

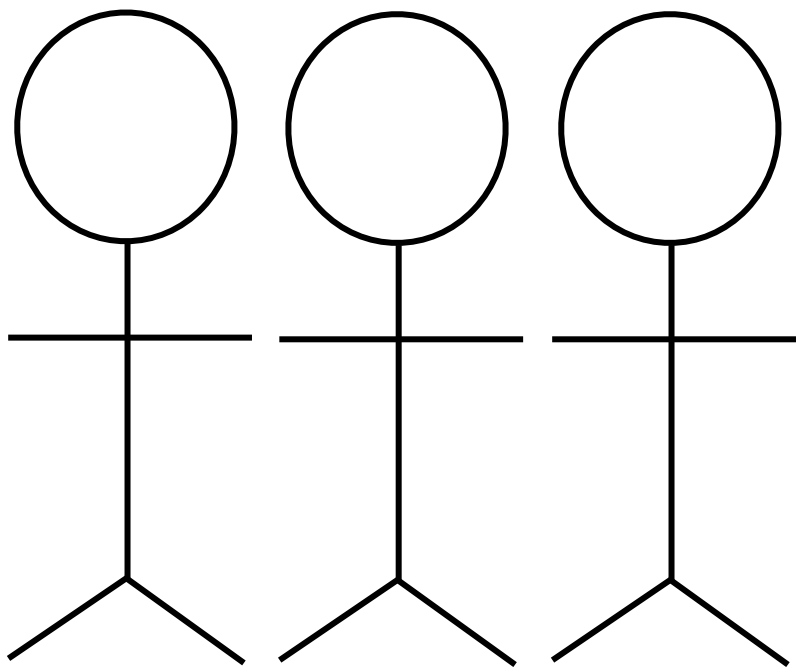
Computing information criteria for multilevel models using different sample sizes

