

# Noisy Circle: End-to-End gflow Workflow

## Overview

This vignette demonstrates a full `gflow` workflow on a noisy circle:

1. Build and inspect ikNN graphs across a `k` range.
2. Select a graph scale using graph diagnostics.
3. Denoise coordinates with `data.smoothen()`.
4. Smooth a noisy response `y` (mixture of two circular Gaussian peaks).
5. Smooth a feature matrix `Z`.
6. Find refined basins/extrema from `y.hat`.
7. Compare Pearson/Spearman and local correlation (`lcor`) analyses.
8. Add permutation-based inference for `lcor` (implemented here in the vignette).
9. Build a feature module graph from local-correlation profiles.

## Helper Functions

```
wrap_angle <- function(theta, mu) {
  atan2(sin(theta - mu), cos(theta - mu))
}

`%||%` <- function(x, y) {
  if (!is.null(x)) x else y
}

as_lcor_matrix <- function(x) {
  if (is.list(x) && !is.null(x$column.coefficients)) {
    return(as.matrix(x$column.coefficients))
  }
  as.matrix(x)
}

assign_from_vertices_list <- function(vertices.list, n) {
  out <- rep(NA_character_, n)
  if (is.null(vertices.list) || length(vertices.list) == 0) {
    return(out)
  }
  for (nm in names(vertices.list)) {
    out[as.integer(vertices.list[[nm]])] <- nm
  }
  out
}

compute_basins_safe <- function(adj.list, edge.length.list, fitted.values, verbose = TRUE) {
  has_both_extrema <- function(obj) {
    if (is.null(obj$summary) || nrow(obj$summary) == 0L || is.null(obj$summary$type)) {
      return(FALSE)
    }
  }
}
```

```

tab <- table(obj$summary$type)
("max" %in% names(tab)) && ("min" %in% names(tab)) &&
  (tab[["max"]] > OL) && (tab[["min"]] > OL)
}

strict <- try(
  compute.refined.basins(
    adj.list = adj.list,
    edge.length.list = edge.length.list,
    fitted.values = fitted.values,
    edge.length.quantile.thld = 0.75,
    min.rel.value.max = 1.1,
    max.rel.value.min = 0.9,
    max.overlap.threshold = 0.15,
    min.overlap.threshold = 0.15,
    p.mean.nbrs.dist.threshold = 0.9,
    p.mean.hopk.dist.threshold = 0.9,
    p.deg.threshold = 0.9,
    min.basin.size = 10,
    expand.basins = TRUE,
    with.trajectories = TRUE,
    verbose = verbose
  ),
  silent = TRUE
)

if (!inherits(strict, "try-error") && has_both_extrema(strict)) {
  return(strict)
}

message("Strict basin settings failed (or removed all maxima/minima); retrying with relaxed settings.")
compute.refined.basins(
  adj.list = adj.list,
  edge.length.list = edge.length.list,
  fitted.values = fitted.values,
  edge.length.quantile.thld = 0.9,
  min.rel.value.max = 1.02,
  max.rel.value.min = 0.98,
  max.overlap.threshold = 0.25,
  min.overlap.threshold = 0.25,
  p.mean.nbrs.dist.threshold = 0.98,
  p.mean.hopk.dist.threshold = 0.98,
  p.deg.threshold = 0.98,
  min.basin.size = 5,
  apply.geometric.filter = FALSE,
  expand.basins = TRUE,
  with.trajectories = TRUE,
  verbose = verbose
)
}

perm_test_lcor <- function(adj.list,
                           edge.length.list,

```

```

        y,
        Z,
        type = "derivative",
        y.diff.type = "difference",
        z.diff.type = "difference",
        n.perm = 200L,
        seed = 1L) {

set.seed(seed)
Z <- as.matrix(Z)
y <- as.numeric(y)

lcor.obs <- as_lcor_matrix(
  lcor(
    adj.list = adj.list,
    weight.list = edge.length.list,
    y = y,
    z = Z,
    type = type,
    y.diff.type = y.diff.type,
    z.diff.type = z.diff.type
  )
)
stat.obs <- colMeans(abs(lcor.obs))

stat.perm <- matrix(NA_real_, nrow = n.perm, ncol = ncol(Z))
colnames(stat.perm) <- colnames(Z)

for (b in seq_len(n.perm)) {
  Zb <- Z[sample.int(nrow(Z)), , drop = FALSE]
  lcor.b <- as_lcor_matrix(
    lcor(
      adj.list = adj.list,
      weight.list = edge.length.list,
      y = y,
      z = Zb,
      type = type,
      y.diff.type = y.diff.type,
      z.diff.type = z.diff.type
    )
  )
  stat.perm[b, ] <- colMeans(abs(lcor.b))
}

p.value <- (1 + colSums(t(t(stat.perm) >= stat.obs))) / (n.perm + 1)
q.value <- stats::p.adjust(p.value, method = "BH")

list(
  stat.obs = stat.obs,
  stat.perm = stat.perm,
  p.value = p.value,
  q.value = q.value,
  table = data.frame(
    feature = colnames(Z),
    stat.obs = stat.obs,

```

```

    p.value = p.value,
    q.value = q.value,
    row.names = NULL
)
)
}

```

## Simulate Noisy Circle and Response

```

n <- 250L
radius <- 1

X.df <- generate.circle.data(
  n = n,
  radius = radius,
  noise = 0.15,
  type = "random",
  noise.type = "normal",
  seed = 11
)
X <- as.matrix(X.df[, c("x", "y")])

if ("angles" %in% colnames(X.df)) {
  theta.obs <- as.numeric(X.df$angles)
} else {
  theta.obs <- atan2(X[, 2], X[, 1])
  theta.obs <- ifelse(theta.obs < 0, theta.obs + 2 * pi, theta.obs)
}

## Two wrapped Gaussian peaks over the circle: one larger, one smaller
mu1 <- 0.80
mu2 <- 3.95
sd1 <- 0.30
sd2 <- 0.48
amp1 <- 1.40
amp2 <- 0.75

comp1 <- amp1 * exp(-0.5 * (wrap_angle(theta.obs, mu1) / sd1)^2)
comp2 <- amp2 * exp(-0.5 * (wrap_angle(theta.obs, mu2) / sd2)^2)
y.true <- comp1 + comp2
y <- y.true + rnorm(n, sd = 0.20)

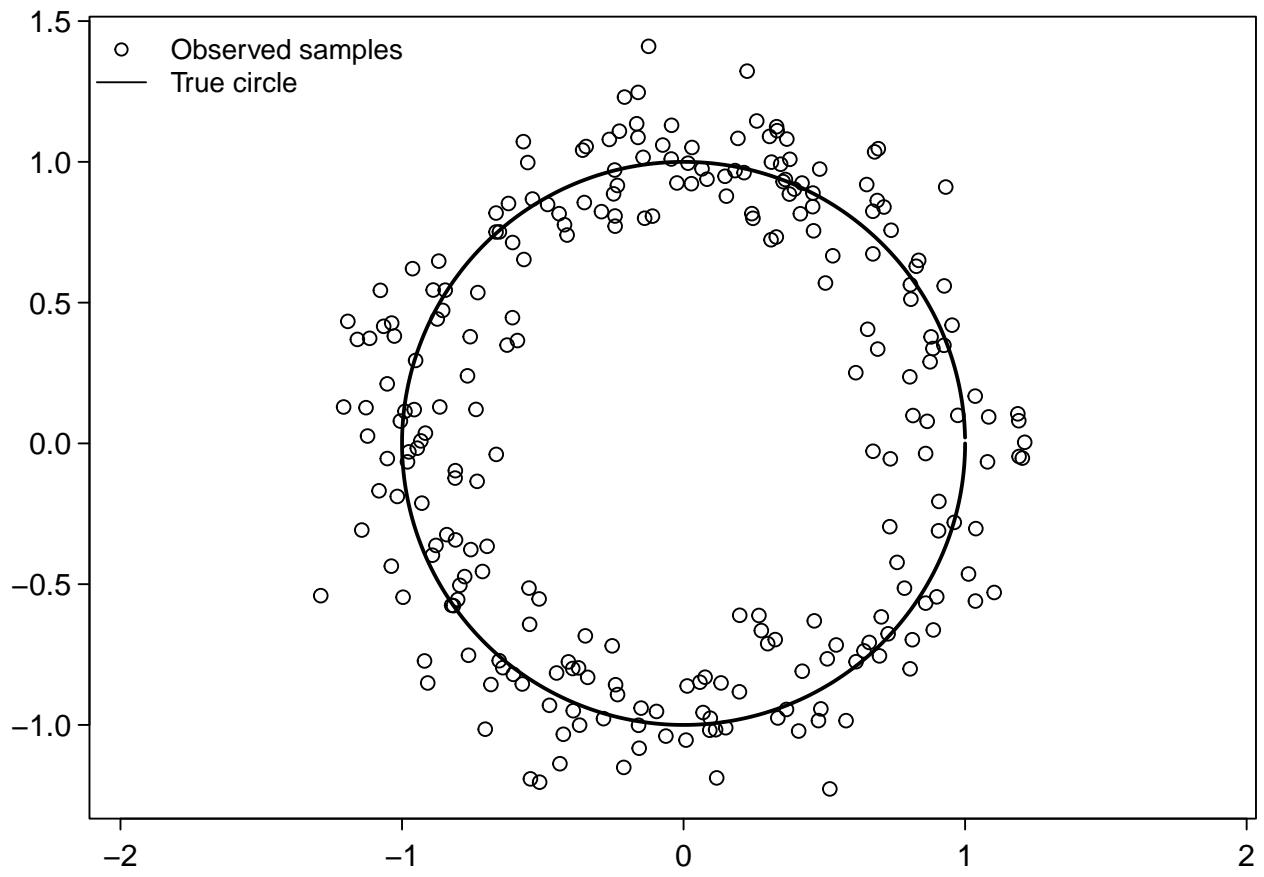
circle.true.df <- generate.circle.data(
  n = 300,
  radius = radius,
  noise = 0,
  type = "uniform",
  noise.type = "normal",
  seed = 1
)
circle.true <- as.matrix(circle.true.df[, c("x", "y")])

