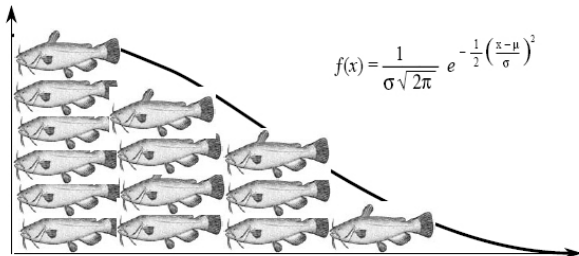


Statistische Beurteilung von Dosiswirkungsbeziehungen

Reinhard Meister

Beuth Hochschule Berlin



Agenda

*algae bacteria bees butterflies bugs cell-cultures cucumber
daphniae dogs ducks fish guinea-pigs hen's eggs humans
limb-buds mice rabbits rats whole-embryo worms yeast
zebrafish-embryos*

Some Construction Lots

- Quantal response – (quasi) complete separation
- Teratology – clustered data
- Time is Information – time-dose-response models
- A general Risk Measure



Data and Model

data $(d_i, r_i, n_i), i = 1, \dots, k; \quad d_1 < d_2 < \dots < d_k$

distribution $r_i \sim \text{binom}(\pi_i, n_i)$

model $\pi_i = p(d_i); \quad F^{-1}(\pi) = \alpha + \beta d = \frac{\mu - d}{\sigma} = \frac{ED_{0.5} - d}{c (ED_{0.5} - ED_q)}$

likelihood $L(\text{model}|\text{data}) = \prod_{i=1}^k \binom{n_i}{r_i} \pi_i^{r_i} (1 - \pi_i)^{n_i - r_i}$



Quantal Response – (quasi) complete separation

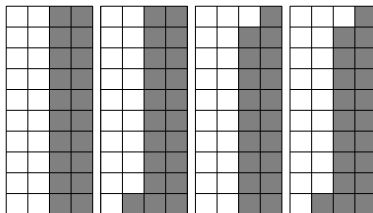
Separated Data

monotonicity $r_1/n_1 \leq r_2/n_2 \leq \dots \leq r_k/n_k$

complete $\max_i r_i (n_i - r_i) = 0$

quasi complete $\exists i^* : \max_{i \neq i^*} r_i (n_i - r_i) = 0, \quad r_{i^*} (n_{i^*} - r_{i^*}) > 0$

complete quasi quasi overlap



Quantal Response – Likelihood

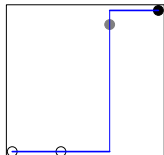
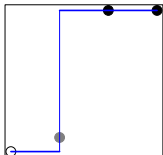
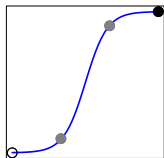
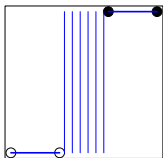
$\sup L$ under (quasi) complete separation

always $L(\pi|data) \leq 1$

complete sep. $\sup L = 1$

quasi complete $\sup L = \text{dbinom}(r_{i^*}, n_{i^*}, r_{i^*}/n_{i^*})$

quasi&complete $\sup L = L(\pi_i = r_i/n_i)$ saturated model



complete	overlap
quasi	quasi



Quantal Response – Benchmarks

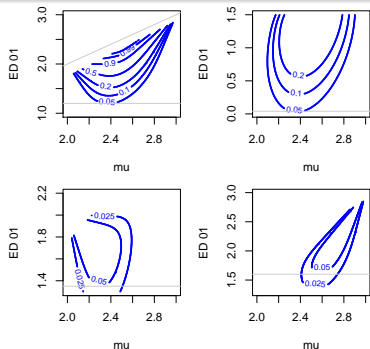
Likelihood intervals under separation

new parameters $(\alpha, \beta) \rightarrow \theta = (ED_q, ED_{0.5})$

LR-statistic $LR(\theta) = L(\theta) / \sup L$

complete sep. $LR(\theta) = L(\theta) = P(\text{data}|\theta)$

quasi complete $Dev(\theta) = -2 \log\{L(\theta) / \text{dbinom}(r_{i^*}, n_{i^*}, r_{i^*}/n_{i^*})\} \sim \chi_{k-2}^2$