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School 1

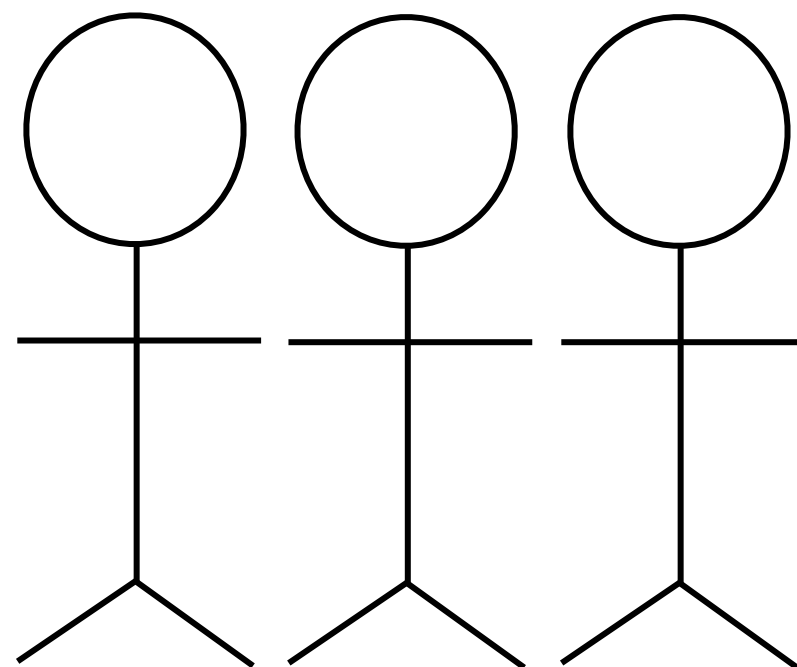
School 2



S1

S2

S3



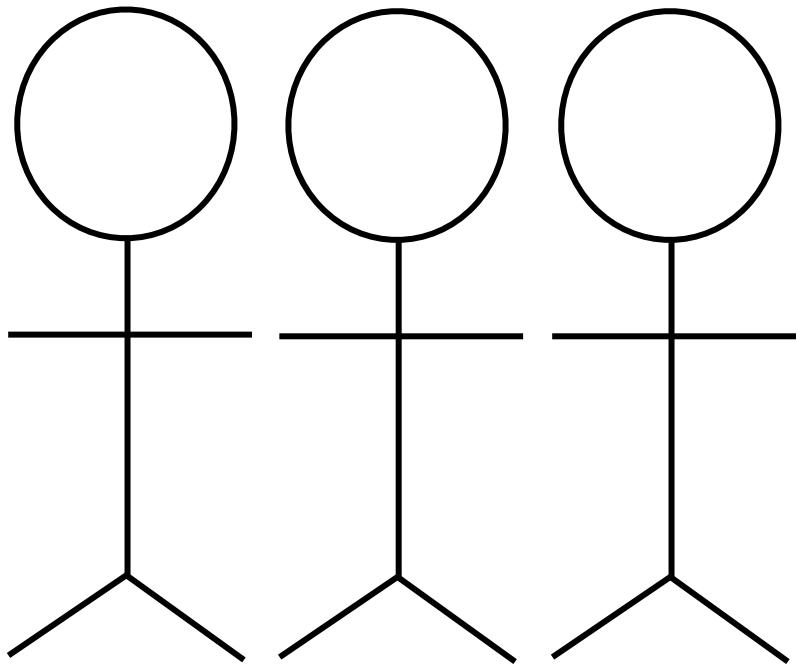
S4

S5

S6

School 1

School 2

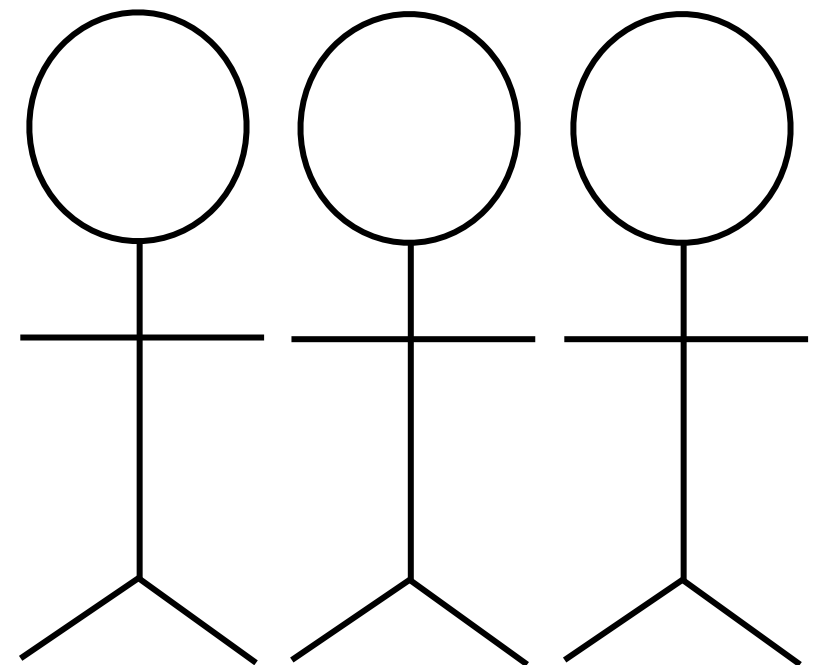


S1

S2

S3

N = 6



S4

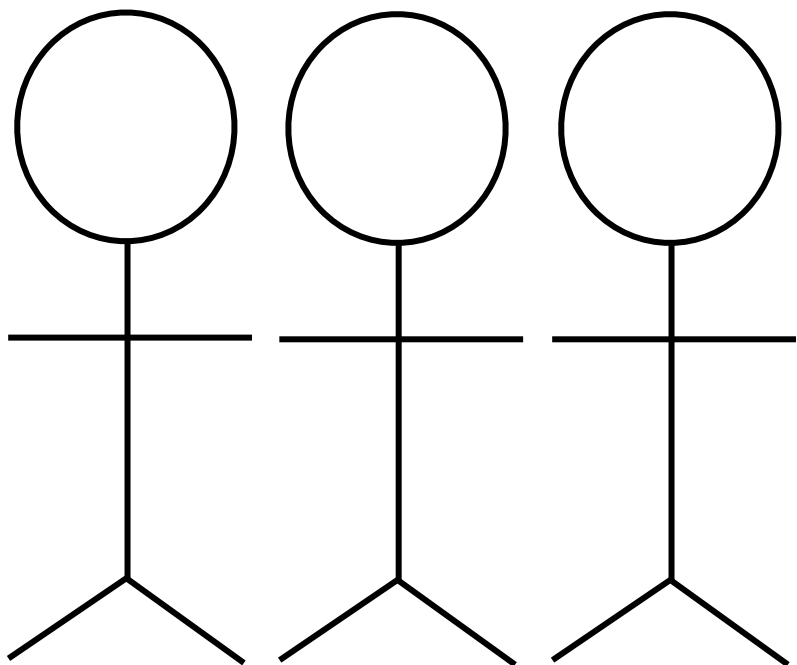
