Analysis: Network of Comorbidity in US Veterans

for the course: Complessità nei sistemi e nelle reti

The Paper

997 138 individuals
US Military Veterans, on average 41 years old, 86% male

95 conditions 17 psychiatric, 78 medical, earliest diagnosis 7 August 2010

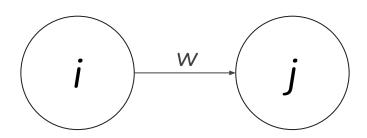
4 main contributions:
relationship between psychiatric and medical conditions
model the temporal aspect of comorbidity
clinical vulnerability score
association between the degree of comorbidity and mortality

The Data

95x95 matrix increased log odds of co-morbidity of condition *i* given condition *j*

95x95 matrix p-values from logistic regression

A curious notation



an arc from i to j represents the influence of currently having condition j on the future development of condition i

if I have condition j my odds of developing condition i increase by w

strong predictors have high in-degree

Data Preparation

start from 8930 (= 95x94) nonzero values

remove 2236 values that are NOT statistically significant (p > 0.05)

remove 2915 negative values they correspond to odds increase between 0 and 1 (negative correlation)

3779 nonzero values left 42.32%

```
# remove values that are not statistically significative
mask = pmat <= 0.05
```

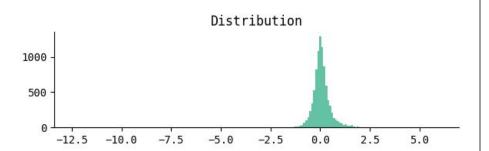
vmat = vmat[mask].fillna(0)

remove negative values

vmat = vmat[vmat > 0].fillna(0)

Before

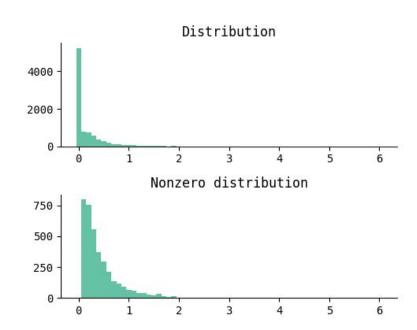
min = -12.36 max = 6.03nonzero = 8930 (= 95x94)

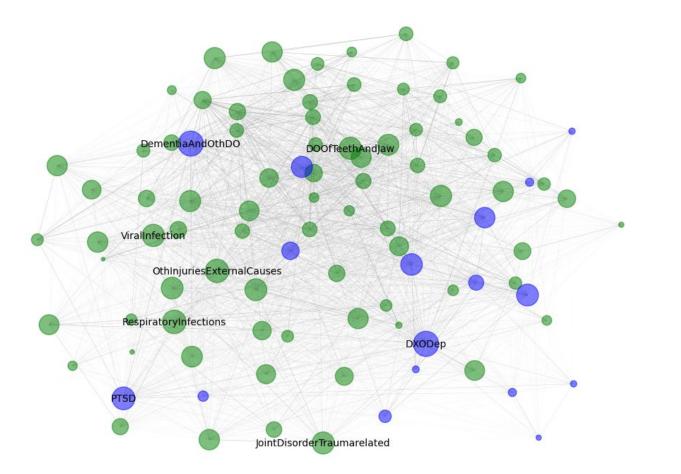


After

$$min = 0.00$$

 $max = 6.03$
 $nonzero = 3779$





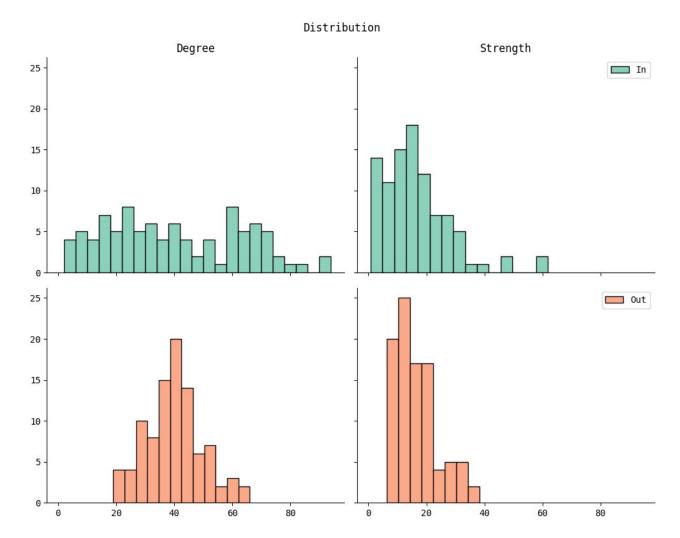
Network Properties

```
directed = True
weighted = True
nodes = 95
edges = 3779
```

clustering = 0.59 diameter = 3 average shortest path = 1.61 assortativity = -0.09

property	min	max	mean
weight	0.02	6.03	0.42
in-degree	2	94	39.78
out-degree	19	66	39.78
in-strength	0.82	61.85	16.60
out-strength	6.26	38.37	16.60

