

Algorithm: Spatial-channel collaborative multi-scale graph interaction deep transfer learning

Input:

- Source domains:  $S = \{S_1, S_2, \dots, S_M\}$
- Target domain:  $T$
- Batch size:  $N_m$
- Number of categories:  $K$
- Number of scales:  $H$
- Spatial-channel refinement parameters

Output:

- Classification predictions for target domain samples

1: // ===== 3.1: Spatial-Channel Collaborative Prototype Extraction Module =====

- 2: for each batch  $(X_{src}, y_{src})$  from  $S$  and  $(X_{tgt}, y_{tgt})$  from  $T$  do
- 3: // Spatial refinement
- 4:  $X_{out} = \beta + \frac{\psi(X - \mu)}{\sqrt{\sigma^2 + \varepsilon}}$  // Eq. (1)
- 5:  $\gamma$  = learnable\_parameter in  $GN$
- 6:  $W_\gamma = \{w_i\} = \gamma_i / \left( \sum_{j=1}^D \gamma_j \right), \quad i, j = 1, 2, \dots, D$  // Eq. (2)
- 7:  $W = \text{Gate}(\text{Sigmoid}(W_\gamma))$
- 8:  $W_1, W_2 = \text{threshold\_gate}(W)$
- 10:  $X^{wl} = X_{11}^w \oplus X_{22}^w$  // Eq. (3)
- 11:  $X^{w2} = X_{21}^w \oplus X_{12}^w$  // Eq. (4)
- 12:  $X^w = \text{Concat}(X^{wl}, X^{w2})$  // Refined spatial feature
- 13: // Channel refinement
- 14:  $(X_{up}, X_{low}) = \text{Split}(X^w, \text{ratio}=\alpha)$  //  $\alpha$  is splitting ratio
- 15:  $Y_1 = X_{up} \left( \lambda^{GWC} + \lambda^{PWC_1} \right)$  // Eq. (5)
- 16:  $Y_2 = \lambda^{PWC_1} X_{low} \cup X_{low}$  // Eq. (6)
- 17:  $O_1, O_2 = \text{GlobalAveragePool}(Y_1), \text{GlobalAveragePool}(Y_2)$
- 18:  $I_1 = \exp(O_1) / (\exp(O_1) + \exp(O_2))$  // Eq. (7)
- 19:  $I_2 = \exp(O_2) / (\exp(O_1) + \exp(O_2))$
- 20:  $Y = I_1 * Y_1 + I_2 * Y_2$  // Eq. (8) channel refined feature
- 21: // Extract prototypes for source domains
- 22: for each source domain  $m = 1$  to  $M$  do

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23:   for each category  $k = 1$  to  $K$  do
24:      $\hat{c}_k^{S_m} = \text{mean}(f_S(x_i))$  for all  $x_i$  in category  $k$  // Eq. (9)
25:   end for
26: end for

27: // ===== Temporarily skip the target domain prototype update and proceed with graph construction and propagation =====
28: // Use the current target domain prototype (obtained from the previous batch update or initial value)
29: // If it is the first iteration, initialize to the source domain mean
30: if first_batch then
31:   for  $k = 1$  to  $K$  do
32:      $\hat{c}_k^T = \text{mean}(\hat{c}_k^{S_m} \text{ for } m=1,2,\dots,M)$ 
33:   end for
34: end if

35: // ===== 3.2: Multi-Scale Graph Interaction Transfer Network =====
36: // Step 1: Build multi-domain knowledge graph
37:  $\mathbf{F} = \begin{bmatrix} \underbrace{c_1^{S_1} c_2^{S_1} \cdots c_K^{S_1}}_{\text{prototypes of } S_1} \cdots \underbrace{c_1^{S_M} c_2^{S_M} \cdots c_K^{S_M}}_{\text{prototypes of } S_M} & \underbrace{c_1^T c_2^T \cdots c_K^T}_{\text{prototypes of } T} \end{bmatrix}^T$ 
38: for  $i,j$  in 1 to  $(M+1)*K$  do
39:    $\mathbf{A}_{i,j} = G(\mathbf{F}_i^T, \mathbf{F}_j^T)$  // Eq. (10)
40: end for

41: // Step 2: Build extended graph with query samples
42: for each query sample  $q$  in batch  $N_m$  do
43:    $\bar{\mathbf{F}} = [\mathbf{F}^T f(q_1) f(q_2) \cdots f(q_{|N_m|})]^T$  // Eq. (11)
44:    $\mathbf{S}_{i,j} = G(\mathbf{F}_i^T, f(q_j))$  // Eq. (12)
45:    $\bar{\mathbf{A}} = \begin{bmatrix} \mathbf{A} & \mathbf{S} \\ \mathbf{S}^T & \mathbf{I} \end{bmatrix}$  // Eq. (13)
46: end for

47: // Step 3: Multi-scale graph interaction
48:  $\mathbf{X\_list} = [\mathbf{X}^{(0)}, \mathbf{X}^{(1)}, \mathbf{X}^{(h)}, \dots, \mathbf{X}^{(H)}]$ 
49: for  $h = 1$  to  $H$  do
50:    $\mathbf{AX}^{(h-1)} \rightarrow \mathbf{X}^{(h)}$ ,  $\forall h=1, 2, \dots, H$  // Eq. (14)
51:    $\mathbf{X\_list.append}(\mathbf{X}^{(h)})$ 

