Exam Answers

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Matched Answers

Question 1: Short answer/objective question.

(a) Convert incidence matrix to adjacency matrix:

Incidence Matrix to Adjacency Matrix

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The incidence matrix provided is: \[
\begin{bmatrix}
1 & 0 & 0 \\
1 & 1 & 1 \\
0 & 1 & 0 \\
0 & 0 & 1 \\
\end{bmatrix}
\]
```

The matrix structure has been given the name. Each edge in an incidence matrix corresponds to a row and every node has its designated column. Each entry of value '1' shows that the node has an incidence with the edge. A simple undirected graph contains exactly two adjacent nodes for every single edge within

From the incidence matrix:

- -> Edge 1 connects Node 1 and Node 2
- -> Edge 2 connects Node 2 and Node 3
- -> Edge 3 connects Node 2 and Node 4

The connection relationship between nodes in a graph structure

in adjacency matrix format. The adjacency matrix contains value 1 in position (i, j) and (j, i) when node i and node j having an edge, otherwise both positions contain 0.

The adjacency matrix will represent node to node connection.

| begin{bmatrix}
0 & 1 & 0 & 0 \\
1 & 0 & 1 & 1 \\
0 & 1 & 0 & 0 \\
0 & 1 & 0 & 0 \\
tend{bmatrix}

| end{bmatrix}

(b) Network model for uniform probability:

Ans -> Option-(B) -> Erd&s{Rényi (Random Network) Model.

In the Erdøs{Rényi model the edges connect randomly because every edge formation remains isolated from other edges.

(c) Game theory situation:

Ans -> Option-(C) -> Nash Equilibrium

Players reach a stable position known as Nash Equilibrium when none of them achieve higher rewards through independent strategy modifications.

(d) Tendency for individuals to associate:

Answer -> Option-(B) Assortative Mixing

Under the concept of Assortative Mixing, similar nodes established connects with each other.

(e) Betweenness centrality relevance:

Answer -> Option-(D) Because it quantifies how often a node lie shortest paths between other nodes.

The measurement technique called Betweenness Centrality evaluates nodes based on their capacity to link other nodes because it determines information flow efficiencies.

(f) Scale-free networks property:

Answer -> Option-(C) The presence of many nodes with very high degrees (hubs) that maintain connectivity.

Scale-free Network (Hubs) operate through central nodes which serve as crucial connections. So their attack vulnerability becomes particularly pronounced.

(g) Community detection for high modularity:

Answer -> Option-(A) The number of intra-community edges is significantly higher than expected in a random network with the same degree sequence.

The goal of Community detection (Modularity) is to identify communities through dense connection within those groups compared to random chance expectation.

(h) Jaccard Coefficient for link prediction:

Answer -> Option-(B) 2/5

Jaccard Coefficient,

Neighbors of X: A, B, C, D Neighbors of Y: C, D, B

Intersection: C, D (2 nodes)
Union: A, B, C, D, E (5 nodes)

Jaccard Coefficient = Intersection / Union = 2/5 = 0.4

The Jaccard's Coefficient calculates set (neighbors) similarity dividing the overlapping intersection by their combined element

(i) ICM vs LTM activation mechanism:

Answer -> Option-(B) -> LTM uses edge probabilities independent

LTM uses a weighted sum of active neighbors compared to a node

ICM operates through independent edge probability, but LTM adopts a threshold-based method which takes the weighted sum of neighbors' impact.

(j) GCN message-passing approach:

Answer -> Option-(B) -> Because aggregating features from dissi neighbors can blur the node's own representational features, ma classification harder.

When heterophily occurs during GCN and heterophily operations it becomes difficult to classify nodes because they lose their individual characteristics through the features aggregation process from dissimilar neighbors.

Question 2: Novel influenza strain strategy:

Influenza Spread and Vaccination

For reducing influenza transmission to 5% level a combined approach of centrality measurement should be used to determine who gets vaccinated first.

A high value of betweenness centrality reveals that the node functions as a "bridge" through which the disease transmits to numerous other connections. The immunization of these specific people breaks transmission pathways between different segments of the network.

High degree centrality helps identify "hubs" which possess numerous connections since they have the potential to infect multiple other people.

A computation between betweenness and degree centrality should be performed for every individual:

- -> A list of ranked people forms based on the centrality measurement result.
- -> The group of people selected for vaccination needs a strong ranking in both centrality measures.
- -> Additional individuals should be included for vaccination until 5% coverage is reached if the top selected candidates do not suffice.

Justification ->

A combined approach delivers better results since betweenness centrality identifies network bridges which link different network sections while degree centrality distinguishes hubs that spread widely.

Vaccinating key hubs and bridges provides maximum efficiency by controlling both small outbreaks and widespread distribution of the virus.

Question 3: Improving 'suggested collaborators' feature:

Suggested Collaborators Feature

Link Prediction -> System utilizes linking prediction algorithms that generate potential collaboration prediction through the analysis of current co-authorship and citation data.

When researchers A and B demonstrate numerous joint authors link with each other scientist a link prediction method could indicate potential collaboration between them.

Through Node2Vec the system creates vector representations of researchers based on network structural information, embedding that align with each other between researchers demonstrates similar research interest and collaboration potential.

Homophily ->

Due to the natural human technology of homophily the system will provide recommendations related to researchers within similar academic fields.

Due to encouraging diverse collaboration the system requires a diversity metric within its recommendation functionality. Although the link prediction score may be slightly lower the system will give priority to research links that combine fields from distinct areas.

Question 4: Girvan-Newman algorithm:

(a) Core idea:

Girvan-Newman Core Idea

The Girvan-Newman algorithm detects communities through a recursive method which successively eliminates edges connecting different communities.

(b) Edge betweenness centrality:

Edge Betweenness Centrality ->

Edge Betweenness Centrality analyzes the number of times an edge lies between all pairs of nodes to discover these edges.

Edges with high betweenness values link separate communities according to this method.

(c) Major computational limitation:

Computational Limitation

The cost of computation increases substantially when calculating edge betweenness centrality on extensive networks especially when they involve frequent edge removal operation.

(d) Louvain method explanation:

Louvain Method

The Louvain method functions as a scalable approach which uses greedy optimization to shift nodes between communities during its iterative process.

Question 5: PageRank algorithm:

(a) Intuition behind PageRank:

PageRank Algorithm

According to PageRank the importance of nodes depends on both the quantity and quality of incoming links passing through them.

Page importance derives from other pages that link to it.

(b) Role of 'damping factor':

Damping Factor (d)

During random web navigation a surfer has a probability explored through the Damping Factor (d) either to pause their link clicks or to move to an arbitrary page.

(c) Problem with dangling nodes:

Dangling Nodes

The lack of outgoing links from Dangling Nodes makes PageRank flow through the network. We handle this condition using an equal distribution among all network nodes.

Each iteration distributes the rank of dangling

nodes across all network nodes in an equal way.

Question 6: Game theory payoff matrix:

(a) Pure Strategy Nash Equilibria:

A strategy pair where players can improve their payoff unilaterally changing their strategy.

Check each pair:

