

Summary of Analysis

Data resolution poor. Caution advised.

Best model found: log_probit

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	8.199	0.085	122.024	76.06	37.657
gamma	nan	nan	nan	nan	nan
weibull	7.962	0.047	124.0	79.082	34.123
log_logistic	7.9	0.048	122.183	0.292	nan
probit	nan	nan	nan	nan	nan
log_probit	7.801	0.05	121.964	0.248	nan
multistage_2	nan	nan	nan	nan	nan
quantal_linear	nan	nan	nan	nan	nan

Scaled Residuals

Model	dose0	dose1	dose2	dose3	dose4	dose5
logistic	1.674024	-2.223854	-0.273798	0.384757	0.475774	-0.039655
gamma	nan	nan	nan	nan	nan	nan
weibull	1.70127	-2.144477	-0.221374	0.423121	0.489094	-0.038064
log_logistic	0.957503	-2.292365	-0.056091	0.784677	1.015534	-0.279343
probit	nan	nan	nan	nan	nan	nan
log_probit	0.87724	-2.290935	-0.041977	0.802612	1.034342	-0.259494
multistage_2	nan	nan	nan	nan	nan	nan
quantal_linear	nan	nan	nan	nan	nan	nan

Dose-response with best fit model (log_probit)