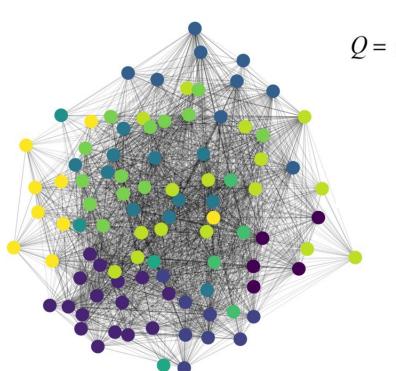
Complementary cumulative distribution Strength Degree 10^{θ} - In - Out 10-1 10-2 10¹ 10² 10⁰ 10¹ 1.0 0.8 0.6 -0.4 -0.2-0.0 20 40 80 20 60 10 30 0 50 60



centrality	min	max	mean	top-3
in-degree	0.02	1.00	0.42	'DXODep' 'DementiaAndOthDO' 'RespiratoryInfections'
out-degree	0.20	0.70	0.42	'DisOfTheUrinarySystem' 'DisOfArteries' 'DisOfTheHeart'
eigenvector	0.00	0.19	0.09	'DXODep' 'DementiaAndOthDO' 'OthInjuriesExternalCauses'
katz	-0.16	0.31	0.01	'PancreaticDO' 'FluidAndElectrolyteDO' 'NoninfectiousGastroenteritO'
in-closeness	0.36	1.00	0.64	'DXODep' 'DementiaAndOthDO' 'RespiratoryInfections'
out-closeness	0.53	0.76	0.62	'DisOfTheUrinarySystem' 'DisOfArteries' 'DisOfTheHeart'
betweenness	0.00	0.02	0.01	'DXDRG' 'OthInjuriesExternalCauses' 'PersonalityDO'

Mesoscale Analysis

Louvain Community Detection Algorithm: 11 communities found.



 $Q = (fraction \ of \ links \ internal \ to \ communities) - (expected \ fraction \ of \ such \ links)$

$$= \frac{1}{2L} \sum_{C_{i}} \sum_{i,j \in C_{i}} \left[a_{ij} - \frac{k_{i}k_{j}}{2L} \right]$$

Modularity: 0.24

Community 1 (5 nodes): {'COPDAndBronchiectasis_DT', 'OthLowerRespiratoryDs_DT', 'OthUpperRespiratoryDs_DT', 'RespiratoryInfections_DT', 'Asthma_DT'}
Community 2 (16 nodes): {'Anemia_DT',

{'OthGIDO_DT', 'cancerdxDT', 'AbdominalHernia DT'. 'UpperGIDO_DT', 'GIHemorrhage_DT', 'LowerGIDO DT', 'DisOfVeins DT', 'NoninfectiousGastroenterit0'} **Community 4** (9 nodes): {'SprainsAndStrains_DT', 'OthBoneDs_DT','EarConditions_DT', 'OthConnectiveTissueDs DT'. 'JointDisorderTraumarelated_DT', 'anomdxDT','AcquiredDeformities_DT', 'Spondylosis_DT', 'NontraumaticJointDO_DT'} Community 5 (11 nodes): ('SuperficialInjury_DT', 'CrushingInjuryOrInternalInO', 'DOOfTeethAndJaw DT'. 'Fractures_DT', 'DisOfMouth_DT', 'OthInjuriesExternalCauses'', 'Burns_DT', 'OpenWounds_DT', 'SkinInfections_DT', 'EyeDO_DT', 'InfectiveArthritis_DT'}

Community 3 (8 nodes):