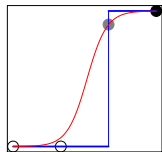
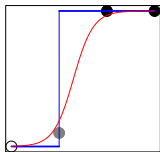
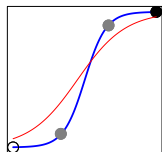
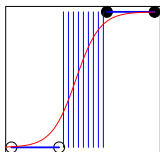
complete overlap
quasi quasi

doses: 1, 2, 3, 4
logit-link

Quantal Response – Benchmarks

Results Example BD01, $\alpha = 0.05$

- assume logistic model with $ED_{0.5} = 2.5$, $ED_{0.01} = 2$ then $P(\cup \text{data}) \approx 1$, $P(\text{overlap}) = 0.008$
- benchmarks appear reasonable
- **ATTENTION!** choice of model is crucial

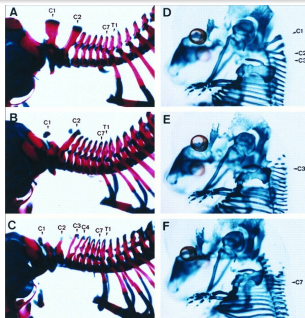


complete (1.2)	overlap (0.04)
quasi(1.35)	quasi (1.6)



Dose-Response Studies

- treat pregnant animals (rodents), randomized to dose groups
- record no. of implantations, resorptions, anomalies, etc per litter
- rather large litters, 8–15 in mice and rats



Yamaguchi et al, PNAS June 23, 1998 vol. 95 no. 13 7491-7496

Decomposition with Structured Noise

- data = signal + noise
- noise caused by **litters** b_i and by **fetuses** e_{ij}
 $i = 1, \dots, l; j = 1, \dots, n_i$ for simplicity assume all n_i equal
- $y_{ij} = \mu + b_i + e_{ij}$, assumptions: b_i, e_{ij} independent,
 $Eb_i = Ee_{ij} = 0, \text{Var}(b_i) = \sigma_b^2, \text{Var}(e_{ij}) = \sigma^2$.
- data **correlated within litters**, no totally independent information

$$\text{Corr}(y_{ij}, y_{ik}) = \sigma_b^2 / (\sigma_b^2 + \sigma^2) = \rho$$

$$\text{Var}(\bar{y}_{..}) = \frac{1}{l} \left(\sigma_b^2 + \frac{\sigma^2}{n} \right) = \frac{1}{l \times n} \left(1 + n \times \frac{\rho}{1 - \rho} \right) \sigma^2$$

Intra-litter correlation causes variance inflation of the mean

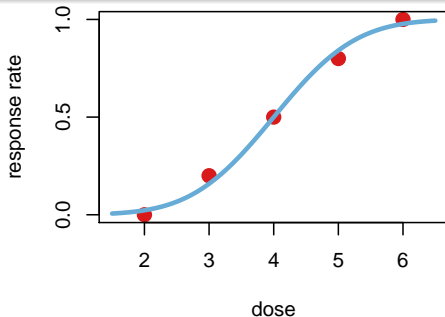
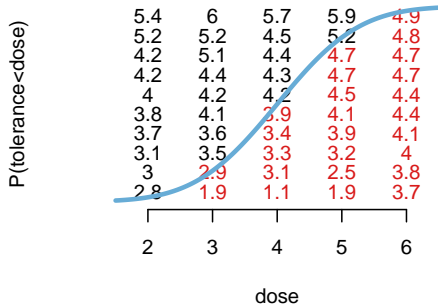


Tolerance and Quantal Data

Distribution of individual tolerances determines response probabilities

Hypothetical tolerances for 50 individuals randomly assigned to 5 dose groups.

