSF12A_S5	TNFRSF1A_S2	TNFRSF1B_S3
TNFSF13B_S3 \					
0	0.115225	0.080005	0.168056	0.252215	0.044209
	0.007273				
1	0.015413	0.100122	0.044841	0.219536	0.089376
	0.005195				
2	0.115225	0.054746	0.121907	0.350215	0.129051
	0.027343				
3	0.017496	0.058073	0.108840	0.273136	0.162439
	0.007791				
4	0.115225	0.114188	0.049923	0.433645	0.040264
	0.019876				

	TOMM40_S5	TPI1_S5	TPM1_S5	TPM2_S5	TPM3_S5	TPRKB_S5
TRA2B_S5 \						
0	0.030447	0.055560	0.085022	0.054904	0.248836	0.224479
	0.370910					
1	0.039365	0.121625	0.019840	0.010281	0.191961	0.427167
	0.307541					
2	0.044709	0.170366	0.031936	0.029378	0.219096	0.172608
	0.150605					
3	0.085939	0.142167	0.048872	0.013824	0.219590	0.089482
	0.170531					
4	0.041386	0.247876	0.027684	0.008869	0.276453	0.239914
	0.384092					

	TUBA4A_S5	TUBG1_S5	UAP1_S5	UBE2J1_S5	UMPS_S5	UQCR10_S5
USPL1_S5 \						
0	0.097268	0.249644	0.107880	0.182237	0.252513	0.037382
	0.201070					
1	0.002220	0.227805	0.080434	0.249096	0.145911	0.072831
	0.190156					
2	0.040278	0.238908	0.078863	0.267850	0.215546	0.063526
	0.178736					
3	0.097268	0.293324	0.038459	0.137453	0.194995	0.042015
	0.102779					
4	0.063399	0.435219	0.129999	0.277074	0.312816	0.022331

0.054357

	VCAN_S2	VDAC1_S5	VSIG4_S3	VTA1_S5	WDHD1_S5	WDR54_S5
WDR77_S5 \						
0	0.072346	0.195607	0.003054	0.269151	0.079282	0.072920
0.102072						
1	0.108592	0.229437	0.013031	0.175206	0.034451	0.188849
0.240956						
2	0.040453	0.227530	0.023227	0.184954	0.094871	0.037184
0.124078						
3	0.018265	0.353334	0.005992	0.102899	0.018202	0.034451
0.221328						
4	0.206705	0.262335	0.017925	0.348754	0.114240	0.052860
0.092856						

	WIPF1_S3	WNT2B_S2	WNT8B_S2	WSB2_S5	ZWILCH_S5	ZYX_S2	Immune Subtype
0	0.089859	0.010692	0.008595	0.124660	0.154684	0.191966	
0							
1	0.067037	0.006442	0.002612	0.089291	0.144194	0.126821	
0							
2	0.091530	0.002030	0.003011	0.142595	0.099884	0.293115	
0							
3	0.026653	0.012035	0.003367	0.134369	0.079119	0.256544	
0							
4	0.126317	0.041299	0.000520	0.167126	0.160216	0.132789	
0							

```
corr_matrix["Immune Subtype"].head()
```

```
ACTL6A_S5    -0.505404
ADAM9_S2     -0.072987
ADAMTS1_S5   -0.066917
ADCY7_S3      0.031620
AIMP2_S5     -0.157777
Name: Immune Subtype, dtype: float64
```

Comparision of different feature reduction methods

PCA VS NMF VS T-SNE

```
# Data preparation
X_full = df_final_normalized.drop(columns=['Immune Subtype'])
y = df_final_normalized['Immune Subtype']

# Scaling
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X_full)
minmax_scaler = MinMaxScaler()
```



```

X_nonneg = minmax_scaler.fit_transform(X_full)

# Dimensionality reductions (create all versions but don't visualize 100D)
## PCA
pca = PCA(n_components=0.90)
X_pca = pca.fit_transform(X_scaled) # For later use
X_pca_2d = PCA(n_components=2).fit_transform(X_scaled)
X_pca_3d = PCA(n_components=3).fit_transform(X_scaled)
X_pca_100d = PCA(n_components=100).fit_transform(X_scaled) # Created but not visualized

## NMF
nmf = NMF(n_components=pca.n_components_, init='nndsvda',
random_state=42)
X_nmf = nmf.fit_transform(X_nonneg) # For later use
X_nmf_2d = NMF(n_components=2, init='nndsvda',
random_state=42).fit_transform(X_nonneg)
X_nmf_3d = NMF(n_components=3, init='nndsvda',
random_state=42).fit_transform(X_nonneg)
X_nmf_100d = NMF(n_components=100, init='nndsvda',
random_state=42).fit_transform(X_nonneg) # Created but not visualized

## t-SNE
X_tsne_2d = TSNE(n_components=2, perplexity=30,
random_state=42).fit_transform(X_scaled)
X_tsne_3d = TSNE(n_components=3, perplexity=30, random_state=42,
method='exact').fit_transform(X_scaled)
# Skip 100D t-SNE as it's not recommended

# Visualization function for 2D plots
def plot_2d(X, y, title, ax):
    sns.scatterplot(x=X[:, 0], y=X[:, 1], hue=y,
                    palette='viridis', alpha=0.8, s=60, ax=ax)
    ax.set_title(title)
    ax.set_xlabel(f'{title.split()[0]}1')
    ax.set_ylabel(f'{title.split()[0]}2')

# Visualization function for 3D plots
def plot_3d(X, y, title, ax):
    scatter = ax.scatter(X[:, 0], X[:, 1], X[:, 2],
                        c=y.astype('category').cat.codes,
                        cmap='viridis', alpha=0.7, s=40)
    ax.set_title(title)
    ax.set_xlabel(f'{title.split()[0]}1')
    ax.set_ylabel(f'{title.split()[0]}2')
    ax.set_zlabel(f'{title.split()[0]}3')
    return scatter

# Create 2D comparison plot

```

```

fig_2d, axes_2d = plt.subplots(1, 3, figsize=(24, 8))
plot_2d(X_pca_2d, y, 'PCA 2D', axes_2d[0])
plot_2d(X_nmf_2d, y, 'NMF 2D', axes_2d[1])
plot_2d(X_tsne_2d, y, 't-SNE 2D', axes_2d[2])
axes_2d[0].legend(title='Immune Subtype', bbox_to_anchor=(1, 1))
axes_2d[1].legend().remove()
axes_2d[2].legend().remove()
plt.suptitle('2D Dimensionality Reduction Comparison', y=1.02)
plt.tight_layout()
plt.show()

```

Create 3D comparison plot

```

fig_3d = plt.figure(figsize=(24, 10))
ax1 = fig_3d.add_subplot(1, 3, 1, projection='3d')
plot_3d(X_pca_3d, y, 'PCA 3D', ax1)

```

```

ax2 = fig_3d.add_subplot(1, 3, 2, projection='3d')
plot_3d(X_nmf_3d, y, 'NMF 3D', ax2)

```

```

ax3 = fig_3d.add_subplot(1, 3, 3, projection='3d')
plot_3d(X_tsne_3d, y, 't-SNE 3D', ax3)

```

Add single legend

```

legend_labels = y.unique()
for i, label in enumerate(legend_labels):
    ax1.scatter([], [], [], label=label,
                color=sns.color_palette('viridis')[i])
fig_3d.legend(title='Immune Subtype', bbox_to_anchor=(1.1, 1),
              loc='upper left')

```

```

plt.suptitle('3D Dimensionality Reduction Comparison', y=1.02)
plt.tight_layout()
plt.show()

```

All transformed datasets are available for later use:

X_pca, X_pca_100d, X_nmf, X_nmf_100d (but not visualized)

```

/usr/local/lib/python3.10/dist-packages/sklearn/decomposition/_nmf.py:1665: ConvergenceWarning: Maximum number of iterations 200 reached. Increase it to improve convergence.

```

```

warnings.warn(

```

```

/usr/local/lib/python3.10/dist-packages/sklearn/decomposition/_nmf.py:1665: ConvergenceWarning: Maximum number of iterations 200 reached. Increase it to improve convergence.

```

```

warnings.warn(

```

```

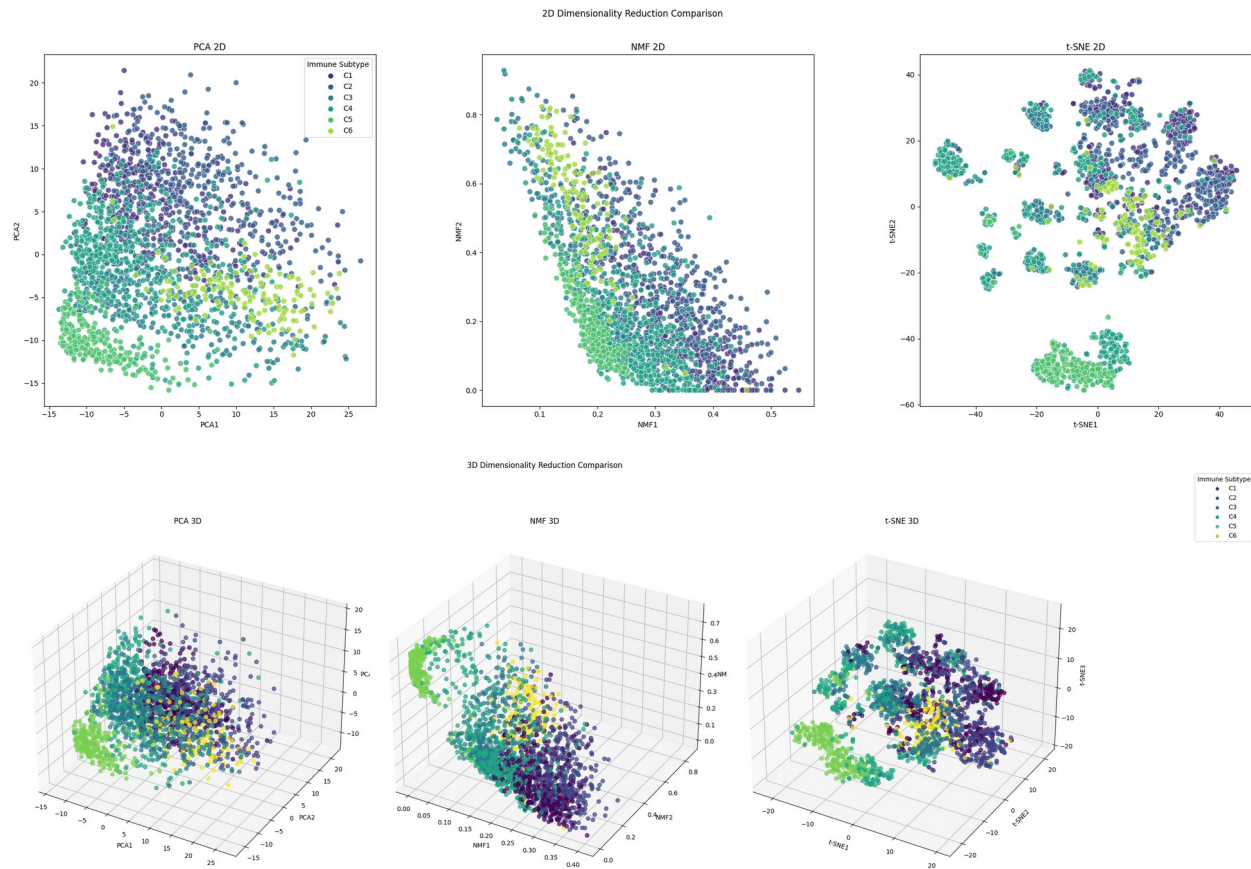
/usr/local/lib/python3.10/dist-packages/sklearn/decomposition/_nmf.py:1665: ConvergenceWarning: Maximum number of iterations 200 reached. Increase it to improve convergence.

```

```

warnings.warn(

```



Best Feature Reduction:

t-SNE → Best separation, distinct clusters.

PCA → Retains global structure but overlaps.

NMF → More structured than PCA but less separation than t-SNE.

For interpretability → PCA/NMF For classification → t-SNE

EDA

```
pca_df = pd.DataFrame(X_pca_3d)

y = le.fit_transform(df_final_normalized['Immune Subtype']) # Encode
categorical labels

# Convert y to DataFrame
y_df = pd.DataFrame(y, columns=['Immune Subtype'])

# Drop target from X

# Reset indices and concatenate
#X = X.reset_index(drop=True)
#y_df = y_df.reset_index(drop=True)
```

```

df = pd.concat([pca_df, y_df], axis=1)
import sweetviz as sv

# Generate a report
report = sv.analyze(pd.DataFrame(X_pca))

# Display inline in Kaggle
report.show_html("report.html") # Save if needed
report.show_notebook() # Inline display

{"model_id":"13b099be6ed44d488c8a96267550b88f","version_major":2,"version_minor":0}

/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py:219: PerformanceWarning: DataFrame is highly
fragmented. This is usually the result of calling `frame.insert` many
times, which has poor performance. Consider joining all columns at
once using pd.concat(axis=1) instead. To get a de-fragmented frame,
use `newframe = frame.copy()`
  new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
  new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
  new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
  new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
  new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
  new_dataframe[feature] = pd.Series(dtype=float)

```

```
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
```

```
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
```

```
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
```

```
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
```



```

usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)
/usr/local/lib/python3.10/dist-packages/sweetviz/graph_associations.py
:219: PerformanceWarning: DataFrame is highly fragmented. This is
usually the result of calling `frame.insert` many times, which has
poor performance. Consider joining all columns at once using
pd.concat(axis=1) instead. To get a de-fragmented frame, use `newframe
= frame.copy()`
    new_dataframe[feature] = pd.Series(dtype=float)

```

Report report.html was generated! NOTEBOOK/COLAB USERS: the web browser MAY not pop up, regardless, the report IS saved in your notebook/colab files.

<IPython.core.display.HTML object>

```

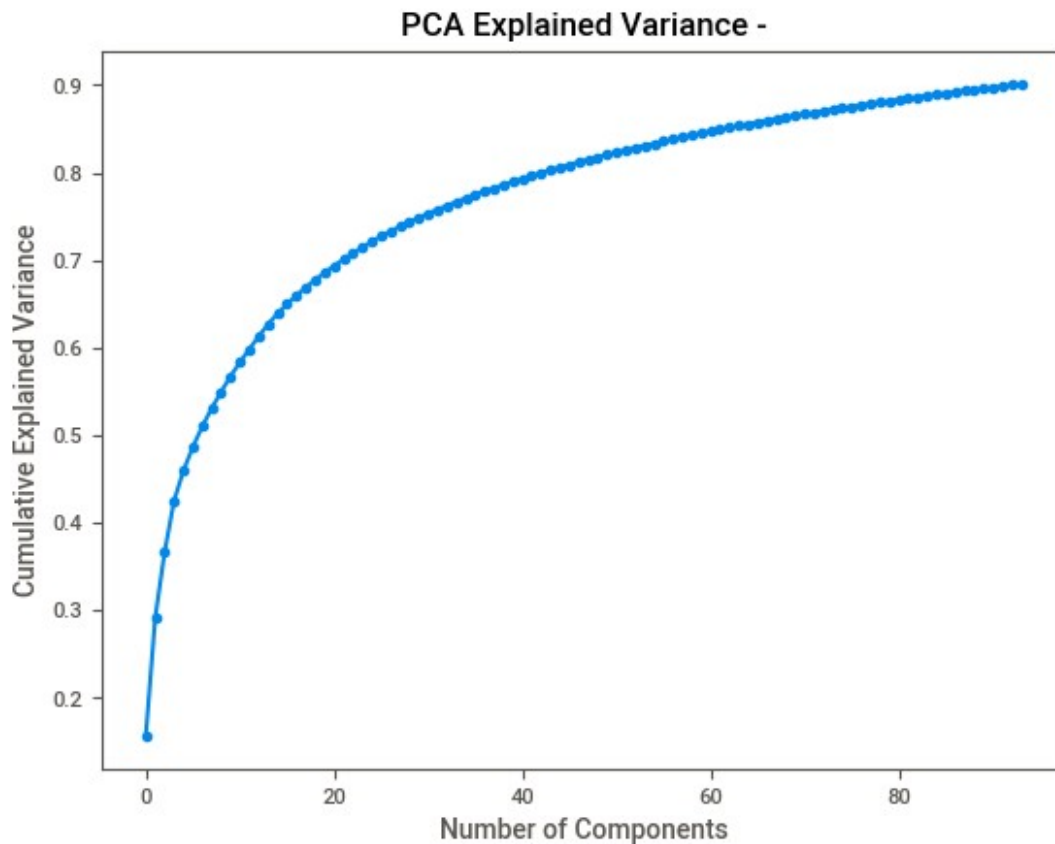
# X_full: Full dataset without dimensionality reduction
X_full = df_final_normalized.drop(columns=['Immune Subtype']) #
Assuming 'Immune Subtype' is the target variable
# X_pca: PCA-reduced dataset
from sklearn.decomposition import PCA

pca = PCA(n_components=0.90)
X_pca = pca.fit_transform(X_full)

explained_var_ratio = np.cumsum(pca.explained_variance_ratio_)

```

```
plt.plot(explained_var_ratio, marker='o')
plt.xlabel("Number of Components")
plt.ylabel("Cumulative Explained Variance")
plt.title(f"PCA Explained Variance -")
plt.show()
```



Section 2 - Classification Model Development

```
# Label Encoding
y = LabelEncoder().fit_transform(df_final_normalized['Immune
Subtype'])

# PCA Transformation (90% Variance Retention)
#pca = PCA(n_components=0.90)
#X_pca = pca.fit_transform(X_full)
#X_FE=df_features.drop(columns=['Immune Subtype'])

# PCA and Kernel PCA
pca = PCA(n_components=0.90)
kPCA = KernelPCA(n_components=50, kernel='rbf')
```

```

X_pca = pca.fit_transform(X_full)
X_kpca = kpca.fit_transform(X_full)
datasets = {
    "NMF": X_nmf,
    "X_pca": X_pca,
    "X_kpca": X_kpca,
    # "X_FE": X_FE,
    "X_nmf_100d": X_nmf_100d,
    # "tsne_100d": tsne_100d,
    "X_full": X_full,
    "X_tsne_3d": X_tsne_3d,
    # "Top_20_Correlated_Features": X_selected
}

# Scale Data
scaler = StandardScaler()

def train_and_evaluate_model(model, model_name, X_train, X_test,
y_train, y_test):
    model.fit(X_train, y_train)
    y_pred = model.predict(X_test)
    accuracy = accuracy_score(y_test, y_pred)

    print(f'\n{model_name} Accuracy: {accuracy:.4f}')
    print(classification_report(y_test, y_pred,
target_names=LabelEncoder().fit(df_final_normalized['Immune
Subtype']).classes_))

    # Confusion Matrix
    cm = confusion_matrix(y_test, y_pred)
    plt.figure(figsize=(6, 5))
    sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
    plt.xlabel('Predicted')
    plt.ylabel('Actual')
    plt.title(f'Confusion Matrix - {model_name}')
    plt.show()

    return accuracy

def create_nn(input_dim):
    model = Sequential([
        Dense(64, activation='relu', input_shape=(input_dim,)),
        Dropout(0.3),
        Dense(32, activation='relu'),
        Dropout(0.3),
        Dense(len(set(y)), activation='softmax')
    ])
    model.compile(optimizer=Adam(learning_rate=0.001),
loss='sparse_categorical_crossentropy', metrics=['accuracy'])
    return model

