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import numpy as np
import matplotlib.pyplot as plt
from scipy.optimize import golden

# -----
# Problem definition (1D test)
# -----
np.random.seed(42)

Omega = (0.0, 1.0)
Gamma = 0.25
Omega1 = (0.0, Gamma)
Omega2 = (Gamma, 1.0)

def beta(x):
    x = np.array(x)
    return np.where(x < Gamma, 1.0, 10.0)

def u_exact(x):
    x = np.array(x)
    return np.where(x <= Gamma, np.sin(2*np.pi*x), np.cos(2*np.pi*x))

def du_exact(x):
    x = np.array(x)
    return np.where(x <= Gamma, 2*np.pi*np.cos(2*np.pi*x),
-2*np.pi*np.sin(2*np.pi*x))

# f(x) = -(beta u')' in subdomains where beta constant -> -beta * u''.

def f_rhs(x):
    x = np.array(x)
    left = -1.0 * (1.0) * (-2*np.pi)**2 * np.sin(2*np.pi*x) # =
(2π)^2 sin
    right = -1.0 * (10.0) * (-2*np.pi)**2 * np.cos(2*np.pi*x) # =
10*(2π)^2 cos

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    return np.where(x < Gamma, left, right)

# interface jumps h1 and h2
def h1_val():
    return u_exact(Gamma - 1e-12) - u_exact(Gamma + 1e-12)

def h2_val():
    #  $\beta_{\text{left}} * u'_{\text{left}} - \beta_{\text{right}} * u'_{\text{right}}$ 
    du_left = 2*np.pi*np.cos(2*np.pi*Gamma)
    du_right = -2*np.pi*np.sin(2*np.pi*Gamma)
    return 1.0 * du_left - 10.0 * du_right

# boundary g
def g_val(x):
    return u_exact(x)

# -----
# Multi-TransNet parameters
# -----
# covering ball centers and radii for each subdomain
x_c1, R1 = 0.125, 0.135
x_c2, R2 = 0.625, 0.385

# neurons per subdomain
M1, M2 = 5, 5

# activation
def sigma(z):
    return np.tanh(z)

# hidden-layer randomized params
a1 = np.random.choice([-1.0, 1.0], size=M1)
a2 = np.random.choice([-1.0, 1.0], size=M2)
r1 = np.random.uniform(0.0, R1, size=M1)
r2 = np.random.uniform(0.0, R2, size=M2)

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# -----
# Basis and operator builders
# -----

def build_basis(X, x_c, a, r, gamma, M):
    X = np.atleast_1d(X)
    N = len(X)
    B = np.ones((N, M+1))
    for m in range(M):
        z = gamma * (a[m] * (X - x_c) + r[m])
        B[:, m+1] = sigma(z)
    return B

def build_L_basis(X, x_c, a, r, gamma, M, beta_func):
    """
    Build matrix for L(phi) = - (beta phi')' approximated by -beta * phi'' (inside subdomains).
    Use centered finite difference for phi'' with small h.
    """
    X = np.atleast_1d(X)
    N = len(X)
    B = np.zeros((N, M+1))
    h = 1e-5
    for m in range(M):
        # evaluate phi at x-h, x, x+h vectorized
        z_c = gamma * (a[m] * (X - x_c) + r[m])
        z_p = gamma * (a[m] * (X + h - x_c) + r[m])
        z_m = gamma * (a[m] * (X - h - x_c) + r[m])
        phi_c = sigma(z_c)
        phi_p = sigma(z_p)
        phi_m = sigma(z_m)
        d2phi = (phi_p - 2.0*phi_c + phi_m) / (h*h)
        B[:, m+1] = - beta_func(X) * d2phi
    # constant column stays zero (second derivative of const = 0)
    return B

def build_J_basis(X, x_c, a, r, gamma, M, beta_func):

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"""
Build matrix for flux J(phi) = beta * phi' (normal in 1D).
Use centered finite difference for phi' with small h.
"""

X = np.atleast_1d(X)
N = len(X)
B = np.zeros((N, M+1))
h = 1e-6
for m in range(M):
    z_p = gamma * (a[m] * (X + h - x_c) + r[m])
    z_m = gamma * (a[m] * (X - h - x_c) + r[m])
    phi_p = sigma(z_p)
    phi_m = sigma(z_m)
    dphi = (phi_p - phi_m) / (2.0*h)
    B[:, m+1] = beta_func(X) * dphi
return B

# -----
# Sampling configuration (modified per your request)
# -----
# interior collocation points used in compute_loss (large sampling)
N_points = 10000
X1 = np.linspace(Omega1[0] + 0.01, Omega1[1] - 0.01, N_points//2) # 500
X2 = np.linspace(Omega2[0] + 0.01, Omega2[1] - 0.01, N_points//2) # 500

# boundary: 100 at x=0 and 100 at x=1 -> total 200 rows
n_bnd_each = 100
Xg_left = np.array([0.0] * n_bnd_each)
Xg_right = np.array([1.0] * n_bnd_each)

