

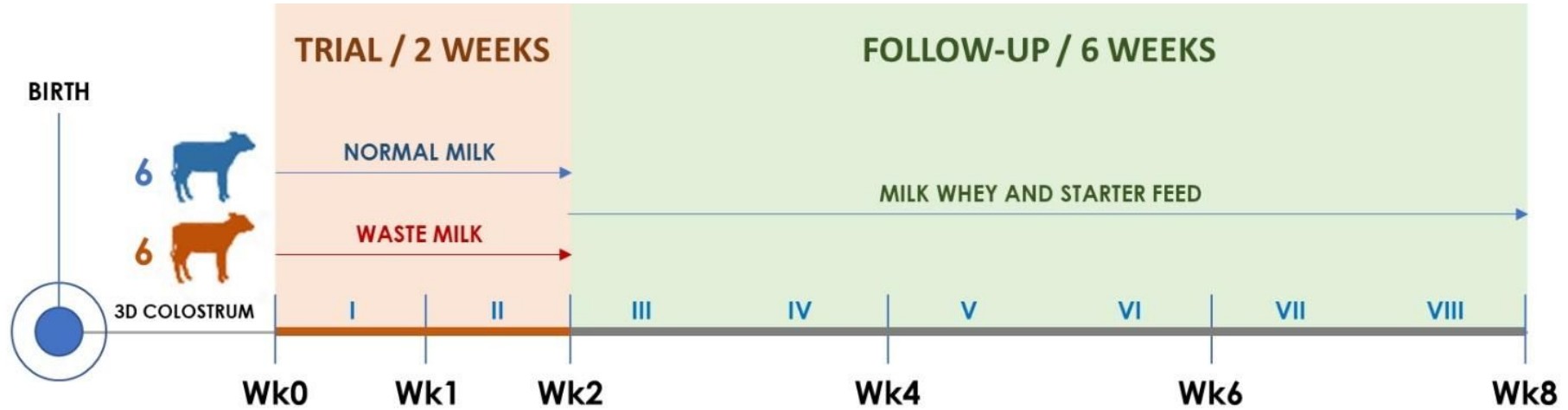
The basic experimental setting: treatments and timepoints

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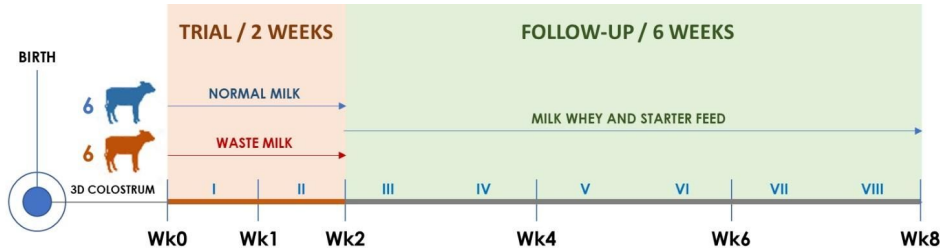
Treatments and **timepoints**



From: [Penati et al. 2021](#) (Front. Vet. Sci., 08 July 2021)



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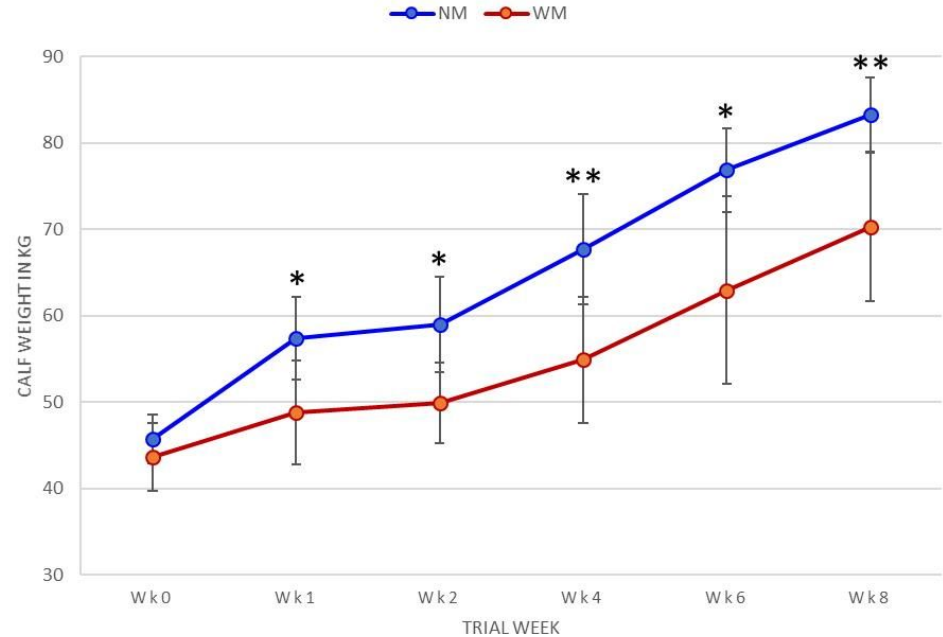
- treatments: groups
- timepoints: longitudinal (time) component