```

```

op <- par(mar = c(2.5, 2.5, 0.5, 0.5), mgp = c(2.5, 0.5, 0), tcl = -0.3)
plot(X, asp = 1, las = 1, xlab = "", ylab = "")
lines(circle.true, lwd = 2)
legend("topleft",
       legend = c("Observed samples", "True circle"),
       pch = c(1, NA),
       lty = c(NA, 1),
       col = c("black", "black"),
       bty = "n", cex = 0.9)

```



```
par(op)
```

## Build ikNN Graphs Across k

```

k.min <- 5L
k.max <- 12L

X.graphs <- create.iknn.graphs(
  X,
  kmin = k.min,
  kmax = k.max,
  max.path.edge.ratio.deviation.thld = 0.1,
  n.cores = 1L,
  verbose = TRUE
)
#> Using 1 OpenMP thread

```

```

#> Processing k values from 5 to 12 for 250 vertices
#> Requested parallel.mode: auto
#> Parallel mode: serial
#> Starting graph processing
#> [compute_knn] running...[compute_knn] 0.001s
#> [graph_build/geometric_prune/isize_prune] running...[graph_build/geometric_prune/isize_prune] 0.361s
#> Graph processing completed (0.361)
#> [compute_knn] 0.001s
#> [graph_build] 0.013s
#> [geometric_prune] 0.348s
#> [isize_prune] skipped (with.isize.pruning=FALSE)
#> Creating return list objects ... DONE (0.000)
#> Total elapsed time (0.362)
summary(X.graphs)
#> Summary of iknn_graphs object
#> -----
#> Number of vertices: 250
#> k range: 5 to 12
#> Max path-edge ratio deviation threshold: 0.1
#> Path-edge ratio percentile: 0.5
#> Graph type: geometrically pruned
#>
#>   idx  k n_ccomp edges mean_degree min_degree max_degree sparsity
#>   1   5      1   877    7.02        2       15  0.97182
#>   2   6      1  1056    8.45        3       17  0.96607
#>   3   7      1  1262   10.10        3       22  0.95945
#>   4   8      1  1437   11.50        3       23  0.95383
#>   5   9      1  1613   12.90        3       29  0.94818
#>   6  10      1  1766   14.13        3       29  0.94326
#>   7  11      1  1927   15.42        3       32  0.93809
#>   8  12      1  2047   16.38        3       33  0.93423

```

## Alternative k Selection Utility

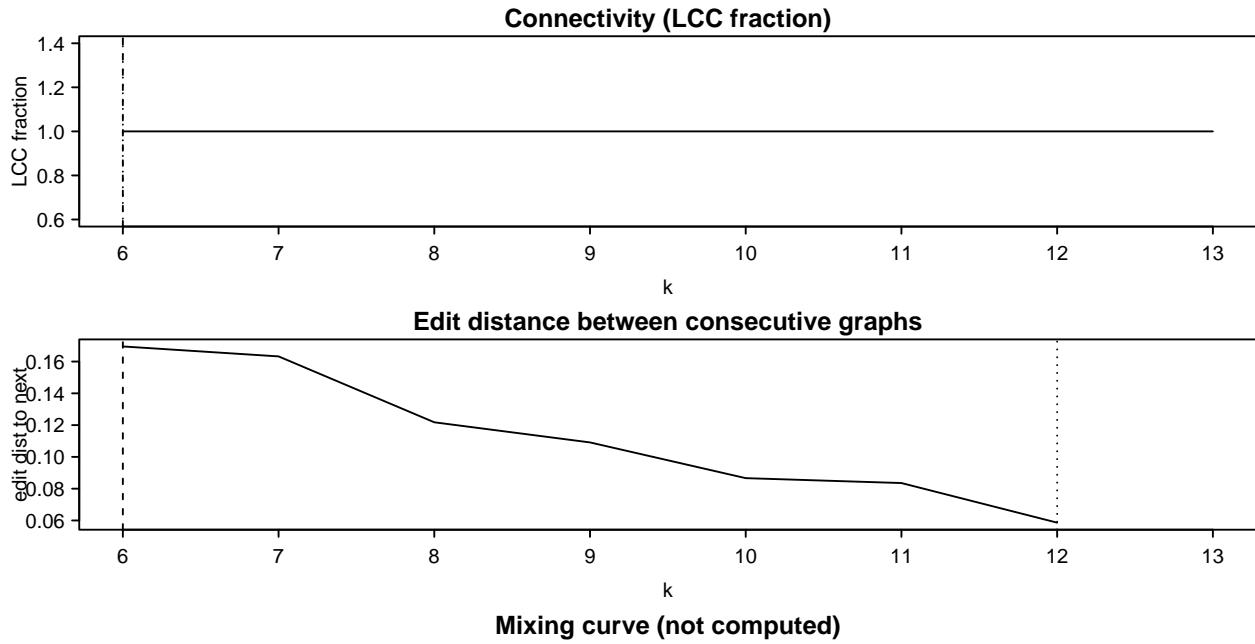
```

X.graphs2 <- build.iknn.graphs.and.selectk(
  X,
  kmin = k.min,
  kmax = k.max,
  method = "edit",
  n.cores = 1L,
  verbose = TRUE
)
#> Using 1 OpenMP thread
#> Processing k values from 5 to 12 for 250 vertices
#> Requested parallel.mode: auto
#> Parallel mode: serial
#> Starting graph processing
#> [compute_knn] running...[compute_knn] 0.001s
#> [graph_build/geometric_prune/isize_prune] running...[graph_build/geometric_prune/isize_prune] 0.371s
#> Graph processing completed (0.371)
#> [compute_knn] 0.001s
#> [graph_build] 0.013s
#> [geometric_prune] 0.357s

```

```
#> [isize_prune] skipped (with.isize.pruning=FALSE)
#> Creating return list objects ... DONE (0.000)
#> Total elapsed time (0.371)
```

```
X.graphs2$k.opt.edit
#> [1] 12
plot(X.graphs2)
```



## Visualize One Graph Layout

```
k.values <- if (!is.null(colnames(X.graphs$k_statistics)) &&
               "k" %in% colnames(X.graphs$k_statistics)) {
  as.integer(X.graphs$k_statistics[, "k"])
} else {
  seq_len(length(X.graphs$geom_pruned_graphs))
}

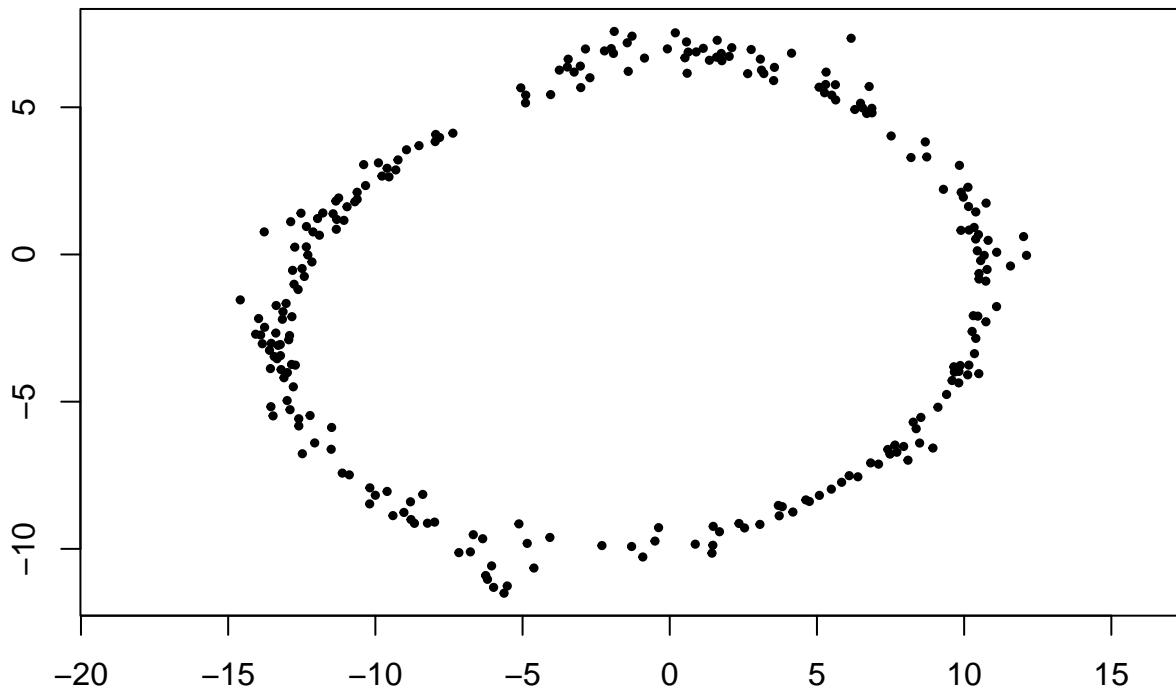
sel.k <- as.integer(X.graphs2$k.opt.edit %||% k.values[1])
k.idx <- which(k.values == sel.k)[1]
if (is.na(k.idx)) k.idx <- 1L

g.sel <- X.graphs$geom_pruned_graphs[[k.idx]]
layout.2d <- graph.embedding(
  adj.list = g.sel$adj_list,
  weights.list = g.sel$weight_list,
  invert.weights = TRUE,
  dim = 2,
  method = "fr"
)

plot(layout.2d, pch = 19, cex = 0.5, asp = 1,
     main = sprintf("2D graph embedding for selected k = %d", sel.k),
```

```
xlab = "", ylab = "")
```

