return membership_list(lines) if output_type == 'clusters': return clusters_list(lines) def membership_list(lines): #returns a list of membership (to a cluster) for the nodes (lines) membership_list=[] if isinstance(lines, str): lines = iter(lines.split('\n')) lines = iter([line.rstrip('\n') for line in lines]) while lines: try: l = next(lines) except: # EOF break if 1.lower().startswith("*vertices"): 1, nnodes = 1.split() for vertice in range(int(nnodes)): 1 = next(lines) membership_list.append(int(1)) else: break return membership_list def clusters_list(lines): #return a list of the clusters if isinstance(lines, str): lines = iter(lines.split('\n')) lines = iter([line.rstrip('\n') for line in lines]) while lines: try: l = next(lines) except: # EOF break if 1.lower().startswith("*vertices"): 1, nnodes = 1.split() communities = defaultdict(list) for vertice in range(int(nnodes)): 1 = next(lines) community = int(1)communities.setdefault(community, []).append(vertice) else: break return [set(v) for k, v in dict(communities).items()] In [54]: def clusters_to_membership(c): #converts a list of clusters into a membership list membership_list=[] clusters=[sorted(list(s)) for s in c] for i,cluster in enumerate(clusters): for node in range(len(cluster)): membership_list.append(i) return membership_list def write_clu(membership_list,name): #to store a membership list as a .clu file textfile = open('clu_files/'+name+'.clu', "w") textfile.write('*Vertices {} \n'.format(len(membership_list))) for element in membership_list: textfile.write(str(element) + "\n") textfile.close() In [3]: | #import and store all the networks nx_nets={} ig_nets={} clu ={} #membership clu_clusters = {} clu_names=[] nets_folder_path="A3-networks/" with os.scandir(nets_folder_path) as folders_list: for folder in folders_list: if folder.is_dir(): networks=os.listdir(nets_folder_path+folder.name) **for** net **in** networks: if net.endswith(".net"): file=nets_folder_path+folder.name+'/'+net $nx_nets[('{})'.format(net[:-4]))]=nx.Graph(nx.read_pa$ jek(file)) ig_nets[('{}'.format(net[:-4]))]=ig.read(file,format ="pajek").simplify() elif net.endswith(".clu"): clu_names.append(net[:-4]) file=nets_folder_path+folder.name+'/'+net clu[('{}'.format(net[:-4]))]= read_clu(file, 'member ship') clu_clusters[('{}'.format(net[:-4]))]= read_clu(file , 'clusters') In [104]: #keys_to_clu={'graph3+1+3':'graph3+1+3', '20x2+5x2':'20x2+5x2', 'graph4+ 4':'graph4+4', # 'star':'star', 'cat_cortex_sim':'cat_cortex_sim', # 'zachary_unwh':'zachary_unwh-real', 'dolphins':'dolphins-r eal', 'football':'football-conferences', '256_4_4_4_13_18_p':'256 _4_4_4_13_18_p', 'rb125':'rb125-1', '256_4_4_2_15_18_p':'256_4_4_2_15_18_p' #rb125 can be sent also to 'rb125-2' or 'rb125-3' clu_to_nets={'graph3+1+3': 'graph3+1+3', '20x2+5x2': '20x2+5x2', 'graph4+4': 'graph4+4', 'star': 'star', 'cat_cortex_sim': 'cat_cortex_sim', 'zachary_unwh-real': 'zachary_unwh', 'dolphins-real': 'dolphins', 'football-conferences': 'football', '256_4_4_4_13_18_p': '256_4_4_4_13_18_p', 'rb125-1': 'rb125', 'rb125-2': 'rb125', 'rb125-3': 'rb125', '256_4_4_2_15_18_p': '256_4_4_2_15_18_p'} Clustering algorithms In [250]: def comm_fastgreedy(name): f=deepcopy(ig_nets[name]) v=f.community_fastgreedy() clusters = v.as_clustering() return clusters def comm_optimal_modularity(name): f=deepcopy(ig_nets[name]) v=f.community_optimal_modularity() return v def comm_leading_eigenvector(name): f=deepcopy(ig_nets[name]) v=f.community_leading_eigenvector() return v def comm_label_propagation(name): f=deepcopy(ig_nets[name]) v=f.community_label_propagation() return v def plot_color_coded(name, membership, target=None): f=deepcopy(ig_nets[name]) pal=ig.drawing.colors.ClusterColoringPalette(len(membership)) f.vs['color']=pal.get_many(membership) if 'x' in ig_nets[name].vs.attributes(): ig.plot(f, bbox=(0, 0, 400, 200), vertex_size=5, edge_width= 0.5, target=target) else: ig.plot(f, bbox=(0, 0, 400, 200), layout='kamada_kawai', vertex_size=5, edge_width=0.5, target=target) else: ig.plot(f, bbox=(0, 0, 400, 200), layout='kamada_kawai', vertex_size=5, edge_width=0.5) In [276]: def fastgreedy(labels_true, name): labels_pred=comm_fastgreedy(name).membership write_clu(labels_pred, name+'-fastgreedy') return ['fastgreedy', jaccard_index(labels_true, labels_pred), ig.compare_communities(labels_true,labels_pred, method='nmi' normalized_vi(labels_true, labels_pred)] def optimal_modularity(labels_true, name): labels_pred=comm_optimal_modularity(name).membership write_clu(labels_pred, name+'-optimal_modularity') return ['optimal_modularity', jaccard_index(labels_true, labels_pred), ig.compare_communities(labels_true,labels_pred, method='nmi'), normalized_vi(labels_true, labels_pred)] def leading_eigenvector(labels_true, name): labels_pred=comm_leading_eigenvector(name).membership write_clu(labels_pred, name+'-lead_eig') return ['leading_eigenvector', jaccard_index(labels_true, labels_pred), ig.compare_communities(labels_true,labels_pred, method='nmi'), normalized_vi(labels_true, labels_pred)] def label_prop(labels_true, name): labels_pred= comm_label_propagation(name).membership write_clu(labels_pred, name+'-label_prop') return ['label_propagation', jaccard_index(labels_true, labels_pred), ig.compare_communities(labels_true,labels_pred, method='nmi'), normalized_vi(labels_true, labels_pred)] In [301]: def jaccard_index(C1,C2): #C1, C2 are membership lists of the clusters C=pair_confusion_matrix(C1,C2) den=(C[1,1]+C[1,0]+C[0,1])if den==0: return 'inf' index=C[1,1]/den return index from math import log def normalized_vi(C1, C2): #C1, C2 are membership lists of the clusters X=list(ig.Clustering(C1)) Y=list(ig.Clustering(C2)) n = float(sum([len(x) for x in X])) S = 0.0for x in X: p = len(x) / nfor y in Y: q = len(y) / nr = len(set(x) & set(y)) / n**if** r>0.0: S += r * (log(r / p, 2) + log(r / q, 2))return abs(S)/log(n) Comparison measures In [305]: | comparison={} for clu_name in clu_to_nets: name=clu_to_nets[clu_name] comparison[name] = pd.DataFrame(columns=('algorithm', 'Jaccard', 'NM I', 'NVI')) labels_true=clu[clu_name] comparison[name].loc[0]=fastgreedy(labels_true, name) comparison[name].loc[1]=leading_eigenvector(labels_true, name) comparison[name].loc[2]=label_prop(labels_true, name) #comparison[name].loc[3]=optimal_modularity(labels_true, name) In [306]: **for** name **in** comparison: print('') print(name) print(comparison[name]) dfi.export(comparison[name].style.set_caption(name),'comparison_meas ures/'+name+'.png') graph3+1+3 algorithm Jaccard NVI fastgreedy 0.666667 0.80954 0.238237 1 leading_eigenvector 0.666667 0.80954 0.238237 label_propagation 0.666667 0.80954 0.238237 20x2+5x2 algorithm NMI NVI Jaccard 0 fastgreedy 0.941176 0.938345 0.051124 leading_eigenvector 0.941176 0.938345 0.051124 label_propagation 1.000000 1.000000 0.000000 graph4+4 algorithm Jaccard NMI NVI fastgreedy 1.0 1.0 0.0 1 leading_eigenvector 1.0 1.0 0.0 label_propagation 1.0 1.0 0.0 star algorithm Jaccard NMI NVI 0 fastgreedy 1.0 1.00.0 leading_eigenvector 1.0 1.0 0.0 1 label_propagation 1.0 1.0 0.0 cat_cortex_sim algorithm Jaccard NMI fastgreedy 0.542169 0.656873 0.296602 leading_eigenvector 0.547872 0.618651 0.332598 label_propagation 0.257239 0.000000 0.481994 zachary_unwh algorithm Jaccard NMI NVI 0 fastgreedy 0.683274 0.692467 0.217697 leading_eigenvector 0.505495 0.677092 0.269804 label_propagation 0.462845 0.225967 0.351184 dolphins algorithm Jaccard NMI fastgreedy 0.504125 0.572700 0.271613 leading_eigenvector 0.329314 0.448914 0.423804 label_propagation 0.943044 0.888836 0.049311 football NMI NVI algorithm Jaccard fastgreedy 0.362153 0.697732 0.385871 leading_eigenvector 0.350324 0.698670 0.407185 label_propagation 0.545143 0.848038 0.210807 256_4_4_4_13_18_p Jaccard algorithm NMI NVI fastgreedy 1.000000 1.000000 0.000000 leading_eigenvector 1.000000 1.000000 0.000000 label_propagation 0.269841 0.680851 0.338132 rb125 algorithm NMI NVI Jaccard fastgreedy 0.281143 0.825443 0.287653 leading_eigenvector 0.031742 0.000000 0.967777 label_propagation 0.324100 0.887932 0.198671 256_4_4_2_15_18_p algorithm Jaccard fastgreedy 0.483871 0.869708 0.166303 leading_eigenvector 0.542431 0.924071 0.102676 label_propagation 1.000000 1.000000 0.000000 **Modularity** In [69]: modularity_nets = pd.DataFrame(columns=('fastgreedy', 'leading_eig', 'la bel_prop')) for name in ig_nets: modularity_nets.loc[name]=[comm_fastgreedy(name).modularity, comm_leading_eigenvector(name).modularity, comm_label_propagation(name).modularity] In [121]: modularity_nets Out[121]: fastgreedy leading_eig label_prop 0.367188 0.367188 graph3+1+3 0.367188 20x2+5x2 0.542579 0.542579 0.541586 0.423077 0.423077 0.423077 graph4+4 grid-p-6x6 0.401235 0.000000 0.277778 0.000000 0.000000 0.000000 star 0.255355 cat_cortex_sim 0.260436 0.000000 0.380671 0.393409 0.364809 zachary_unwh dolphins 0.495491 0.491199 0.392073 airports_UW 0.662490 0.639231 0.516750 football 0.549741 0.492606 0.540977 256_4_4_4_13_18_p 0.696773 0.664728 0.696773 0.752151 256_4_4_2_15_18_p 0.765660 0.781804 0.608733 0.000000 0.583748 rb125 In [111]: modularity_clu = pd.DataFrame(columns=('modularity',)) for clu_name in clu_to_nets: name=clu_to_nets[clu_name] vc=ig.VertexClustering(ig_nets[name], membership=clu[clu_name]) modularity_clu.loc[clu_name]=[vc.modularity] In [112]: modularity_clu Out[112]: modularity graph3+1+3 0.351562 0.541586 20x2+5x2 0.423077 graph4+4 0.000000 0.245996 cat_cortex_sim zachary_unwh-real 0.371466 0.373482 dolphins-real football-conferences 0.553973 256_4_4_4_13_18_p 0.696773 rb125-1 0.600595 rb125-2 0.558144 rb125-3 0.553147 256_4_4_2_15_18_p 0.781804 In [310]: dfi.export(modularity_nets.style.set_caption('partitions found'),'modula rity_nets.png') dfi.export(modularity_clu.style.set_caption('reference'), 'modularity_cl u.png') Color-coded plots In [190]: def plots_alg(name): plot_color_coded(name, comm_fastgreedy(name).membership, 'color_plot s/'+name+'-fastgreedy.png') plot_color_coded(name, comm_label_propagation(name).membership, 'col or_plots/'+name+'-label_prop.png') plot_color_coded(name, comm_leading_eigenvector(name).membership, 'c olor_plots/'+name+'-lead_eig.png') In [251]: #store the plots in color_plots directory for name in ig_nets: plots_alg(name) **NetworkX community detection** from networkx.algorithms.community import greedy_modularity_communities, k_clique_communities, label_propagation_communities def greedy_mod_nx(labels_true, name): c = greedy_modularity_communities(nx_nets[name]) labels_pred=clusters_to_membership(c) write_clu(labels_pred, 'nx_'+name+'-greedy_mod') return ['greedy_modularity', jaccard_index(labels_true, labels_pred), metrics.normalized_mutual_info_score(labels_true, labels_pre d), normalized_vi(labels_true, labels_pred)] #def k_clique_nx(labels_true, name, k): c=k_clique_communities(nx_nets[name], k) labels_pred=clusters_to_membership(c) write_clu(labels_pred, 'nx_'+name+'-k_clique') return ['k-clique', jaccard_index(labels_true, labels_pred), metrics.normalized_mutual_info_score(labels_true, labels_pr ed), normalized_vi(labels_true, labels_pred)] def label_prop_nx(labels_true, name): c=label_propagation_communities(nx_nets[name]) labels_pred=clusters_to_membership(c) write_clu(labels_pred, 'nx_'+name+'-label_prop') return ['label_propagation', jaccard_index(labels_true, labels_pred), metrics.normalized_mutual_info_score(labels_true, labels_pre d), normalized_vi(labels_true, labels_pred)] In [307]: comparison_nx={} for clu_name in clu_to_nets: name=clu_to_nets[clu_name] comparison_nx[name] = pd.DataFrame(columns=('algorithm', 'Jaccard', 'NMI', 'NVI')) labels_true=clu[clu_name] comparison_nx[name].loc[0]=greedy_mod_nx(labels_true, name) #comparison_nx[name].loc[1]=k_clique_nx(labels_true, name, k) comparison_nx[name].loc[2]=label_prop_nx(labels_true, name) In [308]: for name in comparison_nx: print('') print(name) print(comparison_nx[name]) dfi.export(comparison_nx[name].style.set_caption(name), 'comparison_m easures/nx_'+name+'.png') graph3+1+3 algorithm Jaccard NMI NVI greedy_modularity 0.666667 0.80954 0.238237 2 label_propagation 0.666667 0.80954 0.238237 20x2+5x2 NVI Jaccard NMI algorithm 0 greedy_modularity 0.941176 0.938345 0.051124 2 label_propagation 1.000000 1.000000 graph4+4 algorithm Jaccard NMI NVI greedy_modularity 1.0 1.0 2 label_propagation 1.0 1.0 0.0 star algorithm Jaccard NMI greedy_modularity 1.0 1.0 0.0 2 label_propagation 1.0 1.0 0.0 cat_cortex_sim Jaccard algorithm NMI NVI 0.257299 greedy_modularity 0.571930 0.702341 label_propagation 0.257239 0.000000 0.481994 zachary_unwh algorithm Jaccard NMI NVI greedy_modularity 0.379009 0.289122 0.503219 label_propagation 0.403361 0.254134 0.494754 dolphins NVI algorithm Jaccard NMI 0 greedy_modularity 0.245068 0.010911 0.628715

