**Codes**

%%% main %%

clear;

clc;

for k=1:1

 k;

globalposition=FRMCLDA\_LOOCV\_DATA1();

overallauc1(k)=positiontooverallauc\_loocv\_DATA1();

%

% globalposition=FRMCLDA\_LOOCV\_DATA2();

% overallauc2(k)=positiontooverallauc\_loocv\_DATA2();

%

% globalposition=FRMCLDA\_LOOCV\_DATA3();

% overallauc3(k)=positiontooverallauc\_loocv\_DATA3();

end

save FRMCLDA\_LOOCV\_DATA1\_COSINE;

% a1=mean(overallauc1); % mean of auc values

% b1=std(overallauc1); % standard deviation of auc values

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

function [ result ] = cosSim( data )

%COSSIM Summary of this function goes here

% Detailed explanation goes here

rows=size(data,1);

result=zeros(rows,rows);

 for i=1:rows

 for j=1:i

 if (norm(data(i,:))\*norm(data(j,:))==0)

 result(i,j)=0;

 else

 result(i,j)=dot(data(i,:),data(j,:))/(norm(data(i,:))\*norm(data(j,:)));

 %%n=norm(A) % Returns the maximum singular value of A

 %%dot(a,b) % he dot product of vectors a and b

 end

 result(j,i)=result(i,j); %% symmetric matrix

 end

end

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

function globalposition=FRMCLDA\_LOOCV\_DATA1()

load data1

[m1,n2]=size(LD);

Adj=[LS,LD;LD',DS];

% color block matrix

% figure(1);

% image(Adj,'CDataMapping','scaled');

% colorbar;

interaction=LD;

save interaction1 interaction;

index=find(LD);

[pp,qq]=size(index);

A=zeros(pp,2);

for i=1:pp

[lncrna,disease]=ind2sub(size(LD),index(i));

A(i,1)=lncrna;

A(i,2)=disease;

end

save('knownlinks1.mat','A');

%implement leave-one-out cross validation

for cv=1:pp

 cv

 % obtain training sample

 load interaction1;

 interaction(A(cv,1),A(cv,2))=0;

% %%%%%%%%%% similarity computation %%%%%%%%%%%%%%%%%%%

 w1=0.7;

 w2=0.9;

 Slcos = cosSim(interaction);

 Sdcos = cosSim(interaction');

 LS=w1\*LS+(1-w1)\*Slcos;

 DS=w2\*DS+(1-w2)\*Sdcos;

%%%%%%%%% matrix completion %%%%%%%%%%%%%%%%%%%%%%%%%%

 Adj1=[LS,interaction;interaction',DS];

 SP=sparsity(Adj1)

 M = sparse(Adj1);

 t = cputime;

 [X] = fastSVT\_Q(M, 0.4, [0 1], 100, 10); %%tol=0.4, ran=[0,1], r\_reuse=100,q\_reuse=10

 t\_SVT = cputime - t;

 F=X(1:m1,(m1+1):(m1+n2)); % F is the matrix of predicted scores

finalscore=F(A(cv,1),A(cv,2));

for i=1:m1

 for j=1:n2

 if interaction(i,j)==1

 F(i,j)=-10000;

 end

 end

end

% obtain the position of tested disease-microbe interaction as variable globalposition(1,cv),

[ll1,mm1]=size(find(F>=finalscore));

[ll2,mm2]=size(find(F>finalscore));

globalposition(1,cv)=ll2+1+(ll1-ll2-1)/2;

end

 save('globalposition\_LOOCV\_DATA1\_COSINE.mat','globalposition');

end

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

 function overallauc=positiontooverallauc\_loocv\_DATA1()

load globalposition\_LOOCV\_DATA1\_COSINE.mat;

load interaction1;

[n,m]=size(interaction);

sID=importdata('knownlinks1.mat');

[pp,qq]=size(sID);

for i=1:pp

if globalposition(i)>m\*n-pp+1

globalposition(i)=m\*n-pp+1;

end

end

 pre\_y=[];

 rec\_y=[];

for k=1:m\*n-pp+1

 tp=0;

 for t=1:pp

 if globalposition(1,t)<=k

 tp=tp+1;

 end

 end

 tpr(1,k)=tp/pp;

 fp=k\*pp-tp;

 fpr(1,k)=fp/(pp\*(m\*n-pp));

 fn=pp-tp;

 Precision(1,k)=tp/(k\*pp);

 recall(1,k)=tp/(tp+fn);

 if mod(k,10)==0

 pre\_y(1,k)=Precision(1,k);

 rec\_y(1,k)=recall(1,k); %calculate the recall

 end

 %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

end

 Spe=1-fpr;

 Sen=recall;

 AUC=abs(trapz(1-Spe,Sen));

 AUPR=abs(trapz(Sen,Precision));

 overallauc=AUC;

% plot(fpr,tpr);

% str = ['AUC= ',num2str(AUC),'AUPR=',num2str(AUPR)];

% title('FMCLDA ROC curve');

% text(0.3,0.2,str);

% axis([-0.01 1.00 0 1.01]);

% xlabel('FPR');

% ylabel('TPR');

% figure;

% plot(recall,Precision);

% %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

% figure;

% bar(pre\_y,5);

% xlabel('top-k');

% ylabel('precision');

% title('FMCLDA precision-rank');

% axis([0 300 0 0.02]);

% set(gca,'XTickLabel',{'','','top-20','','top-40','','top-60','','top-80','','top-100'})

% figure;

% bar(rec\_y,5);

% xlabel('top-k')

% ylabel('recall')

% title('FMCLDA recall-rank');

% axis([0 300 0 0.5]);

% set(gca,'XTickLabel',{'','top-20','','top-40','','top-60','','top-80','','top-100'})

end