## 2D graph embedding for selected k = 12



Denoise X with `data.smoother()`

```
X.denoise.res <- data.smoother(  
  X,  
  kmin = k.min,  
  kmax = k.max,  
  n.cores = 1L,  
  proxy.response = "pc1",  
  max.iterations = 8L,  
  n.eigenpairs = 100L,  
  filter.type = "heat_kernel",  
  t.scale.factor = 0.5,  
  beta.coef.factor = 0.1,  
  verbose = TRUE  
)  
#> Building ikNN graphs (geom pruned) for k in [5, 12]  
#> Using 1 OpenMP thread  
#> Processing k values from 5 to 12 for 250 vertices  
#> Requested parallel.mode: auto  
#> Parallel mode: serial  
#> Starting graph processing  
#> [compute_knn] running... [compute_knn] 0.001s  
#> [graph_build/geometric_prune/isize_prune] running... [graph_build/geometric_prune/isize_prune] 0.368s  
#> Graph processing completed (0.368)  
#> [compute_knn] 0.001s  
#> [graph_build] 0.013s
```

```

#> [geometric_prune] 0.355s
#> [isize_prune] skipped (with.isize.pruning=FALSE)
#> Creating return list objects ... DONE (0.000)
#> Total elapsed time (0.369)
#> Summary of iknn_graphs object
#> -----
#> Number of vertices: 250
#> k range: 5 to 12
#> Max path-edge ratio deviation threshold: 0.1
#> Path-edge ratio percentile: 0.5
#> Graph type: geometrically pruned
#>
#>   idx  k n_ccomp edges mean_degree min_degree max_degree sparsity
#>   1   5     1    877    7.02        2       15  0.97182
#>   2   6     1   1056    8.45        3       17  0.96607
#>   3   7     1   1262   10.10        3       22  0.95945
#>   4   8     1   1437   11.50        3       23  0.95383
#>   5   9     1   1613   12.90        3       29  0.94818
#>   6  10     1   1766   14.13        3       29  0.94326
#>   7  11     1   1927   15.42        3       32  0.93809
#>   8  12     1   2047   16.38        3       33  0.93423
#> Proxy response: PC1 score (fraction variance explained = 0.5565)
#> Fitting fit.rdgraph.regression() across k values and extracting GCV
#> Selected k = 5 (min GCV = 1.212e-05)

X.smoothed <- X.denoise.res$X.smoothed
X.denoise.res$k.best
#> [1] 5

if (isTRUE(X.denoise.res$trimmed)) {
  kept.rows <- X.denoise.res$kept.rows
  X.use <- X[kept.rows, , drop = FALSE]
  theta.use <- theta.obs[kept.rows]
  y.use <- y[kept.rows]
  y.true.use <- y.true[kept.rows]
  comp1.use <- comp1[kept.rows]
  comp2.use <- comp2[kept.rows]
} else {
  X.use <- X
  theta.use <- theta.obs
  y.use <- y
  y.true.use <- y.true
  comp1.use <- comp1
  comp2.use <- comp2
}

op <- par(mfrow = c(1, 2),
          mar = c(2.75, 2.75, 0.5, 0.5),
          mgp = c(2.5, 0.5, 0),
          tcl = -0.3)

plot(X.use, las = 1, asp = 1, xlab = "", ylab = "")
lines(circle.true, lwd = 2)
legend("topleft",

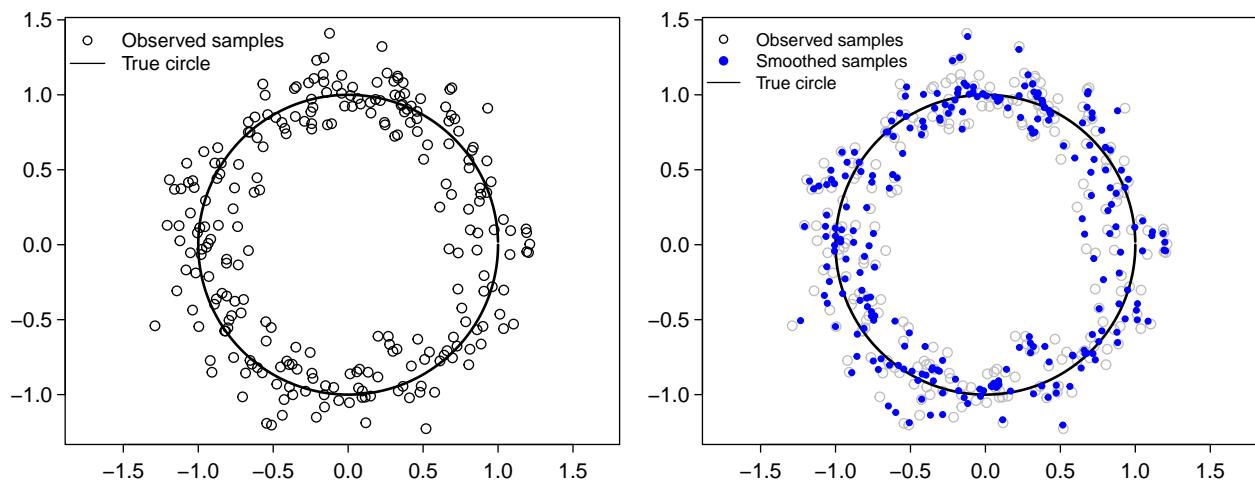
```

```

legend = c("Observed samples", "True circle"),
pch = c(1, NA),
lty = c(NA, 1),
col = c("black", "black"),
bty = "n", cex = 0.9)

plot(X.use, las = 1, asp = 1, col = "gray", xlab = "", ylab = "")
lines(circle.true, lwd = 2)
points(X.smoothed, col = "blue", pch = 19, cex = 0.6)
legend("topleft",
      legend = c("Observed samples", "Smoothed samples", "True circle"),
      pch = c(1, 19, NA),
      lty = c(NA, NA, 1),
      col = c("black", "blue", "black"),
      bty = "n", cex = 0.85)

```



```
par(op)
```

```

rad.obs <- sqrt(rowSums(X.use^2))
rad.sm <- sqrt(rowSums(X.smoothed^2))

rmse.obs <- sqrt(mean((rad.obs - radius)^2))
rmse.sm <- sqrt(mean((rad.sm - radius)^2))

data.frame(
  metric = c("RMSE radius (observed)", "RMSE radius (smoothed)"),
  value = c(rmse.obs, rmse.sm)
)
#>           metric      value
#> 1 RMSE radius (observed) 0.1472329
#> 2 RMSE radius (smoothed) 0.1321032

```

## Smooth y on the Selected Graph

We reuse the graph/spectral structure selected by `data.smoother()` and apply it to `y` via `refit.rdggraph.regression()`.

```

fit.seed <- X.denoise.res$fit.best

y.fit <- refit.rdggraph.regression(

```

```

fitted.model = fit.seed,
y.new = as.double(y.use),
per.column.gcv = FALSE,
n.cores = 1L,
verbose = FALSE
)

y.hat <- as.double(y.fit$fitted.values)

data.frame(
  metric = c("cor(y.hat, y.true)", "RMSE(y.hat, y.true)"),
  value = c(cor(y.hat, y.true.use),
            sqrt(mean((y.hat - y.true.use)^2)))
)
#>           metric    value
#> 1  cor(y.hat, y.true) 0.941873
#> 2 RMSE(y.hat, y.true) 0.128056

```

## Optional 3D Layout and Surface Coloring

```

X.denoise.graph.layout.3d <- graph.embedding(
  adj.list = fit.seed$graph$adj.list,
  weights.list = fit.seed$graph$edge.length.list,
  invert.weights = TRUE,
  dim = 3,
  method = "fr"
)

if (requireNamespace("rgl", quietly = TRUE)) {
  plot3D.cont(
    X.denoise.graph.layout.3d,
    y.hat,
    radius = 0.03,
    legend.title = "y.hat"
  )
}

if (requireNamespace("rgl", quietly = TRUE) &&
  requireNamespace("htmlwidgets", quietly = TRUE)) {
  plot3D.cont.html(
    X.denoise.graph.layout.3d,
    y.hat,
    legend.title = "y.hat",
    output.file = NULL
  )
}

```

## Build Feature Matrix Z and Smooth It

Features are built so some correlate with selected arcs of the response, while others are pure noise.

```

set.seed(1002)

window.1 <- as.numeric(abs(wrap_angle(theta.use, mu1)) <= 1.1)

```

```

window.2 <- as.numeric(abs(wrap_angle(theta.use, mu2)) <= 1.1)

z.global <- y.true.use + rnorm(length(theta.use), sd = 0.08)
z.peak1.arc <- window.1 * y.true.use + rnorm(length(theta.use), sd = 0.08)
z.peak2.arc <- window.2 * y.true.use + rnorm(length(theta.use), sd = 0.08)
z.peak.contrast <- (comp1.use - comp2.use) + rnorm(length(theta.use), sd = 0.08)

z.noise1 <- rnorm(length(theta.use))
z.noise2 <- rnorm(length(theta.use))
z.noise3 <- rnorm(length(theta.use))

Z <- cbind(
  z_global = z.global,
  z_peak1_arc = z.peak1.arc,
  z_peak2_arc = z.peak2.arc,
  z_peak_contrast = z.peak.contrast,
  z_noise1 = z.noise1,
  z_noise2 = z.noise2,
  z_noise3 = z.noise3
)
Z <- scale(Z)

Z.fit <- refit.rdggraph.regression(
  fitted.model = fit.seed,
  y.new = Z,
  per.column.gcv = FALSE,
  n.cores = 1L,
  verbose = FALSE
)
Z.sm <- as.matrix(Z.fit$fitted.values)
summary(Z.fit)
#>
#> Summary: Refitted Riemannian Graph Regression
#> -----
#>
#> Observations: 250
#> Responses: 7
#>
#> Fit quality summary across 7 responses:
#>   R-sq: min=0.3491, Q1=0.3768, med=0.9406, Q3=0.9621, max=0.9775
#>   RMSE: min=0.1499, Q1=0.1941, med=0.2433, Q3=0.7879, max=0.8051