```

```

results = {}

for name, X in datasets.items():
    print(f"\n===== Training Models on {name} Dataset =====")

    X_train, X_test, y_train, y_test = train_test_split(X, y,
stratify=y, test_size=0.2, random_state=42)
    print(f"Training set: {X_train.shape}, Test set: {X_test.shape}")
    X_train = scaler.fit_transform(X_train)
    X_test = scaler.transform(X_test)

    # Neural Network Training
    nn_model = create_nn(X_train.shape[1])
    history = nn_model.fit(X_train, y_train, validation_data=(X_test,
y_test), epochs=50, batch_size=16, verbose=0)

    plt.plot(history.history['accuracy'], label='Train')
    plt.plot(history.history['val_accuracy'], label='Validation')
    plt.title(f'Neural Network Accuracy - {name}')
    plt.legend()
    plt.show()

    y_pred_nn = nn_model.predict(X_test).argmax(axis=1)
    print(f"\nNeural Network - {name} Classification Report:\n",
classification_report(y_test, y_pred_nn))

    # Random Forest Classifier
    rf_clf = RandomForestClassifier(n_estimators=200, max_depth=10,
random_state=42)
    results[f"RandomForest_{name}"] = train_and_evaluate_model(rf_clf,
"Random Forest", X_train, X_test, y_train, y_test)

    # XGBoost with Hyperparameter Tuning
    xgb_params = {'max_depth': [1,2], 'learning_rate': [0.01, 0.1],
'n_estimators': [100, 200]}
    xgb_clf = GridSearchCV(XGBClassifier(eval_metric='logloss',
random_state=42), xgb_params, cv=3)
    xgb_clf.fit(X_train, y_train)
    print(f"Best XGBoost Parameters for {name}:
{xgb_clf.best_params_}")
    results[f"XGBoost_{name}"] =
train_and_evaluate_model(xgb_clf.best_estimator_, "XGBoost", X_train,
X_test, y_train, y_test)

    # SVM with Hyperparameter Tuning
    svm_params = {
        'C': [0.1, 1, 10, 50, 100],
        'gamma': ['scale', 'auto', 0.001, 0.01, 0.1],
        'kernel': ['linear', 'rbf', 'poly'],

```

```

        'class_weight': ['balanced', None]
    }
    svm_clf = GridSearchCV(SVC(), svm_params, cv=3, verbose=1,
n_jobs=-1)
    svm_clf.fit(X_train, y_train)
    print(f"Best SVM Parameters for {name}: {svm_clf.best_params_}")
    results[f"SVM_{name}"] =
train_and_evaluate_model(svm_clf.best_estimator_, "SVM", X_train,
X_test, y_train, y_test)

    # Bagging SVM
    bagging_svm =
BaggingClassifier(base_estimator=SVC(**svm_clf.best_params_),
n_estimators=10, random_state=42)

    results[f"BaggingSVM_{name}"] =
train_and_evaluate_model(bagging_svm, "Bagging SVM", X_train, X_test,
y_train, y_test)

    # Logistic Regression
    log_reg = LogisticRegression(max_iter=500)
    results[f"LogisticRegression_{name}"] =
train_and_evaluate_model(log_reg, "Logistic Regression", X_train,
X_test, y_train, y_test)

```

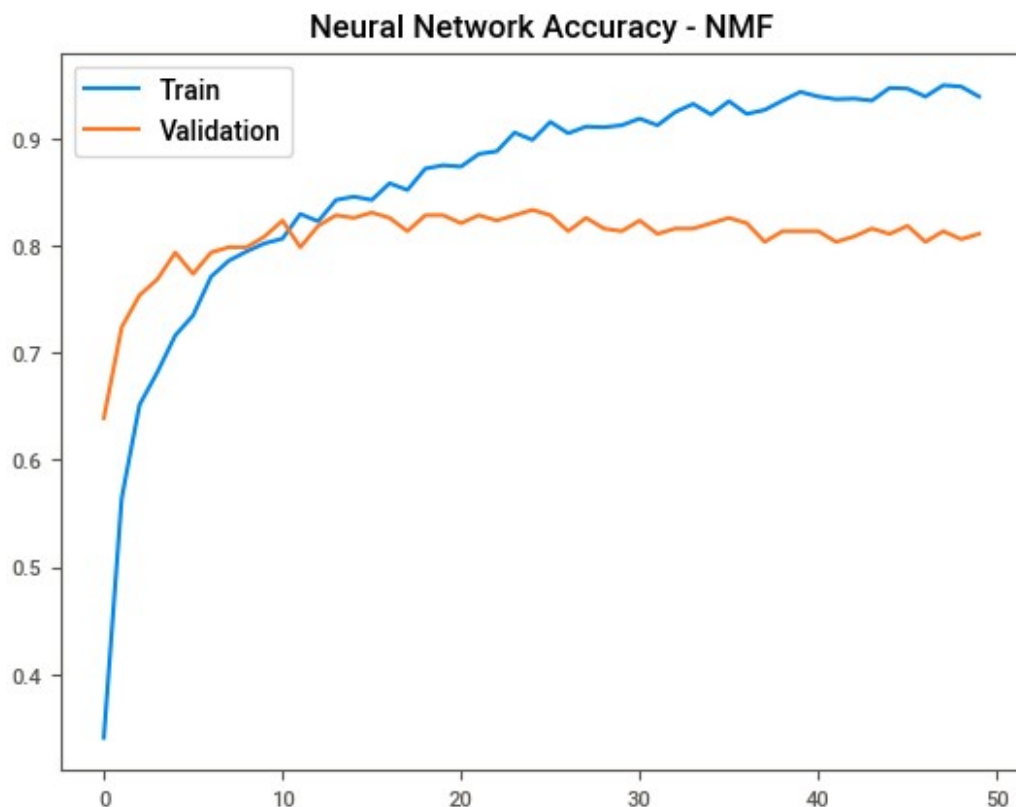
==== Training Models on NMF Dataset ====

Training set: (1607, 138), Test set: (402, 138)

```

/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/
dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim`
argument to a layer. When using Sequential models, prefer using an
`Input(shape)` object as the first layer in the model instead.
    super().__init__(activity_regularizer=activity_regularizer,
**kwargs)

```



13/13 ————— 0s 9ms/step

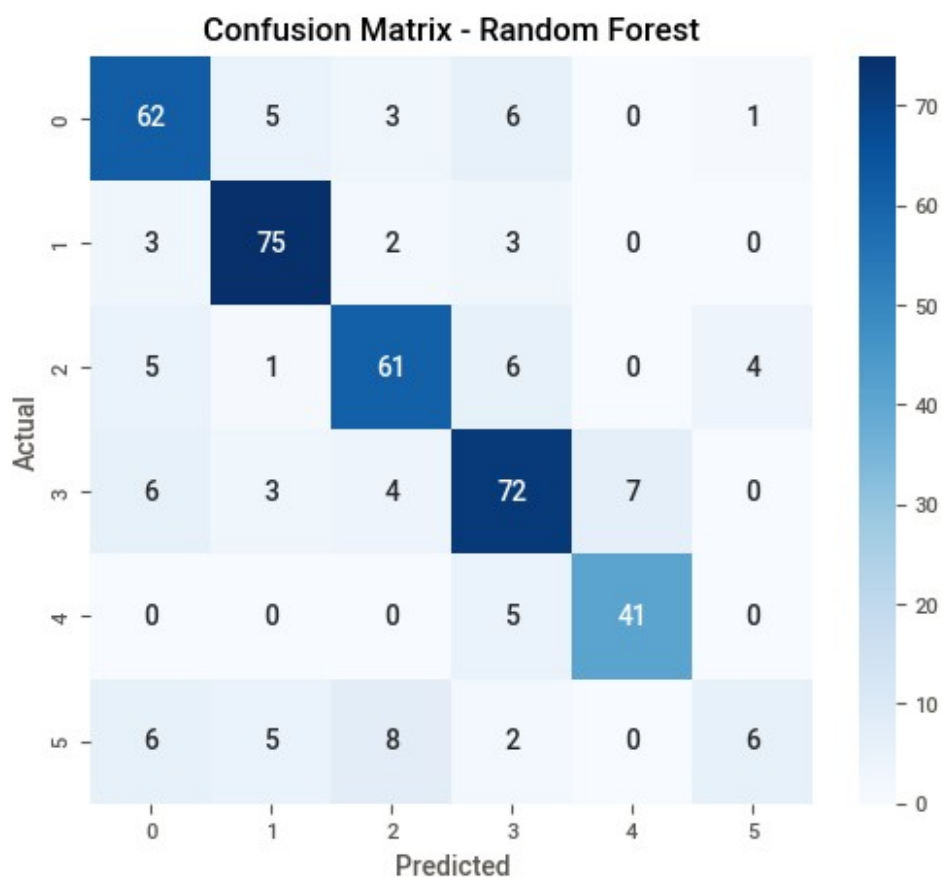
Neural Network - NMF Classification Report:

	precision	recall	f1-score	support
0	0.76	0.87	0.81	77
1	0.89	0.92	0.90	83
2	0.81	0.75	0.78	77
3	0.81	0.75	0.78	92
4	0.83	0.93	0.88	46
5	0.65	0.48	0.55	27
accuracy			0.81	402
macro avg	0.79	0.78	0.78	402
weighted avg	0.81	0.81	0.81	402

Random Forest Accuracy: 0.7886

	precision	recall	f1-score	support
C1	0.76	0.81	0.78	77
C2	0.84	0.90	0.87	83
C3	0.78	0.79	0.79	77
C4	0.77	0.78	0.77	92
C5	0.85	0.89	0.87	46

C6	0.55	0.22	0.32	27
accuracy			0.79	402
macro avg	0.76	0.73	0.73	402
weighted avg	0.78	0.79	0.78	402

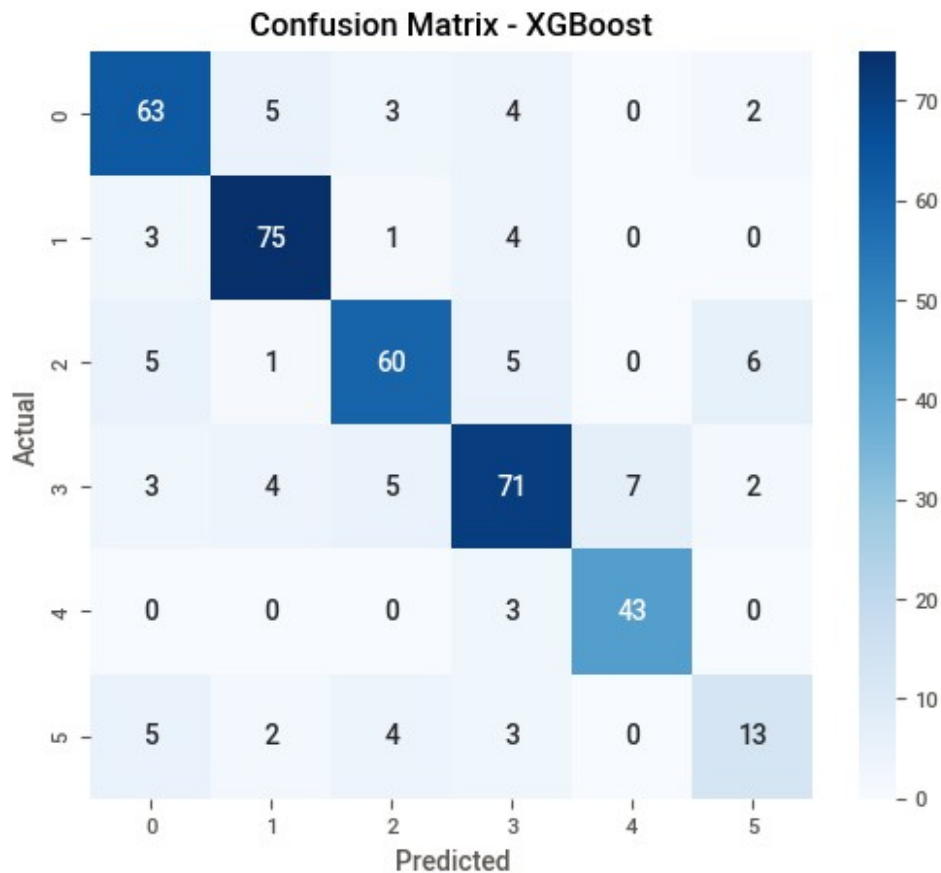


Best XGBoost Parameters for NMF: {'learning_rate': 0.1, 'max_depth': 2, 'n_estimators': 200}

XGBoost Accuracy: 0.8085

	precision	recall	f1-score	support
C1	0.80	0.82	0.81	77
C2	0.86	0.90	0.88	83
C3	0.82	0.78	0.80	77
C4	0.79	0.77	0.78	92
C5	0.86	0.93	0.90	46
C6	0.57	0.48	0.52	27
accuracy			0.81	402
macro avg	0.78	0.78	0.78	402

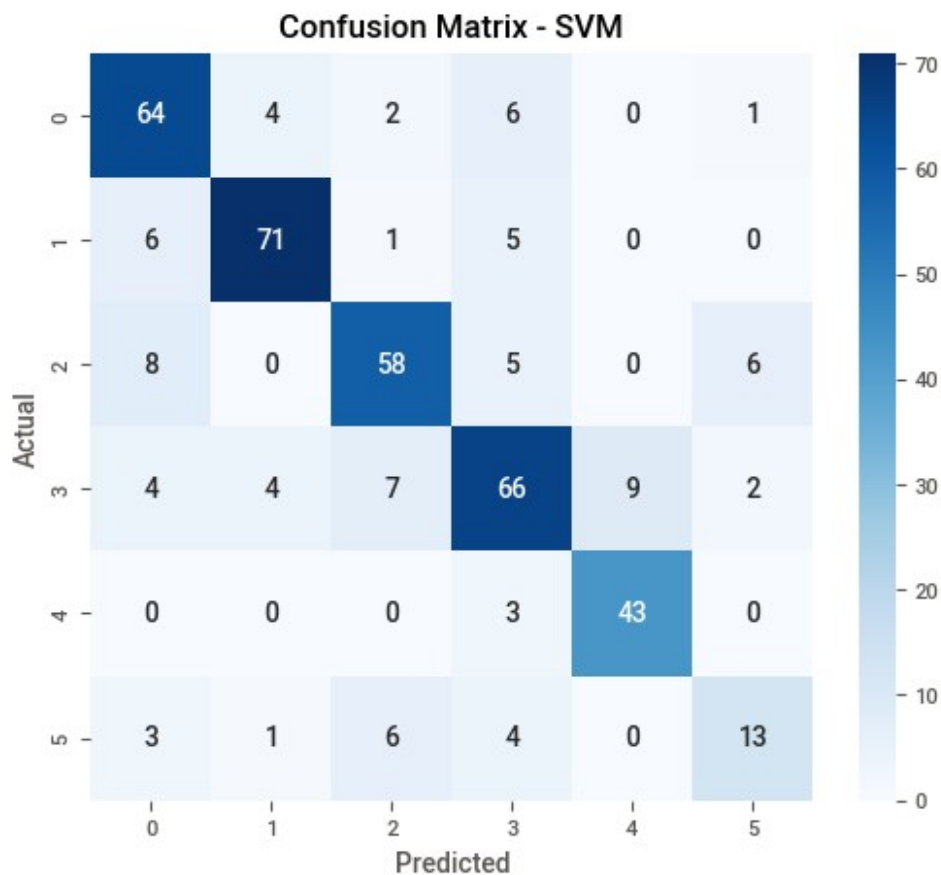
weighted avg	0.81	0.81	0.81	402
--------------	------	------	------	-----



Fitting 3 folds for each of 150 candidates, totalling 450 fits
 Best SVM Parameters for NMF: {'C': 10, 'class_weight': 'balanced', 'gamma': 'scale', 'kernel': 'rbf'}

SVM Accuracy: 0.7836

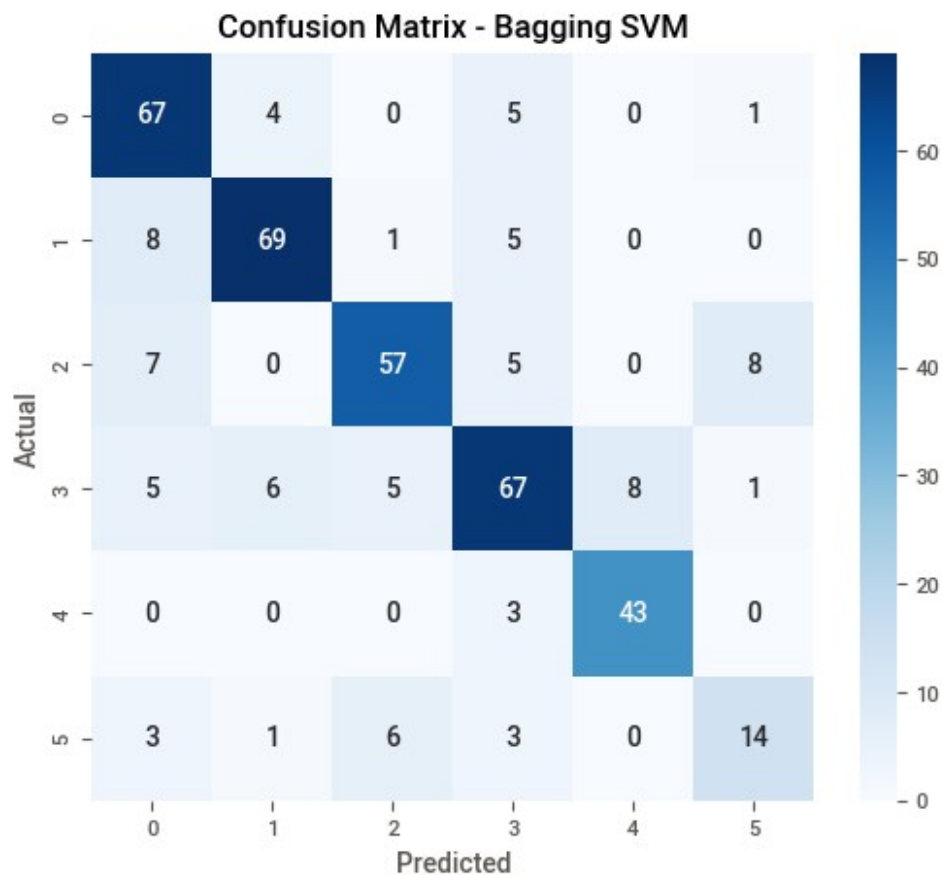
	precision	recall	f1-score	support
C1	0.75	0.83	0.79	77
C2	0.89	0.86	0.87	83
C3	0.78	0.75	0.77	77
C4	0.74	0.72	0.73	92
C5	0.83	0.93	0.88	46
C6	0.59	0.48	0.53	27
accuracy			0.78	402
macro avg	0.76	0.76	0.76	402
weighted avg	0.78	0.78	0.78	402



```
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
warnings.warn(
```

