

**Some statistical models and corresponding SAS software code (Normal [left] or binomial [right])**

Design	LMM (Normal cond. dist.)	SAS code (LMM; normal)	GLMM (Binomial cond. dist.)	SAS code (GLMM; binomial)
Completely randomized (CR), one factor	$\mu_i = \eta_i = \theta + \alpha_i$ $Y_{ij} \sim N(\mu_i, \sigma^2)$  <i>j is an index for the j-th experimental unit of the treatment (such as the plot)</i>	<pre>class alpha ; model y = alpha;</pre>	<b>Naïve GLMM</b> $\text{logit}(p_i) = \eta_i = \theta + \alpha_i$ $Y_i \sim \text{Bin}(p_i, n)$	<pre>class alpha ; model y/n = alpha / dist=bin link=logit;</pre>
			<b>Cond. GLMM (w/ unit effect)</b> $\text{logit}(p_{ij}) = \eta_{ij} = \theta + \alpha_i + v_{ij}$ $Y_{ij}   v_{ij} \sim \text{Bin}(p_{ij}, n)$ $v_{ij} \sim N(0, \sigma_v^2)$ experimental unit (j) within treatment	<pre>class alpha ; model y/n = alpha / dist=bin link=logit; random plot(alpha);</pre> <p><i>plot is a code to identify each unique experimental unit receiving a treatment</i></p>
			<b>Quasi-likelihood</b> $\text{logit}(p_i) = \eta_i = \theta + \alpha_i$ $Y_i \sim \text{quasi-Bin}(p_i, n; \phi)$	<pre>class alpha ; model y/n = alpha / dist=bin link=logit; random _residual_;</pre>
Randomized complete block (RCB), one factor	$\mu_{ij} = \eta_{ij} = \theta + \alpha_i + b_j$ $Y_{ij}   b_j \sim N(\mu_{ij}, \sigma_e^2)$ $b_j \sim N(0, \sigma_b^2)$	<pre>class alpha block; model y = alpha; random block;</pre>	<b>Naïve GLMM</b> $\text{logit}(p_{ij}) = \eta_{ij} = \theta + \alpha_i + b_j$ $Y_{ij}   b_j \sim \text{Bin}(p_{ij}, n)$ $b_j \sim N(0, \sigma_b^2)$	<pre>class alpha block; model y/n = alpha / dist=bin link=logit; random block;</pre>
			<b>Cond. GLMM (w/ unit effect)</b> $\text{logit}(p_{ij}) = \eta_{ij} = \theta + \alpha_i + b_j + v_{ij}$ $Y_{ij}   b_j, v_{ij} \sim \text{Bin}(p_{ij}, n)$ $b_j \sim N(0, \sigma_b^2)$ $v_{ij} \sim N(0, \sigma_v^2)$	<pre>class alpha block; model y/n = alpha / dist=bin link=logit; random block block*alpha;</pre>
			<b>Quasi-likelihood</b> $\text{logit}(p_{ij}) = \eta_{ij} = \theta + \alpha_i + b_j$ $Y_{ij}   b_j \sim \text{quasi-Bin}(p_{ij}, n; \phi)$ $b_j \sim N(0, \sigma_b^2)$	<pre>class alpha block; model y/n = alpha / dist=bin link=logit; random block; random _residual_;</pre>
RCB, two factors	$\mu_{ijk} = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k$ $Y_{ijk}   b_k \sim N(\mu_{ijk}, \sigma_e^2)$	<pre>class alpha beta block; model y = alpha beta; random block;</pre>	<b>Naïve GLMM</b> $\text{logit}(p_{ijk}) = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k$ $Y_{ijk}   b_k \sim \text{Bin}(p_{ijk}, n)$	<pre>class alpha beta block; model y/n = alpha beta / dist=bin link=logit; random block;</pre>