# interface: 100 points total, choose 50 left + 50 right near Gamma
n_if_total = 100
n_if_each_side = n_if_total // 2
eps = 1e-3

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# distribute points moving away from Gamma: small offsets
left_offsets = np.linspace(eps/n_if_each_side, eps, n_if_each_side)
right_offsets = np.linspace(eps/n_if_each_side, -eps, n_if_each_side)
Xr_left = Gamma - left_offsets
Xr_right = Gamma + right_offsets
Xr = np.concatenate([Xr_left, Xr_right]) # length 100

# -----
# Loss / assembly for given gammas
# -----

def compute_loss(g1):
    # predict g2 from g1 using the relation used in your code pattern
    g2 = g1 * (R1 / R2) * (M2 / M1)

    # build basis and operator matrices
    Phi1 = build_basis(X1, x_c1, a1, r1, g1, M1)
    Phi2 = build_basis(X2, x_c2, a2, r2, g2, M2)
    L1 = build_L_basis(X1, x_c1, a1, r1, g1, M1, beta)
    L2 = build_L_basis(X2, x_c2, a2, r2, g2, M2, beta)

    # boundary basis (many repeated rows)
    Phi_g1 = build_basis(Xg_left, x_c1, a1, r1, g1, M1) # shape
    (100, M1+1)
    Phi_g2 = build_basis(Xg_right, x_c2, a2, r2, g2, M2) # shape
    (100, M2+1)

    # interface bases and flux
    Phi_r1 = build_basis(Xr, x_c1, a1, r1, g1, M1) # (100, M1+1)
    Phi_r2 = build_basis(Xr, x_c2, a2, r2, g2, M2) # (100, M2+1)
    J_r1 = build_J_basis(Xr, x_c1, a1, r1, g1, M1, beta)
    J_r2 = build_J_basis(Xr, x_c2, a2, r2, g2, M2, beta)

    # count rows
    n1 = len(X1)
    n2 = len(X2)
    nb = len(Phi_g1) + len(Phi_g2) # 200

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ni = len(Xr)                                # 100 (each produces 2 equations
=> 200 rows)
total_rows = n1 + n2 + nb + 2*ni
total_cols = M1 + M2 + 2 # alpha1..alpha_M1, alpha0_1,
alpha2..alpha_M2, alpha0_2

F = np.zeros((total_rows, total_cols))
T = np.zeros(total_rows)

# PDE residuals Omega1
F[0:n1, :M1+1] = L1
T[0:n1] = f_rhs(X1)

# PDE residuals Omega2
F[n1:n1+n2, M1+1:] = L2
T[n1:n1+n2] = f_rhs(X2)

# boundary: left (x=0) -> fill rows
idx_b_left_start = n1 + n2
idx_b_left_end = idx_b_left_start + len(Phi_g1)
F[idx_b_left_start:idx_b_left_end, :M1+1] = Phi_g1
T[idx_b_left_start:idx_b_left_end] = g_val(0.0)

# boundary: right (x=1)
idx_b_right_start = idx_b_left_end
idx_b_right_end = idx_b_right_start + len(Phi_g2)
F[idx_b_right_start:idx_b_right_end, M1+1:] = Phi_g2
T[idx_b_right_start:idx_b_right_end] = g_val(1.0)

# interface rows start
idx_if_start = idx_b_right_end

# for each interface sampling point, add [u] equation: phi_left -
phi_right = h1
for i in range(ni):
    row = idx_if_start + i

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# left phi (M1+1), right phi (M2+1)
F[row, :M1+1] = Phi_r1[i]
F[row, M1+1:] = -Phi_r2[i]
T[row] = h1_val()

# for each interface sampling point, add flux jump equation:
beta_left phi'_left - beta_right phi'_right = h2
for i in range(ni):
    row = idx_if_start + ni + i
    F[row, :M1+1] = J_r1[i]
    F[row, M1+1:] = -J_r2[i]
    T[row] = h2_val()

# solve least squares and return residual norm
try:
    alpha, residuals, rank, s = np.linalg.lstsq(F, T, rcond=None)
    if len(residuals) > 0:
        return residuals[0]
    else:
        return np.linalg.norm(F @ alpha - T)
except Exception as e:
    # fallback large penalty if solver fails
    return 1e12

# -----
# Golden-section search for gamma1
# -----
print("Starting golden-section search for gamma1 (this may take some time)...")
gamma1_opt = golden(lambda g: compute_loss(g), brack=(0.1, 10.0),
                    tol=1e-3, full_output=False)
gamma2_opt = gamma1_opt * (R1 / R2) * (M2 / M1)
print(f"Found gamma1_opt = {gamma1_opt:.6e}, gamma2_opt = {gamma2_opt:.6e}")

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# Final solve using gamma_opt with denser final collocation (kept
moderate)
# -----
# final collocation inside each subdomain (smaller for speed but can
be increased)
N_final = 1000
X1_final = np.linspace(Omega1[0] + 0.001, Omega1[1] - 0.001,
N_final//2)
X2_final = np.linspace(Omega2[0] + 0.001, Omega2[1] - 0.001,
N_final//2)

