

## Summary of Analysis

Data resolution poor. Caution advised.

Best model found: gamma

Model fit might be unreliable.

p-val for chi-squared statistic was  $< 0.1$  for all converged models.

# Model Predictions

| Model          | Chi-squared | p-val | AIC    | BMD10 | BMDL10 |
|----------------|-------------|-------|--------|-------|--------|
| logistic       | 12.182      | 0.002 | 69.888 | 0.498 | 0.383  |
| gamma          | 3.906       | 0.048 | 66.638 | 0.922 | 0.781  |
| weibull        | 3.906       | 0.048 | 66.638 | 0.997 | nan    |
| log_logistic   | 3.906       | 0.048 | 66.638 | 0.967 | 0.764  |
| probit         | 12.967      | 0.002 | 71.437 | 0.444 | 0.342  |
| log_probit     | nan         | nan   | nan    | nan   | nan    |
| multistage_2   | 12.803      | 0.0   | 81.68  | 0.353 | 0.329  |
| quantal_linear | nan         | nan   | nan    | nan   | nan    |

## Scaled Residuals

| Model          | dose0     | dose1    | dose2    | dose3     |
|----------------|-----------|----------|----------|-----------|
| logistic       | -2.948703 | 1.008324 | 1.470854 | -0.553864 |
| gamma          | -1.541835 | 1.20078  | 0.295322 | 2.9e-05   |
| weibull        | -1.541889 | 1.200763 | 0.295298 | -0.000289 |
| log_logistic   | -1.54186  | 1.200772 | 0.295328 | -2.2e-05  |
| probit         | -2.87114  | 1.066479 | 1.715497 | -0.80198  |
| log_probit     | nan       | nan      | nan      | nan       |
| multistage_2   | -0.061898 | 1.933044 | 2.931828 | 0.683003  |
| quantal_linear | nan       | nan      | nan      | nan       |

Dose-response with best fit model (gamma)

