

Choosing representative BRT-models from sets of replicates

Supplementary Information
Cape vs SWA publication

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A representative model from a set of replicates is one which falls within the 5% quantile bound about the set's median values for *all* the various model-quality statistics (nt , R_{pseudo}^2 , R_{E-O}^2). When there is more than one model that meets this criterion, one is chosen at random.

```
is_between <- function(x, a, b) {  
  (x >= a) & (x <= b)  
}  
  
determine_bounds <- function(reps, bound_quantile = 0.05) {  
  columns <- c("nt", "pseudo_r2", "pred_obs_r2", "pred_obs_r2_exp")  
  summarise_at(reps, columns, funs(  
    # E.g. 0.025 + 0.025 = 5% quantile about the median  
    lwr = quantile(., (0.5 - (bound_quantile / 2))),  
    upr = quantile(., (0.5 + (bound_quantile / 2)))  
  ))  
}  
  
filter_reps_by_bounds <- function(reps, bounds) {  
  bounds %$% filter(reps,  
    is_between(nt,          nt_lwr,          nt_upr),  
    is_between(pseudo_r2,  pseudo_r2_lwr,  pseudo_r2_upr),  
    is_between(pred_obs_r2, pred_obs_r2_lwr, pred_obs_r2_upr)  
  )  
}
```

```
reps <- read.csv(...)  
bounds <- determine_bounds(reps)  
filter_reps_by_bounds(reps, bounds)
```