```
(U, A): Payoff (3, 2)
```

- \rightarrow Player 1: Switch to L \rightarrow 2 < 3, no improve
- -> Player 2: Switch to B -> 1 < 2, no improve
- -> Nash Equilibrium.

(U, B): Payoff (0, 1)

- \rightarrow Player 1: Switch L \rightarrow 2 > 0, improve
- -> Not Nash

(L, B): Payoff (2, 3)

- \rightarrow Player 1: Switch to U \rightarrow 0 < 2, no improve
- -> Player 2: Switch to A -> 0 < 3, no improvement
- -> Nash Equilibrium.

There are two pure strategy Nash equilibria: (U, A) & (L, B).

(b) Expected payoff calculation:

Player 1 plays U with probability p, L with probability 1-p.

Player 2's strategy:

Expected payoff for Player 2 if choosing Strategy A
\[
E[A] = p \times 2 + (1-p) \times 0 = 2p
\]

Expected payoff for Player 2 if choosing Strategy B $\[\]$

(c) Expected outcome for p = 0.77:

For Player 2:

Since $\langle (E[B] > E[A] \rangle$, Player 2 could choose Strategy B.

For Player 1: (0.7 U, 0.3 L)Expected payoff = $0.7 \text{ \times } 0 + 0.3 \text{ \times } 2$ = 0.6 when Player 2 chooses B

The expected outcome for Player 1 receives a payoff of 0.6 if Player 2 receives a payoff of 1.6.

Question 7: Graph Neural Network feature update:

Graph Neural Network Feature Update

Given,

Directed edge A -> B C -> B

D -> B

Weighted matrix W =
\[
\begin{bmatrix}

0.5 & 0 \\ 0.1 & 0.2 \\

```
\end{bmatrix}
\]
Step 1: Aggregate neighbor features.
] \
h_A^{(0)} + h_C^{(0)} + h_D^{(0)} =
\begin{bmatrix}
1 \\
\end{bmatrix}
\begin{bmatrix}
3 \\
\end{bmatrix}
\begin{bmatrix}
3 \\
\end{bmatrix}
\]
h_{N(B)}^{(0)} = \frac{1}{3}
\begin{bmatrix}
7 \\
6 \\
\end{bmatrix}
\begin{bmatrix}
1 \\
2 \\
\end{bmatrix}
\]
Step 2: Transform
] \
\begin{bmatrix}
1 \\
2 \\
\end{bmatrix}
```

```
\begin{bmatrix}
0.5 & 0 \\
0.1 & 0.2 \\
\end{bmatrix}
\begin{bmatrix}
1 \\
2 \\
\end{bmatrix}
/]
1/
\begin{bmatrix}
0.5 \\
0.1 \times 1 + 0.2 \times 2 \\
\end{bmatrix}
\]
] \
\begin{bmatrix}
0.5 \\
0.1 + 0.4 \\
\end{bmatrix}
\]
1/
\begin{bmatrix}
0.5 \\
0.5 \\
\end{bmatrix}
\]
Step 3: Activate
ReLU
\begin{bmatrix}
0.5, 0.5 \\
\end{bmatrix}
```

```
begin{bmatrix}

0.5, 0.5 \\
\end{bmatrix}

Final Answer \(h_B^{(1)}\) =
\begin{bmatrix}

0.5, 0.5 \\
\end{bmatrix}
```