Guidelines for the supervised learning of species interactions

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- 1. The prediction of species interaction networks is gaining momentum as a way to circumvent limitations in data volume. Yet, ecological networks are challenging to predict because they are typically small and sparse. Dealing with extreme class imbalance is a challenge for most binary classifiers, and there are currently no guidelines as to how predictive models can be trained.
- 2. Using simple mathematical arguments and numerical experiments in which a variety of classifiers (for supervised learning) are trained on simulated networks, we develop a series of guidelines related to the choice of measures to use for model selection, and the degree of unbiasing to apply to the training dataset.
- 3. Classifier accuracy and the ROC-AUC are not informative measures for the performance of interaction prediction. PR-AUC is a fairer assessment of performance. In some cases, even standard measures can lead to selecting a more biased classifier because the effect of connectance is strong. The amount of correction to apply to the training dataset depends as a function of the classifier and the network connectance.
- 4. These results reveal that training machines to predict networks is a challenging task, and that in virtually all cases, the composition of the training set needs to be experimented on before performing the actual training. We discuss these consequences in the context of the low volume of data.

- example on diagnostic test: rare events are hard to detect even with really good models
- summary of model challenges for networks
- Strydom et al. (2021) importance of drawing on traits + validation is challenging + comparing
- 4 across space
- 5 Binary classifiers are usually assessed by measuring properties of their confusion matrix, i.e. the
- 6 contingency table reporting true/false positive/negative hits. A confusion matrix is laid out as

$$\begin{pmatrix} tp & fp \\ fn & tn \end{pmatrix}$$

- 7 wherein tp is the number of interactions predicted as positive, tn is the number of non-interactions
- 8 predicted as negative, fp is the number of non-interactions predicted as positive, and fn is the number of
- 9 interactions predicted as negative. Almost all measures based on the confusion matrix express rates of
- error or success as proportions, and therefore the values of these components matter in a relative way. At a
- coarse scale, a classifier is accurate when the trace of the matrix divided by the sum of the matrix is close
- to 1, with other measures focusing on different ways in which the classifer is wrong.
- 13 The same approach is used to evaluate e.g. species distribution models (SDMs). Indeed, the training and
- evaluation of SDMs as binary classifiers suffers from the same issue of low prevalence. In a previous work,
- Allouche et al. (2006) suggested that κ was a better test of model performance than the True Skill Statistic
- (TSS), which we will refer to as Youden's informedness (or J); these conclusions were later criticized by
- Somodi et al. (2017), who emphasized that informedness' relationship to prevalence depends on
- assumptions about bias in the model, and therefore recommend the use of κ as a validation of
- 19 classification performance. Although this work offers recommendations about the comparison of models,
- 20 it doesn't establishes baselines or good practices for training on imbalanced ecological data. Within the
- 21 context of networks, there are three specific issues that need to be adressed. First, what values of
- performance measures are we expecting for a classifier that has poor performance? This is particularly
- 23 important as it can evaluate whether low prevalence can lull us into a false sense of predictive accuracy.
- Second, independently of the question of model evaluation, is low prevalence an issue for training, and
- 25 can we remedy it? Finally, because the low amount of data on interaction makes a lot of imbalance
- ²⁶ correction methods (see e.g. Branco et al., 2015) hard to apply, which indicators can be optimized with the

- 27 least amount of positive interaction data?
- 28 In addition to the literature on SDMs, most of the research on machine learning application to life
- sciences is focused on genomics (which has very specific challenges, see a recent discussion by Whalen et
- al., 2021); this sub-field has generated largely different recommendations. Chicco & Jurman (2020)
- suggest using Matthews correlation coefficient (MCC) over F_1 , as a protection against over-inflation of
- predicted results; Delgado & Tibau (2019) advocate against the use of Cohen's κ , again in favor of MCC, as
- the relative nature of κ means that a worse classifier can be picked over a better one; similarly, Boughorbel
- et al. (2017) recommend MCC over other measures of performance for imbalanced data, as it has more
- desirable statistical properties. More recently, Chicco et al. (2021) temper the apparent supremacy of the
- MCC, by suggesting it should be replaced by Youden's informedness (also known as J, bookmaker's
- accuracy, and the True-Skill Statistic) when the imbalance in the dataset may not be representative
- ³⁸ (Jordano, 2016a, which is the case as networks are under-sampled; 2016b), when classifiers need to be
- 39 compared across different datasets (for example when predicting a system in space, where undersampling
- varies locally; McLeod et al., 2021), and when comparing the results to a no-skill (baseline) classifier is
- important. As these conditions are likely to be met with network data, there is a need to evaluate which
- measures of classification accuracy respond in a desirable way.
- 43 A lot of binary classifiers are built by using a regressor (whose task is to guess the value of the interaction,
- 44 amd can therefore return somethins considered to be a pseudo-probability); in this case, the optimal value
- below which predicitions are assumed to be negative (i.e. the interaction does not exist) can be determined
- by picking a threshold maximizing some value on the ROC curve or the PR curve. The area under these
- curves (ROC-AUC and PR-AUC henceforth) give ideas on the overall goodness of the classifier. Saito &
- 48 Rehmsmeier (2015) established that the ROC-AUC is biased towards over-estimating performance for
- 49 imbalanced data; on the contrary, the PR-AUC is able to identify classifiers that are less able to detect
- 50 positive interactions correctly, with the additional advantage of having a baseline value equal to
- prevalence. Therefore, it is important to assess whether these two measures return different results when
- ⁵² applied to ecological network prediction.
- 53 We establish that due to the low prevalence of interactions, even poor classifiers applied to food web data
- will reach a high accuracy; this is because the measure is dominated by the accidental correct predictions
- of negatives. The F_1 score and positive predictive values are less sensitive to bias, but **TODO**

56 Baseline values

57 Definition of the performance measures

58 K

59 F_{β}

60 informedness

61 MCC

62 Confusion matrix with skill and bias

In this section, we will assume a network of connectance ρ , *i.e.* having ρS^2 interactions (where S is the species richness), and $(1-\rho)S^2$ non-interactions. Therefore, the vector describing the *true* state of the network is a column vector $\mathbf{o}^T = [\rho(1-\rho)]$ (we can safely drop the S^2 terms, as we will work on the confusion matrix, which ends up expressing *relative* values).

In order to write the values of the confusion matrix for a hypothetical classifier, we need to define two characteristics: its skill, and its bias. Skill, here, refers to the propensity of the classifier to get the correct answer (*i.e.* to assign interactions where they are, and to not assign them where they are not). A no-skill classifier guesses at random, *i.e.* it will guess interactions with a probability ρ . The predictions of a no-skill classifier can be expressed as a row vector $\mathbf{p} = [\rho(1-\rho)]$. The confusion matrix \mathbf{M} for a no-skill classifier is given by the element-wise product of these vectors $\mathbf{o} \odot \mathbf{p}$, *i.e.*

$$\mathbf{M} = \begin{pmatrix} \rho^2 & \rho(1-\rho) \\ (1-\rho)\rho & (1-\rho)^2 \end{pmatrix}.$$

In order to regulate the skill of this classifier, we can define a skill matrix **S** with diagonal elements equal to s, and off-diagonal elements equal to (1-s), and re-express the skill-adjusted confusion matrix as **M** \odot **S**, *i.e*.

$$\begin{pmatrix} \rho^2 & \rho(1-\rho) \\ (1-\rho)\rho & (1-\rho)^2 \end{pmatrix} \odot \begin{pmatrix} s & (1-s) \\ (1-s) & s \end{pmatrix}.$$

- Note that when s = 0, $Tr(\mathbf{M}) = 0$ (the classifier is *always* wrong), when s = 0.5, the classifier is no-skill and guesses at random, and when s = 1, the classifier is perfect.
- The second element we can adjust in this hypothetical classifier is its bias, specifically its tendency to over-predict interactions. Like above, we can do so by defining a bias matrix **B**, where interactions are
- over-predicted with probability b, and express the final classifier confusion matrix as $\mathbf{M} \odot \mathbf{S} \odot \mathbf{B}$, *i.e.*

$$\begin{pmatrix} \rho^2 & \rho(1-\rho) \\ (1-\rho)\rho & (1-\rho)^2 \end{pmatrix} \odot \begin{pmatrix} s & (1-s) \\ (1-s) & s \end{pmatrix} \odot \begin{pmatrix} b & b \\ (1-b) & (1-b) \end{pmatrix}.$$

The final expression for the confusion matrix in which we can regulate the skill and the bias is

$$\mathbf{C} = \begin{pmatrix} s \times b \times \rho^2 & (1-s) \times b \times \rho(1-\rho) \\ (1-s) \times (1-b) \times (1-\rho)\rho & s \times (1-b) \times (1-\rho)^2 \end{pmatrix}.$$

In all further simulations, the confusion matrix $\bf C$ is transformed so that it sums to 1.