```

## Compute Refined Basins from y.hat

```

y.basins <- compute_basins_safe(
  adj.list = fit.seed$graph$adj.list,
  edge.length.list = fit.seed$graph$edge.length.list,
  fitted.values = y.hat,
  verbose = TRUE
)
#> Step 1: Computing initial basins of attraction...
#>   Found 17 maxima and 22 minima
#>   initial.summary:

```

```

#>   label vertex      value  rel.value type hop.idx basin.size
#> 1    m1     245 -0.26630000 -0.81112583 min     2       4
#> 2    m2     120 -0.26183263 -0.79751862 min     2       7
#> 3    m3      92 -0.19902809 -0.60622164 min     2      10
#> 4    m4     230 -0.18650066 -0.56806424 min     3      13
#> 5    m5      78 -0.18240414 -0.55558661 min     4      14
#> 6    m6     208 -0.15682252 -0.47766730 min     1       3
#> 7    m7      93 -0.12477777 -0.38006185 min     3       6
#> 8    m8     250 -0.12346690 -0.37606906 min     1       6
#> 9    m9     101 -0.10990627 -0.33476461 min     1       6
#> 10   m10    6 -0.06511288 -0.19832799 min     6      15
#> 11   m11    203 0.02142402 0.06525565 min     8      25
#> 12   m12    211 0.02177421 0.06632230 min     2       6
#> 13   m13     64 0.04779295 0.14557303 min     5      30
#> 14   m14     72 0.05842157 0.17794685 min     1       4
#> 15   m15    132 0.12752894 0.38844168 min     3      15
#> 16   m16    146 0.18143515 0.55263513 min     1       6
#> 17   m17     55 0.26844958 0.81767328 min     3      14
#> 18   m18    180 0.37570701 1.14436974 min     4       6
#> 19   m19    162 0.56169281 1.71086575 min     1       3
#> 20   m20    164 0.62733373 1.91080209 min     2       7
#> 21   m21     33 0.97835737 2.97998852 min     2       5
#> 22   m22     23 1.01710762 3.09801830 min     0       1
#> 23   M1      30 1.46221755 4.45378309 max     8      39
#> 24   M2      28 1.32545340 4.03721181 max     2       7
#> 25   M3     148 0.96668007 2.94442051 max     0       1
#> 26   M4     163 0.90997849 2.77171260 max     2       3
#> 27   M5     153 0.85694374 2.61017354 max     2       8
#> 28   M6     190 0.54820939 1.66979647 max     4      12
#> 29   M7     217 0.35249067 1.07365485 max     1       6
#> 30   M8      60 0.34274227 1.04396212 max     0       1
#> 31   M9     103 0.33509074 1.02065624 max     2       7
#> 32   M10     8 0.29976846 0.91306776 max     3       5
#> 33   M11     77 0.29952927 0.91233921 max     4      18
#> 34   M12    129 0.27586066 0.84024674 max     1       3
#> 35   M13    216 0.25715592 0.78327380 max     1       4
#> 36   M14    247 0.24114239 0.73449802 max     3       8
#> 37   M15     71 0.22591942 0.68813024 max     0       1
#> 38   M16    233 0.16328801 0.49736059 max     2      11
#> 39   M17    102 0.12545658 0.38212947 max     2       7
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1          0.788          0.912  4 0.956
#> 2          0.100          0.512  5 0.896
#> 3          0.168          0.092  6 0.684
#> 4          0.696          0.364  7 0.504
#> 5          0.404          0.648  6 0.684
#> 6          0.780          0.964  5 0.896
#> 7          0.812          0.720  5 0.896
#> 8          0.016          0.380  5 0.896
#> 9          0.012          0.632  5 0.896
#> 10         0.496          0.224  5 0.896
#> 11         0.928          0.916  4 0.956
#> 12         0.732          0.572  6 0.684

```

```

#> 13      0.164      0.080    9 0.252
#> 14      0.464      0.592    5 0.896
#> 15      0.748      0.480    6 0.684
#> 16      0.172      0.412    6 0.684
#> 17      0.348      0.176    7 0.504
#> 18      0.664      0.976    3 0.980
#> 19      0.860      0.848    5 0.896
#> 20      0.764      0.440    5 0.896
#> 21      0.624      0.952    5 0.896
#> 22      0.996      0.996    4 0.956
#> 23      0.328      0.208    3 0.980
#> 24      0.096      0.616    6 0.684
#> 25      1.000      1.000    3 0.980
#> 26      0.628      0.288    2 1.000
#> 27      0.580      0.524    7 0.504
#> 28      0.920      0.764    4 0.956
#> 29      0.304      0.388    6 0.684
#> 30      0.972      0.676    2 1.000
#> 31      0.548      0.472    5 0.896
#> 32      0.568      0.048    3 0.980
#> 33      0.408      0.232    5 0.896
#> 34      0.244      0.008    2 1.000
#> 35      0.444      0.904    6 0.684
#> 36      0.032      0.504    5 0.896
#> 37      0.988      0.992    2 1.000
#> 38      0.240      0.096    5 0.896
#> 39      0.020      0.756    5 0.896
#>
#> Step 2: Filtering by relative values (max >= 1.10, min <= 0.90)...
#> Retained 6 maxima and 17 minima
#> relvalue.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1     245 -0.26630000 -0.81112583 min     2       4
#> 2   m2     120 -0.26183263 -0.79751862 min     2       7
#> 3   m3      92 -0.19902809 -0.60622164 min     2      10
#> 4   m4     230 -0.18650066 -0.56806424 min     3      13
#> 5   m5      78 -0.18240414 -0.55558661 min     4      14
#> 6   m6     208 -0.15682252 -0.47766730 min     1       3
#> 7   m7      93 -0.12477777 -0.38006185 min     3       6
#> 8   m8     250 -0.12346690 -0.37606906 min     1       6
#> 9   m9     101 -0.10990627 -0.33476461 min     1       6
#> 10  m10     6 -0.06511288 -0.19832799 min     6      15
#> 11  m11    203  0.02142402  0.06525565 min     8      25
#> 12  m12    211  0.02177421  0.06632230 min     2       6
#> 13  m13     64  0.047779295 0.14557303 min     5      30
#> 14  m14     72  0.05842157  0.17794685 min     1       4
#> 15  m15    132  0.12752894  0.38844168 min     3      15
#> 16  m16    146  0.18143515  0.55263513 min     1       6
#> 17  m17     55  0.26844958  0.81767328 min     3      14
#> 18  M1      30  1.46221755  4.45378309 max     8      39
#> 19  M2      28  1.32545340  4.03721181 max     2       7
#> 20  M3     148  0.96668007  2.94442051 max     0       1
#> 21  M4     163  0.90997849  2.77171260 max     2       3

```

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#> 22      M5      153  0.85694374  2.61017354  max      2      8
#> 23      M6      190  0.54820939  1.66979647  max      4     12
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1       0.788      0.912  4 0.956
#> 2       0.100      0.512  5 0.896
#> 3       0.168      0.092  6 0.684
#> 4       0.696      0.364  7 0.504
#> 5       0.404      0.648  6 0.684
#> 6       0.780      0.964  5 0.896
#> 7       0.812      0.720  5 0.896
#> 8       0.016      0.380  5 0.896
#> 9       0.012      0.632  5 0.896
#> 10      0.496      0.224  5 0.896
#> 11      0.928      0.916  4 0.956
#> 12      0.732      0.572  6 0.684
#> 13      0.164      0.080  9 0.252
#> 14      0.464      0.592  5 0.896
#> 15      0.748      0.480  6 0.684
#> 16      0.172      0.412  6 0.684
#> 17      0.348      0.176  7 0.504
#> 18      0.328      0.208  3 0.980
#> 19      0.096      0.616  6 0.684
#> 20      1.000      1.000  3 0.980
#> 21      0.628      0.288  2 1.000
#> 22      0.580      0.524  7 0.504
#> 23      0.920      0.764  4 0.956
#>
#> Step 3: Clustering maxima (overlap threshold = 0.15)...
#> Found 6 clusters (6 singletons, 0 to be merged)
#> Merging clustered maxima...
#> Result: 6 maxima after merging
#> merged.max.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min      2      4
#> 2   m2    120 -0.26183263 -0.79751862 min      2      7
#> 3   m3     92 -0.19902809 -0.60622164 min      2     10
#> 4   m4    230 -0.18650066 -0.56806424 min      3     13
#> 5   m5     78 -0.18240414 -0.55558661 min      4     14
#> 6   m6    208 -0.15682252 -0.47766730 min      1      3
#> 7   m7     93 -0.12477777 -0.38006185 min      3      6
#> 8   m8    250 -0.12346690 -0.37606906 min      1      6
#> 9   m9    101 -0.10990627 -0.33476461 min      1      6
#> 10  m10     6 -0.06511288 -0.19832799 min      6     15
#> 11  m11    203  0.02142402  0.06525565 min      8     25
#> 12  m12    211  0.02177421  0.06632230 min      2      6
#> 13  m13     64  0.04779295  0.14557303 min      5     30
#> 14  m14     72  0.05842157  0.17794685 min      1      4
#> 15  m15    132  0.12752894  0.38844168 min      3     15
#> 16  m16    146  0.18143515  0.55263513 min      1      6
#> 17  m17     55  0.26844958  0.81767328 min      3     14
#> 18  M1     30  1.46221755  4.45378309 max      8     39
#> 19  M2     28  1.32545340  4.03721181 max      2      7
#> 20  M3    148  0.96668007  2.94442051 max      0      1

```