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	$b_j \sim N(0, \sigma_b^2)$	--or, equivalently-- random int / sub=block;	$b_k \sim N(0, \sigma_b^2)$	
			<b>Cond. GLMM (w/ unit effect)</b> $\text{logit}(p_{ijk}) = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + v_{ijk}$ $Y_{ijk} / b_k, v_{ijk} \sim \text{Bin}(p_{ijk}, n)$ $b_k \sim N(0, \sigma_b^2)$ $v_{ijk} \sim N(0, \sigma_v^2)$	<pre>class alpha beta block; model y/n = alpha beta / dist=bin link=logit; random block block*alpha*beta;</pre>
			<b>Quasi-likelihood</b> $\text{logit}(p_{ijk}) = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k$ $Y_{ijk} / b_k \sim \text{quasi-Bin}(p_{ijk}, n; \phi)$ $b_k \sim N(0, \sigma_b^2)$	<pre>class alpha beta block; model y/n = alpha beta / dist=bin link=logit; random block; random _residual_;</pre>
<u>RCB, Split plot</u> Whole-plot ( $\alpha$ ) Sub-plot ( $\beta$ )	$\mu_{ijk} = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + d_{ik}$ $Y_{ijk} / b_k, d_{ik} \sim N(\mu_{ijk}, \sigma_e^2)$ <b>Conditional dist.</b> $b_k \sim N(0, \sigma_b^2)$ <b>block</b> $d_{ik} \sim N(0, \sigma_d^2)$ <b>whole-plot</b> <b>(=b x <math>\alpha</math>)</b>	<pre>class alpha beta block; model y = alpha beta; random block block*alpha;</pre> --or, equivalently-- random int alpha / sub=block;	<b>Naïve GLMM</b> $\text{logit}(p_{ijk}) = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + d_{ik}$ $Y_{ijk} / b_k, d_{ik} \sim \text{Bin}(p_{ijk}, n)$ $b_k \sim N(0, \sigma_b^2)$ $d_{ik} \sim N(0, \sigma_d^2)$	<pre>class alpha beta block; model y = alpha beta / dist=bin link=logit; random block block*alpha;</pre> --or, equivalently-- random int alpha / sub=block;
			<b>Cond. GLMM (w/ unit effect)</b> $\text{logit}(p_{ijk}) = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + d_{ik} + v_{ijk}$ $Y_{ijk} / b_k, d_{ik}, v_{ijk} \sim \text{Bin}(p_{ijk}, n)$ $b_k \sim N(0, \sigma_b^2)$ $d_{ik} \sim N(0, \sigma_d^2)$ $v_{ijk} \sim N(0, \sigma_v^2)$	<pre>class alpha beta block; model y = alpha beta / dist=bin link=logit; random block block*alpha block*alpha*beta;</pre> --or, equivalently-- random int alpha <b>alpha*beta</b> / sub=block;
			<b>Quasi-likelihood</b> $\text{logit}(p_{ijk}) = \eta_{ijk} = \theta + \alpha_i + \beta_j + (\alpha\beta)_{ij} + b_k + d_{ik}$ $Y_{ijk} / b_k, d_{ik} \sim \text{quasi-Bin}(p_{ijk}, n; \phi)$	<pre>class alpha beta block; model y = alpha beta / dist=bin link=logit; random block block*alpha; random _residual_;</pre>

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			$b_k \sim N(0, \sigma_b^2)$ $d_{ik} \sim N(0, \sigma_d^2)$	--or, equivalently-- <pre>random int alpha / sub=block; random _residual_;</pre>
<u>RCB, split-split plot</u> Whole-plot ( $\alpha$ ) Sub-plot ( $\beta$ ) Sub-sub plot ( $\gamma$ )	$\mu_{ijkl} = \eta_{ijkl} = \theta + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + f_{ijk}$ $Y_{ijkl}/b_k, d_{ik}, f_{ijk} \sim N(\mu_{ijkl}, \sigma_e^2)$ $b_k \sim N(0, \sigma_b^2) \text{ block}$ $d_{ik} \sim N(0, \sigma_d^2) \text{ whole-plot (=b x } \alpha)$ $f_{ijk} \sim N(0, \sigma_f^2) \text{ sub-plot (=b x } \alpha \text{ x } \beta)$	<pre>class alpha beta gamma block; model y= alpha beta gamma ; random block block*alpha block*alpha*beta;</pre> --or, equivalently-- <pre>random int alpha alpha*beta /sub=block;</pre>	<b>Naïve GLMM</b> $\text{logit}(p_{ijkl}) = \eta_{ijkl} = \theta + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + f_{ijk}$ $Y_{ijkl}/b_k, d_{ik}, f_{ijk} \sim \text{Bin}(p_{ijkl}, n)$ $b_k \sim N(0, \sigma_b^2) \text{ block}$ $d_{ik} \sim N(0, \sigma_d^2) \text{ whole-plot (=b x } \alpha)$ $f_{ijk} \sim N(0, \sigma_f^2) \text{ sub-plot (=b x } \alpha \text{ x } \beta)$	<pre>class alpha beta gamma block; model y= alpha beta gamma / dist=bin link=logit; random block block*alpha block*alpha*beta;</pre> --or, equivalently-- <pre>random int alpha alpha*beta /sub=block;</pre>
			<b>Cond. GLMM (w/ unit effect)</b> $\text{logit}(p_{ijkl}) = \eta_{ijkl} = \theta + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + f_{ijk} + v_{ijkl}$ $Y_{ijkl}/b_k, d_{ik}, f_{ijk}, v_{ijkl} \sim \text{Bin}(p_{ijkl}, n)$ $b_k \sim N(0, \sigma_b^2)$ $d_{ik} \sim N(0, \sigma_d^2)$ $f_{ijk} \sim N(0, \sigma_f^2)$ $v_{ijkl} \sim N(0, \sigma_v^2)$	<pre>class alpha beta gamma block; model y= alpha beta gamma / dist=bin link=logit; random block block*alpha block*alpha*beta block*alpha*beta*gamma;</pre> --or, equivalently-- <pre>random int alpha alpha*beta alpha*beta*gamma / sub=block;</pre>
			<b>Quasi-likelihood</b> $\text{logit}(p_{ijkl}) = \eta_{ijkl} = \theta + \alpha_i + \beta_j + \gamma_l + (\alpha\beta)_{ij} + (\alpha\gamma)_{il} + (\beta\gamma)_{jl} + (\alpha\beta\gamma)_{ijl} + b_k + d_{ik} + f_{ijk}$ $Y_{ijkl}/b_k, d_{ik}, f_{ijk} \sim \text{quasi-Bin}(p_{ijkl}, n; \phi)$	<pre>class alpha beta gamma block; model y= alpha beta gamma / dist=bin link=logit; random block block*alpha block*alpha*beta; random _residual_ ;</pre> --or, equivalently--