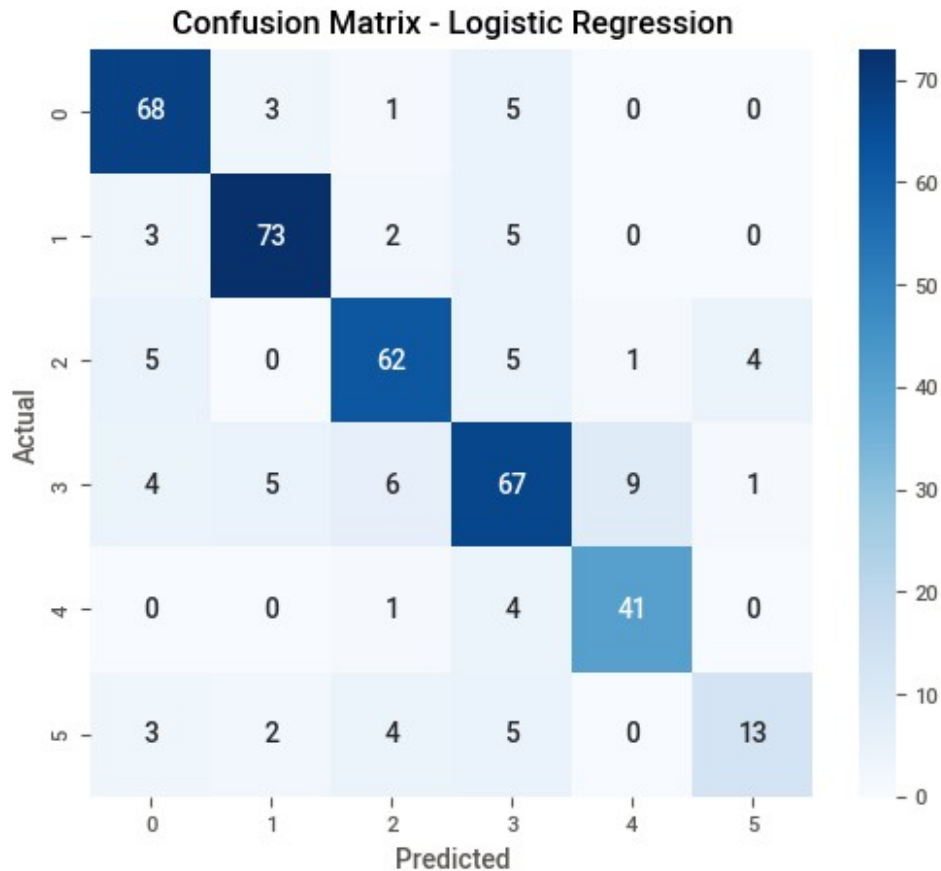
Bagging SVM Accuracy: 0.7886

	precision	recall	f1-score	support
C1	0.74	0.87	0.80	77
C2	0.86	0.83	0.85	83
C3	0.83	0.74	0.78	77
C4	0.76	0.73	0.74	92
C5	0.84	0.93	0.89	46
C6	0.58	0.52	0.55	27
accuracy			0.79	402
macro avg	0.77	0.77	0.77	402
weighted avg	0.79	0.79	0.79	402



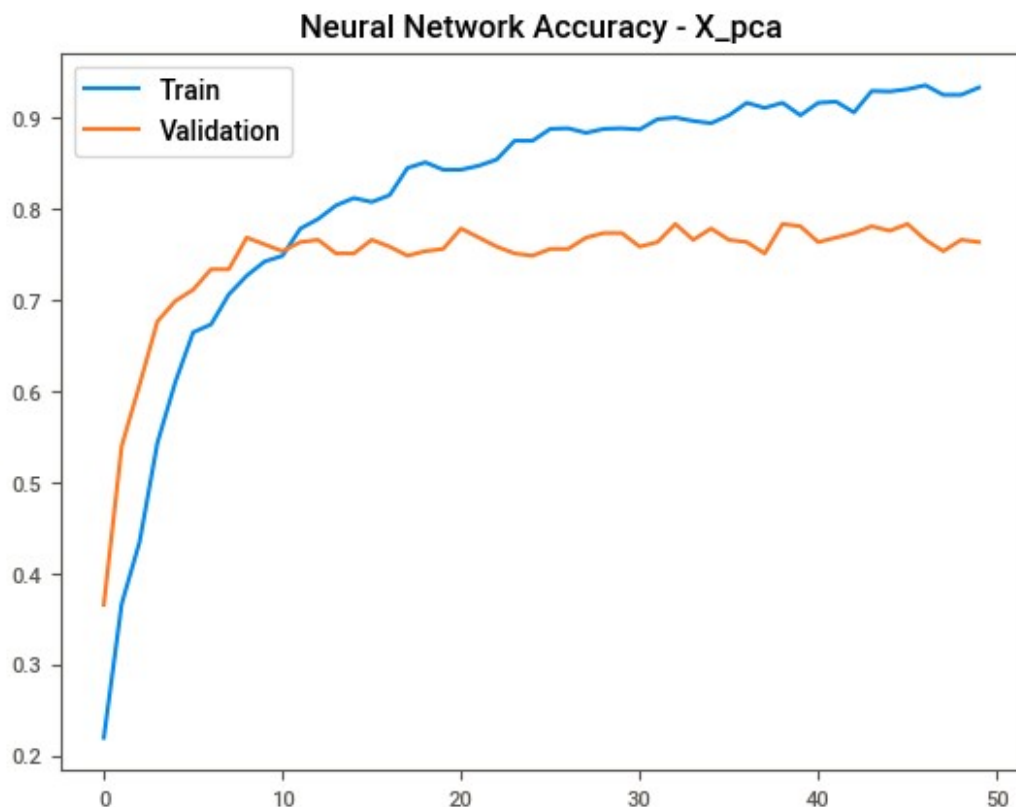
Logistic Regression Accuracy: 0.8060

	precision	recall	f1-score	support
C1	0.82	0.88	0.85	77
C2	0.88	0.88	0.88	83
C3	0.82	0.81	0.81	77
C4	0.74	0.73	0.73	92
C5	0.80	0.89	0.85	46
C6	0.72	0.48	0.58	27
accuracy			0.81	402
macro avg	0.80	0.78	0.78	402
weighted avg	0.80	0.81	0.80	402



```
===== Training Models on X_pca Dataset =====  
Training set: (1607, 94), Test set: (402, 94)
```

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/  
dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim`  
argument to a layer. When using Sequential models, prefer using an  
`Input(shape)` object as the first layer in the model instead.  
super().__init__(activity_regularizer=activity_regularizer,  
**kwargs)
```



13/13 ————— 0s 8ms/step

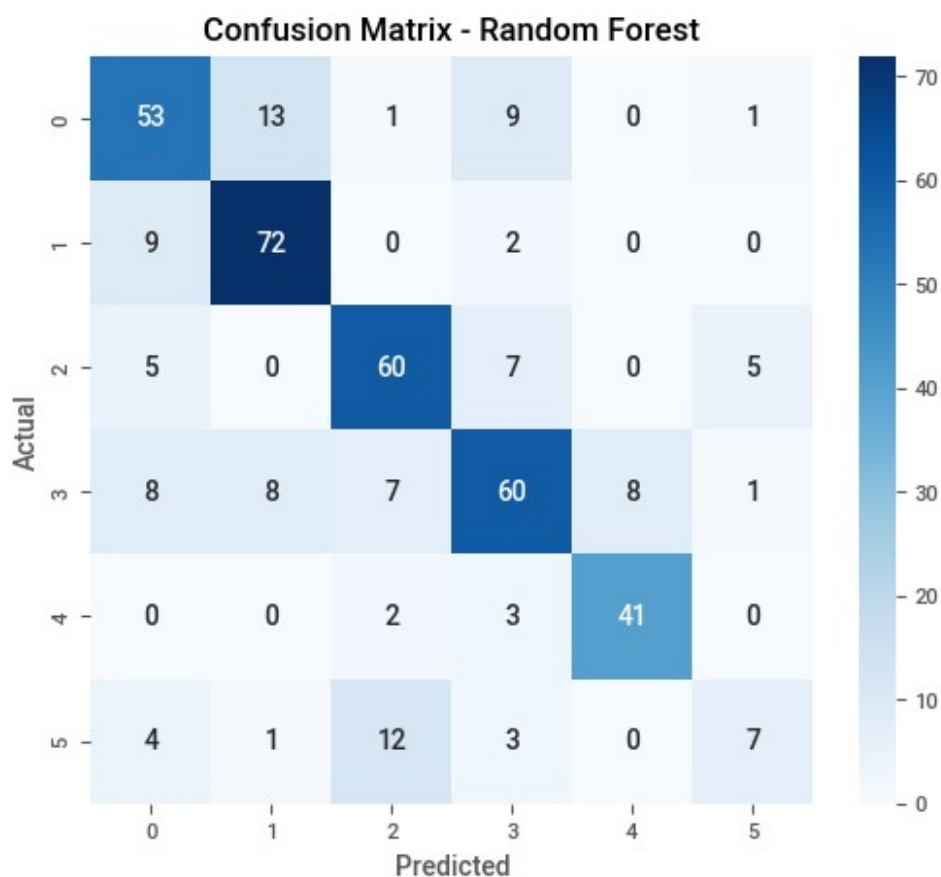
Neural Network - X_pca Classification Report:

	precision	recall	f1-score	support
0	0.74	0.69	0.71	77
1	0.81	0.87	0.84	83
2	0.79	0.70	0.74	77
3	0.69	0.76	0.73	92
4	0.88	0.91	0.89	46
5	0.67	0.59	0.63	27
accuracy			0.76	402
macro avg	0.76	0.75	0.76	402
weighted avg	0.76	0.76	0.76	402

Random Forest Accuracy: 0.7289

	precision	recall	f1-score	support
C1	0.67	0.69	0.68	77
C2	0.77	0.87	0.81	83
C3	0.73	0.78	0.75	77
C4	0.71	0.65	0.68	92
C5	0.84	0.89	0.86	46

C6	0.50	0.26	0.34	27
accuracy			0.73	402
macro avg	0.70	0.69	0.69	402
weighted avg	0.72	0.73	0.72	402

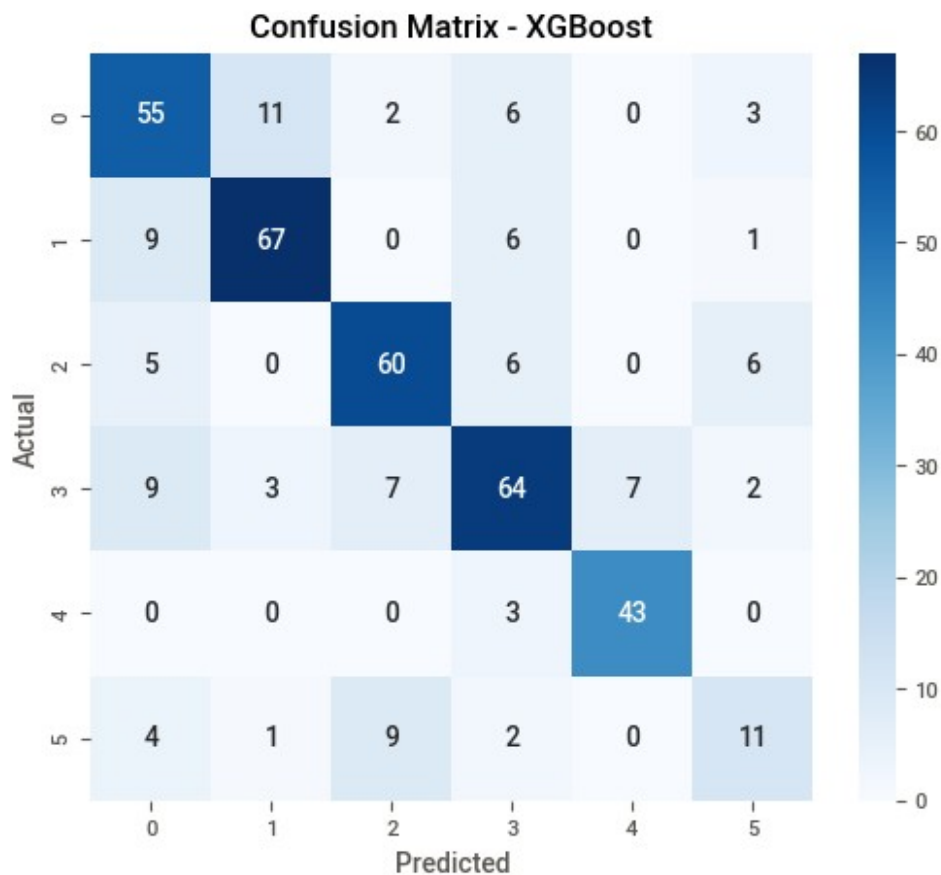


Best XGBoost Parameters for X_pca: {'learning_rate': 0.1, 'max_depth': 2, 'n_estimators': 200}

XGBoost Accuracy: 0.7463

	precision	recall	f1-score	support
C1	0.67	0.71	0.69	77
C2	0.82	0.81	0.81	83
C3	0.77	0.78	0.77	77
C4	0.74	0.70	0.72	92
C5	0.86	0.93	0.90	46
C6	0.48	0.41	0.44	27
accuracy			0.75	402
macro avg	0.72	0.72	0.72	402

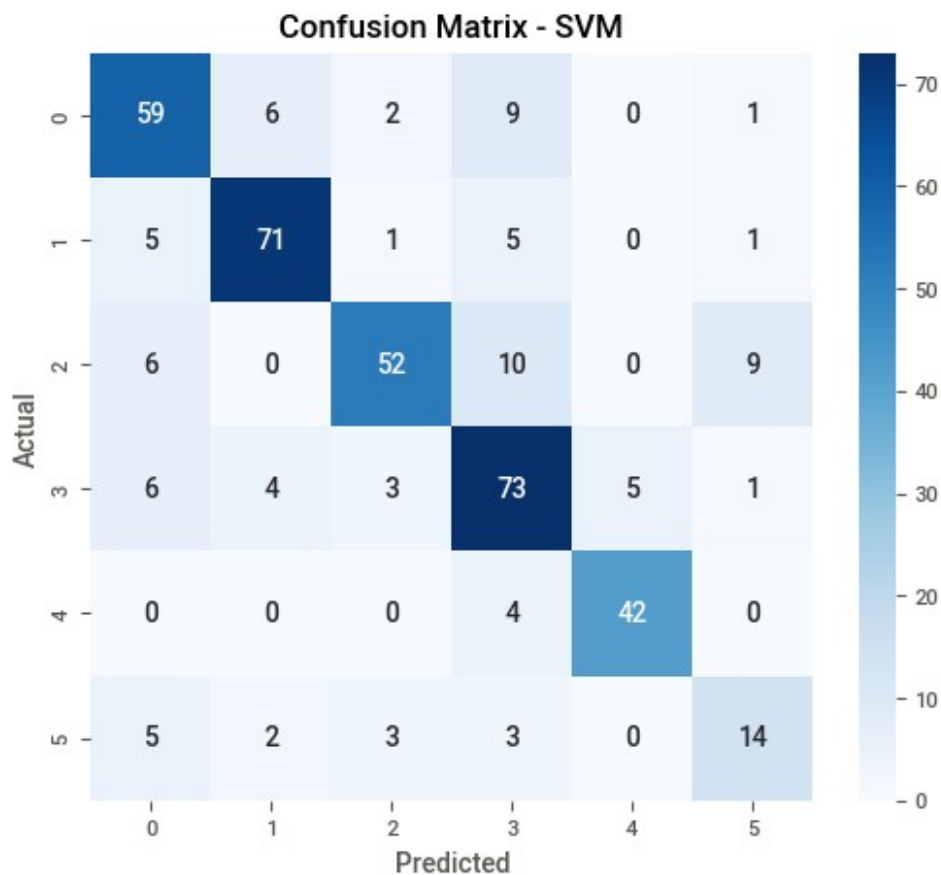
weighted avg	0.74	0.75	0.74	402
--------------	------	------	------	-----



Fitting 3 folds for each of 150 candidates, totalling 450 fits
 Best SVM Parameters for X_pca: {'C': 10, 'class_weight': None, 'gamma': 0.01, 'kernel': 'rbf'}

SVM Accuracy: 0.7736

	precision	recall	f1-score	support
C1	0.73	0.77	0.75	77
C2	0.86	0.86	0.86	83
C3	0.85	0.68	0.75	77
C4	0.70	0.79	0.74	92
C5	0.89	0.91	0.90	46
C6	0.54	0.52	0.53	27
accuracy			0.77	402
macro avg	0.76	0.75	0.76	402
weighted avg	0.78	0.77	0.77	402



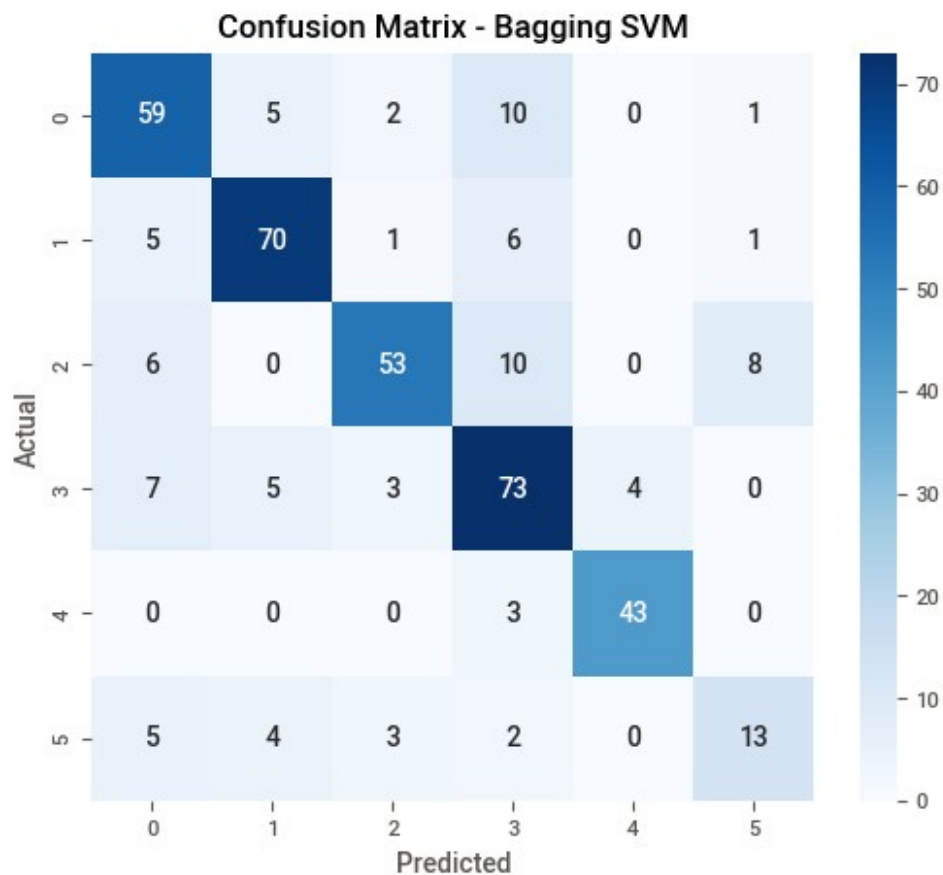
```