```

#> 21 M4 163 0.90997849 2.77171260 max 2 3
#> 22 M5 153 0.85694374 2.61017354 max 2 8
#> 23 M6 190 0.54820939 1.66979647 max 4 12
#>   p.mean.nbros.dist p.mean.hopk.dist deg p.deg
#> 1      0.788      0.912 4 0.956
#> 2      0.100      0.512 5 0.896
#> 3      0.168      0.092 6 0.684
#> 4      0.696      0.364 7 0.504
#> 5      0.404      0.648 6 0.684
#> 6      0.780      0.964 5 0.896
#> 7      0.812      0.720 5 0.896
#> 8      0.016      0.380 5 0.896
#> 9      0.012      0.632 5 0.896
#> 10     0.496      0.224 5 0.896
#> 11     0.928      0.916 4 0.956
#> 12     0.732      0.572 6 0.684
#> 13     0.164      0.080 9 0.252
#> 14     0.464      0.592 5 0.896
#> 15     0.748      0.480 6 0.684
#> 16     0.172      0.412 6 0.684
#> 17     0.348      0.176 7 0.504
#> 18     0.328      0.208 3 0.980
#> 19     0.096      0.616 6 0.684
#> 20     1.000      1.000 3 0.980
#> 21     0.628      0.288 2 1.000
#> 22     0.580      0.524 7 0.504
#> 23     0.920      0.764 4 0.956
#>
#> Step 4: Clustering minima (overlap threshold = 0.15)...
#> Found 16 clusters (15 singletons, 1 to be merged)
#> Merging clustered minima...
#> Result: 16 minima after merging
#> merged.min.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min   2      4
#> 2   m2    120 -0.26183263 -0.79751862 min   2      7
#> 3   m3     92 -0.19902809 -0.60622164 min   2     10
#> 4   m4    230 -0.18650066 -0.56806424 min   3     13
#> 5   m5     78 -0.18240414 -0.55558661 min   4     14
#> 6   m6    208 -0.15682252 -0.47766730 min   1      3
#> 7   m7     93 -0.12477777 -0.38006185 min   3      6
#> 8   m8    250 -0.12346690 -0.37606906 min   1      6
#> 9   m9    101 -0.10990627 -0.33476461 min   1      6
#> 10  m10     6 -0.06511288 -0.19832799 min   6     15
#> 11  m11    203 0.02142402 0.06525565 min   8     25
#> 12  m12    211 0.02177421 0.06632230 min   2      6
#> 13  m13     64 0.04779295 0.14557303 min   5     31
#> 14  m14     72 0.05842157 0.17794685 min   1      4
#> 15  m15    132 0.12752894 0.38844168 min   3     15
#> 16  m16    146 0.18143515 0.55263513 min   1      6
#> 17  M1     30 1.46221755 4.45378309 max   8     39
#> 18  M2     28 1.32545340 4.03721181 max   2      7
#> 19  M3    148 0.96668007 2.94442051 max   0      1

```

```

#> 20 M4 163 0.90997849 2.77171260 max 2 3
#> 21 M5 153 0.85694374 2.61017354 max 2 8
#> 22 M6 190 0.54820939 1.66979647 max 4 12
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1      0.788      0.912 4 0.956
#> 2      0.100      0.512 5 0.896
#> 3      0.168      0.092 6 0.684
#> 4      0.696      0.364 7 0.504
#> 5      0.404      0.648 6 0.684
#> 6      0.780      0.964 5 0.896
#> 7      0.812      0.720 5 0.896
#> 8      0.016      0.380 5 0.896
#> 9      0.012      0.632 5 0.896
#> 10     0.496      0.224 5 0.896
#> 11     0.928      0.916 4 0.956
#> 12     0.732      0.572 6 0.684
#> 13     0.164      0.080 9 0.252
#> 14     0.464      0.592 5 0.896
#> 15     0.748      0.480 6 0.684
#> 16     0.172      0.412 6 0.684
#> 17     0.328      0.208 3 0.980
#> 18     0.096      0.616 6 0.684
#> 19     1.000      1.000 3 0.980
#> 20     0.628      0.288 2 1.000
#> 21     0.580      0.524 7 0.504
#> 22     0.920      0.764 4 0.956
#>
#> Step 5: Filtering by geometric characteristics and basin size:
#>   p.mean.nbrs.dist < 0.90, p.mean.hopk.dist < 0.90, p.deg < 0.90, basin.size >= 10 ...
#>   Maxima: 6 -> 0 (removed 6)
#> geom.summary:
#>   label vertex      value    rel.value type hop.idx basin.size
#> 1   m1 245 -0.26630000 -0.81112583 min 2 4
#> 2   m2 120 -0.26183263 -0.79751862 min 2 7
#> 3   m3 92 -0.19902809 -0.60622164 min 2 10
#> 4   m4 230 -0.18650066 -0.56806424 min 3 13
#> 5   m5 78 -0.18240414 -0.55558661 min 4 14
#> 6   m6 208 -0.15682252 -0.47766730 min 1 3
#> 7   m7 93 -0.12477777 -0.38006185 min 3 6
#> 8   m8 250 -0.12346690 -0.37606906 min 1 6
#> 9   m9 101 -0.10990627 -0.33476461 min 1 6
#> 10  m10 6 -0.06511288 -0.19832799 min 6 15
#> 11  m11 203 0.02142402 0.06525565 min 8 25
#> 12  m12 211 0.02177421 0.06632230 min 2 6
#> 13  m13 64 0.04779295 0.14557303 min 5 31
#> 14  m14 72 0.05842157 0.17794685 min 1 4
#> 15  m15 132 0.12752894 0.38844168 min 3 15
#> 16  m16 146 0.18143515 0.55263513 min 1 6
#> 17  M1 30 1.46221755 4.45378309 max 8 39
#> 18  M2 28 1.32545340 4.03721181 max 2 7
#> 19  M3 148 0.96668007 2.94442051 max 0 1
#> 20  M4 163 0.90997849 2.77171260 max 2 3
#> 21  M5 153 0.85694374 2.61017354 max 2 8

```

```

#> 22      M6     190  0.54820939  1.66979647  max        4          12
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1       0.788       0.912  4 0.956
#> 2       0.100       0.512  5 0.896
#> 3       0.168       0.092  6 0.684
#> 4       0.696       0.364  7 0.504
#> 5       0.404       0.648  6 0.684
#> 6       0.780       0.964  5 0.896
#> 7       0.812       0.720  5 0.896
#> 8       0.016       0.380  5 0.896
#> 9       0.012       0.632  5 0.896
#> 10      0.496       0.224  5 0.896
#> 11      0.928       0.916  4 0.956
#> 12      0.732       0.572  6 0.684
#> 13      0.164       0.080  9 0.252
#> 14      0.464       0.592  5 0.896
#> 15      0.748       0.480  6 0.684
#> 16      0.172       0.412  6 0.684
#> 17      0.328       0.208  3 0.980
#> 18      0.096       0.616  6 0.684
#> 19      1.000       1.000  3 0.980
#> 20      0.628       0.288  2 1.000
#> 21      0.580       0.524  7 0.504
#> 22      0.920       0.764  4 0.956
#>   Minima: 16 -> 6 (removed 10)
#>
#> Generating final summary (hop.k = 2)...
#> Populating extrema labels in basins object...
#> final.summary:
#>   label vertex      value rel.value type hop.idx basin.size p.mean.nbrs.dist
#> 1   m1    92 -0.19902809 -0.6062216 min     2         10      0.168
#> 2   m2   230 -0.18650066 -0.5680642 min     3         13      0.696
#> 3   m3    78 -0.18240414 -0.5555866 min     4         14      0.404
#> 4   m4     6 -0.06511288 -0.1983280 min     6         15      0.496
#> 5   m5    64  0.04779295  0.1455730 min     5         31      0.164
#> 6   m6   132  0.12752894  0.3884417 min     3         15      0.748
#>   p.mean.hopk.dist deg p.deg
#> 1       0.092  6 0.684
#> 2       0.364  7 0.504
#> 3       0.648  6 0.684
#> 4       0.224  5 0.896
#> 5       0.080  9 0.252
#> 6       0.480  6 0.684
#>   Assigned 0 maximum labels
#>   Assigned 6 minimum labels
#> Storing graph structure in basins object...
#> Constructing basin vertices lists...
#> Computing final overlap distance matrices...
#>
#> Step 6: Expanding basins to cover all vertices...
#>   Minima basins: 90 -> 250 vertices covered
#>
#> Refinement complete!

```