Fit of quantal response model: *Maximum Likelihood for binomial data.*

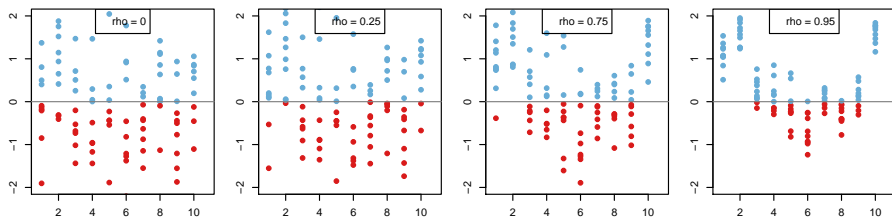


Quantal response allows indirect estimation of tolerance distribution

Litter Effects for Tolerances

How intra-litter correlation changes pattern of reactions

Tolerances with litter effects, 10 fetuses for 10 litters. Increasing intra-litter correlation.



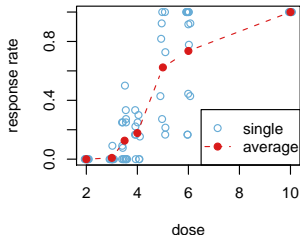
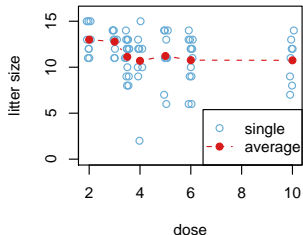
Litter effects increase dissimilarities of response rates between litters



MNU Cleft palate induced by MNU

Raw data from Platzcek et al

dose	reactions/litter size	$\sum r_i / \sum n_i$	litter effect
2	0/11 0/11 0/12 0/12 0/13 0/13 0/15 0/15 0/15	0	
3	0/11 1/11 0/12 0/13 0/13 0/13 0/14 0/14 0/14	0.009	-
3.5	0/8 2/8 0/9 3/9 5/10 1/11 3/11 0/12 0/12 3/12 3/12 0/13 0/13 1/13 0/14	0.13	**
4	0/2 2/9 0/10 3/10 0/12 2/12 2/12 3/12 2/13 5/15	0.18	-
5	1/6 7/7 3/11 8/11 9/11 12/13 3/14 6/14 14/14	0.62	**
6	6/6 6/6 4/9 7/9 10/10 11/11 2/12 2/12 11/12 12/13 13/13 13/13 6/14	0.74	**
10	7/7 8/8 9/9 11/11 12/12 12/12 13/13 14/14	1	



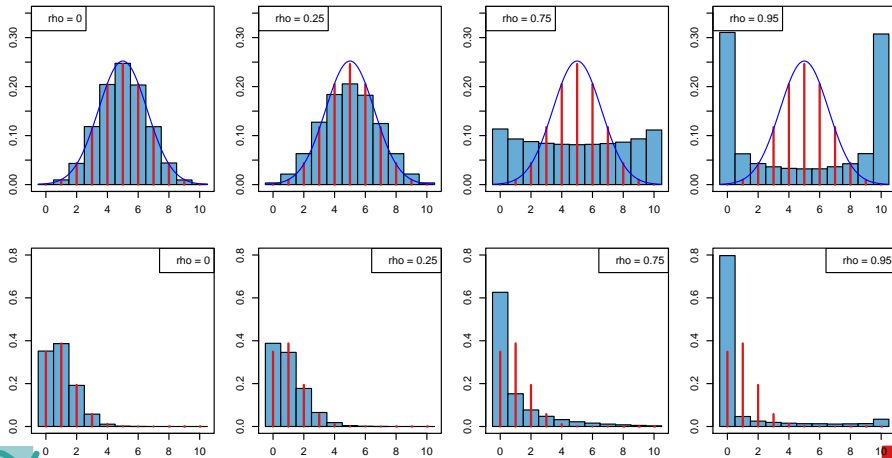
Results: slightly decreasing litter size, increase in response rate