/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(

```

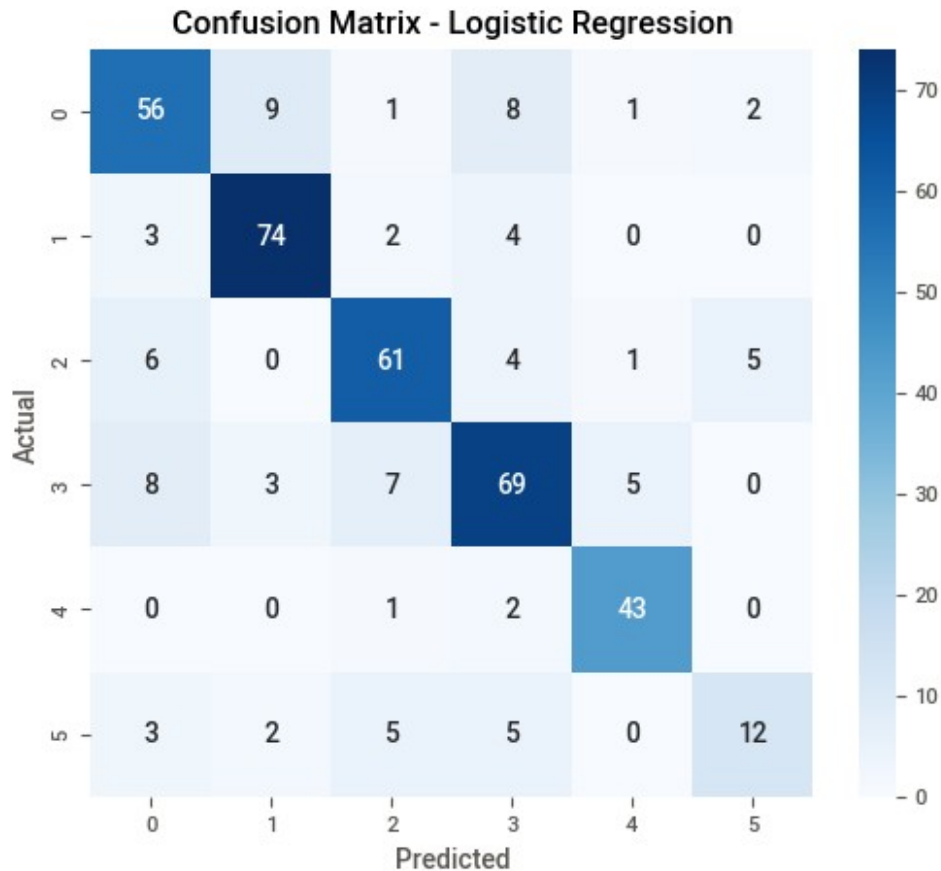
Bagging SVM Accuracy: 0.7736

	precision	recall	f1-score	support
C1	0.72	0.77	0.74	77
C2	0.83	0.84	0.84	83
C3	0.85	0.69	0.76	77
C4	0.70	0.79	0.74	92
C5	0.91	0.93	0.92	46
C6	0.57	0.48	0.52	27
accuracy			0.77	402
macro avg	0.76	0.75	0.76	402
weighted avg	0.78	0.77	0.77	402



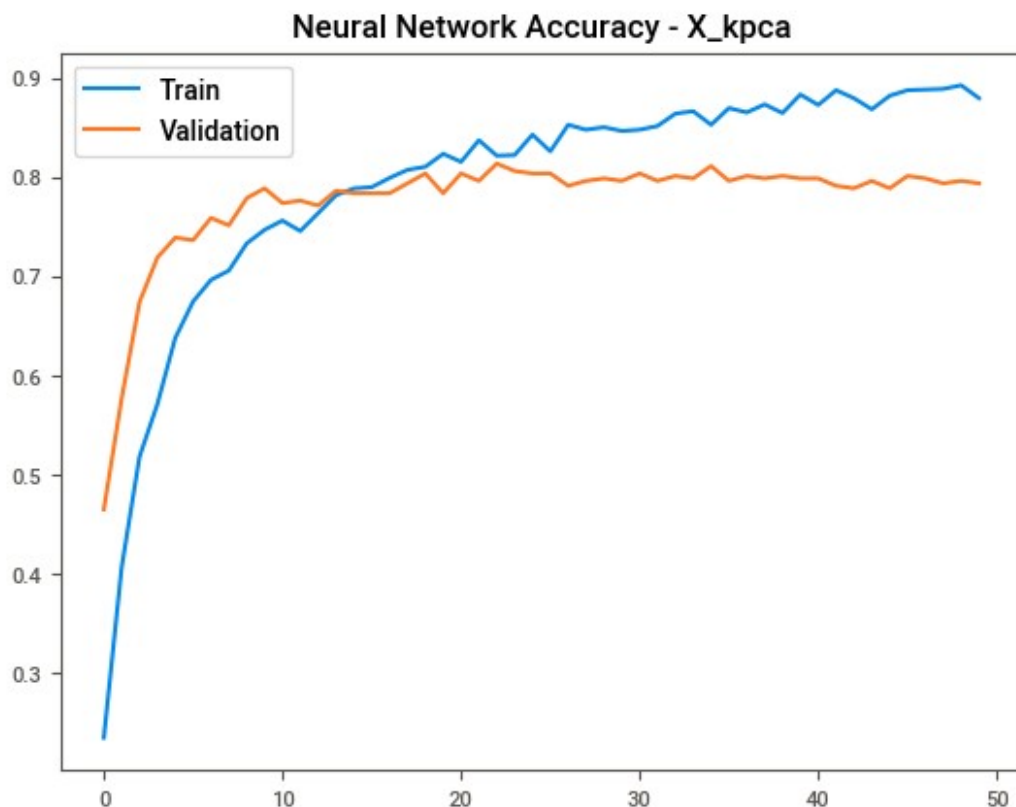
Logistic Regression Accuracy: 0.7836

	precision	recall	f1-score	support
C1	0.74	0.73	0.73	77
C2	0.84	0.89	0.87	83
C3	0.79	0.79	0.79	77
C4	0.75	0.75	0.75	92
C5	0.86	0.93	0.90	46
C6	0.63	0.44	0.52	27
accuracy			0.78	402
macro avg	0.77	0.76	0.76	402
weighted avg	0.78	0.78	0.78	402



```
===== Training Models on X_kpca Dataset =====  
Training set: (1607, 50), Test set: (402, 50)
```

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/  
dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim`  
argument to a layer. When using Sequential models, prefer using an  
`Input(shape)` object as the first layer in the model instead.  
    super().__init__(activity_regularizer=activity_regularizer,  
    **kwargs)
```



13/13 ————— 0s 10ms/step

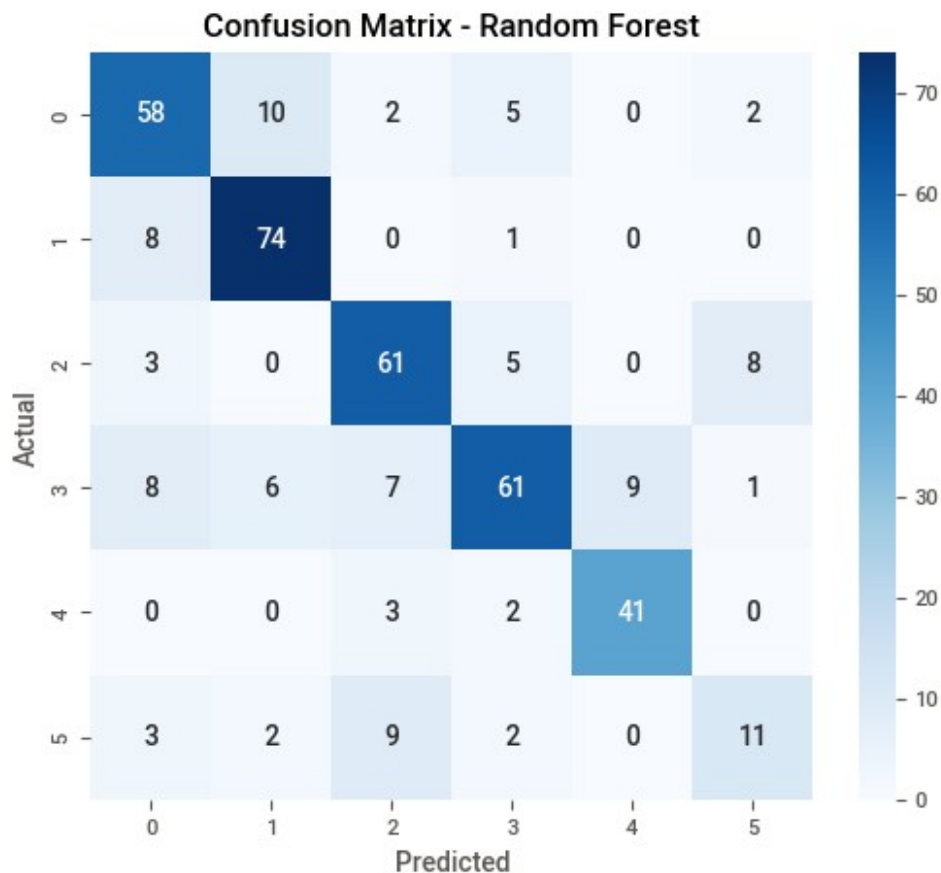
Neural Network - X_kpca Classification Report:

	precision	recall	f1-score	support
0	0.81	0.82	0.81	77
1	0.89	0.86	0.87	83
2	0.84	0.74	0.79	77
3	0.71	0.76	0.73	92
4	0.81	0.93	0.87	46
5	0.62	0.56	0.59	27
accuracy			0.79	402
macro avg	0.78	0.78	0.78	402
weighted avg	0.80	0.79	0.79	402

Random Forest Accuracy: 0.7612

	precision	recall	f1-score	support
C1	0.72	0.75	0.74	77
C2	0.80	0.89	0.85	83
C3	0.74	0.79	0.77	77
C4	0.80	0.66	0.73	92
C5	0.82	0.89	0.85	46

C6	0.50	0.41	0.45	27
accuracy			0.76	402
macro avg	0.73	0.73	0.73	402
weighted avg	0.76	0.76	0.76	402

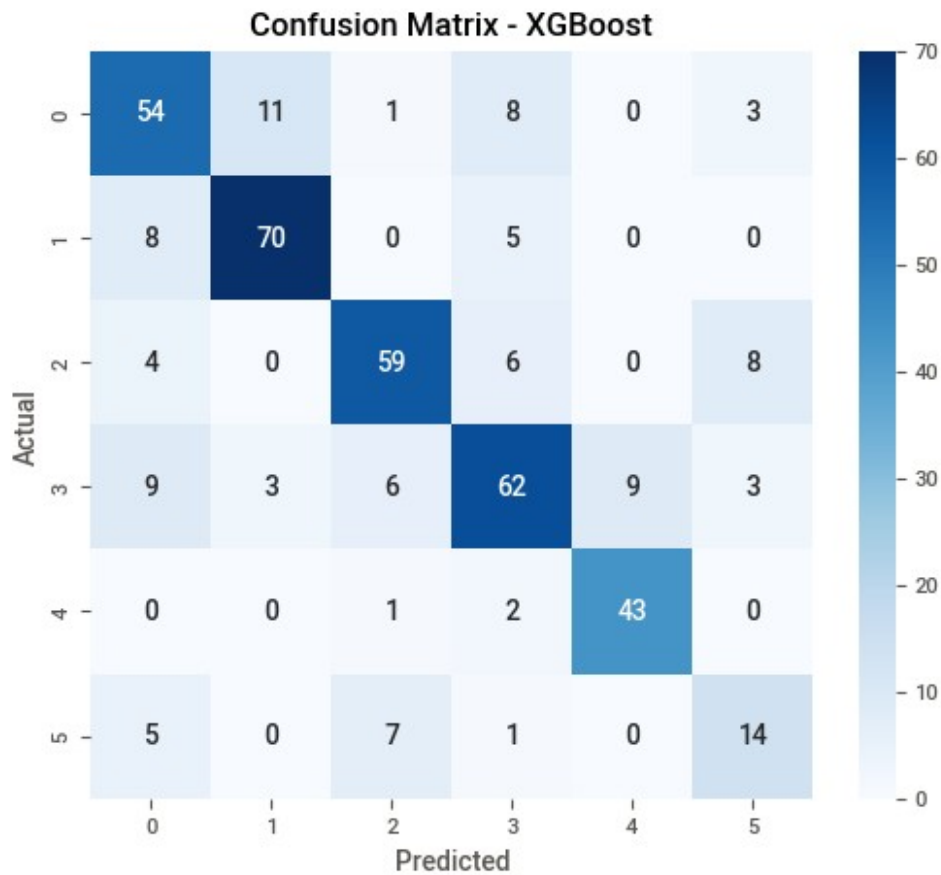


Best XGBoost Parameters for X_kpca: {'learning_rate': 0.1, 'max_depth': 2, 'n_estimators': 200}

XGBoost Accuracy: 0.7512

	precision	recall	f1-score	support
C1	0.68	0.70	0.69	77
C2	0.83	0.84	0.84	83
C3	0.80	0.77	0.78	77
C4	0.74	0.67	0.70	92
C5	0.83	0.93	0.88	46
C6	0.50	0.52	0.51	27
accuracy			0.75	402
macro avg	0.73	0.74	0.73	402

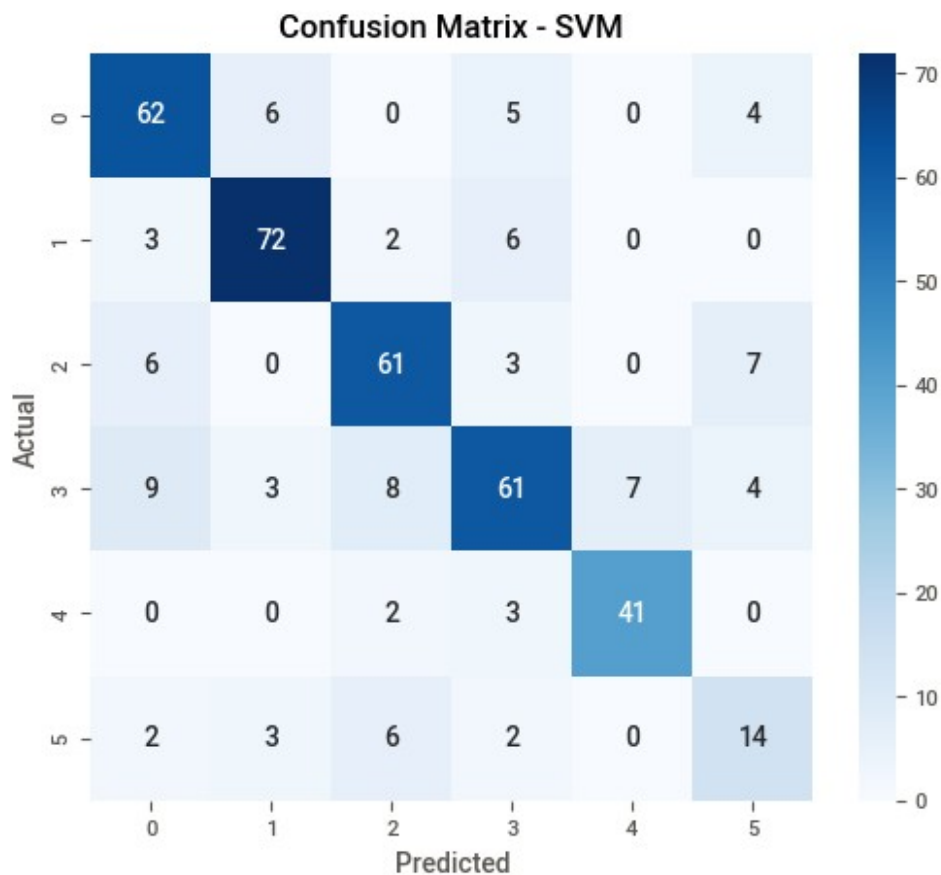
weighted avg	0.75	0.75	0.75	402
--------------	------	------	------	-----



Fitting 3 folds for each of 150 candidates, totalling 450 fits
 Best SVM Parameters for X_kpca: {'C': 50, 'class_weight': None, 'gamma': 0.001, 'kernel': 'rbf'}

