## TESTING MODELS

- The Assumptions are valid
- Model Structure is sound
- Model Parameters are believable
- Model Predictions match observation

Important to remember that the model is an approximation - ignore less important features, "random errors" are expected.

## Assumptions

- Consider what the assumptions really are
- What should they be?
e.g. Do we want a linear relationship or any increasing relationship? Assumptions are often models and may need testing as such!



## Model Structure

How sensitive is the model to changes in model structure?

Quantitive changes (predicted value changes)

Qualitative changes (nature of prediction changes)
Some structures that can change outputs:
Stochasticity
Non-linearity
Modelling physical space explicitly
...!
Only qualitative (different behaviour) changes matter at this stage!

## Example: Changing Model Structure

Lotka-Volterra Predator Prey model:

Deterministic model gives stable cycles... Adding ANY noise produces unstable cycles!


## Parameters

Question: How sensitive is the model to the parameters?
Another: how do you find the "best" parameters?
Need Goodness of fit measure.
Most commonly used are:

- Sum-squared error
- Likelihood


## Estimating model parameters

## Deterministic models

- Minimise mean square error
- Assume normal and uncorrelated errors $\rightarrow$ std. errors
- Pretend model is true, discrepancy is observation error
- Likelihood $L(D \mid p)$ : probability that the observed errors could happen Likelihood $=\exp (-$ sum of [squared error/standard deviation] )

Stochastic models

- Likelihood $L(D \mid p)$ is a natural concept
- Only use model assumptions $\rightarrow$ parameter distributions
- Missing observations must be averaged out


## Estimating model parameters

To obtain "good" model parameters:

Maximise Likelihood (or minimise errors)
Many methods to do this:

- Solve for maximum (usually impossible!)
- Numerically find "Maximum Likelihood" (easy, but only local maxima found)
- Simulated Annealing (hard, should find global maxima)
- MCMC (hard, should find global maxima AND give correct parameter distribution around it)


## Sensitivity Analysis

(Sensitivity Analysis eds. Saltelli, Chan, Scott.)


## Sensitivity Analysis Methods

LOCAL
(directional) derivative:
$S_{i}=\frac{x_{j}}{y_{i}} \frac{\partial y_{i}}{\partial x_{j}}$
i.e. the relative change from making a small change to each parameter

GLOBAL

- Sampling based methods (Monte Carlo, Latin Hypercube Sampling)
- Sensitivity Indices (Importance Measures, SOBOL)
- FAST sampling (Fourier methods)

These are all implemented in the R package "sensitivity".

## Sensitivity Analysis Example



Relative change in output from each input

## Sensitivity Analysis Example 2

HeathMod Grazing System (MLURI)



## Prediction

Test model prediction against NEW and INDEPENDENT data! Why:

- We used the old data to fit the model - avoid overfitting
- Want general prediction - not for specific case
e.g. Fit model for a farm. Better to test on different farm, rather than new data for old farm. Otherwise only get good model for one specific farm!

Sometimes hard to get new data; instead, resample current data.

- Split data into chunks (e.g. leave one out)
- Randomize over which data used to fit model, which to predict from model


## Reasons for Prediction Errors

A) "Random" variation
B) Excluded effects, i.e. incomplete model
C) Wrong model:

- poor model structure
- poor parameter estimates


## Summary Statistics

Bias

$$
B=\sum_{i}\left(O_{i}-P_{i}\right) / n
$$

Standard deviation

$$
S D=\left\{\sum_{i}\left[\left(O_{i}-P_{i}\right)-B\right]^{2} / n\right\}^{\frac{1}{2}}
$$

Prediction mean square error

$$
P M S E=\sum_{i}\left(O_{i}-P_{i}\right)^{2} / n=B^{2}+S D^{2}
$$

## Comparison of Models

Issues to consider include:

- generality
- sensitivity
- predictive ability

NB equations with different functional forms can give similar predictions

Subjective element - what is it you want from the model?

## Information criterion

With Likelihood $L$, and using $k$ parameters for the model, consider:

- AIC (Akaike Information Criterion)
- BIC (Bayesian Information Criterion)
- DIC (Deviance Information Criterion)

Each has a slightly different form and is built on different assumptions - but usually agree

Many (stats) programs provide them routinely
Take the form $-2 \log (L)+f(n)$ ( $f=2 k$ for $\operatorname{AIC}, k \log (n)$ for BIC, where $n$ is the number of observations)

Choose minimum IC value; difference of 10 is very significant, $5-10$ is strong, $0-5$ means both models could be right

## Example: AIC

Model: linear model with 4 components: Each a factor a less important than the last, with noise.
Linear model: $y_{i}=\sum_{j=1: 4}\left(a^{j} x_{i j}+\sigma_{i j}\right)$



Use lowest AIC for predictions: Only use 2 variables. WHY? The noise swamps out the other 2 and so it isnt worth the extra complexity.