Simulations Litter Effect and Quantal Data

How intra-litter correlation changes distribution

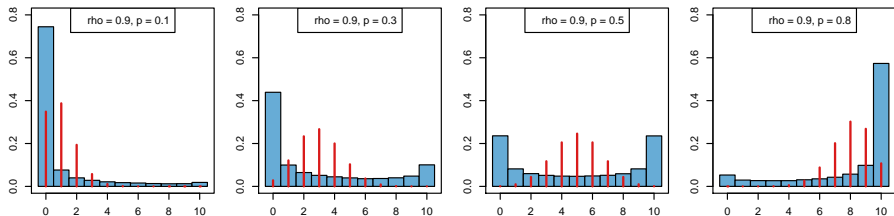
$p = 0.5, 0.1$



Intra-litter correlation results in overdispersed number of reactions per litter

Simulations Litter Effect and Quantal Data

How response probability changes distribution



probability p	0.1	0.3	0.5	0.8
mean	1	3	5	8
variance	0.52	1.32	1.60	0.98
binomial variance	0.09	0.21	0.25	0.16
over-dispersion	5.73	6.30	6.41	6.11

Intra-litter correlations induces nearly constant overdispersion

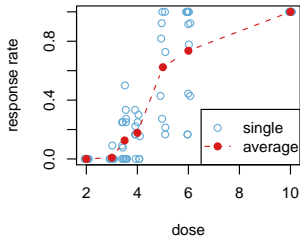
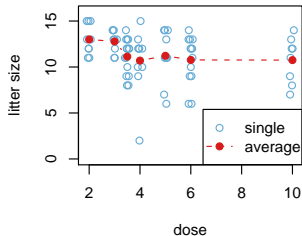


MNU Cleft palate induced by MNU

Raw data from Platzeck et al(1988)

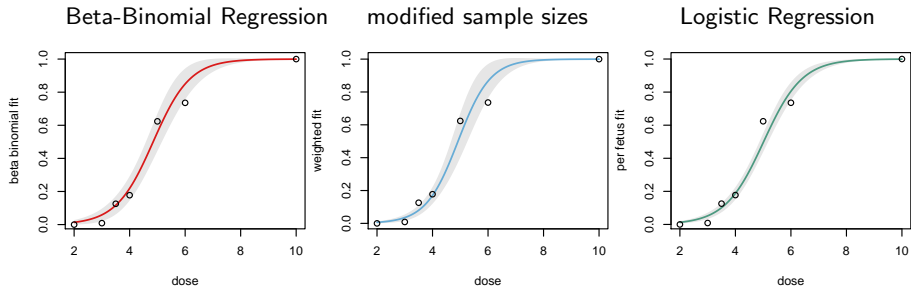
The Teratogenic Potency of MNU in Mice. Arch.Toxicol. 62: 411-423

dose	reactions/litter size	$\sum r_i / \sum n_i$	litter effect
2	0/11 0/11 0/12 0/12 0/13 0/13 0/15 0/15 0/15	0	
3	0/11 1/11 0/12 0/13 0/13 0/13 0/14 0/14 0/14	0.009	-
3.5	0/8 2/8 0/9 3/9 5/10 1/11 3/11 0/12 0/12 3/12 3/12 0/13 0/13 1/13 0/14	0.13	**
4	0/2 2/9 0/10 3/10 0/12 2/12 2/12 3/12 2/13 5/15	0.18	-
5	1/6 7/7 3/11 8/11 9/11 12/13 3/14 6/14 14/14	0.62	**
6	6/6 6/6 4/9 7/9 10/10 11/11 2/12 2/12 11/12 12/13 13/13 13/13 6/14	0.74	**
10	7/7 8/8 9/9 11/11 12/12 12/12 13/13 14/14	1	



Results: slightly decreasing litter size, increase in response rate

How different treatment of litter-effect changes fit



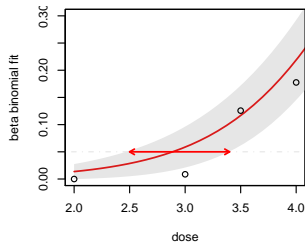
litter effects cause weighted fit to data
ignoring litter effects gives too narrow confidence bands



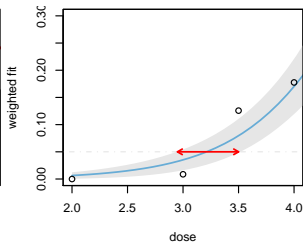
MNU Cleft palate induced by MNU

How different treatment of litter-effect changes benchmarks

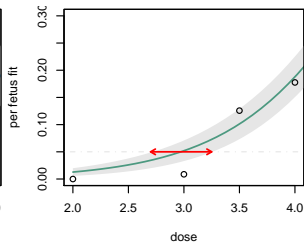
Beta-Binomial Regression



modified sample sizes



Logistic Regression



Selected method influences value **and** width of confidence interval
beta binomial fit recommended

