# Evaluation 2: get table 1

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13/03/2023

### Introduction

In our manuscript, we present the summary of the simulation study in Table 1. In part, this was done before, but here I want to present it a bit nicer. This includes for all analyzed n over all tested k with at least one phylogenetic set:

- Prevalence of phylogenetic decisive sets in simulation (overall, min, max)
- True Positive Rate (TPR, sensitivity, power) (overall, min, max)
- Negative Predictive Value (NPV) (overall, min, max)
- Minimal set size satisfying the 4-way partition property (4WPP) in the simulation
- Minimal set size which is fixing taxon traceable (FTT) in the simulation

In addition, I want a second table with more number, not necessarily out of the simulation

- Minimal set size satisfying the 4-way partition property
- Maximal set size not satisfying the 4-way partition property
- Minimal set size which is fixing taxon traceable
- Maximal set size which satisfy the 4-way partition property but is not fixing taxon traceable

## Initialize

I use a file names *SourceFile.R* that contains all relevant R packages and user-/server-specific path to the R library. If using this code, you must make all the necessary changes within the template source file.

```
rm(list = ls())
time0<-Sys.time()

source(".../SourceFile.R")
source(".../helperFunctions/TestHelpRFunction.R")

x_lowerBound = c()
x_upperBound = c()

for(i in 6:10){
    #i=7
    y = i %% 6

    if(y==0){</pre>
```

```
min_quad = 0.25 * choose(i,3) + i/6
  }else if(y %in% c(2,4)){
   min_quad = 0.25 * choose(i,3)
  }else{
   min_quad = (1/6) * ((i-1)/2) * choose(i,2) + i/12
   min_quad = ceiling(min_quad)
 x_lowerBound = c(x_lowerBound,min_quad)
 max_quad = choose(i,4) - (i-4)
 x_upperBound = c(x_upperBound,max_quad)
}
load("../results/02_SimulationResults_n06.RData")
load("../results/02_SimulationResults_n07.RData")
load("../results/02_SimulationResults_n08.RData")
load("../results/02_SimulationResults_n09.RData")
load("../results/02_SimulationResults_n10.RData")
SimulationResults_n06[,n := 6]
SimulationResults_n07[,n := 7]
SimulationResults_n08[,n := 8]
SimulationResults_n09[,n := 9]
SimulationResults_n10[,n := 10]
sim n6 = SimulationResults n06[k>=x lowerBound[1] & k<=x upperBound[1]]
sim_n7 = SimulationResults_n07[k>=x_lowerBound[2] & k<=x_upperBound[2]]</pre>
sim_n8 = SimulationResults_n08[k>=x_lowerBound[3] & k<=x_upperBound[3]]</pre>
sim_n9 = SimulationResults_n09[k>=x_lowerBound[4] & k<=x_upperBound[4]]
sim_n10 = SimulationResults_n10[k>=x_lowerBound[5] & k<=x_upperBound[5]]
sim = rbind(sim_n6,sim_n7,sim_n8,sim_n9,sim_n10)
table(sim$n)
   6 7 8 9 10
#>
#> 7 20 52 96 174
```

# Get Table 1

```
#> 2: 0.4541400 1 0.9728737 0.9664861 1 7
#> 4: 0.5146885 1 0.9877019 0.9882595 1 9
dumTab3 = foreach(i = 6:10)%do%{
 # i=6
 mySim = copy(sim)
 mySim = mySim[n == i,]
 dumTab4 = foreach(k = 1:dim(mySim)[1])%do%{
   # k=1
   myRow = copy(mySim)
   myRow = myRow[k,]
   stats_k = TestHelpRFunction(P = myRow$NR_PhyloDec,
                         N = myRow$NR NotPhyloDec,
                         PP = myRow$NR_FTT)
   stats_k[,n :=i]
   stats_k[,k := myRow$k]
   stats k
 }
 myStats_k = rbindlist(dumTab4)
 myStats_k
myStats_k = rbindlist(dumTab3)
x1 = myStats_k[Prevalence!=0,min(k),by = n]
x2 = myStats_k[Prevalence!=0 & TPR>0,min(k),by = n]
tab1 = copy(x1)
setnames(tab1,"V1","k_min_4WPP")
tab1[,k_min_FTT := x2$V1]
filt_TPR = !is.na(myStats_k$TPR)
tab6 = myStats_k[n==6 & filt_TPR,summary(TPR)]
tab7 = myStats_k[n==7 & filt_TPR,summary(TPR)]
tab8 = myStats_k[n==8 & filt_TPR,summary(TPR)]
tab9 = myStats_k[n==9 & filt_TPR,summary(TPR)]
tab10 = myStats_k[n==10 & filt_TPR, summary(TPR)]
tab_TPR = rbind(tab6,tab7,tab8,tab9,tab10)
tab1[,TPR := signif(tab_TPR[,3],2)]
tab1[,TPR_IQR := paste0("[",signif(tab_TPR[,2],2),",",signif(tab_TPR[,5],2),"]")]
filt_NPV = !is.na(myStats_k$NPV)
tab6 = myStats_k[n==6 & filt_NPV,summary(NPV)]
tab7 = myStats_k[n==7 & filt_NPV,summary(NPV)]
tab8 = myStats_k[n==8 & filt_NPV,summary(NPV)]
tab9 = myStats_k[n==9 & filt_NPV,summary(NPV)]
tab10 = myStats_k[n==10 & filt_NPV,summary(NPV)]
tab NPV = rbind(tab6,tab7,tab8,tab9,tab10)
tab1[,NPV := signif(tab_NPV[,3],2)]
```

```
tab1[,NPV_IQR := paste0("[",signif(tab_NPV[,2],2),",",signif(tab_NPV[,5],2),"]")]
tab6 = myStats_k[n==6 & Prevalence!=0,summary(Prevalence)]
tab7 = myStats_k[n==7 & Prevalence!=0, summary(Prevalence)]
tab8 = myStats_k[n==8 & Prevalence!=0,summary(Prevalence)]
tab9 = myStats_k[n==9 & Prevalence!=0,summary(Prevalence)]
tab10 = myStats_k[n==10 & Prevalence!=0,summary(Prevalence)]
tab Prev = rbind(tab6,tab7,tab8,tab9,tab10)
tab1[,Prev := signif(tab_Prev[,3],2)]
tab1[,Prev_IQR := paste0("[",signif(tab_Prev[,2],2),",",signif(tab_Prev[,5],2),"]")]
tab1
      n k_min_4WPP k_min_FTT TPR TPR_IQR NPV NPV_IQR Prev
#>
                                                               Prev_IQR
#> 1: 6
                          10 0.97 [0.71,1] 1 [0.98,1] 0.62 [0.36,0.83]
                9
#> 2: 7
                17
                          20 0.99 [0.65,1] 1 [0.97,1] 0.74 [0.25,0.96]
#> 3: 8
                30
                          35 1.00 [0.8,1] 1 [0.98,1] 0.85 [0.29,0.99]
#> 4: 9
                46
                          56 1.00 [0.91,1] 1 [0.99,1] 0.91
                                                             [0.32,1]
#> 5: 10
                          84 1.00 [0.97,1] 1 [1,1] 0.95
                72
                                                                [0.44,1]
knitr::kable(tab1)
```