- the effect of the treatment(s) is not expected at T0
- we may observe the effect of the treatment later in time, progressively stronger
- the effect may appear and then subside
- there may be no effect of the treatment
- etc.



Treatments and timepoints

Effect of treatment apparent from after one week, then constant or intensifies



* p < 0.05

** p < 0.005

From: [Penati et al. 2021](#) (Front. Vet. Sci., 08 July 2021)

Treatments and timepoints

- Is there an effect of treatment on the response variable?
 - e.g. drug vs placebo for disease response
- How strong is the effect of treatment?
 - e.g. how fast / how much does the drug reduce symptoms vs placebo?
- How accurately can we predict future/unobserved response variables?
 - e.g. date of remission based on treatment; fish growth based on diet; etc.
- Does the effect of treatment change with time?
 - e.g. lag of drug to combat disease; faster effect at first, decreasing returns with time; reversed effect of antimicrobial if continued for too long; etc.



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Q: can you think of additional examples from your own research? Let's discuss them!



Simple **linear regression**

$$y = \beta_0 + \beta_1 x + e$$

Q: are you familiar with this equation?



Simple **linear** regression

$$y = \beta_0 + \beta_1 x + e$$



$$\text{response} = \beta_0 + \beta_1 \text{treatment} + e$$



Simple **linear** regression

$$y = \beta_0 + \beta_1 x + e$$

Q: how can we estimate the model coefficients (β_0 , β_1)?



Least squares

Minimise the residuals → Q: how do we calculate the residuals of the model?
→ Q: what happens to the sum of the residuals?
→ Q: what do we do then?



Least squares

Minimise the residuals → Q: how do we calculate the residuals of the model?
→ Q: what happens to the sum of the residuals?
→ Q: what do we do then?

residuals = $(y - (b_0 + b_1x))$
 $\text{sum}(y - (b_0 + b_1x)) = 0$

$$\text{RSS} = \sum_{i=1}^N (y_i - (\beta_0 + \beta_1 x_i))^2 \quad \left. \vphantom{\sum_{i=1}^N} \right\} \text{to be minimised}$$



Least squares: **the normal equations**

$$\beta_1 = \frac{\sum (x_i - \bar{x})(y_i - \bar{y})}{\sum (x_i - \bar{x})^2}$$

$$\beta_0 = \bar{y} - \beta_1 \bar{x}$$

We can use a more compact matrix notation:

$$\mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{e}$$

where \mathbf{y} , \mathbf{b} , and \mathbf{e} are vectors of responses, coefficients and residuals; \mathbf{X} is the design matrix

e.g. $\mathbf{b} = [\mathbf{b}_0, \mathbf{b}_1]$

$$\mathbf{X}'\mathbf{X}\mathbf{b} = \mathbf{X}'\mathbf{y}$$

$$\mathbf{b} = (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}'\mathbf{y}$$

(this is generalizable to multiple linear regression)