83 What are the baseline values of performance measures?

In this section, we will change the values of b, s, and ρ , and report how the main measures discussed in the introduction (MCC, F_1 , κ , and informedness) are responding to issues with the classifier. Before we do so, it is important to explain why we will not focus on accuracy too much. Accuracy is the number of correct predictions (Tr(C)) divided by the sum of the confusion matrix. For a no-skill, no-bias classifier, accuracy is equal to $\rho^2 + (1-\rho)^2$; for $\rho = 0.05$, this is ≈ 0.90 , and for $\rho = 0.01$, this is equal to ≈ 0.98 . In other words, the values of accuracy are expected to be so high that they are not really informatived (this is simply explained by the fact that for ρ small, $\rho^2 \ll (1-\rho)^2$). More concerning is the fact that introducing bias changes the response of accuracy in unexpected ways. Assuming a no-skill classifier, the numerator of accuracy becomes $b\rho^2 + (1-b)(1-\rho)^2$, which increases when b is low, which specifically means that at equal skill, a classifier that under-predicts interactions will have higher accuracy than an un-biased classifier. These issues are absent from balanced accuracy, but should nevertheless lead us to not report accuracy as the primary measure of network prediction success; moving forward, we will focus on other measures.

In order to examine how MCC, F_1 , κ , and informedness change w.r.t. the imbalance, skill, and bias, we performed a grid exploration of the values of logit(s) and logit(b) linearly from -10 to 10, of ρ linearly in [0,0.5], which is within the range of usually observed connectance values for empirical food webs. Note that at this point, there is no food web model to speak of; rather, the confusion matrix we discuss can be obtained for any classification task. Based on the previous discussion, the desirable properties for a measure of classifier success should be: an increase with classifier skill, especially at low bias; a hump-shaped response to bias, especially at high skill, and ideally center around logit(b) = 0; an increase with prevalence up until equiprevalence is reached.

[Figure 1 about here.]

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In fig. 1, we show that none of the four measures satisfy all the considerations at once: F_1 increases with 106 skill, and increases monotonously with bias; this is because F_1 does not account for true negatives, and the 107 increase in positive detection masks the over-prediction of interactions. Informedness varies with skill, reaching 0 for a no-skill classifier, but is entirely unsensitive to bias. Both MCC and κ have the same 109 behavior, whereby they increase with skill. κ peaks at increasing values of biass for increasing skill, i.e. is 110 likely to lead to the selection of a classifier that over-predicts interactions. By contract, MCC peaks at the 111 same value, regardless of skill, but this value is not logit(b) = 0: unless at very high classifier skill, MCC 112 risks leading to a model that over-predicts interactions. In fig. 2, we show that all measures except F_1 give 113 a value of 0 for a no-skill classifier, and are forced towars their correct maximal value when skill changes (i.e. a more connected networks will have higher values for a skilled classifierd, and lower values for a classifier making mostly mistakes).

[Figure 2 about here.]

These two analyses point to the following recommendations: MCC is indeed more appropriate than κ , as although sensitive to bias, it is sensitive in a consistent way. Informedness is appropriate at discriminating between different skills, but confounded by bias. F_1 is increasing with bias, and should not be prioritized to evalue the performance of the model. The discussion of sensitivity to bias should come with a domain-specific caveat: although it is likely that interactions documented in ecological networks are correct, a lot of non-interactions are simply unobserved; as predictive models are used for data-inflation

(*i.e.* the prediction of new interactions), it is not necessarilly a bad thing to select models that predict more interactions than the original dataset, because the original dataset misses some interactions.