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52: end for

53: // Dynamic interaction flow construction
54: for each node  $i = 1$  to  $|\bar{V}|$  do
55:    $\mathbf{Z}_i = \xi(\mathbf{X}_i^{(1)} \parallel \mathbf{X}_i^{(2)} \parallel \dots \parallel \mathbf{X}_i^{(H)})$            // Eq. (15)
56:   for each propagation step  $l$  do
57:      $\mathbf{X}_i^{(l)} = \mathbf{X}_i^{(l)} \parallel \mathbf{Z}_i$                                 // Eq. (16)
58:      $\tilde{w}_i(l) = \text{Sig}(\mathbf{X}_i^{(l)} \cdot s)$                          // Eq. (17)
59:   end for
60:   // Normalize over scales
61:   for  $l = 1$  to  $H$  do
62:      $w_i(l) = \frac{e^{\tilde{w}_i(l)}}{\sum_{h=0}^H e^{\tilde{w}_i(h)}}$                 // Eq. (18)
63:   end for
64: end for

65: // Build weight matrix and fuse features
66: for  $h = 1$  to  $H$  do
67:    $\eta_h[i] = w_i(h) \quad 1 \leq i \leq n \quad \text{for all nodes } i$           // Eq. (19)
68:    $\mathbf{W}_h = \text{Diag}(\eta_h)$                                          // Eq. (20)
69: end for
70:  $\mathbf{U}_x = \sum_{h=0}^H \mathbf{W}_h \mathbf{X}^{(h)}$                            // Eq. (21)

71: // Step 4: Cross-domain diagnosis
72:  $\mathbf{P} = f_H(\bar{\mathbf{F}}, \bar{\mathbf{A}})$                                // Eq. (22) - classification probabilities
73: // Extract prototypes for target domain using confidence-thresholded pseudo-labeling
74: Now proceed with updating the target domain prototype (based on graph network output)
75: Extracting the classification probability of target samples from  $\mathbf{P}$ 
76: HC = empty_set // High confidence sample set
77: for  $j = 1$  to N_target_samples do
78:   // Node index of the target sample in the extended graph
79:   node_idx =  $|F| + j$ 
80:    $p_t = \mathbf{P}[\text{node\_idx}]$  // The classification probability of the target sample

81: // Calculate entropy value
82: entropy_t =  $-\sum_{k=1}^K p_t[k] * \log(p_t[k])$ 

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83:
84: if entropy_t < threshold_entropy then
85:   pseudo_label = argmax( $p_i$ )
86:   // Store sample features and pseudo labels
87:   HC.add
88: end if
89: end for

90: // Update target domain prototype
91: for each category  $k = 1$  to  $K$  do
92:   // Collect all high confidence sample features for this category
93:   features_k = { $z \mid (z, \text{label})$  in HC and label =  $k$ }
94:
95: if features_k not empty then
96:   // Update the target domain prototype using high confidence samples
97:    $\hat{c}_k^T = \text{mean}(\text{features}_k)$ 

98: else
99:   // If there are no high confidence samples, maintain the original value or use the source domain mean
100:  //  $\hat{c}_k^T$  remain unchanged

101: end if
102: end for

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103: // ===== 3.3: Category Constraint Loss =====

104: // Global constraint loss

$$105: L_R^g = \frac{1}{(M+1)^4} \sum_{m,n,i,j=1}^{M+1} \| \mathbf{A}_{i,j} - \mathbf{A}_{m,n} \|_F \quad // \text{Eq. (23)}$$

106: // Local constraint loss

$$107: L_R^l = \frac{1}{|N_m|} \sum_{k=1}^K \left( \sum_{(x_i^T, y_i^T) \in T_k} \| f(x_i^T) - c_k^T \|_2^2 + \sum_{m=1}^M \sum_{(x_i^{S_m}, y_i^{S_m}) \in \mathcal{S}_m^k} \| f(x_i^{S_m}) - c_k^{S_m} \|_2^2 \right) \quad // \text{Eq. (24)}$$

108: // Classification losses

$$109: L_C^s = \frac{\sum_{m=1}^M \left( \mathbb{E}_{(x_i^{S_m}, y_i^{S_m}) \in \mathcal{S}_m} L_{ce}(p(x_i^{S_m}), y_i^{S_m}) \right)}{M} \quad // \text{Eq. (25)}$$

$$110: L_C^p = \frac{\sum_{k=1}^K L_{ce}(p(c_k^T), k) + \sum_{m=1}^M \sum_{k=1}^K L_{ce}(p(c_k^{S_m}), k)}{K(M+1)} \quad // \text{Eq. (26)}$$

$$111: L_C^t = -\mathbb{E}_{(x_i^T, \hat{y}_i^T) \in T} \sum_{k=1}^K p(\hat{y}_i^T = k | x_i^T) \log p(\hat{y}_i^T = k | x_i^T) \quad // \text{Eq. (27)}$$

112: // Total loss

$$113: L_{CCL} = L_C + L_R, \quad // \text{Eq. (28)}$$

114: // Optimization

115: Update  $f_S, f_H, \lambda, s, \gamma$ , etc. via backpropagation

116: // Inference on target domain

117: for each target sample  $x_t$  in  $T$  do

118:    Compute extended graph with  $x_t$

$$119: p_t = f_H(\bar{\mathbf{F}}, \bar{\mathbf{A}})$$

$$120: y_{\text{pred}} = \text{argmax}(p_t)$$

121: end for

122: return  $y_{\text{pred}}$  for all target samples