

Summary of Analysis

Data resolution poor. Caution advised.

Best model found: logistic

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 | 6.797 | 0.079 | 76.38 | 1.47 | 1.075 |
| gamma | nan | nan | nan | nan | nan |
| weibull | 5.978 | 0.05 | 78.154 | 1.746 | 1.468 |
| log_logistic | 5.982 | 0.05 | 78.154 | 1.457 | 0.567 |
| probit | 7.062 | 0.07 | 76.497 | 1.307 | 0.999 |
| log_probit | nan | nan | nan | nan | nan |
| multistage_2 | 6.237 | 0.044 | 81.267 | 1.389 | 1.168 |
| quantal_linear | nan | nan | nan | nan | nan |

Scaled Residuals

| Model | dose0 | dose1 | dose2 | dose3 | dose4 |
|----------------|-----------|----------|----------|-----------|-----------|
| logistic | -2.197069 | 1.276571 | 0.496345 | 0.305343 | -0.032457 |
| gamma | nan | nan | nan | nan | nan |
| weibull | -1.964123 | 1.345153 | 0.557752 | -0.005726 | 9.9e-05 |
| log_logistic | -1.965909 | 1.344578 | 0.556282 | -0.001658 | 0.000103 |
| probit | -2.253322 | 1.260823 | 0.482026 | 0.400974 | -0.043827 |
| log_probit | nan | nan | nan | nan | nan |
| multistage_2 | -0.755929 | 1.825762 | 1.219745 | 0.897559 | 0.196126 |
| quantal_linear | nan | nan | nan | nan | nan |

Dose-response with best fit model (logistic)