Estimating coefficients: alternatives

1. Maximise the likelihood function

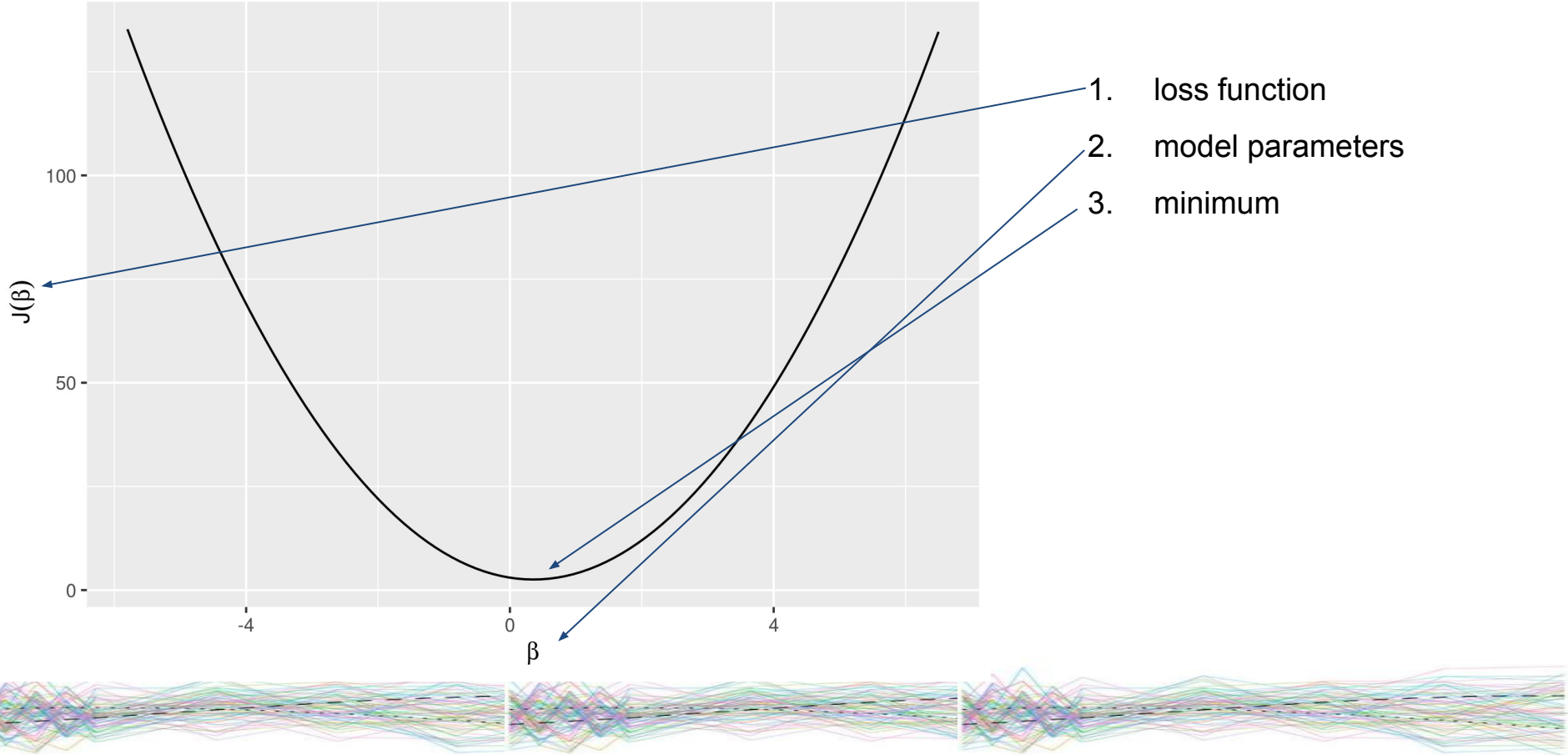
$$\mathcal{L}(\beta, \sigma^2, y, X) = (2\pi\sigma^2)^{-\frac{N}{2}} \exp\left(-\frac{1}{2\sigma^2} \sum (y_i - x_i\beta)^2\right)$$