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Community 6 (2 nodes):
{'PathologicalFracture_D',
'Osteoporosis'}
Community 7 (2 nodes):
{'OthInfections', 'ImmunityDO'}
Community 8 (4 nodes):
{'Mycoses_DT', 'OthSkinDO_DT', 'DXALC_DT', 'ANXgen_DT',
'ViralInfection DT'.
'OthInflamCondOfSkin'}
Community 9 (13 nodes):
{'IntracranialInjury_DT',
'Paralysis_DT',
'ComaBrainDamage_DT',
'OthNervousSystemDO_DT',
'Headache_DT',
'NSConditions_DT',
'EpilepsyConvulsions_DT',
'CerebrovascularDs_DT',
'DevelopmentalDO_DT',
'CNSInfection_DT',
'SpinalCordInjury_DT',
'ChronicUlcerOfSkin DT'.
'DementiaAndOthDO_DT'}
```

'ImpulseControlDONEC_DT', 'SCHZ_DT', 'AdjustmentDO DT'. 'AFBPDX_DT', 'DXDRG_DT', 'PersonalityDO_DT', 'PTSD_DT', 'DXODep_DT', 'AttentionDeficitDO DT'. 'DOInChildhood_DT', 'SuicideAndSelfInjury_DT', 'Poisoning_DT'} Community 11 (9 nodes): {'GoutAndOth DT'. 'OthNutritionalEndocrineDO_D',

'DOOfLipidMetabolism_DT',

'OthEndocrineDO DT'.

'ThyroidDO_DT', 'genitaldxDT',

'dmdxDT', 'Hypertension_DT',

'NutritionalDeficiencies_DT'}

Community 10 (16 nodes):

{'ANXunsp_DT', 'MDD_DT',

What do communities represent?

Community_1 : Respiratory diseases

Community_2 : HEM/CIRC

Community_3: gastrointestinal

Community_4: MSK

Community_5: Mixed Injury and Disorders

Community_6 : Bone diseases

Community_7: Infections

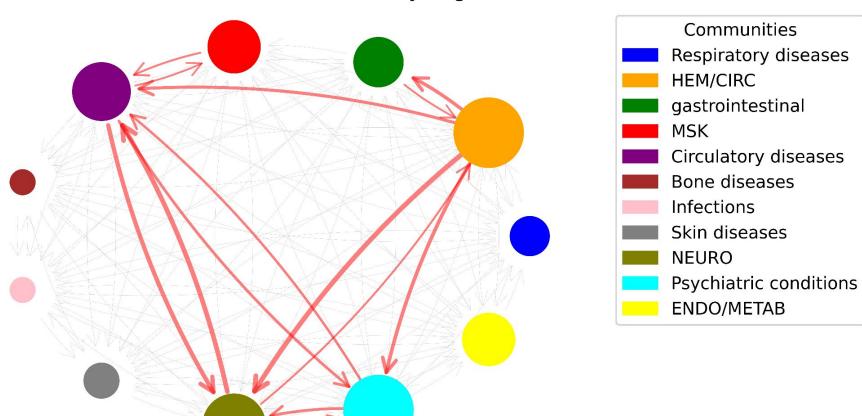
Community_8: Skin diseases

Community_9: NEURO

Community_10 : Psychiatric conditions

Community_11 : ENDO/METAB

Communities and inter-community edges



Who are the predictors?

Weighted edges
entering Community_10:

Weighted edges entering Community_5:

Weighted edges entering Community_9:

Community_7: 3.94 Community_8: 8.56 Community_1: 8.83 Community_4: 9.67

Community_6: 2.21

Community_4: 9.67 Community_3: 12.10 Community_11: 16.29

Community_5: 32.72 Community_9: 34.86 Community_2: 41.86

In-strengths for Community_10: 171.03

Community_11: 3.05 Community_7: 5.95 Community_1: 7.18 Community_3: 7.86 Community_6: 8.50

Community_8: 10.01 Community_4: 23.06 Community_10: 28.87 Community_2: 35.76

Community_9: 54.14

In-strengths for Community_5: 184.37

Community_7: 2.76 Community_8: 3.24

Community_1: 3.64 Community_11: 7.66

Community_3: 9.72 Community_4: 12.28 Community_6: 13.60

Community_10: 32.01 Community_5: 49.64 Community_2: 55.58

In-strengths for Community_9:

190.12

Reference and possible improvements

the paper http://dx.doi.org/10.1098/rspa.2019.0790

the data https://github.com/aaronab/comorbidity_networks/

our code https://github.com/CristinaLiccia/co-morbidity-network-analysis