_								
n	$k\_min\_4WPP$	$k\_min\_FTT$	TPR	TPR_IQR	NPV	NPV_IQR	Prev	Prev_IQR
6	9	10	0.97	[0.71,1]	1	[0.98,1]	0.62	[0.36, 0.83]
7	17	20	0.99	[0.65,1]	1	[0.97,1]	0.74	[0.25, 0.96]
8	30	35	1.00	[0.8,1]	1	[0.98,1]	0.85	[0.29, 0.99]
9	46	56	1.00	[0.91,1]	1	[0.99,1]	0.91	[0.32,1]
10	72	84	1.00	[0.97,1]	1	[1,1]	0.95	[0.44,1]

## Get Table 2

- Minimal set size satisfying the 4-way partition property
- Maximal set size not satisfying the 4-way partition property
- Minimal set size which is fixing taxon traceable
- Maximal set size which satisfy the 4-way partition property but is not fixing taxon traceable

```
x1 = myStats_k[Prevalence!=0,min(k),by = n]
y1 = myStats_k[Prevalence<1,max(k),by = n]
x2 = myStats_k[Prevalence!=0 & TPR>0,min(k),by = n]
y2 = sim[posRate<1,max(k),by = n]
y3 = c()
for(i in 6:10){
    #i=6
    dum = choose(i,4) - 3*i +13
    y3[i-5]=dum
}

tab2 = copy(x1)
setnames(tab2,"V1","k_min_4WPP")
tab2[,k_max_4WPP_sim := y1$V1]
tab2[,k_max_4WPP_theo := x_upperBound-1]
tab2[,k_min_FTT := x2$V1]</pre>
```

```
tab2[,k_max_diff_sim := y2$V1]
tab2[,k_max_diff_theo := y3]

tab2 = t(tab2)
knitr::kable(tab2)
```

6	7	8	9	10
9	17	30	46	72
12	31	62	113	183
12	31	65	120	203
10	20	35	56	84
10	26	50	89	146
10	27	59	112	193
	9 12 12 10 10	9 17 12 31 12 31 10 20 10 26	9 17 30 12 31 62 12 31 65 10 20 35 10 26 50	9 17 30 46 12 31 62 113 12 31 65 120 10 20 35 56

### Session Info

```
sessionInfo()
#> R version 4.2.2 (2022-10-31 ucrt)
#> Platform: x86_64-w64-mingw32/x64 (64-bit)
#> Running under: Windows 10 x64 (build 22621)
#> Matrix products: default
#>
#> locale:
#> [1] LC_COLLATE=German_Germany.utf8 LC_CTYPE=German_Germany.utf8
#> [3] LC MONETARY=German Germany.utf8 LC NUMERIC=C
#> [5] LC_TIME=German_Germany.utf8
#> attached base packages:
           stats graphics grDevices utils datasets methods
#> [1] grid
#> [8] base
#>
#> other attached packages:
#> [1] cowplot_1.1.1
                       gtable\_0.3.1
                                         ggplot2_3.4.1
#> [4] FixingTaxonTraceR_0.0.1 foreach_1.5.2
                                          data.table_1.14.8
#>
#> loaded via a namespace (and not attached):
#> [17] yaml_2.3.7
#> [25] rmarkdown_2.20 compiler_4.2.2 pillar_1.8.1 scales_1.2.1
#> [29] pkgconfig_2.0.3
message("\nTOTAL TIME : " ,round(difftime(Sys.time(),time(),units = "mins"),3)," minutes")
#> TOTAL TIME : 0.024 minutes
```