2. Minimise the loss function

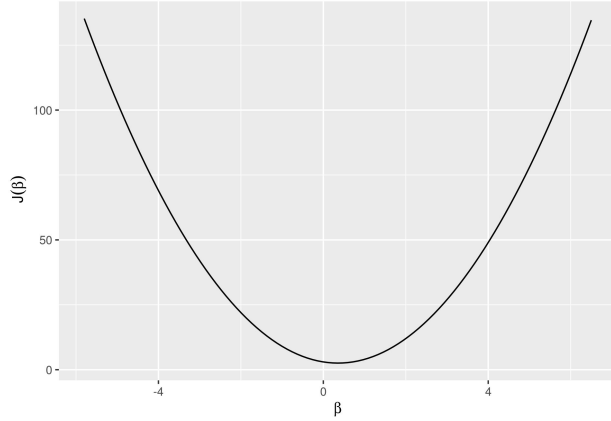
$$J(\beta) = \frac{1}{2n} \sum_{i=1}^n (\beta_i X_i - y_i)^2$$



Minimise loss (maximise likelihood)



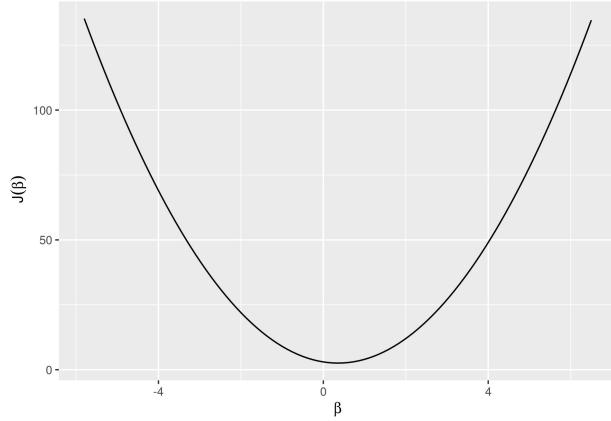
Minimise loss (maximise likelihood)



1. Start with initial values for β : (initialisation)
2. Change β in the direction of reducing $J(\beta)$ / increasing $L(\beta)$: (descent/ascent)
3. Stop when the minimum/maximum is reached : (min/maximisation)



Minimise loss (maximise likelihood)



This is an iterative process that progressively updates the coefficients (unlike least squares that offers a closed form solution)!

1. Start with initial values for β : (initialisation)
2. Change β in the direction of reducing $J(\beta)$ / increasing $L(\beta)$: (descent/ascent)
3. Stop when the minimum/maximum is reached : (min/maximisation)



How reliable are the estimated coefficients?



- Standard errors of the coefficients

$$SE(\hat{\beta}_1) = \sqrt{\frac{\sigma^2 = Var(e)}{\sum (x_i - \bar{x})^2}} = \frac{\sigma}{\sqrt{\sum (x_i - \bar{x})^2}}$$

- Confidence intervals

$$\hat{\beta}_1 \pm z \cdot SE(\hat{\beta}_1)$$

- Hypothesis test

$$t = \frac{\hat{\beta}_1 - 0}{SE(\hat{\beta}_1)} \longrightarrow \text{p-value}$$



Model accuracy

$$\text{RSE} = \sqrt{\frac{1}{n-2} \sum (y_i - \hat{y}_i)^2}$$

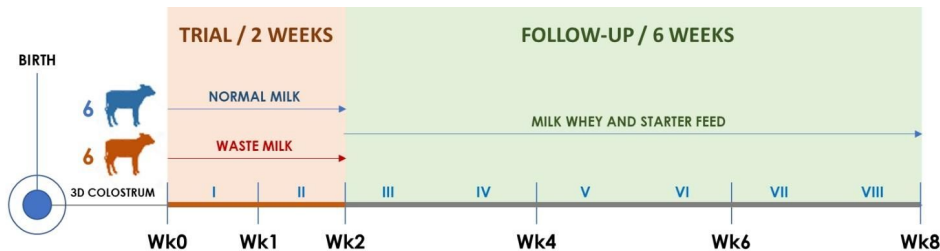
Residual standard error: Q: what value do we wish for it?

$$R^2 = 1 - \frac{\text{RSS}}{\text{TSS}} = 1 - \frac{\sum (y_i - \hat{y}_i)^2}{\sum (y_i - \bar{y})^2}$$

Coefficient of determination: normalized RSE Q: what value do we wish for it?