# final boundary: still 100 each
Phi_g1_final = build_basis(Xg_left, x_c1, a1, r1, gamma1_opt, M1)
Phi_g2_final = build_basis(Xg_right, x_c2, a2, r2, gamma2_opt, M2)

# final interface points (same Xr used)
Phi_r1_final = build_basis(Xr, x_c1, a1, r1, gamma1_opt, M1)
Phi_r2_final = build_basis(Xr, x_c2, a2, r2, gamma2_opt, M2)
J_r1_final = build_J_basis(Xr, x_c1, a1, r1, gamma1_opt, M1, beta)
J_r2_final = build_J_basis(Xr, x_c2, a2, r2, gamma2_opt, M2, beta)

L1_final = build_L_basis(X1_final, x_c1, a1, r1, gamma1_opt, M1,
beta)
L2_final = build_L_basis(X2_final, x_c2, a2, r2, gamma2_opt, M2,
beta)

# assemble final F and T
n1f = len(X1_final)
n2f = len(X2_final)
nbff = len(Phi_g1_final) + len(Phi_g2_final) # 200
nif = len(Xr) # 100
rows_final = n1f + n2f + nbff + 2*nif
cols_final = M1 + M2 + 2

Ff = np.zeros((rows_final, cols_final))
Tf = np.zeros(rows_final)

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# PDE
Ff[0:n1f, :M1+1] = L1_final
Tf[0:n1f] = f_rhs(X1_final)
Ff[n1f:n1f+n2f, M1+1:] = L2_final
Tf[n1f:n1f+n2f] = f_rhs(X2_final)

# boundary
b_left_s, b_right_s = n1f + n2f, n1f + n2f + len(Phi_g1_final)
Ff[b_left_s:b_right_s, :M1+1] = Phi_g1_final
Tf[b_left_s:b_right_s] = g_val(0.0)
Ff[b_right_s:b_right_s+len(Phi_g2_final), M1+1:] = Phi_g2_final
Tf[b_right_s:b_right_s+len(Phi_g2_final)] = g_val(1.0)

# interface rows
start_if = b_right_s + len(Phi_g2_final)
for i in range(nif):
    Ff[start_if + i, :M1+1] = Phi_r1_final[i]
    Ff[start_if + i, M1+1:] = -Phi_r2_final[i]
    Tf[start_if + i] = h1_val()
for i in range(nif):
    Ff[start_if + nif + i, :M1+1] = J_r1_final[i]
    Ff[start_if + nif + i, M1+1:] = -J_r2_final[i]
    Tf[start_if + nif + i] = h2_val()

# solve final least squares
alpha_f, residuals_f, rank_f, s_f = np.linalg.lstsq(Ff, Tf,
rcond=None)
print("Final LS residual (sum squares):", residuals_f if
len(residuals_f)>0 else np.linalg.norm(Ff @ alpha_f - Tf)**2)

# split coefficients
alpha1 = alpha_f[:M1+1]
alpha2 = alpha_f[M1+1:]

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# Evaluate the multi-transnet solution on a dense test grid
# -----
x_test = np.linspace(0.001, 0.999, 1000)
u_pred = np.zeros_like(x_test)
for i, xx in enumerate(x_test):
    if xx <= Gamma:
        B = build_basis([xx], x_c1, a1, r1, gamma1_opt, M1) # shape
        (1, M1+1)
        u_pred[i] = B.dot(alpha1)
    else:
        B = build_basis([xx], x_c2, a2, r2, gamma2_opt, M2)
        u_pred[i] = B.dot(alpha2)

u_true = u_exact(x_test)

# -----
# Plot results
# -----
plt.figure(figsize=(12, 8))

plt.subplot(2,1,1)
plt.plot(x_test, u_true, 'b-', linewidth=2, label='Exact Solution')
plt.plot(x_test, u_pred, 'r--', linewidth=1.5, label='Multi-TransNet
Prediction')
plt.axvline(Gamma, color='k', linestyle='--', alpha=0.7,
label='Interface')
plt.xlabel('x'); plt.ylabel('u(x)')
plt.title('Solution Comparison')
plt.legend(); plt.grid(alpha=0.3)

plt.subplot(2,1,2)
err = np.abs(u_pred - u_true)
plt.semilogy(x_test, err, 'g-', linewidth=1.2)
plt.axvline(Gamma, color='k', linestyle='--', alpha=0.7)
plt.xlabel('x'); plt.ylabel('Absolute Error (log scale)')
plt.title('Absolute Error (log scale)')

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plt.grid(alpha=0.3)

plt.tight_layout()
plt.show()

# -----
# Compute relative errors
# -----
rel_l2 = np.linalg.norm(u_pred - u_true) / np.linalg.norm(u_true)
rel_inf = np.max(np.abs(u_pred - u_true)) / np.max(np.abs(u_true))
print(f"Relative L2 error: {rel_l2:.6e}")
print(f"Relative L_inf error: {rel_inf:.6e}")
```