

## Checking Error Caused by Using Masked Carrier Risk

```
library(tidyverse)
load("RObjects/summary_tables/summaryTable10000Families.Rdata")
```

### Using All Families

```
# calculate average predicted value
avg_full <- mean(summaryTable$fullCarrierRisk) #adjusted means adjusted via prediction model
avg_masked <- mean(summaryTable$carrierRiskUnaffectedInfoMasked)
avg_true <- mean(as.numeric(summaryTable$probandMLH1Status))

sum_full <- sum(summaryTable$fullCarrierRisk) #adjusted means adjusted via prediction model
sum_masked <- sum(summaryTable$carrierRiskUnaffectedInfoMasked)
sum_true <- sum(as.numeric(summaryTable$probandMLH1Status))

# print average predicted value
print(str_interp("Average full carrier risk score: ${avg_full}"))
## [1] "Average full carrier risk score: 0.08222868608331"
print(str_interp("Average masked (unadjusted) carrier risk score: ${avg_masked}"))
## [1] "Average masked (unadjusted) carrier risk score: 0.138178890715236"
print(str_interp("Average number of true carriers of MLH1: ${avg_true}"))
## [1] "Average number of true carriers of MLH1: 0.0978"

mse_full <- mean((as.numeric(summaryTable$probandMLH1Status) - summaryTable$fullCarrierRisk)**2)
mse_masked <- mean((as.numeric(summaryTable$probandMLH1Status) - summaryTable$carrierRiskUnaffectedInfoMasked)**2)
print(str_interp("Full MSE: ${mse_full}"))
## [1] "Full MSE: 0.0594202717519354"
print(str_interp("Masked MSE: ${mse_masked}"))
## [1] "Masked MSE: 0.0612137433039341"
```

### Using Families with Unaffected First Degree Relatives (U1)

#### Unrounded

```
summaryTableFirstDegUnaff <- summaryTable %>% filter(firstDegreeAffectedFamilyMembersBinary == 0)

# calculate average predicted value
avg_full <- mean(summaryTableFirstDegUnaff$fullCarrierRisk) #adjusted means adjusted via prediction model
avg_masked <- mean(summaryTableFirstDegUnaff$carrierRiskUnaffectedInfoMasked)
avg_true <- mean(as.numeric(summaryTableFirstDegUnaff$probandMLH1Status))

# print average predicted value
print(str_interp("Average full carrier risk score: ${avg_full}"))
## [1] "Average full carrier risk score: 0.0332079222774842"
print(str_interp("Average masked carrier risk score: ${avg_masked}"))
## [1] "Average masked carrier risk score: 0.0952801099097885"
```

```

print(str_interp("Average number of true carriers of MLH1: ${avg_true}"))
## [1] "Average number of true carriers of MLH1: 0.046973803071364"

mse_full <- mean((as.numeric(summaryTableFirstDegUnaff$probandMLH1Status) - summaryTableFirstDegUnaff$fullCarrierRisk))
mse_masked <- mean((as.numeric(summaryTableFirstDegUnaff$probandMLH1Status) - summaryTableFirstDegUnaff$carrierRiskUnaffectedInfoMasked))
print(str_interp("Full MSE: ${mse_full}"))
## [1] "Full MSE: 0.0369271312819115"
print(str_interp("Masked MSE: ${mse_masked}"))
## [1] "Masked MSE: 0.0389315218152308"

```

## Using Families with Affected First Degree Relatives (A1)

```

summaryTableFirstDegAff <- summaryTable %>% filter(firstDegreeAffectedFamilyMembersBinary == 1)

# calculate average predicted value
avg_full <- mean(summaryTableFirstDegAff$fullCarrierRisk) #adjusted means adjusted via prediction model
avg_masked <- mean(summaryTableFirstDegAff$carrierRiskUnaffectedInfoMasked)
avg_true <- mean(as.numeric(summaryTableFirstDegAff$probandMLH1Status))
# print average predicted value
print(str_interp("Average full carrier risk score: ${avg_full}"))
## [1] "Average full carrier risk score: 0.461711102398339"
print(str_interp("Average masked carrier risk score: ${avg_masked}"))
## [1] "Average masked carrier risk score: 0.47026945261475"
print(str_interp("Average number of true carriers of MLH1: ${avg_true}"))
## [1] "Average number of true carriers of MLH1: 0.491258741258741"

mse_full <- mean((as.numeric(summaryTableFirstDegAff$probandMLH1Status) - summaryTableFirstDegAff$fullCarrierRisk))
mse_masked <- mean((as.numeric(summaryTableFirstDegAff$probandMLH1Status) - summaryTableFirstDegAff$carrierRiskUnaffectedInfoMasked))
print(str_interp("Full MSE: ${mse_full}"))
## [1] "Full MSE: 0.233545492033869"
print(str_interp("Masked MSE: ${mse_masked}"))
## [1] "Masked MSE: 0.233706185178021"