Treatments and timepoints: **linear regression**



From: [Penati et al. 2021](#) (Front. Vet. Sci., 08 July 2021)

Apply linear regression within-time point:

$$y = b_0 + b_1 * \text{treatment} + e$$

replicate the model for each time point

PROS:

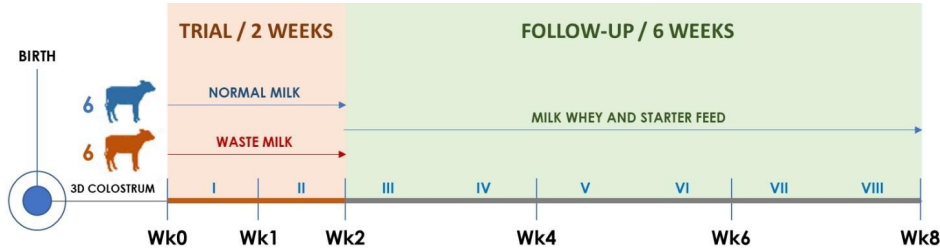
- simple to run
- easy to interpret
- may be your research hypothesis

CONS:

- smaller sample size
- fail to account for changing effect of treatment over time
- may not be your research hypothesis



Across-timepoints analysis



From: [Penati et al. 2021](#) (Front. Vet. Sci., 08 July 2021)

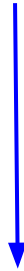
Previously, we saw that we can fit separate regression models for each timepoint

- each separate regression equation ignores the other timepoints: if timepoints are correlated (time trend) this can lead to misleading estimates of the association between treatment and response
- it can be difficult to combine results and interpret results across timepoints (independent models)



Across-timepoints analysis

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + e$$



$$\text{response} = \beta_0 + \beta_1 \text{timepoint} + \beta_2 \text{treatment} + e$$



Across-timepoints analysis

$$\text{response} = \beta_0 + \beta_1 \text{timepoint} + \beta_2 \text{treatment} + e$$

We interpret coefficients “ceteris paribus”:

- b_0 : mean of reference classes (e.g. controls, T0)
- b_1 : $\text{mean}(T1) - \text{mean}(T0)$ [and so on]
- b_2 : $\text{mean}(\text{treated}) - \text{mean}(\text{controls})$
- $b_0 + b_1$: $\text{mean}(\text{controls})$ at T1
- $b_0 + b_2$: $\text{mean}(\text{treated})$ at T0
- $b_0 + b_1 + b_2$: $\text{mean}(\text{treated})$ at T1



Interaction model

Removing the **additive assumption**: x_1 and x_2 are assumed to be independent
But are they?

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 (x_1 \cdot x_2) + e$$

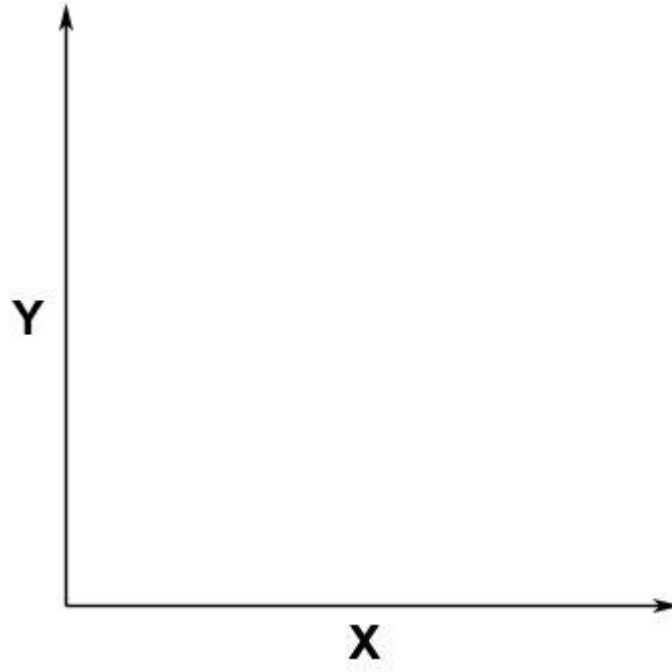


$$\text{resp} = \beta_0 + \beta_1 \text{time} + \beta_2 \text{treat} + \beta_3 (\text{time} \times \text{treat}) + e$$



Interaction model

Interaction plot



Interaction model

hierarchical principle : if we include an interaction in a model we always have to also include the main effects, even if the p-values associated with their coefficients are not significant



(no models with only interaction terms!)