SVM Accuracy: 0.7736

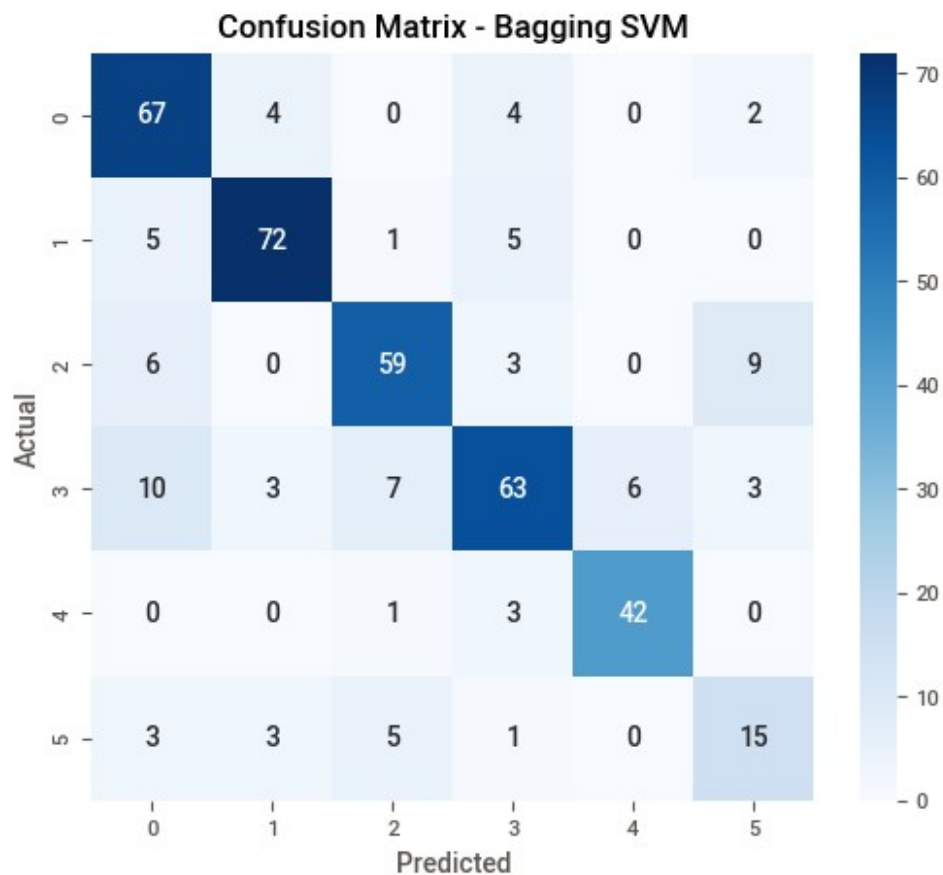
	precision	recall	f1-score	support
C1	0.76	0.81	0.78	77
C2	0.86	0.87	0.86	83
C3	0.77	0.79	0.78	77
C4	0.76	0.66	0.71	92
C5	0.85	0.89	0.87	46
C6	0.48	0.52	0.50	27
accuracy			0.77	402
macro avg	0.75	0.76	0.75	402
weighted avg	0.77	0.77	0.77	402



```
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
warnings.warn(
```

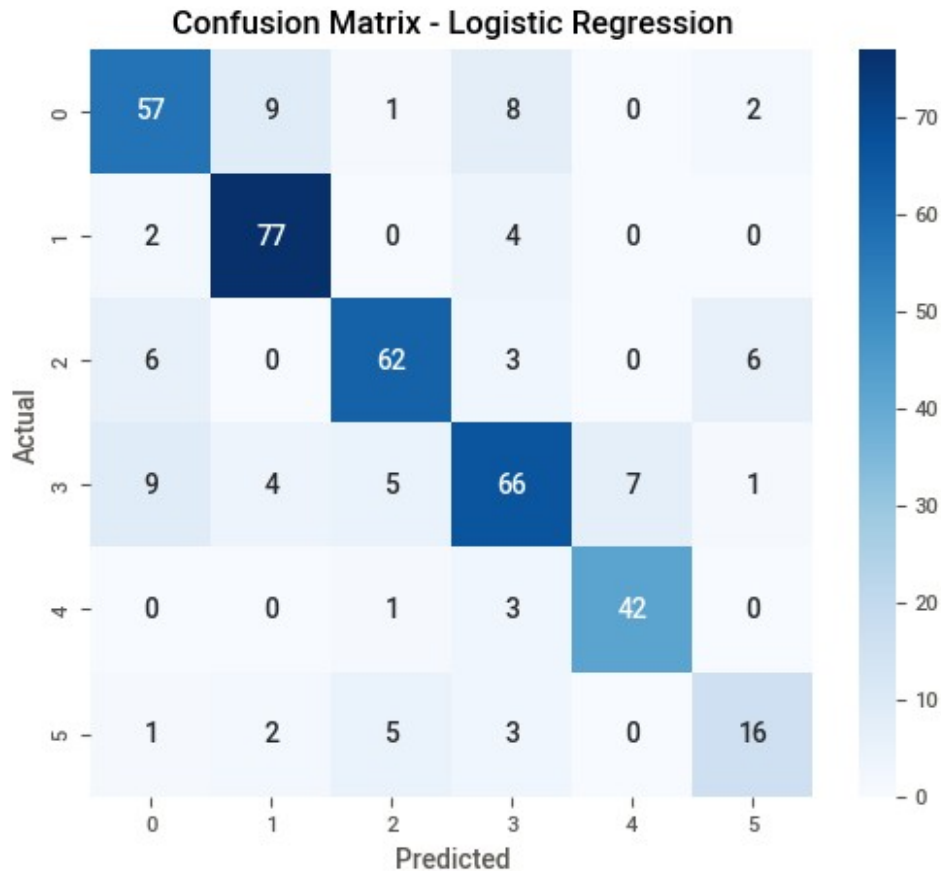
Bagging SVM Accuracy: 0.7910

	precision	recall	f1-score	support
C1	0.74	0.87	0.80	77
C2	0.88	0.87	0.87	83
C3	0.81	0.77	0.79	77
C4	0.80	0.68	0.74	92
C5	0.88	0.91	0.89	46
C6	0.52	0.56	0.54	27
accuracy			0.79	402
macro avg	0.77	0.78	0.77	402
weighted avg	0.79	0.79	0.79	402



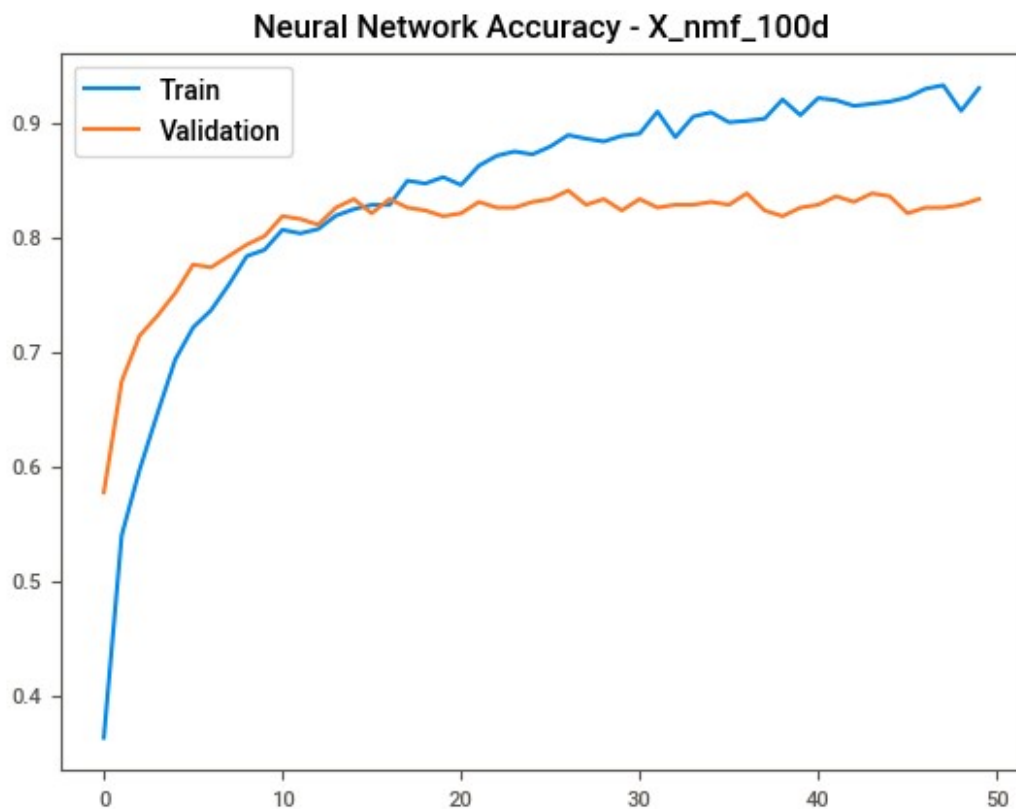
Logistic Regression Accuracy: 0.7960

	precision	recall	f1-score	support
C1	0.76	0.74	0.75	77
C2	0.84	0.93	0.88	83
C3	0.84	0.81	0.82	77
C4	0.76	0.72	0.74	92
C5	0.86	0.91	0.88	46
C6	0.64	0.59	0.62	27
accuracy			0.80	402
macro avg	0.78	0.78	0.78	402
weighted avg	0.79	0.80	0.79	402



```
===== Training Models on X_nmf_100d Dataset =====
Training set: (1607, 100), Test set: (402, 100)
```

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/
dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim`
argument to a layer. When using Sequential models, prefer using an
`Input(shape)` object as the first layer in the model instead.
  super().__init__(activity_regularizer=activity_regularizer,
**kwargs)
```



13/13 ————— 0s 9ms/step

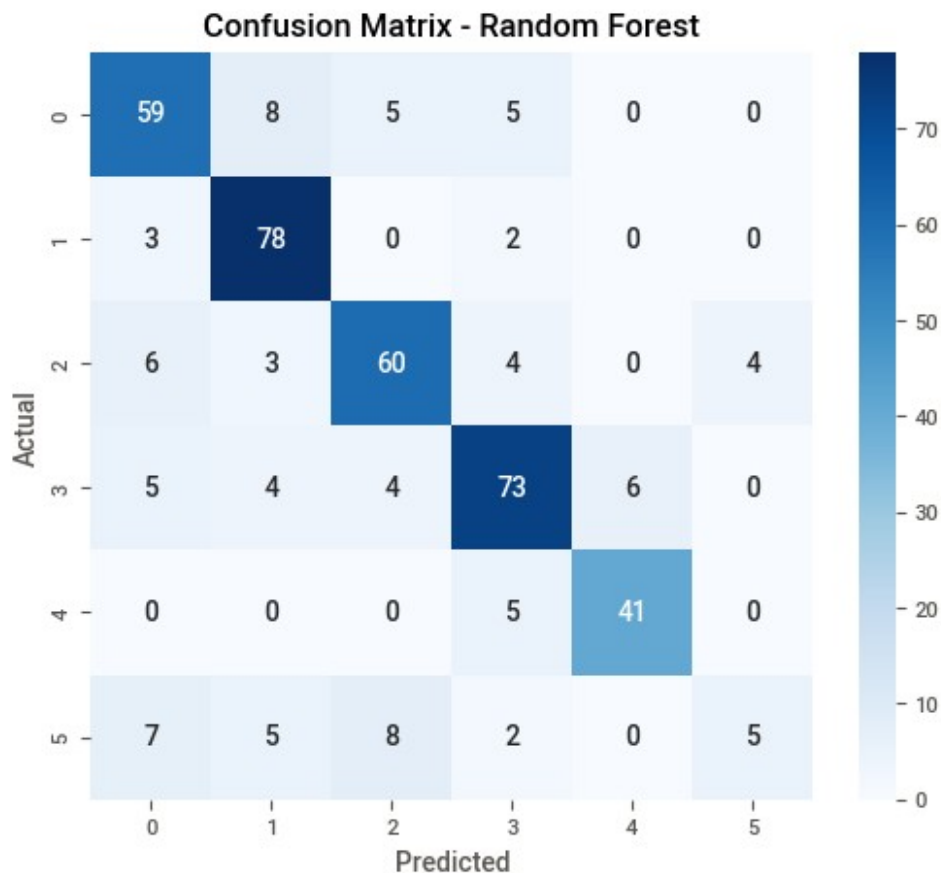
Neural Network - X_nmf_100d Classification Report:

	precision	recall	f1-score	support
0	0.81	0.78	0.79	77
1	0.94	0.89	0.91	83
2	0.85	0.87	0.86	77
3	0.75	0.83	0.79	92
4	0.84	0.93	0.89	46
5	0.83	0.56	0.67	27
accuracy			0.83	402
macro avg	0.84	0.81	0.82	402
weighted avg	0.84	0.83	0.83	402

Random Forest Accuracy: 0.7861

	precision	recall	f1-score	support
C1	0.74	0.77	0.75	77
C2	0.80	0.94	0.86	83
C3	0.78	0.78	0.78	77
C4	0.80	0.79	0.80	92
C5	0.87	0.89	0.88	46

C6	0.56	0.19	0.28	27
accuracy			0.79	402
macro avg	0.76	0.73	0.73	402
weighted avg	0.78	0.79	0.77	402

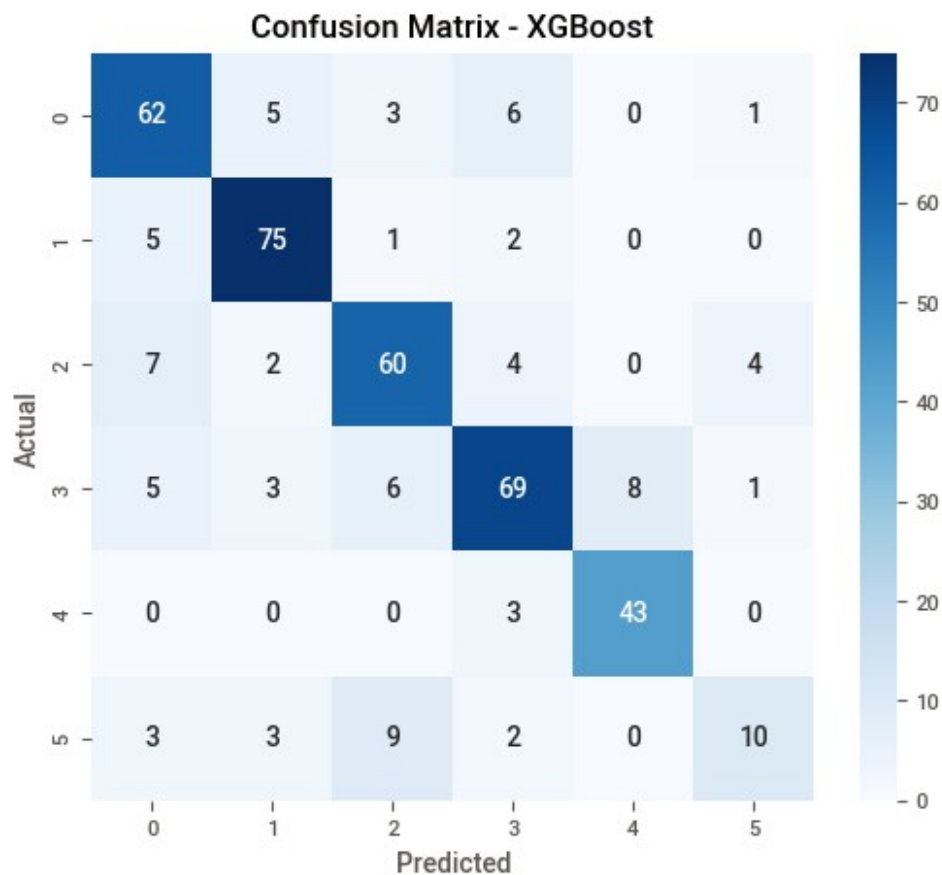


Best XGBoost Parameters for X_nmf_100d: {'learning_rate': 0.1, 'max_depth': 2, 'n_estimators': 200}

XGBoost Accuracy: 0.7935

	precision	recall	f1-score	support
C1	0.76	0.81	0.78	77
C2	0.85	0.90	0.88	83
C3	0.76	0.78	0.77	77
C4	0.80	0.75	0.78	92
C5	0.84	0.93	0.89	46
C6	0.62	0.37	0.47	27
accuracy			0.79	402
macro avg	0.77	0.76	0.76	402

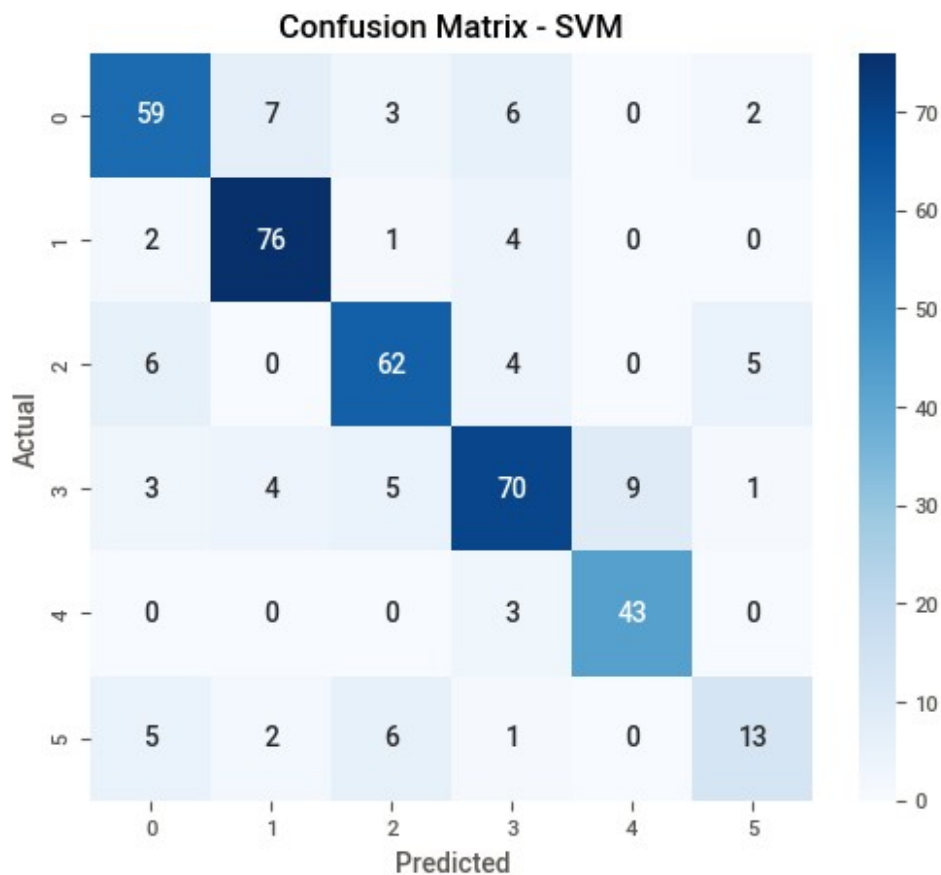
weighted avg	0.79	0.79	0.79	402
--------------	------	------	------	-----



Fitting 3 folds for each of 150 candidates, totalling 450 fits
 Best SVM Parameters for X_nmf_100d: {'C': 50, 'class_weight': 'balanced', 'gamma': 'auto', 'kernel': 'rbf'}

SVM Accuracy: 0.8035

	precision	recall	f1-score	support
C1	0.79	0.77	0.78	77
C2	0.85	0.92	0.88	83
C3	0.81	0.81	0.81	77
C4	0.80	0.76	0.78	92
C5	0.83	0.93	0.88	46
C6	0.62	0.48	0.54	27
accuracy			0.80	402
macro avg	0.78	0.78	0.78	402
weighted avg	0.80	0.80	0.80	402



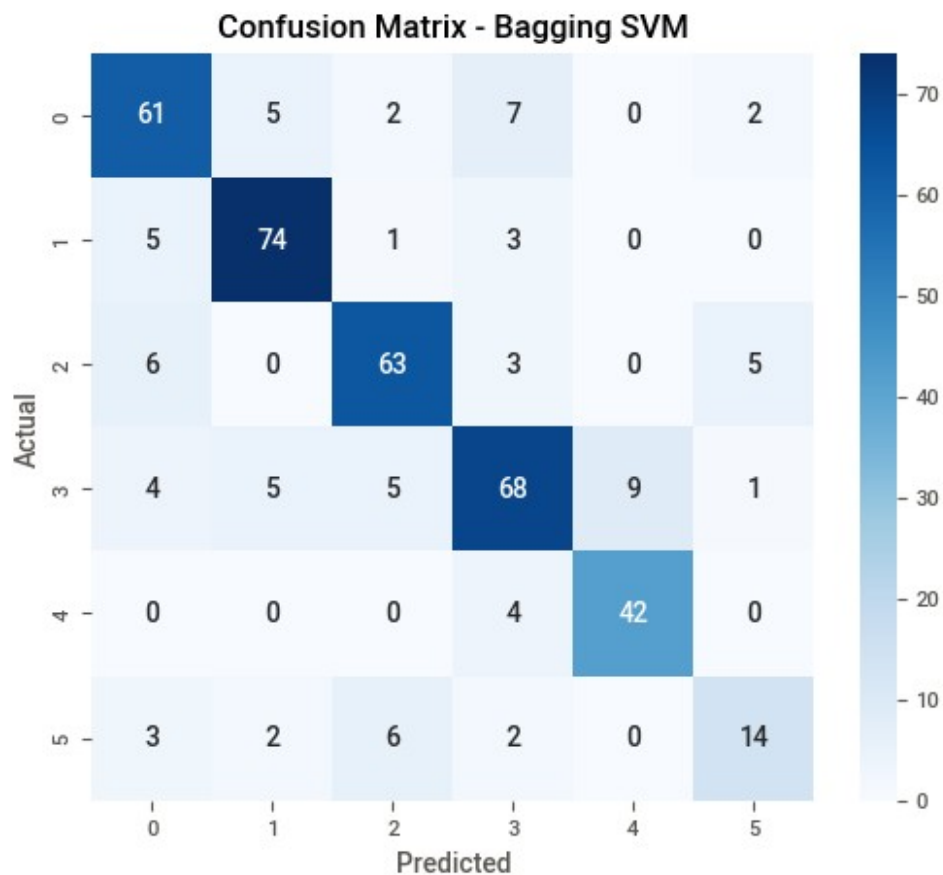
```

