

### Some statistical models and corresponding SAS software code

Completely randomized (CR), one factor	$Y_{ij} = \mu + \alpha_i + e_{ij}, \quad e_{ij} \sim N(0, \sigma^2)$	class alpha ; model y = alpha;
Randomized complete block (RCB), one factor	$Y_{ij} = \mu + \alpha_i + b_j + e_{ij}, \quad b_j \sim N(0, \sigma_b^2),$ $e_{ij} \sim N(0, \sigma_e^2)$	class alpha block; model y = alpha; random block;
CR, two (crossed) factors	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + e_{ijk}$ $e_{ijk} \sim N(0, \sigma_e^2)$	class alpha beta block; model y = alpha beta;
RCB, two factors	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + e_{ijk}$ $b_k \sim N(0, \sigma_b^2), e_{ijk} \sim N(0, \sigma_e^2)$	class alpha beta block; model y = alpha beta; random block;
CR, split plot (in designs in table without a block effect [as here], “block” in SAS code is a label for the replicate experimental unit)  Equivalence of two models holds when $\sigma_d^2$ is positive (same for the RCB, below)	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + d_{ik} + e_{ijk}$ $d_{ik} \sim N(0, \sigma_d^2), e_{ijk} \sim N(0, \sigma_e^2)$  $Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + e_{ijk}$ $e_{ijk} \sim N(0, \mathbf{R})$ <b>R:</b> var.-cov.matrix (fixed corr. or covariance)	class alpha beta block; model y = alpha beta; random block*alpha;  --or, equivalently--  model y = alpha beta; repeated /sub=block*alpha type=cs;
RCB, split plot	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + d_{ik} + e_{ijk}$ $b_k \sim N(0, \sigma_b^2), d_{ik} \sim N(0, \sigma_d^2), e_{ijk} \sim N(0, \sigma_e^2)$  $Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + e_{ijk}$ $b_k \sim N(0, \sigma_b^2), e_{ijk} \sim N(0, \mathbf{R})$ <b>R:</b> var.-cov.matrix (fixed corr. or covariance)	class alpha beta block; model y = alpha beta; random block block*alpha;  --or, equivalently--  model y = alpha beta; random block; repeated /sub=block*alpha type=cs;
CR, <u>repeated measures</u> (time: $\beta$ ), with one experimental factor ( $\alpha$ ); Covariance: CS, AR(1), UN, CSH, ARH(1), etc., etc.	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + e_{ijk}$ $e_{ijk} \sim N(0, \mathbf{R})$  <b>R:</b> var.-cov.matrix (many possibilities)	class alpha beta block; model y = alpha beta / ddfm=kr; repeated /sub=block*alpha type=■;
RCB, <u>repeated measures</u> (time: $\beta$ ), with one experimental factor ( $\alpha$ ); Covariance: CS, AR(1), UN, CSH, ARH(1), etc., etc.	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + e_{ijk}$ $b_k \sim N(0, \sigma_b^2), e_{ijk} \sim N(0, \mathbf{R})$  <b>R:</b> var.-cov.matrix (many possibilities)	class alpha beta block; model y = alpha beta / ddfm=kr; random block; repeated /sub=block*alpha type=■;
RCB, <u>repeated measures</u> (time: $\beta$ ), with one experimental factor ( $\alpha$ ); Covariance: CS+AR(1) [special case]	$Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + d_{ik} + e_{ijk}$ $b_k \sim N(0, \sigma_b^2), d_{ik} \sim N(0, \sigma_d^2), e_{ijk} \sim N(0, \mathbf{R})$  <b>R:</b> var.-cov.matrix (autoregressive error only)	class alpha beta block; model y = alpha beta / ddfm=kr; random block block*alpha; repeated /sub=block*alpha type=ar(1);
RCB, <u>repeated measures</u> (time: $\beta$ ), with <u>two</u> other experimental factors ( $\alpha$ and $\gamma$ ); Covariance: CS, AR(1), UN, CSH, ARH(1), etc., etc.  [take out “random block” if CR]	$Y_{ijk} = \mu + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} +$ $(\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + e_{ijkl},$ $b_k \sim N(0, \sigma_b^2), e_{ijkl} \sim N(0, \mathbf{R})$  <b>R:</b> var.-cov.matrix (many possibilities)	class alpha beta gamma block; model y = alpha beta gamma / ddfm=kr; random block; repeated /sub=block*alpha*gamma type=■;