Numerical experiments

In the following section, we will generate random networks, and train four binary classifiers (as well as an 127 ensemble model using the sum of the outputs) on 30% of the interaction data. Networks are generated by 128 picking random generality g and vulnerability v traits for S = 200 species uniformly on the unit interval, and assigning an interaction from species i to species j if $0.2g_i - \xi \le v_j \le 0.2g_i + \xi$, where ξ is a constant 130 regulating the connectance of the networks, and varies uniformly in $[5 \times 10^{-3}, 10^{-1}]$. This model gives 131 fully interval networks that are close analogues to the niche model (Williams & Martinez, 2000), but has the benefit of only relying on two features (g_i, v_i) , and having the exact same rule for all interactions. It is, 133 therefore, a simple case which most classifiers should be able to learn. 134 The training sample is composed of 30% of the 4×10^4 possible entries in the network, i.e. n = 12000. Out 135 of these interactions, we pick a proportion ν (the training set bias) to be positive, so that the training set 136 has νn interactions, and $(1 - \nu)n$ non-interactions. We vary ν uniformly in]0,1[. This allows to evaluate 137 how the measures of binary classification performance respond to artificially rebalanced dataset for a given network connectance. Note that both ξ and ν are sampled from a distribution rather than being 139 picked on a grid; this is because there is no direct relationship between the value of ξ and the connectance 140 of the simulated network, and therefore the precise value of ξ is not relevant for the analysis of the results. The dataset used for numerical experiments is composed of 20000 such (ξ, ν) pairs, on which four learners 142 are trained: a decision tree regressor, a boosted regression tree, a ridge regressor, and a random forest regressor. All models were taken from the MLJ. jl package (Blaom et al., 2020; Blaom & Vollmer, 2020) in 144 Julia 1.7 (Bezanson et al., 2017). In order to pick the best adjacency matrix for a given learner, we 145 performed a thresholding approach using 500 steps on predictions from the testing set, and picking the 146 threshold that maximized Youden's informedness, which is usually the optimized target for imbalanced classification. During the thresholding step, we measured the area under the receiving-operator 148 characteristic (ROC-AUC) and precision-recall (PR-AUC) curves, as measures of overall performance over 149 the range of returned values. We report the ROC-AUC and PR-AUC, as well as a suite of other measures as introduced in the next section, for the best threshold. The ensemble model was generated by summing the

- $_{152}$ predictions of all component models on the testing set (ranged in [0,1]), then put through the same
- thresholding process. The complete code to run the simulations is given as an appendix.
- After the simulations were completed, we removed all runs (i.e. pairs of ξ and ν) for which at least one of
- the following conditions was met: the accuracy was 0, the true positive or true negative rates were 0, the
- connectance was larger than 0.2. This removes both the obviously failed model runs, and the networks
- that are more densely connected compared to the connectance of empirical food webs (and are therefore
- less difficult to predict, being less imbalanced).

159 Effect of training set bias on performance

[Figure 3 about here.]

[Figure 4 about here.]

Required amount of positives to get the best performance

[Figure 5 about here.]

Guidelines for prediction

55 References

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163

- Allouche, O., Tsoar, A., & Kadmon, R. (2006). Assessing the accuracy of species distribution models:
- Prevalence, kappa and the true skill statistic (TSS). *Journal of Applied Ecology*, 43(6), 1223–1232.
- https://doi.org/10.1111/j.1365-2664.2006.01214.x
- Bezanson, J., Edelman, A., Karpinski, S., & Shah, V. (2017). Julia: A Fresh Approach to Numerical
- 170 Computing. SIAM Review, 59(1), 65–98. https://doi.org/10.1137/141000671
- Blaom, A. D., Kiraly, F., Lienart, T., Simillides, Y., Arenas, D., & Vollmer, S. J. (2020). MLJ: A Julia package
- for composable machine learning. *Journal of Open Source Software*, *5*(55), 2704.
- https://doi.org/10.21105/joss.02704

```
Blaom, A. D., & Vollmer, S. J. (2020, December 31). Flexible model composition in machine learning and its
       implementation in MLJ. http://arxiv.org/abs/2012.15505
175
    Boughorbel, S., Jarray, F., & El-Anbari, M. (2017). Optimal classifier for imbalanced data using Matthews
176
       Correlation Coefficient metric. PloS One, 12(6), e0177678.
177
       https://doi.org/10.1371/journal.pone.0177678
178
    Branco, P., Torgo, L., & Ribeiro, R. (2015, May 13). A Survey of Predictive Modelling under Imbalanced
       Distributions. http://arxiv.org/abs/1505.01658
180
    Chicco, D., & Jurman, G. (2020). The advantages of the Matthews correlation coefficient (MCC) over F1
181
       score and accuracy in binary classification evaluation. BMC Genomics, 21(1), 6.
182
       https://doi.org/10.1186/s12864-019-6413-7
183
    Chicco, D., Tötsch, N., & Jurman, G. (2021). The Matthews correlation coefficient (MCC) is more reliable
184
       than balanced accuracy, bookmaker informedness, and markedness in two-class confusion matrix
185
       evaluation. BioData Mining, 14, 13. https://doi.org/10.1186/s13040-021-00244-z
186
    Delgado, R., & Tibau, X.-A. (2019). Why Cohen's Kappa should be avoided as performance measure in
187
       classification. PloS One, 14(9), e0222916. https://doi.org/10.1371/journal.pone.0222916
188
    Jordano, P. (2016a). Chasing Ecological Interactions. PLOS Biol, 14(9), e1002559.
189
       https://doi.org/10.1371/journal.pbio.1002559
190
    Jordano, P. (2016b). Sampling networks of ecological interactions. Functional Ecology.
191
       https://doi.org/10.1111/1365-2435.12763
192
    McLeod, A., Leroux, S. J., Gravel, D., Chu, C., Cirtwill, A. R., Fortin, M.-J., Galiana, N., Poisot, T., & Wood,
       S. A. (2021). Sampling and asymptotic network properties of spatial multi-trophic networks. Oikos,
194
       n/a(n/a). https://doi.org/10.1111/oik.08650
195
    Saito, T., & Rehmsmeier, M. (2015). The Precision-Recall Plot Is More Informative than the ROC Plot
       When Evaluating Binary Classifiers on Imbalanced Datasets. PLOS ONE, 10(3), e0118432.
197
       https://doi.org/10.1371/journal.pone.0118432
198
    Somodi, I., Lepesi, N., & Botta-Dukát, Z. (2017). Prevalence dependence in model goodness measures with
199
       special emphasis on true skill statistics. Ecology and Evolution, 7(3), 863–872.
200
       https://doi.org/10.1002/ece3.2654
201
```