S5

S6

School 1

$m = 2$

School 2

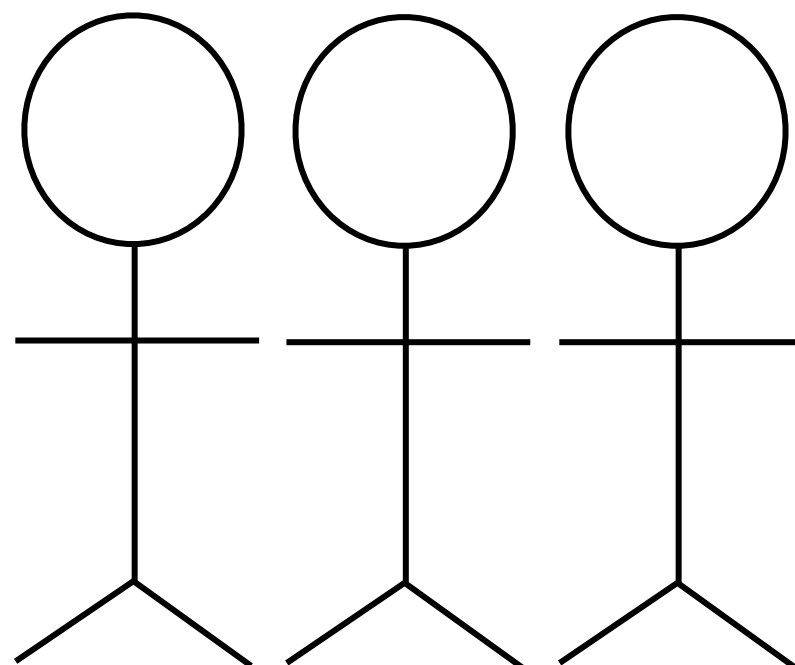


S1

S2

S3

$N = 6$



S4

S5

S6

Model selection using information criteria

①

Model 1

Model 2

Model 3

Model selection using information criteria

①

Model 1

Model 2

Model 3