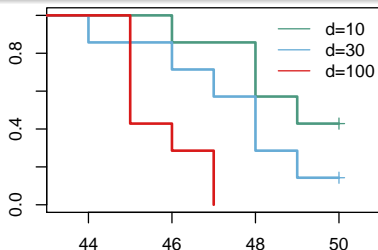


More Information Time-Dose-Response Models

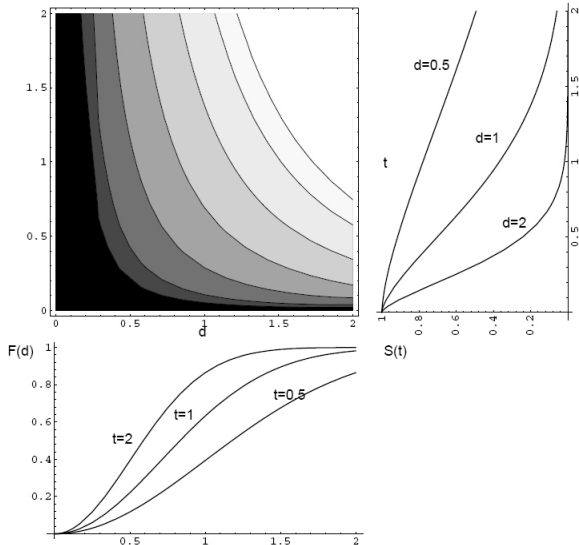
Example Labour Induction in Guinea-Pigs

Dose	Time (days post conception)								Total
	44	45	46	47	48	49	50	> 50	
10	0	0	1	0	2	1	0	3	7
30	1	0	1	1	2	1	0	1	7
100	0	4	1	2	0	0	0	0	7

Time to first abortion by dose of a Progesterone Antagonist (mg/a/d). Data from Elger (1999), pers. commun.



More Information Time-Dose-Response Models



Time-Dose-Response Models

Example: Moskito Fish (Newman/Huscher)

Zeitintervall bis Tod (in h)	Konzentration (in ppt)							Total
	0	10.3	10.8	11.6	13.2	15.8	20.1	
(. - 8]	0	0	0	0	0	9	77	86
(8 - 16]	0	0	0	0	0	26	0	26
(16 - 24]	0	0	0	0	3	28	0	31
(24 - 32]	0	0	0	1	10	11	0	22
(32 - 40]	0	0	2	2	10	2	0	16
(40 - 48]	0	5	6	8	10	2	0	31
(48 - 56]	0	2	2	9	10	0	0	23
(56 - 64]	0	2	5	4	10	0	0	21
(64 - 72]	0	3	3	7	5	0	0	18
(72 - 80]	0	1	0	2	6	0	0	9
(80 - 88]	0	2	0	3	4	0	0	9
(88 - 96]	0	1	4	4	1	0	0	10
<i>zensiert</i>	78	60	57	37	7	0	0	239
Total	78	76	79	77	76	78	77	541

Dörte Huscher (1999) Diploma Thesis TFH Berlin

„Statistische Methoden der Risikoabschätzung bei zeitabhängiger Exposition“

Data: Types of Recording

- C (quasi) continuous
- D discrete
- G grouped (*interval censored*)

Analysis: Types of Models

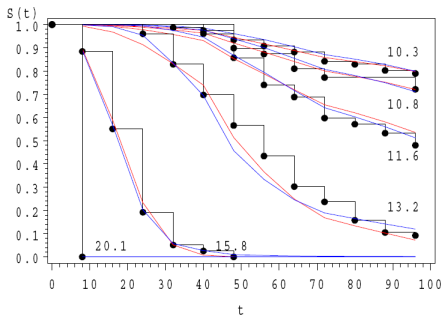
- parametric models C D G
accelerated failure time
glm with $f(\text{time})$ as covariate
- semi parametric models
Cox prop hazard C (D)
glm for ordered categorical G

Choice of model defines type of dose-relation.