/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(

```

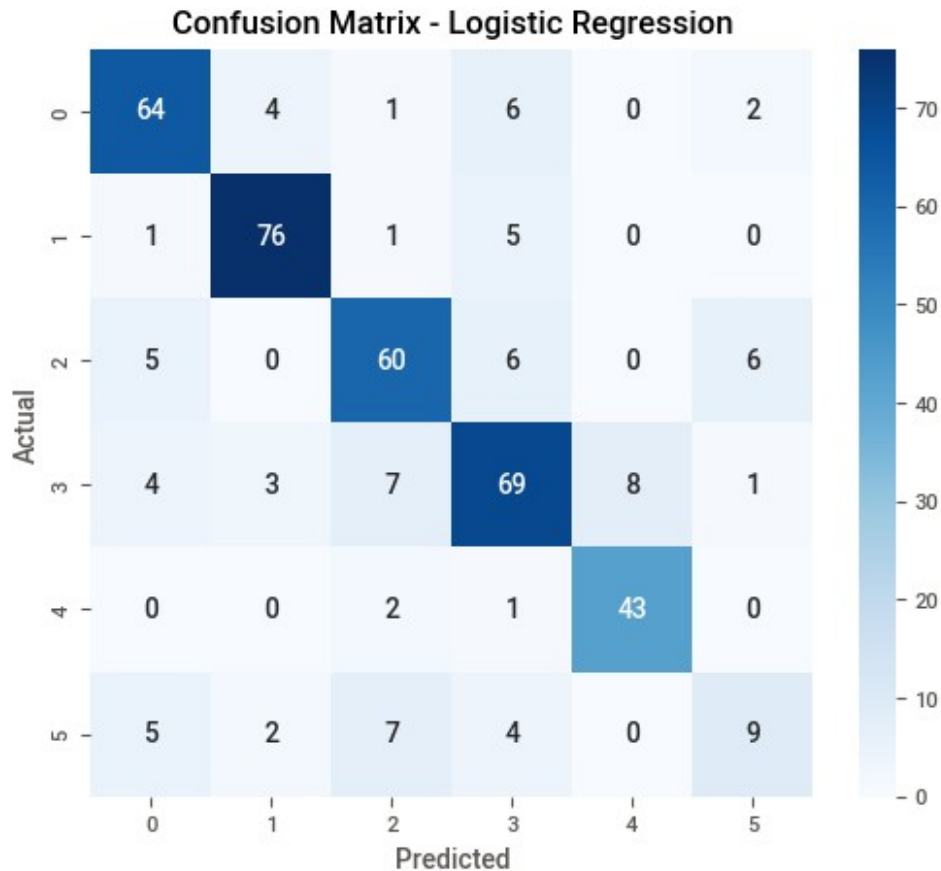
Bagging SVM Accuracy: 0.8010

	precision	recall	f1-score	support
C1	0.77	0.79	0.78	77
C2	0.86	0.89	0.88	83
C3	0.82	0.82	0.82	77
C4	0.78	0.74	0.76	92
C5	0.82	0.91	0.87	46
C6	0.64	0.52	0.57	27
accuracy			0.80	402
macro avg	0.78	0.78	0.78	402
weighted avg	0.80	0.80	0.80	402



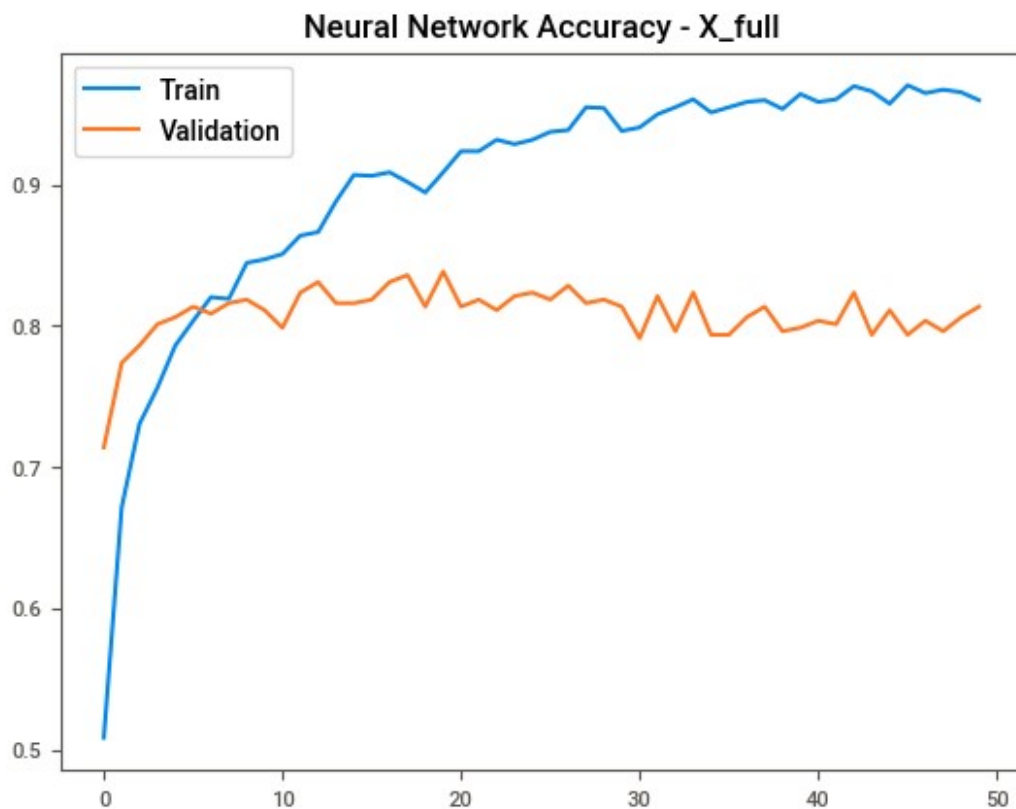
Logistic Regression Accuracy: 0.7985

	precision	recall	f1-score	support
C1	0.81	0.83	0.82	77
C2	0.89	0.92	0.90	83
C3	0.77	0.78	0.77	77
C4	0.76	0.75	0.75	92
C5	0.84	0.93	0.89	46
C6	0.50	0.33	0.40	27
accuracy			0.80	402
macro avg	0.76	0.76	0.76	402
weighted avg	0.79	0.80	0.79	402



```
===== Training Models on X_full Dataset =====  
Training set: (1607, 440), Test set: (402, 440)
```

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/  
dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim`  
argument to a layer. When using Sequential models, prefer using an  
`Input(shape)` object as the first layer in the model instead.  
super().__init__(activity_regularizer=activity_regularizer,  
**kwargs)
```



13/13 ————— 0s 8ms/step

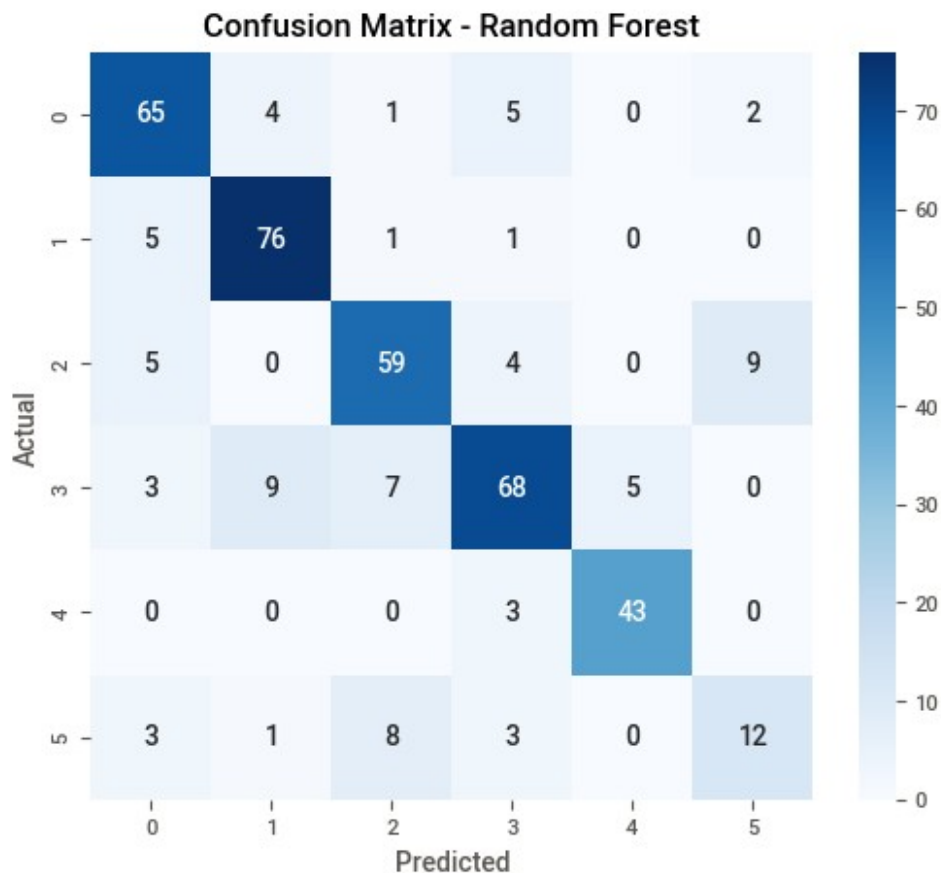
Neural Network - X_full Classification Report:

	precision	recall	f1-score	support
0	0.80	0.86	0.83	77
1	0.88	0.86	0.87	83
2	0.83	0.81	0.82	77
3	0.78	0.79	0.79	92
4	0.84	0.93	0.89	46
5	0.60	0.44	0.51	27
accuracy			0.81	402
macro avg	0.79	0.78	0.78	402
weighted avg	0.81	0.81	0.81	402

Random Forest Accuracy: 0.8035

	precision	recall	f1-score	support
C1	0.80	0.84	0.82	77
C2	0.84	0.92	0.88	83
C3	0.78	0.77	0.77	77
C4	0.81	0.74	0.77	92
C5	0.90	0.93	0.91	46

C6	0.52	0.44	0.48	27
accuracy			0.80	402
macro avg	0.78	0.77	0.77	402
weighted avg	0.80	0.80	0.80	402

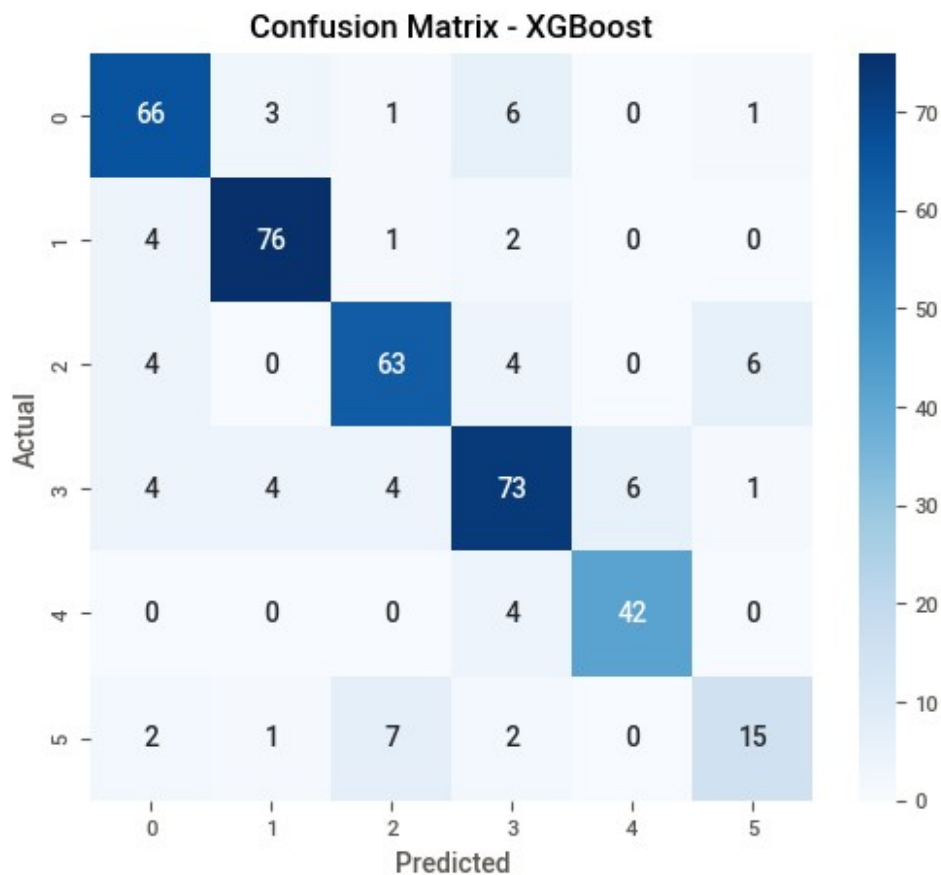


Best XGBoost Parameters for X_full: {'learning_rate': 0.1, 'max_depth': 2, 'n_estimators': 200}

XGBoost Accuracy: 0.8333

	precision	recall	f1-score	support
C1	0.82	0.86	0.84	77
C2	0.90	0.92	0.91	83
C3	0.83	0.82	0.82	77
C4	0.80	0.79	0.80	92
C5	0.88	0.91	0.89	46
C6	0.65	0.56	0.60	27
accuracy			0.83	402
macro avg	0.81	0.81	0.81	402

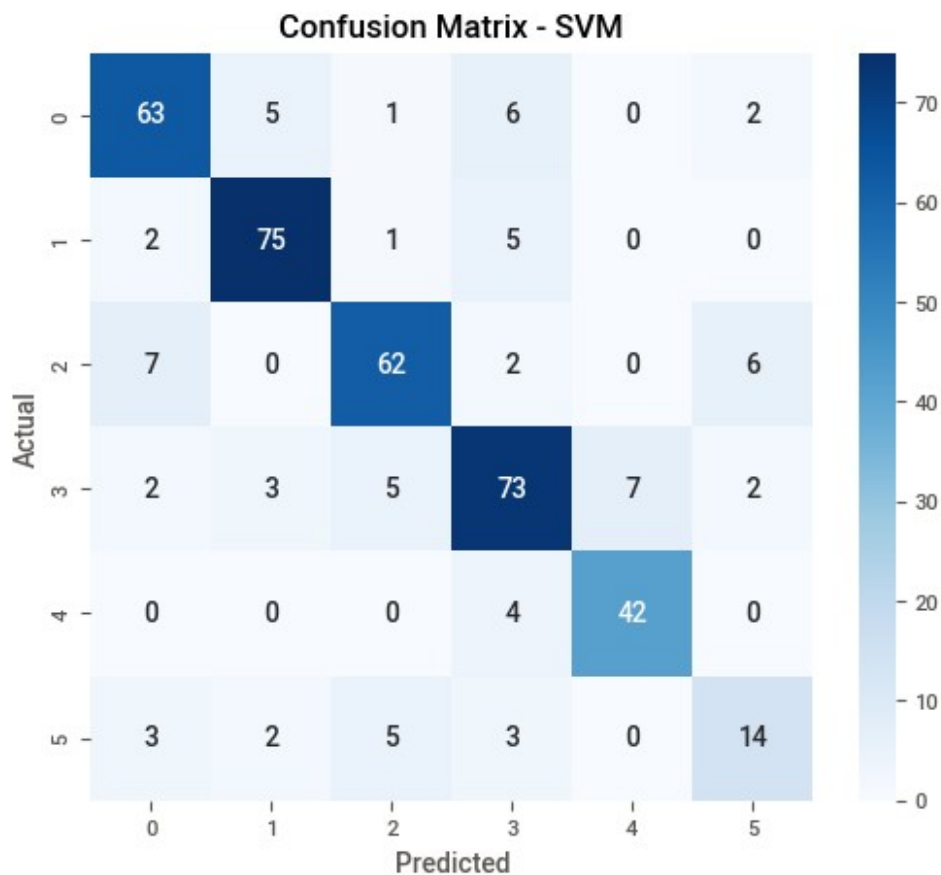
weighted avg	0.83	0.83	0.83	402
--------------	------	------	------	-----