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			$b_k \sim N(0, \sigma_b^2)$ $d_{ik} \sim N(0, \sigma_d^2)$ $f_{ijk} \sim N(0, \sigma_f^2)$	<pre>random int alpha   alpha*beta /sub=block; random _residual_ ;</pre>
<u>RCB, one factor, with sub-sampling</u>  Normal: Linear predictor for <i>ij</i> treatment-block combination, with cond. dist. of <i>Y</i> for <i>ijk</i> treat-block-sampling unit (su) combination	$\mu_{ij} = \eta_{ij} = \theta + \alpha_i + b_j + v_{ij}$ $Y_{ijk} / b_j, v_{ij} \sim N(\mu_{ij}, \sigma_e^2)$ $b_j \sim N(0, \sigma_b^2)$ $v_{ij} \sim N(0, \sigma_v^2)$	<pre>class alpha block; model y = alpha; random block   block*alpha;</pre>	<b>Naïve GLMM</b> $\text{logit}(p_{ij}) = \eta_{ij} = \theta + \alpha_i + b_j + v_{ij}$ $Y_{ijk} / b_j, v_{ij} \sim \text{Bin}(p_{ij}, n)$ $b_j \sim N(0, \sigma_b^2)$ $v_{ij} \sim N(0, \sigma_v^2)$	<pre>class alpha block; model y/n = alpha   / dist=bin link=logit; random block block*alpha;  --or, equivalently-- random int alpha /   sub=block;</pre>
Binomial (with unit-level effect): Linear predictor <i>and</i> cond. dist. of <i>Y</i> for <i>ijk</i> treat-block-sampling unit (su) combination			<b>Cond. GLMM (w/ unit effect)</b> $\text{logit}(p_{ijk}) = \eta_{ij} = \theta + \alpha_i + b_j + v_{ij} + u_{ijk}$  $Y_{ijk} / b_j, v_{ij}, u_{ijk} \sim \text{Bin}(p_{ijk}, n)$ $b_j \sim N(0, \sigma_b^2)$ $v_{ij} \sim N(0, \sigma_v^2)$ $u_{ijk} \sim N(0, \sigma_u^2)$	<pre>class alpha block su; model y/n = alpha   / dist=bin link=logit; random block block*alpha   block*alpha*su;  --or, equivalently-- random int alpha   alpha*su; / sub=block ;</pre>
Quasi-likelihood: Linear predictor for <i>ij</i> treatment-block combination, with cond. quasi-dist. of <i>Y</i> for <i>ijk</i> treat-block-sampling unit (su) combination			<b>Quasi-likelihood</b> $\text{logit}(p_{ij}) = \eta_{ij} = \theta + \alpha_i + b_j + v_{ij}$ $Y_{ijk} / b_j, v_{ij} \sim \text{quasi-Bin}(p_{ij}, n; \phi)$ $b_j \sim N(0, \sigma_b^2)$ $v_{ij} \sim N(0, \sigma_v^2)$	<pre>class alpha block; model y/n = alpha   / dist=bin link=logit; random block block*alpha; random _residual_;  --or, equivalently-- random int alpha /   sub=block; random _residual_;</pre>

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

For other distributions in the exponential family, change `dist=` and `link=` options. For example, for counts, use `dist=Poisson link=log`.

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The quasi-likelihood approach is appropriate for binomial or Poisson conditional distributions (those without a so-called free scale parameter). Note that  $\text{Bin}(\bullet, n; \phi)$  is a customized notation for a quasi-distribution, an overdispersed “binomial” in this case (where  $\phi$  is the overdispersion scale parameter).