```

#> Final structure: 0 maxima and 6 minima
#> Step 1: Computing initial basins of attraction...
#>   Found 17 maxima and 22 minima
#> initial.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min     6    16
#> 2   m2    120 -0.26183263 -0.79751862 min     4    16
#> 3   m3     92 -0.19902809 -0.60622164 min     9    57
#> 4   m4    230 -0.18650066 -0.56806424 min     4    17
#> 5   m5     78 -0.18240414 -0.55558661 min     8    45
#> 6   m6    208 -0.15682252 -0.47766730 min     2     6
#> 7   m7     93 -0.12477777 -0.38006185 min    11    51
#> 8   m8    250 -0.12346690 -0.37606906 min     2     7
#> 9   m9    101 -0.10990627 -0.33476461 min     1     6
#> 10  m10    6 -0.06511288 -0.19832799 min     5    22
#> 11  m11   203 0.02142402 0.06525565 min     8    38
#> 12  m12   211 0.02177421 0.06632230 min     3    10
#> 13  m13    64 0.04779295 0.14557303 min     5    39
#> 14  m14    72 0.05842157 0.17794685 min     1     6
#> 15  m15   132 0.12752894 0.38844168 min     4    22
#> 16  m16   146 0.18143515 0.55263513 min     4    15
#> 17  m17    55 0.26844958 0.81767328 min     4    22
#> 18  m18   180 0.37570701 1.14436974 min     6    11
#> 19  m19   162 0.56169281 1.71086575 min     2     5
#> 20  m20   164 0.62733373 1.91080209 min     2     7
#> 21  m21    33 0.97835737 2.97998852 min     4     9
#> 22  m22    23 1.01710762 3.09801830 min     0     1
#> 23  M1     30 1.46221755 4.45378309 max     9    63
#> 24  M2     28 1.32545340 4.03721181 max     2     8
#> 25  M3     148 0.96668007 2.94442051 max     0     1
#> 26  M4     163 0.90997849 2.77171260 max     2     3
#> 27  M5     153 0.85694374 2.61017354 max     6    25
#> 28  M6     190 0.54820939 1.66979647 max     4    16
#> 29  M7     217 0.35249067 1.07365485 max     3     9
#> 30  M8      60 0.34274227 1.04396212 max     1     2
#> 31  M9     103 0.33509074 1.02065624 max     4    16
#> 32  M10     8 0.29976846 0.91306776 max     2     7
#> 33  M11     77 0.29952927 0.912333921 max     4    26
#> 34  M12    129 0.27586066 0.84024674 max     1     3
#> 35  M13    216 0.25715592 0.78327380 max     5    20
#> 36  M14    247 0.24114239 0.73449802 max     3     8
#> 37  M15     71 0.22591942 0.68813024 max     1     2
#> 38  M16    233 0.16328801 0.49736059 max     2    13
#> 39  M17    102 0.12545658 0.38212947 max     3    11
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1           0.788          0.912    4 0.956
#> 2           0.100          0.512    5 0.896
#> 3           0.168          0.092    6 0.684
#> 4           0.696          0.364    7 0.504
#> 5           0.404          0.648    6 0.684
#> 6           0.780          0.964    5 0.896
#> 7           0.812          0.720    5 0.896
#> 8           0.016          0.380    5 0.896

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```

#> 9      0.012      0.632  5 0.896
#> 10     0.496      0.224  5 0.896
#> 11     0.928      0.916  4 0.956
#> 12     0.732      0.572  6 0.684
#> 13     0.164      0.080  9 0.252
#> 14     0.464      0.592  5 0.896
#> 15     0.748      0.480  6 0.684
#> 16     0.172      0.412  6 0.684
#> 17     0.348      0.176  7 0.504
#> 18     0.664      0.976  3 0.980
#> 19     0.860      0.848  5 0.896
#> 20     0.764      0.440  5 0.896
#> 21     0.624      0.952  5 0.896
#> 22     0.996      0.996  4 0.956
#> 23     0.328      0.208  3 0.980
#> 24     0.096      0.616  6 0.684
#> 25     1.000      1.000  3 0.980
#> 26     0.628      0.288  2 1.000
#> 27     0.580      0.524  7 0.504
#> 28     0.920      0.764  4 0.956
#> 29     0.304      0.388  6 0.684
#> 30     0.972      0.676  2 1.000
#> 31     0.548      0.472  5 0.896
#> 32     0.568      0.048  3 0.980
#> 33     0.408      0.232  5 0.896
#> 34     0.244      0.008  2 1.000
#> 35     0.444      0.904  6 0.684
#> 36     0.032      0.504  5 0.896
#> 37     0.988      0.992  2 1.000
#> 38     0.240      0.096  5 0.896
#> 39     0.020      0.756  5 0.896
#>
#> Step 2: Filtering by relative values (max >= 1.02, min <= 0.98)...
#> Retained 9 maxima and 17 minima
#> relvalue.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min    6    16
#> 2   m2    120 -0.26183263 -0.79751862 min    4    16
#> 3   m3     92 -0.19902809 -0.60622164 min    9    57
#> 4   m4    230 -0.18650066 -0.56806424 min    4    17
#> 5   m5     78 -0.18240414 -0.55558661 min    8    45
#> 6   m6    208 -0.15682252 -0.47766730 min    2     6
#> 7   m7     93 -0.12477777 -0.38006185 min   11    51
#> 8   m8    250 -0.12346690 -0.37606906 min    2     7
#> 9   m9    101 -0.10990627 -0.33476461 min    1     6
#> 10  m10    6 -0.06511288 -0.19832799 min    5    22
#> 11  m11   203  0.02142402  0.06525565 min    8    38
#> 12  m12   211  0.02177421  0.06632230 min    3    10
#> 13  m13    64  0.04779295  0.14557303 min    5    39
#> 14  m14    72  0.05842157  0.17794685 min    1     6
#> 15  m15   132  0.12752894  0.38844168 min    4    22
#> 16  m16   146  0.18143515  0.55263513 min    4    15
#> 17  m17    55  0.26844958  0.81767328 min    4    22

```

```

#> 18   M1    30  1.46221755  4.45378309  max   9    63
#> 19   M2    28  1.32545340  4.03721181  max   2    8
#> 20   M3    148 0.96668007  2.94442051  max   0    1
#> 21   M4    163 0.90997849  2.77171260  max   2    3
#> 22   M5    153 0.85694374  2.61017354  max   6    25
#> 23   M6    190 0.54820939  1.66979647  max   4    16
#> 24   M7    217 0.35249067  1.07365485  max   3    9
#> 25   M8    60  0.34274227  1.04396212  max   1    2
#> 26   M9    103 0.33509074  1.02065624  max   4    16
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1      0.788          0.912  4 0.956
#> 2      0.100          0.512  5 0.896
#> 3      0.168          0.092  6 0.684
#> 4      0.696          0.364  7 0.504
#> 5      0.404          0.648  6 0.684
#> 6      0.780          0.964  5 0.896
#> 7      0.812          0.720  5 0.896
#> 8      0.016          0.380  5 0.896
#> 9      0.012          0.632  5 0.896
#> 10     0.496          0.224  5 0.896
#> 11     0.928          0.916  4 0.956
#> 12     0.732          0.572  6 0.684
#> 13     0.164          0.080  9 0.252
#> 14     0.464          0.592  5 0.896
#> 15     0.748          0.480  6 0.684
#> 16     0.172          0.412  6 0.684
#> 17     0.348          0.176  7 0.504
#> 18     0.328          0.208  3 0.980
#> 19     0.096          0.616  6 0.684
#> 20     1.000          1.000  3 0.980
#> 21     0.628          0.288  2 1.000
#> 22     0.580          0.524  7 0.504
#> 23     0.920          0.764  4 0.956
#> 24     0.304          0.388  6 0.684
#> 25     0.972          0.676  2 1.000
#> 26     0.548          0.472  5 0.896
#>
#> Step 3: Clustering maxima (overlap threshold = 0.25)...
#>   Found 9 clusters (9 singletons, 0 to be merged)
#>   Merging clustered maxima...
#>   Result: 9 maxima after merging
#> merged.max.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min   6    16
#> 2   m2    120 -0.26183263 -0.79751862 min   4    16
#> 3   m3    92  -0.19902809 -0.60622164 min   9    57
#> 4   m4    230 -0.18650066 -0.56806424 min   4    17
#> 5   m5    78  -0.18240414 -0.55558661 min   8    45
#> 6   m6    208 -0.15682252 -0.47766730 min   2     6
#> 7   m7    93  -0.12477777 -0.38006185 min  11    51
#> 8   m8    250 -0.12346690 -0.37606906 min   2     7
#> 9   m9    101 -0.10990627 -0.33476461 min   1     6
#> 10  m10   6  -0.06511288 -0.19832799 min   5    22

```

```

#> 11 m11 203 0.02142402 0.06525565 min 8 38
#> 12 m12 211 0.02177421 0.06632230 min 3 10
#> 13 m13 64 0.04779295 0.14557303 min 5 39
#> 14 m14 72 0.05842157 0.17794685 min 1 6
#> 15 m15 132 0.12752894 0.38844168 min 4 22
#> 16 m16 146 0.18143515 0.55263513 min 4 15
#> 17 m17 55 0.26844958 0.81767328 min 4 22
#> 18 M1 30 1.46221755 4.45378309 max 9 63
#> 19 M2 28 1.32545340 4.03721181 max 2 8
#> 20 M3 148 0.96668007 2.94442051 max 0 1
#> 21 M4 163 0.90997849 2.77171260 max 2 3
#> 22 M5 153 0.85694374 2.61017354 max 6 25
#> 23 M6 190 0.54820939 1.66979647 max 4 16
#> 24 M7 217 0.35249067 1.07365485 max 3 9
#> 25 M8 60 0.34274227 1.04396212 max 1 2
#> 26 M9 103 0.33509074 1.02065624 max 4 16
#> p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1 0.788 0.912 4 0.956
#> 2 0.100 0.512 5 0.896
#> 3 0.168 0.092 6 0.684
#> 4 0.696 0.364 7 0.504
#> 5 0.404 0.648 6 0.684
#> 6 0.780 0.964 5 0.896
#> 7 0.812 0.720 5 0.896
#> 8 0.016 0.380 5 0.896
#> 9 0.012 0.632 5 0.896
#> 10 0.496 0.224 5 0.896
#> 11 0.928 0.916 4 0.956
#> 12 0.732 0.572 6 0.684
#> 13 0.164 0.080 9 0.252
#> 14 0.464 0.592 5 0.896
#> 15 0.748 0.480 6 0.684
#> 16 0.172 0.412 6 0.684
#> 17 0.348 0.176 7 0.504
#> 18 0.328 0.208 3 0.980
#> 19 0.096 0.616 6 0.684
#> 20 1.000 1.000 3 0.980
#> 21 0.628 0.288 2 1.000
#> 22 0.580 0.524 7 0.504
#> 23 0.920 0.764 4 0.956
#> 24 0.304 0.388 6 0.684
#> 25 0.972 0.676 2 1.000
#> 26 0.548 0.472 5 0.896
#>
#> Step 4: Clustering minima (overlap threshold = 0.25)...
#> Found 11 clusters (8 singletons, 3 to be merged)
#> Merging clustered minima...
#> Result: 11 minima after merging
#> merged.min.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1    m1    245 -0.26630000 -0.81112583 min    6    25
#> 2    m2    120 -0.26183263 -0.79751862 min    4    16
#> 3    m3     92 -0.19902809 -0.60622164 min    9    64

```