Fitting 3 folds for each of 150 candidates, totalling 450 fits
 Best SVM Parameters for X_full: {'C': 10, 'class_weight': 'balanced', 'gamma': 'auto', 'kernel': 'rbf'}

SVM Accuracy: 0.8184

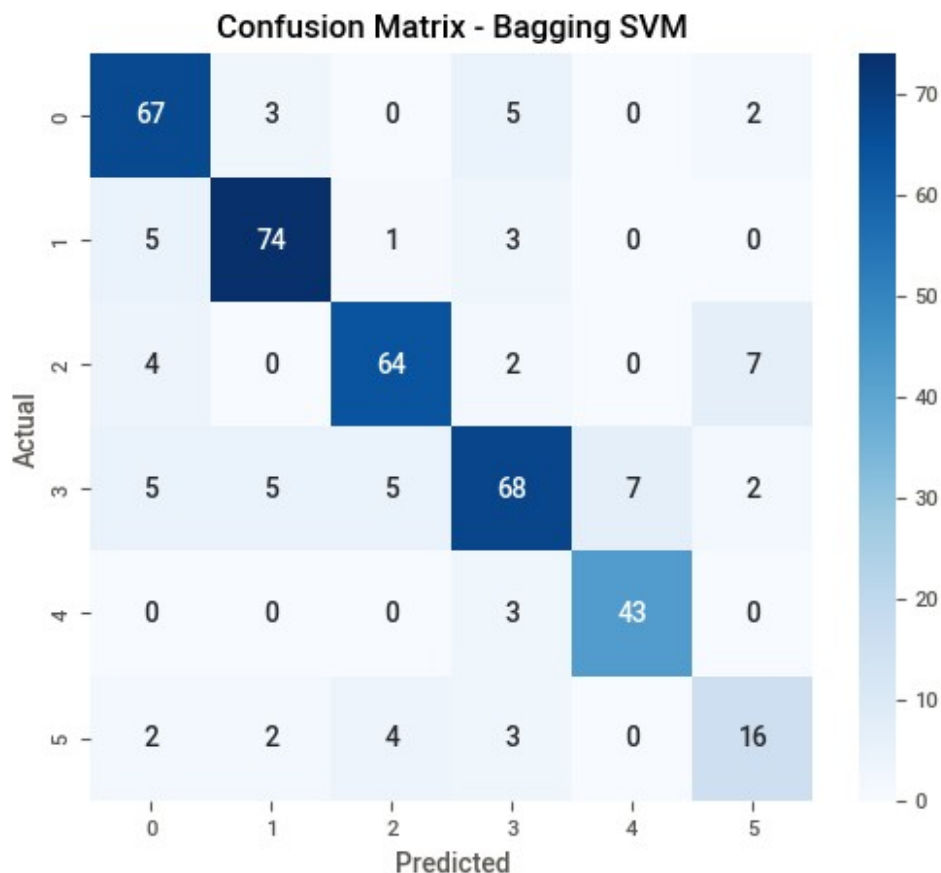
	precision	recall	f1-score	support
C1	0.82	0.82	0.82	77
C2	0.88	0.90	0.89	83
C3	0.84	0.81	0.82	77
C4	0.78	0.79	0.79	92
C5	0.86	0.91	0.88	46
C6	0.58	0.52	0.55	27
accuracy			0.82	402
macro avg	0.79	0.79	0.79	402
weighted avg	0.82	0.82	0.82	402



```
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
warnings.warn(
```

Bagging SVM Accuracy: 0.8259

	precision	recall	f1-score	support
C1	0.81	0.87	0.84	77
C2	0.88	0.89	0.89	83
C3	0.86	0.83	0.85	77
C4	0.81	0.74	0.77	92
C5	0.86	0.93	0.90	46
C6	0.59	0.59	0.59	27
accuracy			0.83	402
macro avg	0.80	0.81	0.81	402
weighted avg	0.83	0.83	0.82	402



```
/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge
(status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
```

Increase the number of iterations (`max_iter`) or scale the data as shown in:

<https://scikit-learn.org/stable/modules/preprocessing.html>

Please also refer to the documentation for alternative solver options:

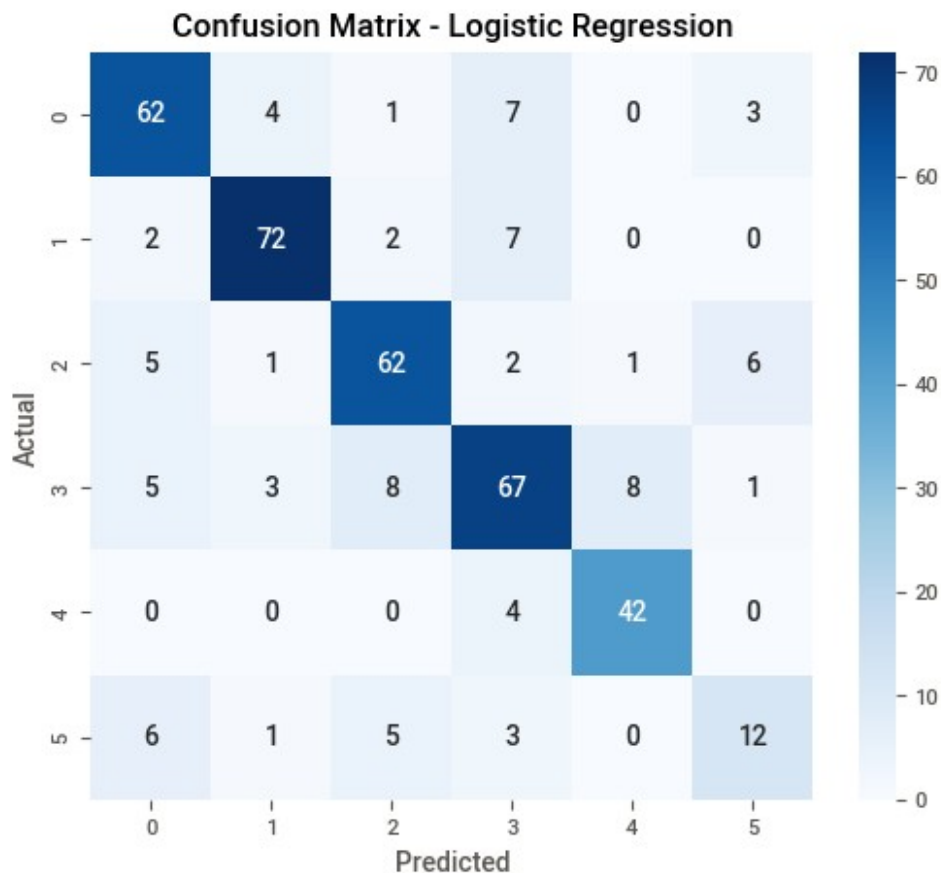
https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression

```
n_iter_i = _check_optimize_result(
```

Logistic Regression Accuracy: 0.7886

	precision	recall	f1-score	support
C1	0.78	0.81	0.79	77
C2	0.89	0.87	0.88	83
C3	0.79	0.81	0.80	77
C4	0.74	0.73	0.74	92
C5	0.82	0.91	0.87	46

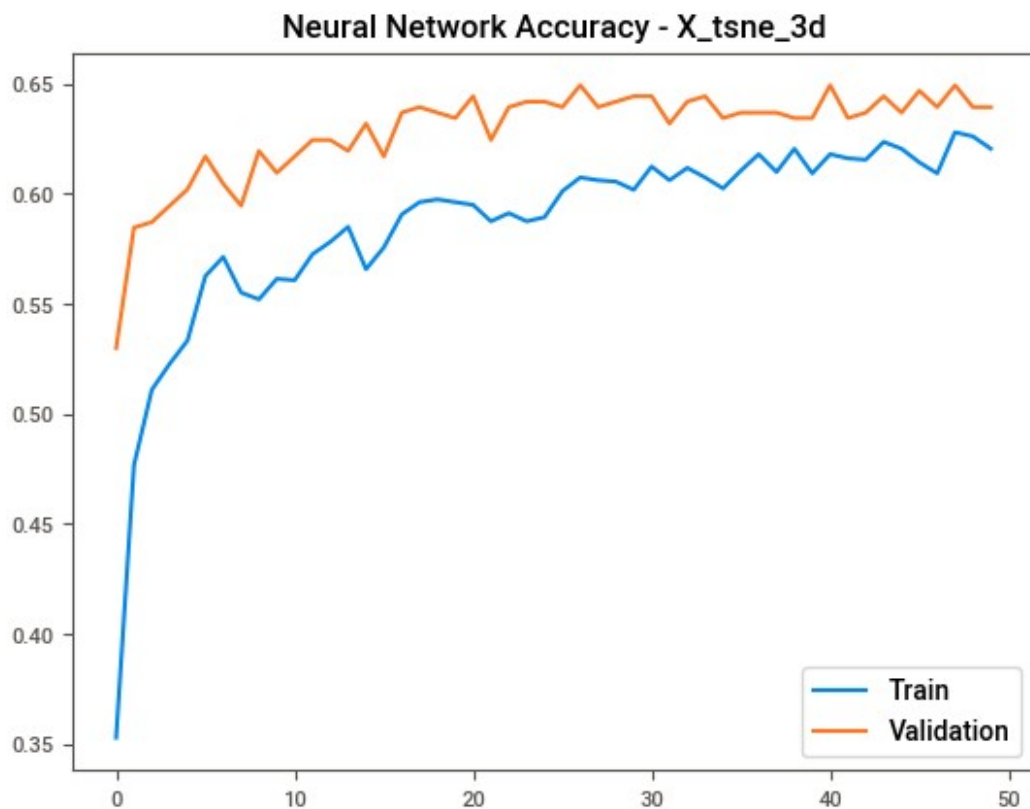
C6	0.55	0.44	0.49	27
accuracy			0.79	402
macro avg	0.76	0.76	0.76	402
weighted avg	0.79	0.79	0.79	402



==== Training Models on X_tsne_3d Dataset =====

Training set: (1607, 3), Test set: (402, 3)

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim` argument to a layer. When using Sequential models, prefer using an `Input(shape)` object as the first layer in the model instead.
  super().__init__(activity_regularizer=activity_regularizer,
**kwargs)
```



13/13 ————— 0s 8ms/step

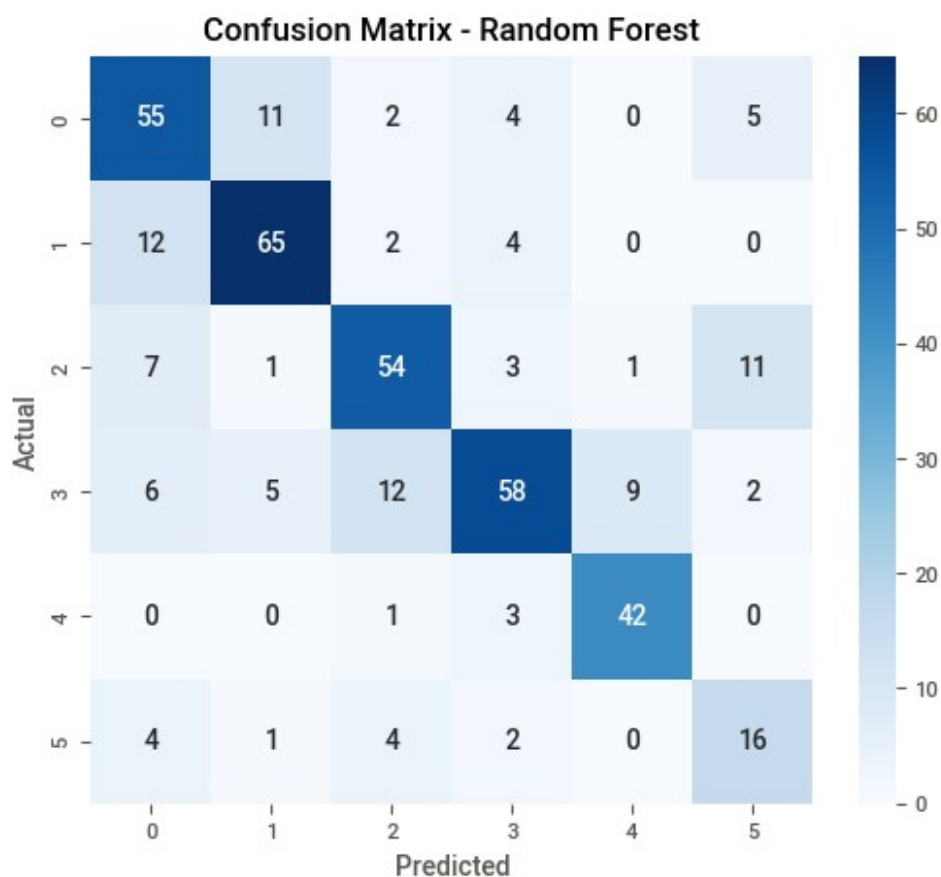
Neural Network - X_tsne_3d Classification Report:

	precision	recall	f1-score	support
0	0.61	0.60	0.61	77
1	0.73	0.71	0.72	83
2	0.57	0.61	0.59	77
3	0.65	0.57	0.60	92
4	0.84	0.89	0.86	46
5	0.34	0.44	0.39	27
accuracy			0.64	402
macro avg	0.62	0.64	0.63	402
weighted avg	0.65	0.64	0.64	402

Random Forest Accuracy: 0.7214

	precision	recall	f1-score	support
C1	0.65	0.71	0.68	77
C2	0.78	0.78	0.78	83
C3	0.72	0.70	0.71	77
C4	0.78	0.63	0.70	92
C5	0.81	0.91	0.86	46

C6	0.47	0.59	0.52	27
accuracy			0.72	402
macro avg	0.70	0.72	0.71	402
weighted avg	0.73	0.72	0.72	402

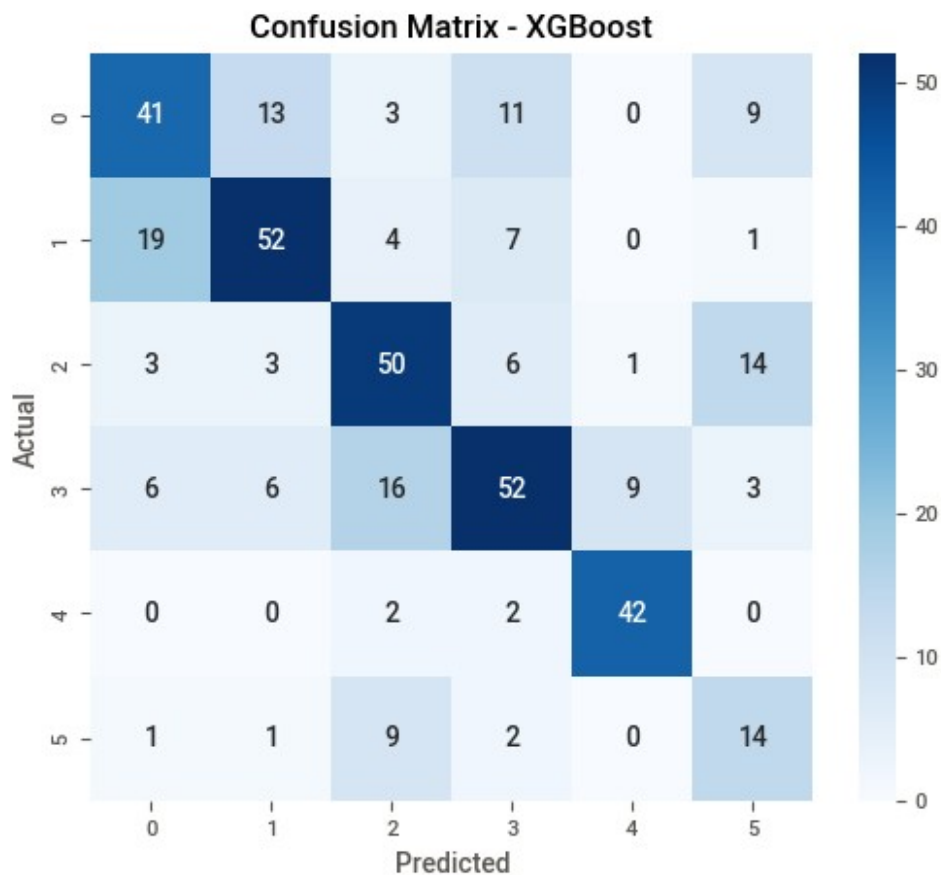


Best XGBoost Parameters for X_tsne_3d: {'learning_rate': 0.1, 'max_depth': 2, 'n_estimators': 200}

XGBoost Accuracy: 0.6244

	precision	recall	f1-score	support
C1	0.59	0.53	0.56	77
C2	0.69	0.63	0.66	83
C3	0.60	0.65	0.62	77
C4	0.65	0.57	0.60	92
C5	0.81	0.91	0.86	46
C6	0.34	0.52	0.41	27
accuracy			0.62	402
macro avg	0.61	0.63	0.62	402

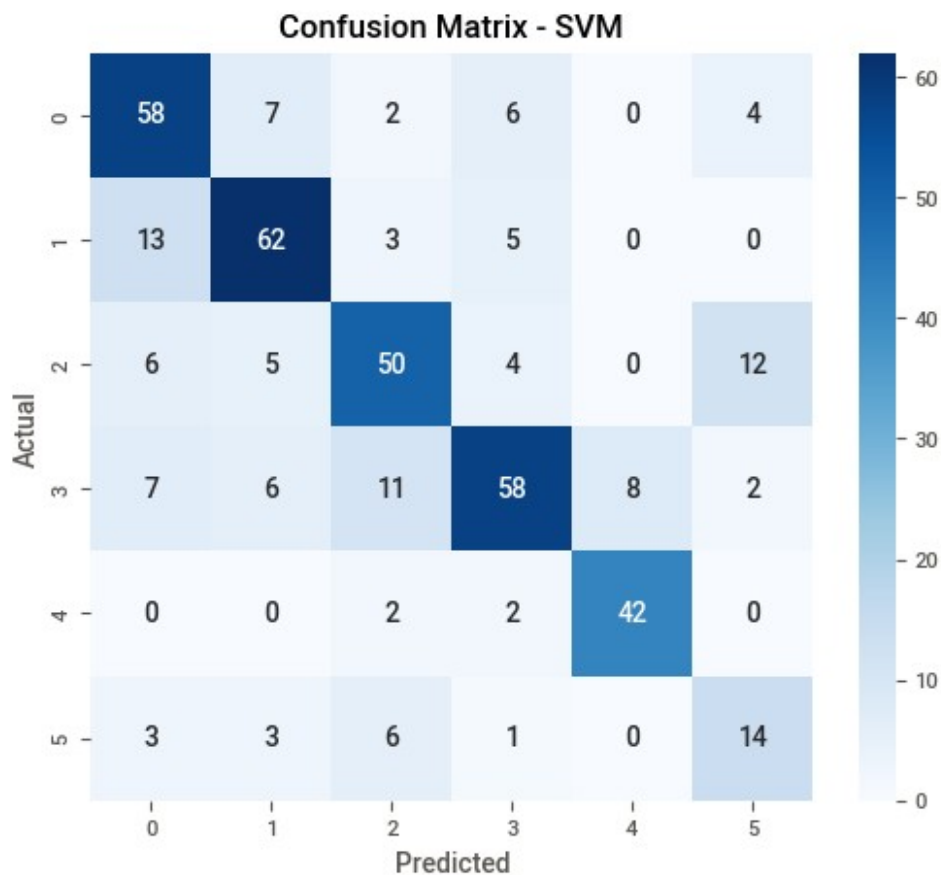
weighted avg	0.63	0.62	0.63	402
--------------	------	------	------	-----



Fitting 3 folds for each of 150 candidates, totalling 450 fits
 Best SVM Parameters for X_tsne_3d: {'C': 100, 'class_weight': None, 'gamma': 'auto', 'kernel': 'rbf'}

SVM Accuracy: 0.7065

	precision	recall	f1-score	support
C1	0.67	0.75	0.71	77
C2	0.75	0.75	0.75	83
C3	0.68	0.65	0.66	77
C4	0.76	0.63	0.69	92
C5	0.84	0.91	0.87	46
C6	0.44	0.52	0.47	27
accuracy			0.71	402
macro avg	0.69	0.70	0.69	402
weighted avg	0.71	0.71	0.71	402



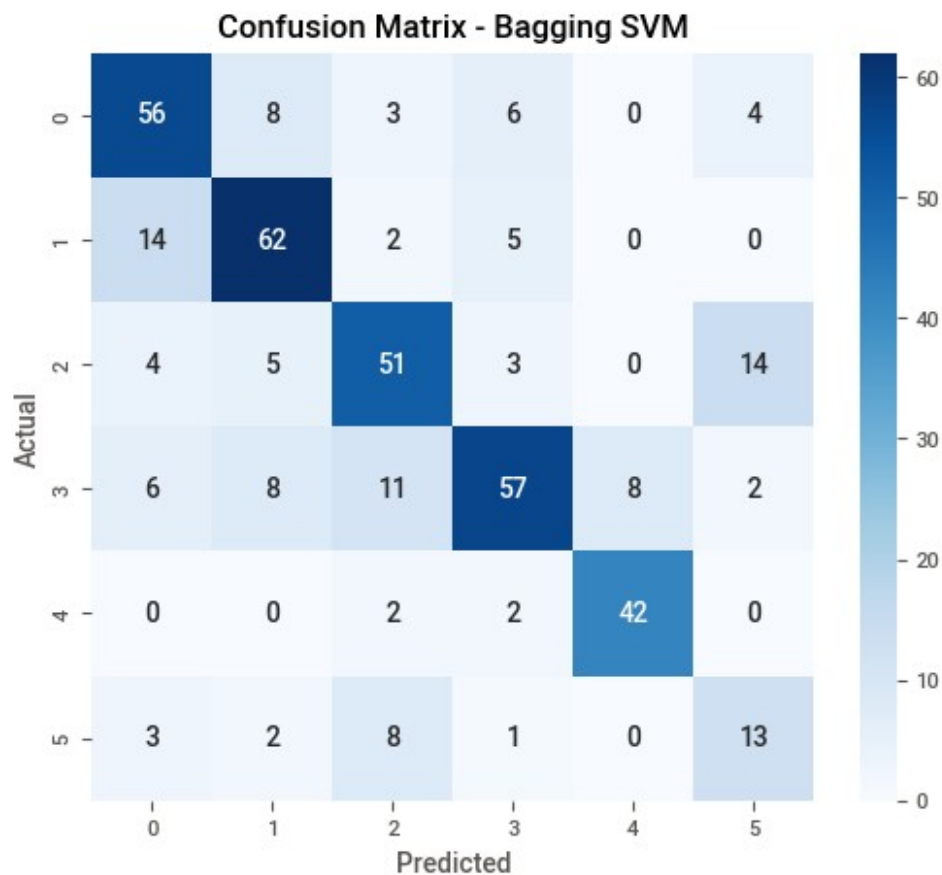
```