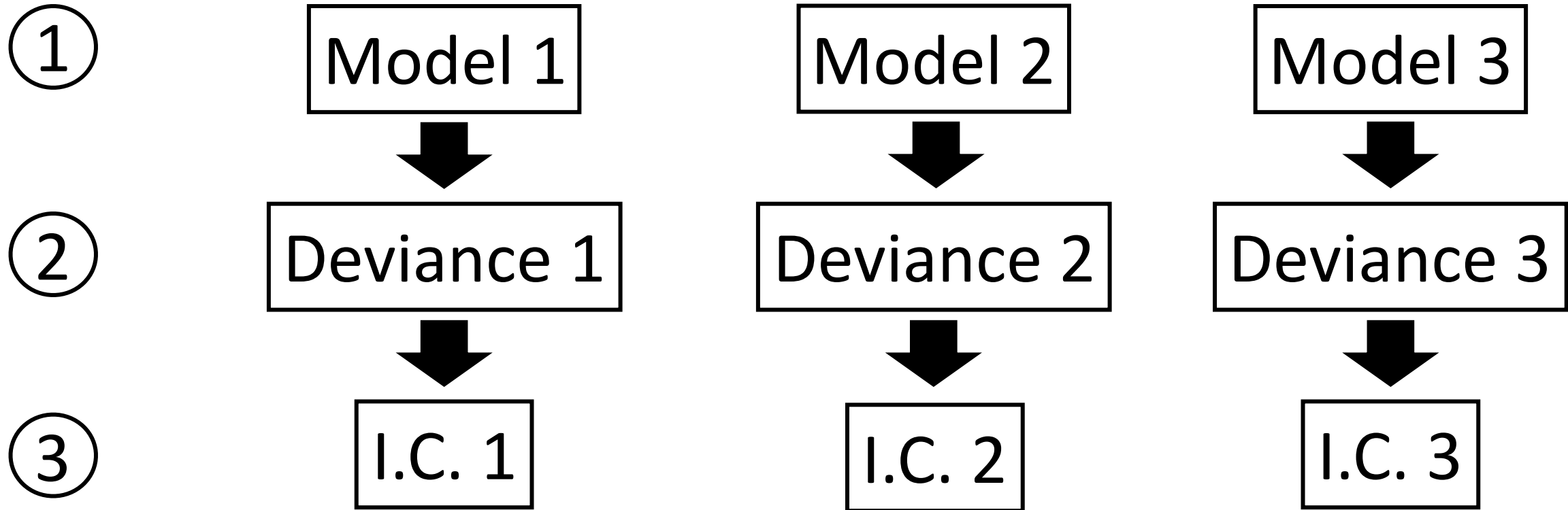
②

Deviance 1

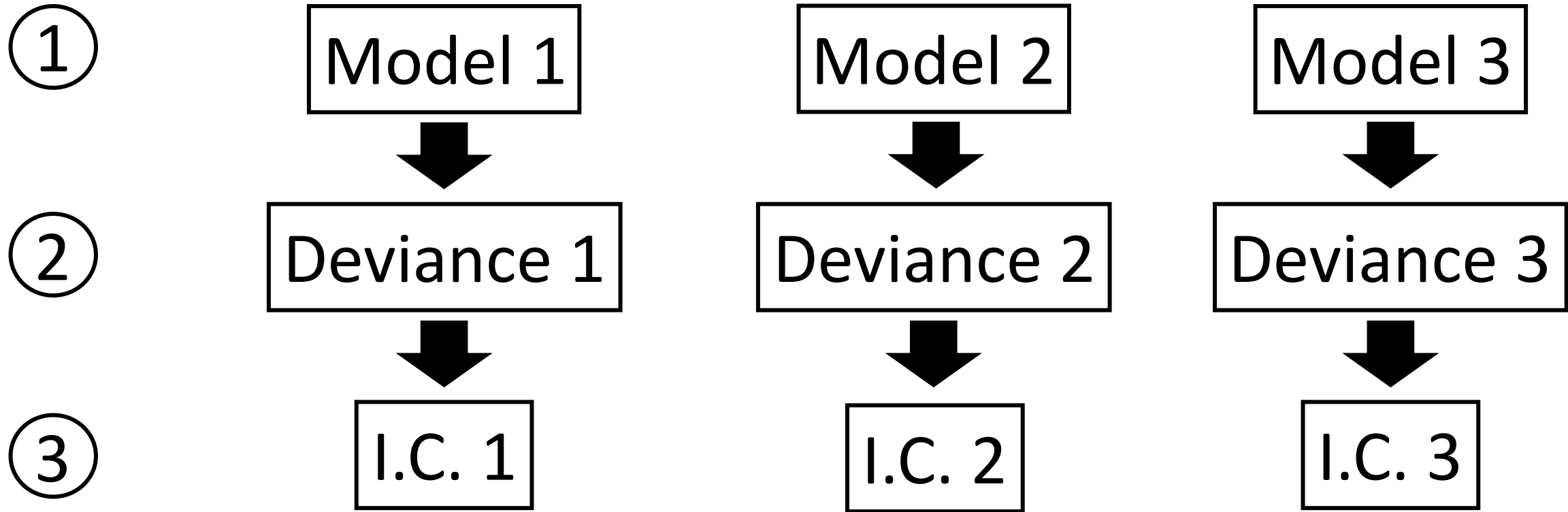
Deviance 2

Deviance 3

Model selection using information criteria



Model selection using information criteria



④ Smallest I.C. value = “best” model

Choice of sample size matters!

AIC

Deviance + $2*k$

BIC

Deviance + $\log(N)*k$

Choice of sample size matters!

AIC

Deviance + $2*k$

BIC

Deviance + $\log(N)*k$

Model deviance



Choice of sample size matters!

AIC

Deviance + $2*k$

BIC

Deviance + $\log(N)*k$

Model deviance

Number of model parameters



Choice of sample size matters!