- Strydom, T., Catchen, M. D., Banville, F., Caron, D., Dansereau, G., Desjardins-Proulx, P., Forero-Muñoz,
- N. R., Higino, G., Mercier, B., Gonzalez, A., Gravel, D., Pollock, L., & Poisot, T. (2021). A roadmap
- towards predicting species interaction networks (across space and time). Philosophical Transactions of
- the Royal Society B: Biological Sciences, 376(1837), 20210063.
- 206 https://doi.org/10.1098/rstb.2021.0063
- Whalen, S., Schreiber, J., Noble, W. S., & Pollard, K. S. (2021). Navigating the pitfalls of applying machine
- learning in genomics. *Nature Reviews Genetics*, 1–13.
- https://doi.org/10.1038/s41576-021-00434-9
- Williams, R., & Martinez, N. (2000). Simple rules yield complex food webs. *Nature*, 404, 180–183.
- http://userwww.sfsu.edu/

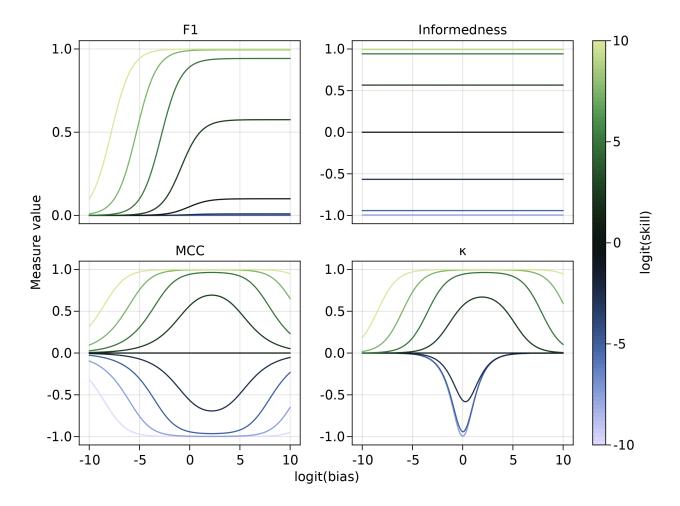


Figure 1: Consequences of changing the classifier skills (s) and bias (s) for a connectance $\rho=0.15$, on accuracy, F_1 , postive predictive value, and κ . Accuracy increases with skill, but also increases when the bias tends towards estimating *fewer* interactions. The F_1 score increases with skill but also increases when the bias tends towards estimating *more* interactions; PPV behaves in the same way. Interestingly, κ responds as expected to skill (being negative whenever s<0.5), and peaks for values of $b\approx0.5$; nevertheless, the value of bias for which κ is maximized in *not* b=0.5, but instead increases with classifier skill. In other words, at equal skill, maximizing κ would lead to select a *more* biased classifier.