2 label_propagation 0.203363 0.041141 0.717362

Jaccard

greedy_modularity 0.058675 0.147975 1.086858 2 label_propagation 0.044610 0.238707 1.113253

Jaccard 0 greedy_modularity 1.000000 1.0000 0.000000 2 label_propagation 0.413333 0.6993 0.263123

Jaccard

Jaccard

greedy_modularity 0.461653 0.845852 0.196748 2 label_propagation 0.245902 0.773341 0.266577

2 label_propagation 0.159132 0.754033 0.420127

NMI

NMI

NMI

NMI

NVI

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NVI

NVI

0.287653

algorithm

algorithm

algorithm

algorithm

0 greedy_modularity 0.281143 0.825443

football

rb125

256_4_4_4_13_18_p

256_4_4_2_15_18_p

COMMUNITY DETECTION

from networkx.algorithms import community

from collections import defaultdict

import networkx.algorithms.community as nx_comm

from sklearn.metrics.cluster import contingency_matrix, pair_confusion_m

import matplotlib.pyplot as plt from matplotlib import gridspec

import dataframe_image as dfi from copy import deepcopy

from sklearn import metrics

In [170]: import pandas as pd

import os

import igraph as ig

import networkx as nx

%matplotlib inline

Useful functions

In [2]: #functions to read the files given

def read_clu(path, output_type): with open(path, 'r') as f: lines = f.readlines()

if output_type == 'membership':