Time-Dose-Response Models

Example: Moskito Fish (Newman/Huscher)



fit: glm ordered categorical

link=cloglog, $c^{2.1}$

→ prop. hazard

link=logit, $c^{2.8}$

→ prop. odds

$$S(t_i) = 1 - \text{link}^{-1}(\alpha_i + \beta c^k); \quad t_i \in \{8, 16, 24, 32, 40, 48, 56, 64, 72, 80, 88, 96\}$$



General Measure of Risk



differential effects

data Let $X_i \sim F_i$, $i = 1, 2$ be independent. Small values indicating *bad* state.

effect
$$p(X_1, X_2) = P(X_2 < X_1) + (1/2)P(X_2 = X_1)$$
$$= \int F_2(x) dF_1(x)$$
modified version of F for discontinuous distributions .

origin Brunner, Akritas *differential treatment effect*

differential effects are invariant under strictly monotonous transformations



differential effects for selected models

normal data $X_i \sim \Phi(\mu_i, \sigma^2), i = 1, 2$
$$p(X_1, X_2) = \Phi\{(\mu_1 - \mu_2)/(\sqrt{2}\sigma)\}$$

prop. hazards $1 - F_2(x) = \{1 - F_1(x)\}^\gamma$
$$p(X_1, X_2) = \gamma/(\gamma + 1)$$

binary data $X_i \sim B(1, q_i), i = 1, 2$
$$p(X_1, X_2) = (1/2) + (1/2)(p_2 - p_1)$$

differential effects are easily expressed as function of parameters



Dose-Response for Differential Effects

data $X_1 = X(0)$ controls, $X_2 = X(d)$ exposed

normalize $\pi(d) = 2\{p(X(0), X(d)) - 1/2\}$

normal data $\pi(d) = 2(\Phi\{(\mu(0) - \mu(d))/(\sqrt{2}\sigma)\} - 1/2)$

prop.hazard $\pi(d) = \{\gamma(d) - 1\}/\{\gamma(d) + 1\}$

binary data $\pi(d) = p(d) - p(0)$

normalized differential effects define risk function



Benchmark Doses for Normalized Differential Effects

parametric models use plug-in estimates of model parameters

ordered categorical use rank-sum estimate:

$$\hat{\pi}(d) = 2[(1/n_d)\{(1/n_0)\sum_{i=1}^{n_0} R_{0,i} - (n_0 + 1)/2\} - (1/2)]$$

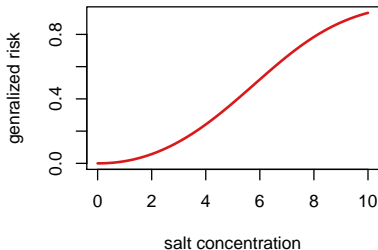
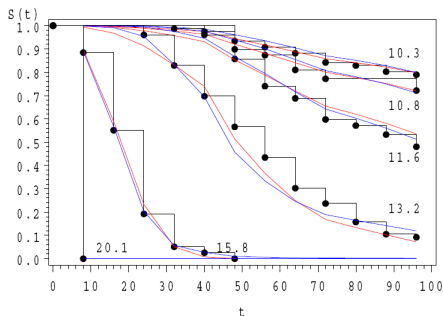
conf. bounds as usual, delta method, profile likelihood, bootstrap

benchmarks can be derived for differential effects



General Measure of Risk: Time-Dose-Response

Example: Moskito Fish (Newman/Huscher)



$$\pi(c) = \frac{\exp(0.0268 \times c^{2.1}) - 1}{\exp(0.0268 \times c^{2.1}) + 1}$$



- Benchmarks can be derived for quantal response data when (quasi) complete separation occurs. **ATTENTION** model dependence
- Simple simulation is a good tool for analyzing clustered binary data. Beta-binomial regression provides reliable models for teratological data.
- Event history modeling extracts **more** and **better** information than quantal response approach.
- **Stochastic order** expressed by normalized differential effects is a suitable general measure of risk.





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