## Comparing models for cattle growth

Two models for prediction of liveweight gain in growing cattle

## Notation:

```
\(M E I=\) metabolisable energy of daily ration (MJ/d)
\(q \quad=\) ration of metabolisable to gross energy in the diet
\(E_{m} \quad=\) energy of maintenance ( \(\mathrm{MJ} / \mathrm{d}\) )
\(k_{m} \quad=\) efficiency of utilisation of dietary ME for maintenance
\(L=M E I * k_{m} / E_{m}\) (level of feeding)
\(E_{g} \quad=\) energy retained in daily weight change ( \(\mathrm{MJ} / \mathrm{d}\) )
\(k_{g} \quad=\) efficiency of utilisation of dietary ME for weight change
\(E V_{g} \quad=\) the energy value of tissue lost or gained ( \(\mathrm{MK} / \mathrm{kg}\) )
\(W \quad=\) liveweight (kg)
\(\Delta W=\) liveweight change ( \(\mathrm{kg} / \mathrm{d}\) )
```


## Comparing models for cattle growth

## General

The daily energy balance in growing cattle may be represented as follows:

$$
\begin{equation*}
M E I=\frac{E_{m}}{k_{m}}+\frac{E_{g}}{k_{g}} \tag{1}
\end{equation*}
$$

writing $E_{g}=E V_{g} \times \Delta W$ we obtain

$$
\begin{equation*}
M E I=\frac{E_{m}}{k_{m}}+\frac{E V_{g} \times \Delta W}{k_{g}} \tag{2}
\end{equation*}
$$

## Comparing models for cattle growth

 MODEL 1: (AFRC 1980)$$
\begin{aligned}
& E_{m}=0.53(W / 1.08)^{0.67}+0.0043 W \\
& k_{m}=0.35 q+0.503 \\
& k_{f}=0.78 q+0.006 \\
& k_{g}=\frac{k_{f}}{L-1} \\
& E V_{g}= \frac{\left(4.1+0.0332 W-0.000009 W^{2}\right)}{1-0.1475 \Delta W} \\
& \Delta W=\frac{E_{g}}{4.1+0.0332 W-0.000009 W 2+0.1475 E_{g}} \\
& E_{g}=k_{g} \times\left(M E I-\frac{E_{m}}{k_{m}}\right) .
\end{aligned}
$$

## Comparing models for cattle growth MODEL 2: (TB33)

$$
\begin{aligned}
E_{m} & =5.67+0.061 W \\
k_{m} & =0.72 \\
k_{g} & =0.9 q \\
E V_{g} & =6.28+0.3 E_{g}+0.0188 W
\end{aligned}
$$

Rearranging (2) and substituting for $E V_{g}$ gives

$$
\Delta W=E_{g} /\left(6.28+0.0188 W+0.3 E_{g}\right)
$$

where $E_{g}=k_{g} \times\left(M E I-E_{m} / k_{m}\right)$.

## Comparing models for cattle growth



## Comparing models for cattle growth

Table A
Predictions of liveweight gains (g/d) according to the competing models

|  | W $(\mathrm{kg})$ |  | 100 |  |  |  |  |  |  |  | 500 |  |  |
| :--- | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :--- | :---: | :---: | :---: | :---: | :---: |
| $M E I(\mathrm{MJ} / \mathrm{d})$ | 20 |  | 40 |  | 60 | 100 |  |  |  |  |  |  |  |
| $q$ | .46 | .68 | .57 | .68 | .46 | .68 | .57 | .68 |  |  |  |  |  |
| Model 1 | 170 | 351 | 1112 | 1322 | 197 | 408 | 1044 | 1239 |  |  |  |  |  |
| Model 2 | 157 | 226 | 947 | 1070 | 215 | 307 | 1010 | 1137 |  |  |  |  |  |

## Comparing models for cattle growth

Table B
Bias in predicting liveweight gain (g/d) using independent data (standard deviations in parentheses)

|  | Data Set | Mean liveweight gain (g/d) | Model 1 | Model 2 |
| :--- | :---: | :---: | :---: | :---: |
| 1. | (Food, Reading) | 1080 | $210(80)$ | $130(80)$ |
| $2 a$. | (Hinks, Edinburgh) | 660 | $180(100)$ | $150(100)$ |
| $2 b$. |  | 890 | $130(180)$ | $100(170)$ |
| $2 c$. |  | 820 | $70(120)$ | $50(120)$ |
| $2 d$. | 970 | $160(130)$ | $220(130)$ |  |
| 3. | (Drayton EHF) | 730 | $-10(120)$ | $0(120)$ |
| 4. | (MLC, Nottingham) | 910 | $170(220)$ | $140(210)$ |

## Working Party Report

Concludes there is little difference in predictive ability Recommends Model 1 because:
a) no need for linearizing approximations of Model 2
b) includes terms for known effects absent in Model 1
c) better platform for future development

## Comparison of two models via precision of parameter estimates

Likelihoods for biological control parameter


Treatment 1 supports model including the biocontrol parameter $b$
Treatment 2 supports simpler model without $b$

