			χ ² Scaled residuals ^b				_		
			Goodness-	Dose	Dose				
			of-fit	below	above	Overall		BMC ₁₀	BMCL ₁₀
Model	DF	X ²	p-value ^a	BMC	BMC	largest	AIC	(mg/m ³)	(mg/m ³)
Gamma ^c	2	4.41	0.11	0.14	-1.03	1.82	77.98	3.69	2.85
Logistic	3	5.04	0.17	-1.20	-0.37	1.86	77.15	3.78	2.95
LogLogistic ^{d,e}	2	0.02	0.99	0.00	-0.00	0.13	70.45	5.87	4.82
LogProbit ^d	1	0.00	0.99	-0.00	-0.00	-0.00	72.42	5.92	4.73
Multistage (1-degree) ^f	2	18.41	0.00	0.28	-3.28	-3.28	95.80	ND	ND
Multistage (2-degree) ^f	2	2.81	0.24	0.20	-1.21	-1.21	74.57	3.40	2.54
Multistage (3-degree) ^f	2	0.02	0.99	0.01	-0.05	0.15	70.46	4.77	2.39
Probit	2	0.48	0.79	0.49	-0.28	0.49	71.03	4.09	3.12

Table A-5. Model Predictions for Hyaline Degeneration of the Nasal RespiratoryEpithelium in Female Rats Exposed to Molybdenum Trioxide (NTP 1997)

^aValues <0.1 fail to meet conventional goodness-of-fit criteria.

^bScaled residuals at doses immediately below and above the BMC; also the largest residual at any dose.

^cPower restricted to \geq 1.

^dSlope restricted to \geq 1.

eSelected model. BMCLs for models providing adequate fit were sufficiently close (differed by <3-fold). Therefore, the model with the lowest AIC was selected.

^fBetas restricted to ≥ 0 .

AIC = Akaike Information Criterion; BMC = maximum likelihood estimate of the exposure concentration associated with the selected benchmark response; BMCL = 95% lower confidence limit on the BMC (subscripts denote benchmark response: i.e., $_{10}$ = exposure concentration associated with 10% extra risk); DF = degrees of freedom; ND = not determined, goodness-of-fit criteria, p<0.10