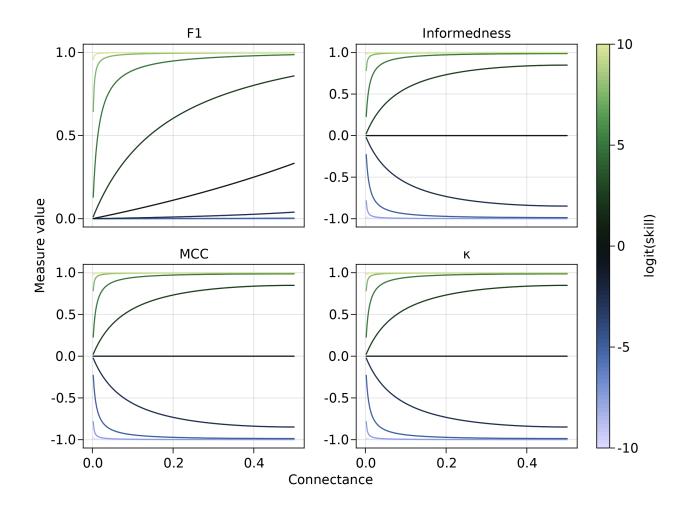


Figure 2: TODO

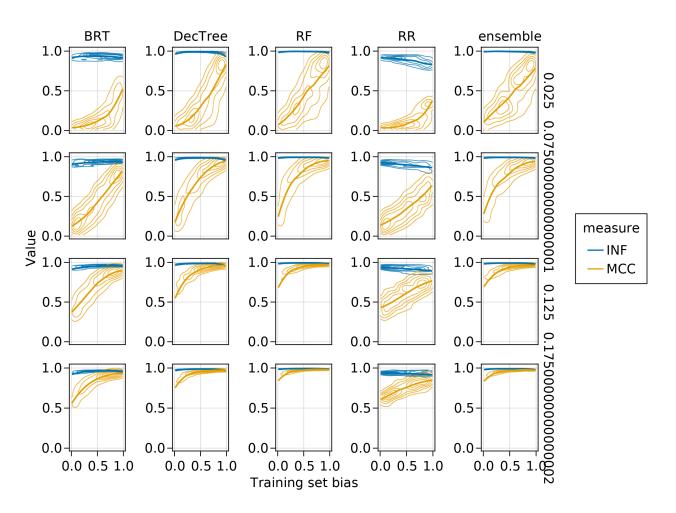


Figure 3: TODO

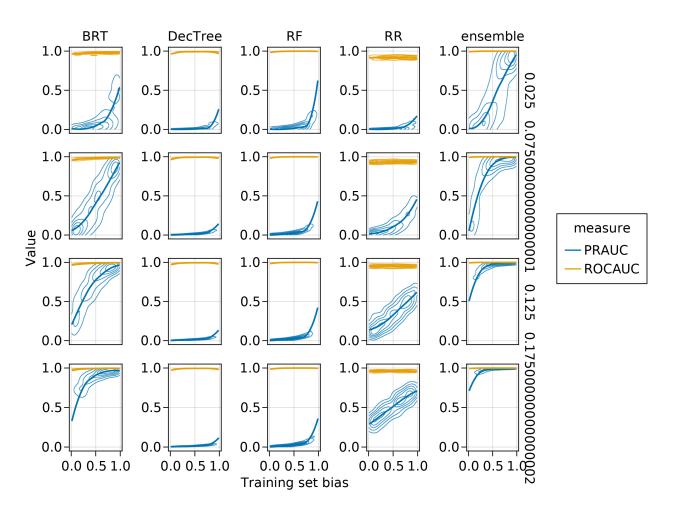


Figure 4: TODO

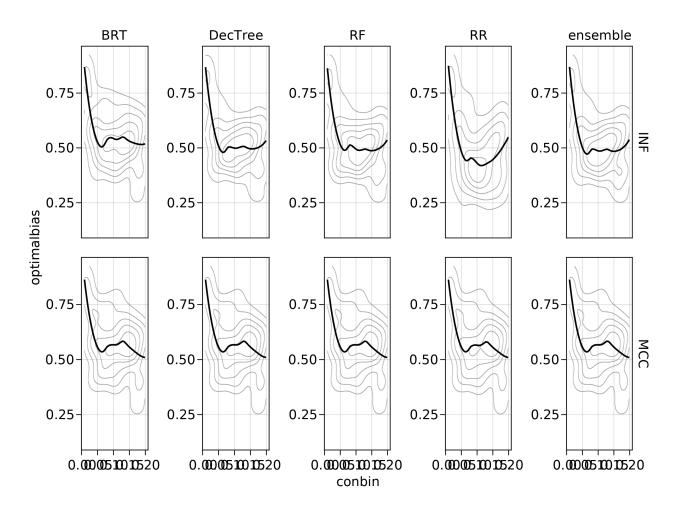


Figure 5: TODO