```

#> 4 m4 230 -0.18650066 -0.56806424 min 4 17
#> 5 m5 208 -0.15682252 -0.47766730 min 2 6
#> 6 m6 250 -0.12346690 -0.37606906 min 2 7
#> 7 m7 101 -0.10990627 -0.33476461 min 1 6
#> 8 m8 203 0.02142402 0.06525565 min 8 40
#> 9 m9 72 0.05842157 0.17794685 min 1 6
#> 10 m10 132 0.12752894 0.38844168 min 4 22
#> 11 m11 146 0.18143515 0.55263513 min 4 15
#> 12 M1 30 1.46221755 4.45378309 max 9 63
#> 13 M2 28 1.32545340 4.03721181 max 2 8
#> 14 M3 148 0.96668007 2.94442051 max 0 1
#> 15 M4 163 0.90997849 2.77171260 max 2 3
#> 16 M5 153 0.85694374 2.61017354 max 6 25
#> 17 M6 190 0.54820939 1.66979647 max 4 16
#> 18 M7 217 0.35249067 1.07365485 max 3 9
#> 19 M8 60 0.34274227 1.04396212 max 1 2
#> 20 M9 103 0.33509074 1.02065624 max 4 16
#> p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1 0.788 0.912 4 0.956
#> 2 0.100 0.512 5 0.896
#> 3 0.168 0.092 6 0.684
#> 4 0.696 0.364 7 0.504
#> 5 0.780 0.964 5 0.896
#> 6 0.016 0.380 5 0.896
#> 7 0.012 0.632 5 0.896
#> 8 0.928 0.916 4 0.956
#> 9 0.464 0.592 5 0.896
#> 10 0.748 0.480 6 0.684
#> 11 0.172 0.412 6 0.684
#> 12 0.328 0.208 3 0.980
#> 13 0.096 0.616 6 0.684
#> 14 1.000 1.000 3 0.980
#> 15 0.628 0.288 2 1.000
#> 16 0.580 0.524 7 0.504
#> 17 0.920 0.764 4 0.956
#> 18 0.304 0.388 6 0.684
#> 19 0.972 0.676 2 1.000
#> 20 0.548 0.472 5 0.896
#>
#> Step 5: Skipping geometric filtering
#>
#> Generating final summary (hop.k = 2)...
#> Populating extrema labels in basins object...
#> final.summary:
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min   6   25
#> 2   m2    120 -0.26183263 -0.79751862 min   4   16
#> 3   m3     92 -0.19902809 -0.60622164 min   9   64
#> 4   m4    230 -0.18650066 -0.56806424 min   4   17
#> 5   m5    208 -0.15682252 -0.47766730 min   2   6
#> 6   m6    250 -0.12346690 -0.37606906 min   2   7
#> 7   m7    101 -0.10990627 -0.33476461 min   1   6
#> 8   m8    203  0.02142402  0.06525565 min   8   40

```

```

#> 9   m9    72  0.05842157  0.17794685 min   1     6
#> 10  m10   132 0.12752894  0.38844168 min   4     22
#> 11  m11   146 0.18143515  0.55263513 min   4     15
#> 12  M1    30  1.46221755  4.45378309 max   9     63
#> 13  M2    28  1.32545340  4.03721181 max   2     8
#> 14  M3    148 0.96668007  2.94442051 max   0     1
#> 15  M4    163 0.90997849  2.77171260 max   2     3
#> 16  M5    153 0.85694374  2.61017354 max   6     25
#> 17  M6    190 0.54820939  1.66979647 max   4     16
#> 18  M7    217 0.35249067  1.07365485 max   3     9
#> 19  M8    60  0.34274227  1.04396212 max   1     2
#> 20  M9    103 0.33509074  1.02065624 max   4     16
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1      0.788          0.912 4 0.956
#> 2      0.100          0.512 5 0.896
#> 3      0.168          0.092 6 0.684
#> 4      0.696          0.364 7 0.504
#> 5      0.780          0.964 5 0.896
#> 6      0.016          0.380 5 0.896
#> 7      0.012          0.632 5 0.896
#> 8      0.928          0.916 4 0.956
#> 9      0.464          0.592 5 0.896
#> 10     0.748          0.480 6 0.684
#> 11     0.172          0.412 6 0.684
#> 12     0.328          0.208 3 0.980
#> 13     0.096          0.616 6 0.684
#> 14     1.000          1.000 3 0.980
#> 15     0.628          0.288 2 1.000
#> 16     0.580          0.524 7 0.504
#> 17     0.920          0.764 4 0.956
#> 18     0.304          0.388 6 0.684
#> 19     0.972          0.676 2 1.000
#> 20     0.548          0.472 5 0.896
#>   Assigned 9 maximum labels
#>   Assigned 11 minimum labels
#> Storing graph structure in basins object...
#> Constructing basin vertices lists...
#> Computing final overlap distance matrices...
#>
#> Step 6: Expanding basins to cover all vertices...
#> Maxima basins: 134 -> 250 vertices covered
#> Minima basins: 201 -> 250 vertices covered
#>
#> Refinement complete!
#> Final structure: 9 maxima and 11 minima

y.basins$summary
#>   label vertex      value  rel.value type hop.idx basin.size
#> 1   m1    245 -0.26630000 -0.81112583 min   6     25
#> 2   m2    120 -0.26183263 -0.79751862 min   4     16
#> 3   m3    92  -0.19902809 -0.60622164 min   9     64
#> 4   m4    230 -0.18650066 -0.56806424 min   4     17
#> 5   m5    208 -0.15682252 -0.47766730 min   2      6

```

```

#> 6   m6   250 -0.12346690 -0.37606906 min    2     7
#> 7   m7   101 -0.10990627 -0.33476461 min    1     6
#> 8   m8   203  0.02142402  0.06525565 min    8     40
#> 9   m9   72   0.05842157  0.17794685 min    1     6
#> 10  m10  132  0.12752894  0.38844168 min    4     22
#> 11  m11  146  0.18143515  0.55263513 min    4     15
#> 12  M1   30   1.46221755  4.45378309 max    9     63
#> 13  M2   28   1.32545340  4.03721181 max    2     8
#> 14  M3   148  0.96668007  2.94442051 max    0     1
#> 15  M4   163  0.90997849  2.77171260 max    2     3
#> 16  M5   153  0.85694374  2.61017354 max    6     25
#> 17  M6   190  0.54820939  1.66979647 max    4     16
#> 18  M7   217  0.35249067  1.07365485 max    3     9
#> 19  M8   60   0.34274227  1.04396212 max    1     2
#> 20  M9   103  0.33509074  1.02065624 max    4     16
#>   p.mean.nbrs.dist p.mean.hopk.dist deg p.deg
#> 1      0.788          0.912  4 0.956
#> 2      0.100          0.512  5 0.896
#> 3      0.168          0.092  6 0.684
#> 4      0.696          0.364  7 0.504
#> 5      0.780          0.964  5 0.896
#> 6      0.016          0.380  5 0.896
#> 7      0.012          0.632  5 0.896
#> 8      0.928          0.916  4 0.956
#> 9      0.464          0.592  5 0.896
#> 10     0.748          0.480  6 0.684
#> 11     0.172          0.412  6 0.684
#> 12     0.328          0.208  3 0.980
#> 13     0.096          0.616  6 0.684
#> 14     1.000          1.000  3 0.980
#> 15     0.628          0.288  2 1.000
#> 16     0.580          0.524  7 0.504
#> 17     0.920          0.764  4 0.956
#> 18     0.304          0.388  6 0.684
#> 19     0.972          0.676  2 1.000
#> 20     0.548          0.472  5 0.896

true.max.label <- ifelse(comp1.use >= comp2.use, "M1.true", "M2.true")

est.max.label <- assign_from_vertices_list(
  y.basins$expanded.max.vertices.list %||% y.basins$max.vertices.list,
  n = length(y.hat)
)

idx.ok <- !is.na(est.max.label)
table(True = true.max.label[idx.ok], Estimated = est.max.label[idx.ok])
#>           Estimated
#> True      M1 M2 M3 M4 M5 M6 M7 M8 M9
#>   M1.true 90  9  0  0  0  0  2  0
#>   M2.true 20  0  1 11 41 29 27  0 20

```

## Basin-Wise Pearson and Spearman Analyses