```

## Round values with clinical significance

```

round <- function(x){
  if(x>=0.2){
    return(1)
  }
  else{
    return(0)
  }
}

summaryTable$roundedFullCarrierRisk <- lapply(summaryTable$fullCarrierRisk, round)
summaryTable$roundedFullCarrierRisk <- as.numeric(summaryTable$roundedFullCarrierRisk)
summaryTable$roundedCarrierRiskUnaffectedInfoMasked <- lapply(summaryTable$carrierRiskUnaffectedInfoMasked, round)
summaryTable$roundedCarrierRiskUnaffectedInfoMasked <- as.numeric(summaryTable$roundedCarrierRiskUnaffectedInfoMasked)
summaryTable$probandMLH1Status <- as.numeric(summaryTable$probandMLH1Status)

```

```
#All families
```

```
#Calibration
```

```
print(str_interp("Sum of rounded full carrier risk: ${sum(summaryTable$roundedFullCarrierRisk)}"))  
## [1] "Sum of rounded full carrier risk: 1167"  
print(str_interp("Sum of rounded masked carrier risk: ${sum(summaryTable$roundedCarrierRiskUnaffectedIn)}"))  
## [1] "Sum of rounded masked carrier risk: 2340"  
print(str_interp("Sum of true MLH1 carriers: ${sum(summaryTable$probandMLH1Status)}"))  
## [1] "Sum of true MLH1 carriers: 978"
```

```
mse_full <- mean((as.numeric(summaryTable$probandMLH1Status) - summaryTable$roundedFullCarrierRisk)**2)  
mse_masked <- mean((as.numeric(summaryTable$probandMLH1Status) - summaryTable$roundedCarrierRiskUnaffectedIn)**2)  
print(str_interp("Full MSE: ${mse_full}"))  
## [1] "Full MSE: 0.0947"  
print(str_interp("Masked MSE: ${mse_masked}"))  
## [1] "Masked MSE: 0.1716"
```

```
#First Degree Unaffected
```

```
summaryTableFirstDegUnaff <- summaryTable %>% filter(firstDegreeAffectedFamilyMembersBinary == 0)
```

```
#Calibration
```

```
print(str_interp("Sum of rounded full carrier risk: ${sum(summaryTableFirstDegUnaff$roundedFullCarrierRisk)}"))  
## [1] "Sum of rounded full carrier risk: 79"  
print(str_interp("Sum of rounded masked carrier risk: ${sum(summaryTableFirstDegUnaff$roundedCarrierRiskUnaffectedIn)}"))  
## [1] "Sum of rounded masked carrier risk: 1223"  
print(str_interp("Sum of true MLH1 carriers: ${sum(summaryTableFirstDegUnaff$probandMLH1Status)}"))  
## [1] "Sum of true MLH1 carriers: 416"
```

```
mse_full <- mean((as.numeric(summaryTableFirstDegUnaff$probandMLH1Status) - summaryTableFirstDegUnaff$roundedFullCarrierRisk)**2)  
mse_masked <- mean((as.numeric(summaryTableFirstDegUnaff$probandMLH1Status) - summaryTableFirstDegUnaff$roundedCarrierRiskUnaffectedIn)**2)  
print(str_interp("Full MSE: ${mse_full}"))  
## [1] "Full MSE: 0.0446025293586269"  
print(str_interp("Masked MSE: ${mse_masked}"))  
## [1] "Masked MSE: 0.129516711833785"
```

```
#First degree affected
```

```
summaryTableFirstDegAff <- summaryTable %>% filter(firstDegreeAffectedFamilyMembersBinary == 1)
```

```
#Calibration
```

```
print(str_interp("Sum of rounded full carrier risk: ${sum(summaryTableFirstDegAff$roundedFullCarrierRisk)}"))  
## [1] "Sum of rounded full carrier risk: 1088"  
print(str_interp("Sum of rounded masked carrier risk: ${sum(summaryTableFirstDegAff$roundedCarrierRiskUnaffectedIn)}"))  
## [1] "Sum of rounded masked carrier risk: 1117"  
print(str_interp("Sum of true MLH1 carriers: ${sum(summaryTableFirstDegAff$probandMLH1Status)}"))  
## [1] "Sum of true MLH1 carriers: 562"
```

```
mse_full <- mean((as.numeric(summaryTableFirstDegAff$probandMLH1Status) - summaryTableFirstDegAff$roundedFullCarrierRisk)**2)  
mse_masked <- mean((as.numeric(summaryTableFirstDegAff$probandMLH1Status) - summaryTableFirstDegAff$roundedCarrierRiskUnaffectedIn)**2)  
print(str_interp("Full MSE: ${mse_full}"))  
## [1] "Full MSE: 0.482517482517482"  
print(str_interp("Masked MSE: ${mse_masked}"))  
## [1] "Masked MSE: 0.497377622377622"
```