/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(

```

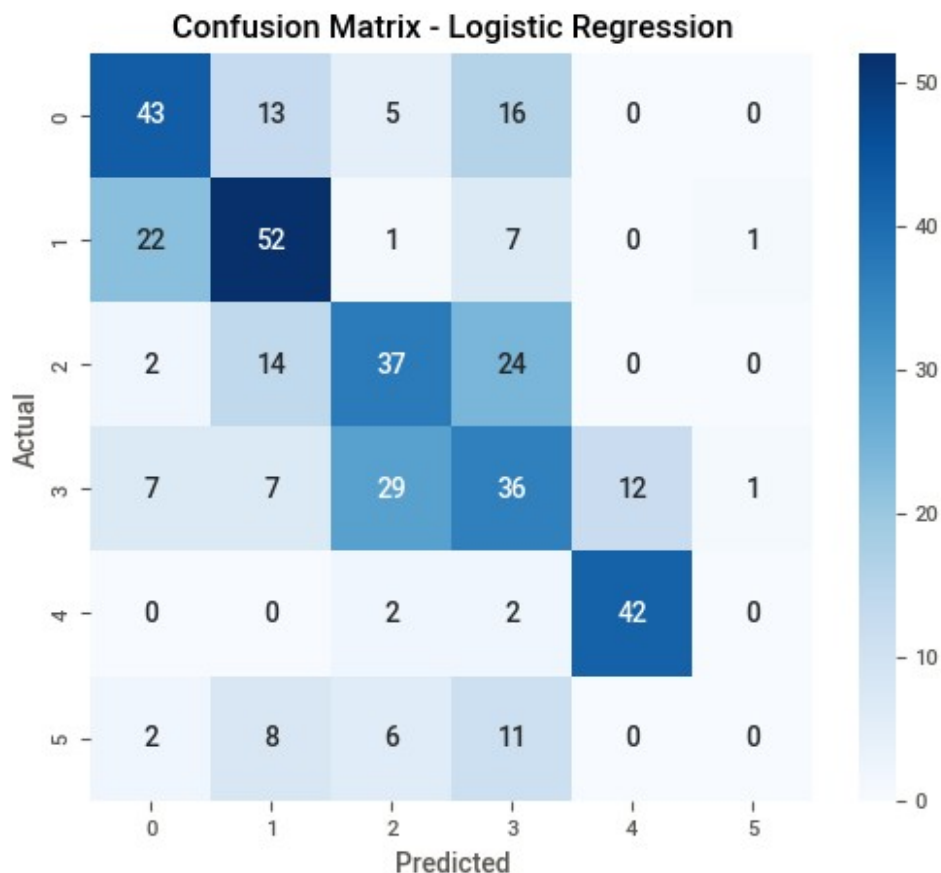
Bagging SVM Accuracy: 0.6990

	precision	recall	f1-score	support
C1	0.67	0.73	0.70	77
C2	0.73	0.75	0.74	83
C3	0.66	0.66	0.66	77
C4	0.77	0.62	0.69	92
C5	0.84	0.91	0.87	46
C6	0.39	0.48	0.43	27
accuracy			0.70	402
macro avg	0.68	0.69	0.68	402
weighted avg	0.71	0.70	0.70	402



Logistic Regression Accuracy: 0.5224

	precision	recall	f1-score	support
C1	0.57	0.56	0.56	77
C2	0.55	0.63	0.59	83
C3	0.46	0.48	0.47	77
C4	0.38	0.39	0.38	92
C5	0.78	0.91	0.84	46
C6	0.00	0.00	0.00	27
accuracy			0.52	402
macro avg	0.46	0.49	0.47	402
weighted avg	0.49	0.52	0.50	402



```
# Summary only
print("\n==== Model Performance Summary =====")
for model, acc in results.items():
    print(f"{model}: {acc:.4f}")
```

```
==== Model Performance Summary =====
```

```
RandomForest_NMF: 0.7886
XGBoost_NMF: 0.8085
SVM_NMF: 0.7836
BaggingSVM_NMF: 0.7886
LogisticRegression_NMF: 0.8060
RandomForest_X_pca: 0.7289
XGBoost_X_pca: 0.7463
SVM_X_pca: 0.7736
BaggingSVM_X_pca: 0.7736
LogisticRegression_X_pca: 0.7836
RandomForest_X_kpca: 0.7612
XGBoost_X_kpca: 0.7512
SVM_X_kpca: 0.7736
BaggingSVM_X_kpca: 0.7910
LogisticRegression_X_kpca: 0.7960
RandomForest_X_nmf_100d: 0.7861
```

```
XGBoost_X_nmf_100d: 0.7935
SVM_X_nmf_100d: 0.8035
BaggingSVM_X_nmf_100d: 0.8010
LogisticRegression_X_nmf_100d: 0.7985
RandomForest_X_full: 0.8035
XGBoost_X_full: 0.8333
SVM_X_full: 0.8184
BaggingSVM_X_full: 0.8259
LogisticRegression_X_full: 0.7886
RandomForest_X_tsne_3d: 0.7214
XGBoost_X_tsne_3d: 0.6244
SVM_X_tsne_3d: 0.7065
BaggingSVM_X_tsne_3d: 0.6990
LogisticRegression_X_tsne_3d: 0.5224
```

2.1 Model Selection and Training

Partitioned the pre-processed dataset into **training (80%)** and **test (20%)** sets using **stratified sampling**, ensuring balanced class distributions. Below are the dataset dimensions:

```
X_train.shape, X_test.shape
((1607, 440), (402, 440))
```

Experimented with multiple classification algorithms to assess their performance on the dataset:

- **Random Forest Classifier**
- **XGBoost Classifier**
- **Support Vector Machine (SVM)**
- **Bagging Classifier with SVM**
- **Logistic Regression**
- **Neural Networks (Feedforward NN)**

Each model was trained on multiple feature representations, including **PCA-transformed**, **Kernel PCA**, **NMF**, **t-SNE**, and the **full feature set**. The goal was to determine the best feature space for classification.

2.2 Optimal Model Development

To enhance model performance, we applied **hyperparameter tuning** via **GridSearchCV** and **cross-validation (k=3)**:

XGBoost Hyperparameter Tuning

```
param_grid = {
    'max_depth': [1, 2],
    'learning_rate': [0.01, 0.1],
```

```
}
    'n_estimators': [100, 200]
```

- Evaluated different combinations using **log-loss** as the evaluation metric.
- Selected the best hyperparameters for each dataset.

SVM Hyperparameter Tuning

```
param_grid = {
    'C': [0.1, 1, 10, 50, 100],
    'gamma': ['scale', 'auto', 0.001, 0.01, 0.1],
    'kernel': ['linear', 'rbf', 'poly'],
    'class_weight': ['balanced', None]
}
```

- Optimized kernel selection, regularization (C), and gamma values for non-linear decision boundaries.

Neural Network Training

- Implemented a **Feedforward Neural Network (FNN)** with dropout for regularization.
- Configured layers as follows:

```
model = Sequential([
    Dense(64, activation='relu', input_shape=(input_dim,)),
    Dropout(0.3),
    Dense(32, activation='relu'),
    Dropout(0.3),
    Dense(len(set(y)), activation='softmax')
])
```

- Used **Adam optimizer** (learning rate = 0.001) and **sparse categorical cross-entropy** as the loss function.

2.3 Justification of the Selected Methods

The chosen models align with the dataset's complexity:

- **Random Forest & XGBoost:** Robust against outliers and handle feature importance well.
- **SVM:** Effective for high-dimensional data, especially with kernel tricks.
- **Bagging Classifier:** Combines multiple weak learners (SVM) to enhance performance.
- **Neural Networks:** Captures non-linear relationships in the dataset, suitable for complex patterns.

Feature engineering was crucial, with **PCA and NMF improving model performance** by reducing redundancy in features.

2.4 Strategies to Address Model Issues

Several strategies were implemented to enhance model generalization and robustness:

- **Class Imbalance Handling:** Applied stratified sampling and `class_weight='balanced'` in models like SVM.
- **Regularization:** Used dropout in NN and `max_depth` tuning in XGBoost to prevent overfitting.
- **Feature Selection:** Evaluated different feature reduction techniques (PCA, NMF, t-SNE) to improve interpretability.
- **Cross-validation:** Ensured models generalize well across different data splits. The following table summarizes the accuracy results obtained for different models and dimensionality reduction techniques:

Model	NMF	PCA	Kernel PCA	NMF (100D)	Full Dataset	t-SNE (3D)
Random Forest	0.7886	0.7289	0.7612	0.7861	0.8035	0.7214
XGBoost	0.8085	0.7463	0.7512	0.7935	0.8333	0.6244
SVM	0.7836	0.7736	0.7736	0.8035	0.8184	0.7065
Bagging SVM	0.7886	0.7736	0.7910	0.8010	0.8259	0.6990
Logistic Regression	0.8060	0.7836	0.7960	0.7985	0.7886	0.5224

Conclusion

The best-performing model was **XGBoost on the full dataset with an accuracy of 83.33%**. Further improvements could be explored using advanced ensemble techniques or deep learning approaches.

Section 3- Model Evaluation

Model Definition and Tuning

We used the XGBoost classifier with hyperparameter tuning via GridSearchCV. The best parameters for the full dataset (using `X_full`) were:

- **Learning Rate:** 0.1
- **Max Depth:** 2
- **Number of Estimators:** 200

After scaling the data and splitting into training and test sets, the model was fit on the training data and predictions were generated on the test set.

Model Performance Metrics

Classification Report

Class	Precision	Recall	F1-Score	Support
C1 (Wound Healing)	0.82	0.86	0.84	77
C2 (IFN-γ Dominant)	0.90	0.92	0.91	83
C3 (Inflammatory)	0.83	0.82	0.82	77
C4 (Lymphocyte Depleted)	0.80	0.79	0.80	92
C5 (Immunologically Quiet)	0.88	0.91	0.89	46
C6 (TGF-β Dominant)	0.65	0.56	0.60	27

Overall Model Performance

- **Accuracy:** 83%
 - **Total Samples:** 402
-

Model Performance Analysis

Overall Performance

- **Accuracy:** The overall model accuracy is **83.33%**, meaning that the classifier correctly labels the majority of samples.
 - **Macro and Weighted Averages:** With a **macro average of ~81%** and a **weighted average of 83%**, performance is reasonably balanced across classes.
-

Per-Class Performance & Implications

1. C1 (Wound Healing)

- **Metrics:** Precision = 0.82, Recall = 0.86, F1-Score = 0.84

- **Implications:**
High recall indicates that most true C1 cases are captured. The slightly lower precision shows that a few samples may be incorrectly predicted as C1.
 - *Reason:* C1 tumors are defined by distinct angiogenic and proliferation signatures, which aids their identification.

2. C2 (IFN- γ Dominant)

- **Metrics:** Precision = 0.90, Recall = 0.92, F1-Score = 0.91
- **Implications:**
With the highest scores among the subtypes, C2 tumors are classified exceptionally well.
 - *Reason:* Strong CD8+ T cell signals and robust M1 macrophage polarization provide a distinct signature, making them easier to classify.

3. C3 (Inflammatory)

- **Metrics:** Precision = 0.83, Recall = 0.82, F1-Score = 0.82
- **Implications:**
Balanced performance suggests that while the inflammatory markers (Th1/Th17) are distinct, there is some overlap with other classes.
 - *Reason:* Some cases may be misclassified as **C1 (Wound Healing)** if angiogenic signals are moderate.

4. C4 (Lymphocyte Depleted)

- **Metrics:** Precision = 0.80, Recall = 0.79, F1-Score = 0.80
- **Implications:**
Performance metrics indicate some misclassification, especially in borderline cases.
 - *Reason:* High M2 macrophage signal and suppressed Th1 response make it similar to **C6 (TGF- β Dominant)**.

5. C5 (Immunologically Quiet)

- **Metrics:** Precision = 0.88, Recall = 0.91, F1-Score = 0.89
- **Implications:**
With high precision and recall, C5 samples are well distinguished.
 - *Reason:* Low lymphocyte and high M2 macrophage features provide a clear signal, making these cases easier to classify.

6. C6 (TGF- β Dominant)

- **Metrics:** Precision = 0.65, Recall = 0.56, F1-Score = 0.60
- **Implications:**
C6 is the most challenging class. The low recall indicates that many true C6 samples are missed, and the low precision suggests a considerable number of false positives.
 - *Reason:*

- **Heterogeneity:** TGF- β Dominant tumors have variable expression and are not restricted to a single cancer type.
 - **Overlapping Features:** Some samples with moderate TGF- β signaling may also exhibit characteristics of **C4 (Lymphocyte Depleted)**, leading to misclassification.
 - **Small Sample Size:** With only 27 instances, statistical noise can significantly affect model performance.
-

Examples of Misclassification and Potential Reasons

1. C6 vs. C4

- **False Negatives:** A true C6 sample might be misclassified as **C4** if it exhibits a strong M2 macrophage signature and only moderate TGF- β expression.
- **False Positives:** Some samples predicted as C6 may not be true TGF- β Dominant cases if they show borderline TGF- β signals, resulting in lower precision.

2. C1 vs. C3 Overlap

- **Borderline Cases:** A sample with features of both **wound healing (C1)** and **inflammatory responses (C3)** might be misclassified.
 - Example: A C1 sample with less prominent angiogenic signals could be predicted as C3 if inflammatory markers are more pronounced.

3. C4 vs. C6 Misclassification

- **Subtle Differences:** The misclassification between these two subtypes emphasizes the subtle differences in their immune microenvironment.
 - **Example:** A sample with moderate TGF- β signaling might be more readily confused with C4 if the overall immune suppression is similar.
-

Summary

The XGBoost model achieved an overall accuracy of **83.33%** on the full dataset, with distinct performance across the six immune subtypes.

- The best performance was observed in **C2 (IFN- γ Dominant)** and **C5 (Immunologically Quiet)**, owing to their well-defined and robust immune signatures.
- In contrast, **C6 (TGF- β Dominant)** posed challenges due to:
 - Its heterogeneous nature

- Overlapping features with C4
- Limited sample size

Specific misclassifications—such as **C6 being confused with C4** and occasional overlaps between **C1 and C3**—highlight the complexity of the tumor immune landscape as described in **Thorsson et al. (2018)**.

References

- Thorsson et al., "The Immune Landscape of Cancer" (2018)