```

max.assign <- assign_from_vertices_list(
  y.basins$expanded.max.vertices.list %||% y.basins$max.vertices.list,
  n = length(y.hat)
)
min.assign <- assign_from_vertices_list(
  y.basins$expanded.min.vertices.list %||% y.basins$min.vertices.list,
  n = length(y.hat)
)
cell.assign <- ifelse(is.na(max.assign) | is.na(min.assign),
                      NA_character_,
                      paste(max.assign, min.assign, sep = " |"))

cor_by_group <- function(yv, zv, g, feature, min.size = 20L) {
  ii <- split(seq_along(g), g)
  ii <- ii[!is.na(names(ii))]
  ii <- ii[!(names(ii) %in% "NA")]
  if (length(ii) == 0L) return(data.frame())
  out <- lapply(names(ii), function(gr) {
    idx <- ii[[gr]]
    if (length(idx) < min.size) return(NULL)
    data.frame(
      feature = feature,
      group = gr,
      n = length(idx),
      pearson = suppressWarnings(cor(yv[idx], zv[idx], method = "pearson")),
      spearman = suppressWarnings(cor(yv[idx], zv[idx], method = "spearman"))
    )
  })
  out <- do.call(rbind, out)
  if (is.null(out)) data.frame() else out
}

basin.cor <- do.call(
  rbind,
  lapply(seq_len(ncol(Z.sm)), function(j) {
    cor_by_group(y.hat, Z.sm[, j], max.assign, colnames(Z.sm)[j], min.size = 20L)
  })
)

if (nrow(basin.cor) > 0) {
  head(basin.cor[order(-abs(basin.cor$spearman)), ], 12)
} else {
  basin.cor
}

#>           feature group   n     pearson     spearman
#> 13      z_peak2_arc    M6  29  0.9171324  0.8926108
#> 18      z_peak_contrast    M6  29 -0.8740734 -0.8472906
#> 3       z_global        M6  29  0.8860806  0.8394089
#> 6       z_peak1_arc     M1 110  0.9632611  0.8263869
#> 1       z_global        M1 110  0.9692105  0.8241688
#> 16     z_peak_contrast    M1 110  0.9658861  0.8131776
#> 2       z_global        M5  41  0.8799458  0.8090592

```

```

#> 17 z_peak_contrast      M5  41 -0.8541682 -0.8083624
#> 12      z_peak2_arc     M5  41  0.8385678  0.8024390
#> 23      z_noise1        M6  29  0.8270852  0.7753695
#> 35      z_noise3        M9  20  0.5967139  0.7127820
#> 33      z_noise3        M6  29 -0.4186189 -0.6842365

```

## Local Correlation with lcor()

```

adj.list <- fit.seed$graph$adj.list
edge.length.list <- fit.seed$graph$edge.length.list

lcor.derivative <- as_lcor_matrix(
  lcor(adj.list, edge.length.list, y.hat, Z, type = "derivative")
)
lcor.unit <- as_lcor_matrix(
  lcor(adj.list, edge.length.list, y.hat, Z, type = "unit")
)
lcor.sign <- as_lcor_matrix(
  lcor(adj.list, edge.length.list, y.hat, Z, type = "sign")
)

summarize_lcor <- function(mat, type) {
  data.frame(
    feature = colnames(mat),
    type = type,
    mean.lcor = colMeans(mat),
    mean.abs.lcor = colMeans(abs(mat)),
    row.names = NULL
  )
}

lcor.summary <- rbind(
  summarize_lcor(lcor.derivative, "derivative"),
  summarize_lcor(lcor.unit, "unit"),
  summarize_lcor(lcor.sign, "sign")
)

lcor.summary[order(lcor.summary$type, -lcor.summary$mean.abs.lcor), ]
#>      feature      type   mean.lcor mean.abs.lcor
#> 1      z_global derivative  0.223553319  0.5408227
#> 4  z_peak_contrast derivative -0.018374965  0.4996884
#> 2  z_peak1_arc derivative   0.172994951  0.4936340
#> 3  z_peak2_arc derivative   0.136151960  0.4776319
#> 7      z_noise3 derivative  -0.005235465  0.4560861
#> 6      z_noise2 derivative   0.001632273  0.4532052
#> 5      z_noise1 derivative   0.014150734  0.4502928
#> 15      z_global      sign  0.247438268  0.4879175
#> 18 z_peak_contrast      sign  0.017436406  0.4683590
#> 16      z_peak1_arc      sign  0.183656190  0.4448450
#> 17      z_peak2_arc      sign  0.111610027  0.3974714
#> 19      z_noise1        sign  0.046876824  0.3869158
#> 21      z_noise3        sign -0.008896981  0.3846748
#> 20      z_noise2        sign -0.012852809  0.3717648

```

```

#> 8      z_global      unit  0.247438268  0.4879175
#> 11     z_peak_contrast  unit  0.017436406  0.4683590
#> 9      z_peak1_arc    unit  0.183656190  0.4448450
#> 10     z_peak2_arc    unit  0.111610027  0.3974714
#> 12     z_noise1       unit  0.046876824  0.3869158
#> 14     z_noise3       unit -0.008896981  0.3846748
#> 13     z_noise2       unit -0.012852809  0.3717648

```

## Permutation Tests for lcor

`lcor()` computes local-correlation coefficients but does not directly return permutation p-values. We add a permutation layer by shuffling vertex labels of  $Z$  and recomputing the chosen `lcor` summary statistic.

```

perm.lcor <- perm_test_lcor(
  adj.list = adj.list,
  edge.length.list = edge.length.list,
  y = y.hat,
  Z = Z,
  type = "derivative",
  n.perm = 200L,
  seed = 2026L
)

perm.lcor$table[order(perm.lcor$table$q.value), ]
#>           feature  stat.obs   p.value   q.value
#> 1      z_global 0.5408227 0.004975124 0.03482587
#> 2      z_peak1_arc 0.4936340 0.009950249 0.03482587
#> 4 z_peak_contrast 0.4996884 0.019900498 0.04643449
#> 3      z_peak2_arc 0.4776319 0.124378109 0.21766169
#> 5      z_noise1 0.4502928 0.835820896 0.83582090
#> 6      z_noise2 0.4532052 0.830845771 0.83582090
#> 7      z_noise3 0.4560861 0.771144279 0.83582090

```

## Build an lcor-Based Feature Module Graph

```

sim <- stats::cor(lcor.derivative, use = "pairwise.complete.obs")
adj.sim <- abs(sim)
diag(adj.sim) <- 0

threshold <- 0.20
adj.sim[adj.sim < threshold] <- 0

if (sum(adj.sim > 0) == 0) {
  message("No edges above threshold. Lower `threshold` to make the module graph denser.")
} else {
  g.mod <- igraph::graph_from_adjacency_matrix(
    adj.sim,
    mode = "undirected",
    weighted = TRUE,
    diag = FALSE
  )

  modules <- igraph::cluster_louvain(g.mod, weights = igraph::E(g.mod)$weight)

```

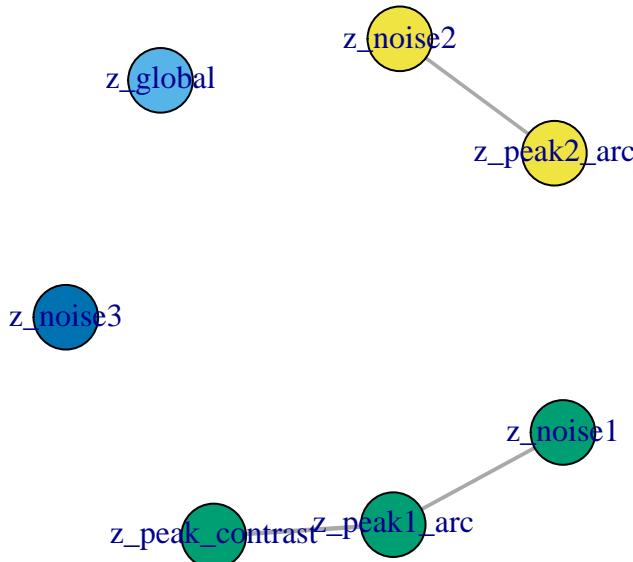
```

igraph::plot.igraph(
  g.mod,
  vertex.label = igraph::V(g.mod)$name,
  vertex.size = 26,
  vertex.color = as.integer(igraph::membership(modules)) + 1,
  edge.width = 1 + 4 * igraph::E(g.mod)$weight,
  main = "Feature modules from local-correlation profiles"
)

data.frame(
  feature = igraph::V(g.mod)$name,
  module = as.integer(igraph::membership(modules)),
  row.names = NULL
)
}

```

## Feature modules from local–correlation profiles



```

#>      feature module
#> 1      z_global     1
#> 2      z_peak1_arc  2
#> 3      z_peak2_arc  3
#> 4      z_peak_contrast 2
#> 5      z_noise1     2
#> 6      z_noise2     3
#> 7      z_noise3     4

```

## Optional grip 3D Layout

```

if (requireNamespace("grip", quietly = TRUE) &&
  requireNamespace("rgl", quietly = TRUE)) {
  g <- X.graphs$geom_pruned_graphs[[k.idx]]
}

```

```

X.graphs.3d <- grip::grip.layout(
  adj_list = g$adj_list,
  weight_list = g$weight_list,
  dim = 3,
  rounds = 50,
  final_rounds = 50,
  num_init = 36,
  num_nbrs = 8,
  seed = 6
)

plot3D.plain(X.graphs.3d, radius = 0.03)
}

```

## Notes

- For publication-quality inference, increase permutation count (`n.perm`), report confidence intervals, and retain `q.value` (FDR-controlled) summaries.
- For larger datasets, use `n.cores > 1` and consider chunk-level caching.
- If basin refinement is unstable for a specific dataset, start with relaxed filtering and tighten thresholds gradually.