<p>Same, but with gamma as <i>fixed</i> location effect (and <i>randomized blocks within locations</i>—note the two subscripts on b)</p>	$Y_{ijk} = \mu + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_{kl} + e_{ijkl},$ $b_{kl} \sim N(0, \sigma_b^2), e_{ijkl} \sim N(0, \mathbf{R})$ <p><b>R</b>: var.-cov.matrix (many possibilities)</p>	<pre>class alpha beta gamma block; model y = alpha beta gamma /       ddfm=kr; random block(gamma); repeated /sub=block*alpha*gamma       type=■;</pre>
<p>Same, but with gamma as <i>random</i> location effect (and randomized blocks within locations)</p> <p>[a random location effect is very different from previous, and here we only consider some random-effect interactions]</p>	$Y_{ijk} = \mu + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + b_{kl} + e_{ijkl},$ $\gamma_l \sim N(0, \sigma_\gamma^2), (\alpha\gamma)_{il} \sim N(0, \sigma_{\alpha\gamma}^2),$ $(\beta\gamma)_{jl} \sim N(0, \sigma_{\beta\gamma}^2), b_{kl} \sim N(0, \sigma_b^2),$ $e_{ijkl} \sim N(0, \mathbf{R})$	<pre>class alpha beta gamma block; model y = alpha beta /       ddfm=kr; random gamma block(gamma) gamma*alpha gamma*beta; repeated /sub=block*alpha*gamma       type=■;</pre>
<p><u>RCB, split-split plot</u></p> <p>Whole-plot (<math>\alpha</math>)</p> <p>Sub-plot (<math>\beta</math>)</p> <p>Sub-sub plot (<math>\gamma</math>)</p> <p>[equivalence when variance terms are positive]</p>	$Y_{ijkl} = \mu + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + f_{ijk} + e_{ijkl},$ $b_k \sim N(0, \sigma_b^2), d_{ik} \sim N(0, \sigma_d^2),$ $f_{ijk} \sim N(0, \sigma_f^2), e_{ijkl} \sim N(0, \sigma_e^2)$ $Y_{ijkl} = \mu + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + e_{ijkl},$ $b_k \sim N(0, \sigma_b^2), d_{ik} \sim N(0, \sigma_d^2),$ $e_{ijkl} \sim N(0, \mathbf{R})$ <p><b>R</b>: var.-cov.matrix (many possibilities)</p>	<pre>class alpha beta gamma block; model y = alpha beta gamma /       ddfm=kr; random block block*alpha       block*alpha*beta;  --or, equivalently--  class alpha beta gamma block; model y = alpha beta gamma /       ddfm=kr; random block block*alpha; repeated / sub=block*alpha*beta       type=cs;</pre>
<p><u>RCB, split-plot+time</u></p> <p>Whole-plot (<math>\alpha</math>)</p> <p>Sub-plot (<math>\beta</math>)</p> <p>Time (<math>\gamma</math>)</p>	$Y_{ijkl} = \mu + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + e_{ijkl},$ $b_k \sim N(0, \sigma_b^2), d_{ik} \sim N(0, \sigma_d^2),$ $e_{ijkl} \sim N(0, \mathbf{R})$ <p><b>R</b>: var.-cov.matrix (many possibilities)</p>	<pre>class alpha beta gamma block; model y = alpha beta gamma /       ddfm=kr; random block block*alpha; repeated / sub=block*alpha*beta       type=■;</pre>

Unless indicated otherwise, Greek letters indicate fixed-effect terms (treatment, factor A [ $\alpha$ ], time or factor B [ $\beta$ ], etc.), and  $b$  is the random block effect.

For all models before the RCB split-split-plot, the beta ( $\beta$ ) term always refers to time, which is clustered within the subjects identified by block\*alpha, or for more factors, block\*alpha\*gamma.