Call: lavaan::sem(model = model, data = data, group = "Group", group.label = c("Control", "Clinical"), missing = "fiml", estimator = "ML")

Free parameters   1000

Number of observations   Control   Clinical

Number of missing patterns   Control   Clinical

Model Test User Model:

Test statistic   NaN

Degrees of freedom   NaN

P-value (Chi-square)   NaN

Test statistic for each group:

Control   NaN

Clinical   NaN

Model Test Baseline Model:

Test statistic   NaN

Degrees of freedom   NaN

P-value   NaN

User Model versus Baseline Model:

Comparative Fit Index (CFI)   NaN

Tucker-Lewis Index (TLI)   NaN

Loglikelihood and Information Criteria:

Loglikelihood user model (H0)   NaN

Loglikelihood unrestricted model (H1)   NaN

Akaike (AIC)   NaN

Bayesian (BIC)   NaN

Sample-size adjusted Bayesian (SABIC)   NaN

Root Mean Square Error of Approximation:

RMSEA   NaN

90 Percent confidence interval - lower   NaN

90 Percent confidence interval - upper   NaN

P-value RMSEA <= 0.05   NaN

Standardized Root Mean Square Residual:

SRMR   NaN

Parameter Estimates:

Standard errors   Standard

Information   Expected

Information saturated (h1) model   Structured

Group 1 [Control]:

Latent Variables:

Group 2 [Clinical]:

Latent Variables: