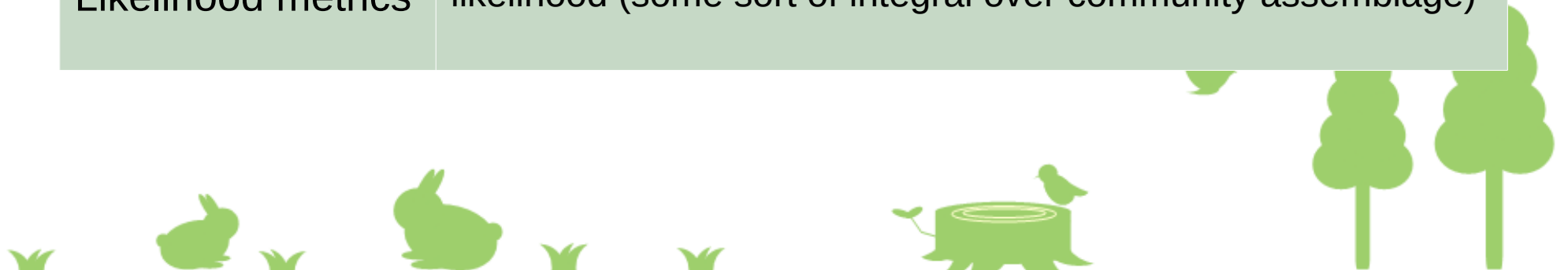


# Evaluation Metrics for sjSDM (JSDM)

Threshold-independent metrics	AUC, RMSE, Spearman rank correlation coefficient
Community dissimilarity indices	Bray-Curtis dissimilarity, Jaccard distance
Species richness metrics	Species richness difference
Likelihood metrics	Independent log-likelihood (Nagelkerke's $R^2$ ), joint log-likelihood (some sort of integral over community assemblage)



Spearman rank correlation coefficient	$\frac{\text{cov}(r_{\text{obs}}, r_{\text{pred}})}{\sigma_{r_{\text{obs}}} \sigma_{r_{\text{pred}}}}$	-1-1 >0 more correct predictions <0 more incorrect predictions
Bray-Curtis dissimilarity	$\frac{A+B-2J}{A+B}$ A: # obs presences, B: # pred, J: # correctly pred	0-1 0 = same composition 1 = opposite composition
Jaccard distance	$\frac{2D}{1+D}$ D: Bray-Curtis	Lower = more correct assemblage prediction
Species richness difference	pred richness – obs richness	$-\infty - \infty$ >0 overpredicts species richness <0 underpredicts species richness
Nagelkerke's R <sup>2</sup>	$L_p = \prod_i (p_i y_i + (1-p_i)(1-y_i))$ $\frac{1 - (L_0/L_p)^{2/N}}{1 - L_0^{2/N}}$	$R^2 = \frac{\log L_p - \log L_0}{1 - \log L_0}$