AIC

Deviance + $2 * k$

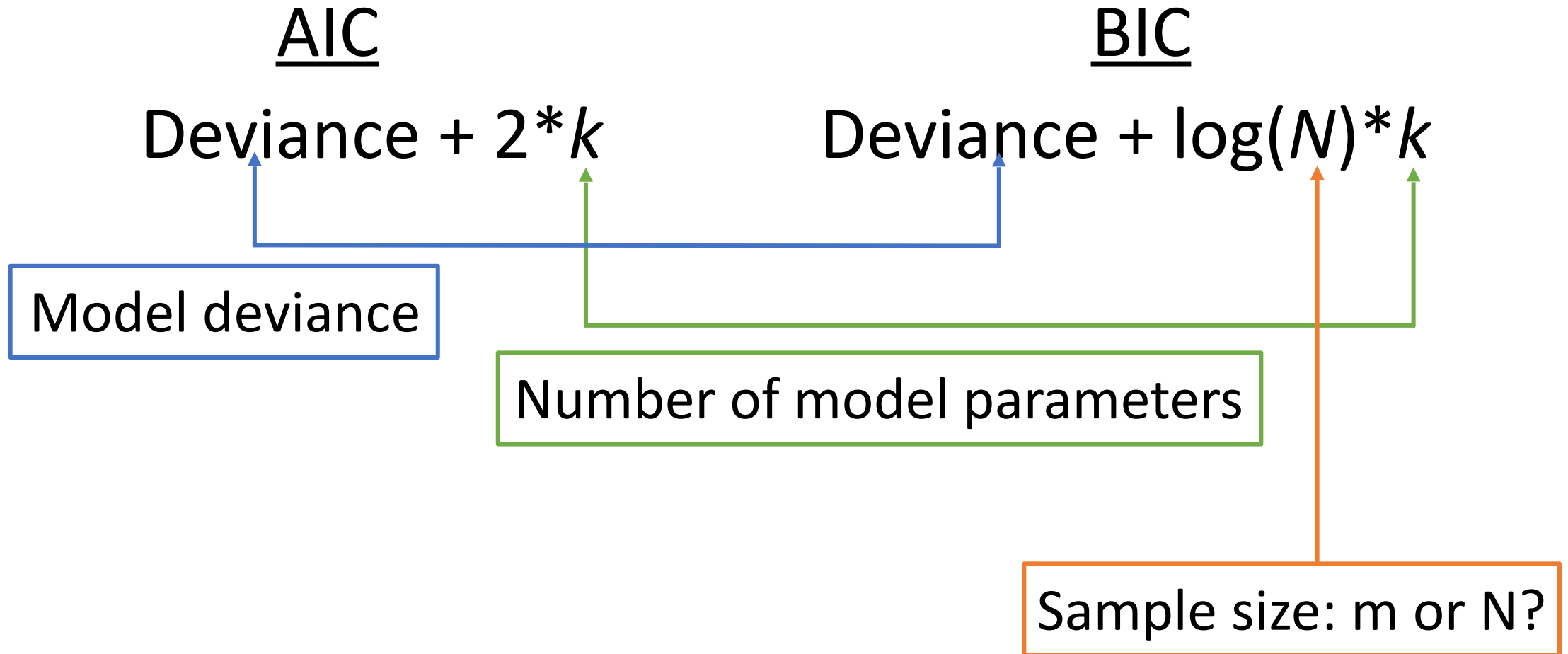
BIC

Deviance + $\log(N) * k$

Model deviance

Number of model parameters

Sample size: m or N?



Example using the nlme package^{[1][2]}

Information criteria

Parameters

N

m

```
Linear mixed-effects model fit by maximum likelihood
Data: alcohol1
      AIC      BIC    logLik
676.156 686.672 -335.078

Random effects:
Formula: ~1 | id
      (Intercept)  Residual
StdDev:   0.7509085 0.7494974

Fixed effects: alcuse ~ 1
              Value Std.Error DF t-value p-value
(Intercept) 0.9219549 0.09590253 164 9.613458      0

Standardized Within-Group Residuals:
      Min       Q1       Med       Q3      Max
-1.8865485 -0.3075977 -0.3066575  0.6136653  2.8567495

Number of Observations: 246
Number of Groups: 82
```

[1] Singer & Willett (2003): *Applied Longitudinal Data Analysis*

[2] UCLA Statistical Consulting Group: <https://stats.idre.ucla.edu/r/examples/alda/r-applied-longitudinal-data-analysis-ch-4/>

Computing BIC using m (number of clusters)

```
str(model) ←  
  
logLikelihood <- model$logLik  
deviance <- -2*logLikelihood  
  
groups <- head(model$dim$ngrps,n=1)  
m <- unname(groups, force = FALSE)  
  
N <- model$dim$N  
  
BIC_m <- deviance+log(m)*3  
  
BIC_N <- deviance+log(N)*3
```

Examine to find necessary
elements from output

Computing BIC using m (number of clusters)

```
str(model)
```

```
logLikelihood <- model$logLik  
deviance <- -2*logLikelihood
```

```
groups <- head(model$dims$ngrps, n=1)  
m <- unname(groups, force = FALSE)
```

```
N <- model$dims$N
```

```
BIC_m <- deviance+log(m)*3
```

```
BIC_N <- deviance+log(N)*3
```

Isolate model log-likelihood



Compute model deviance

Computing BIC using m (number of clusters)

```
str(model)
```

```
logLikelihood <- model$logLik
```

```
deviance <- -2*logLikelihood
```

```
groups <- head(model$dims$ngrps, n=1)
```

```
m <- unname(groups, force = FALSE)
```

```
N <- model$dims$N
```

```
BIC_m <- deviance+log(m)*3
```

```
BIC_N <- deviance+log(N)*3
```



Obtain number of clusters

Computing BIC using m (number of clusters)

```
str(model)

logLikelihood <- model$logLik
deviance <- -2*logLikelihood

groups <- head(model$dim$ngrps,n=1)
m <- unname(groups, force = FALSE)

N <- model$dim$N

BIC_m <- deviance+log(m)*3
BIC_N <- deviance+log(N)*3
```

Compute BIC(m)

683.376

Computing BIC using N (total observations)

```
str(model)

logLikelihood <- model$logLik
deviance <- -2*logLikelihood

groups <- head(model$dims$ngrps, n=1)
m <- unname(groups, force = FALSE)

N <- model$dims$N

BIC_m <- deviance+log(m)*3

BIC_N <- deviance+log(N)*3
```

Matches output!

686.6719...

AIC	BIC	logLik
676.156	686.672	-335.078

Thanks!

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Code available at: <https://github.com/wendychristensen>