

# R Notebook

## Load data

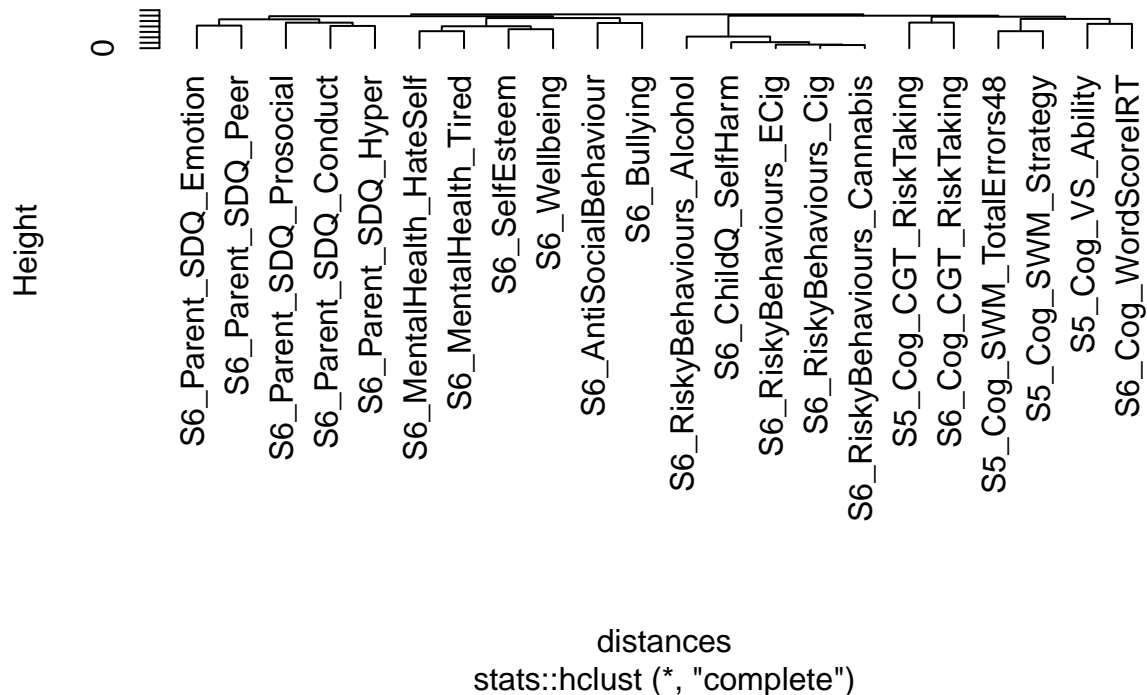
```
rm(list=ls())
source("Script P2S0 - Load Analysis Environment.R")
load(file.path("Output R Data", "cca_boot.Rdata"))
load(file.path("Output R Data", "cca_splithalf.Rdata"))
load(file.path("Output R Data", "cca_boot_teach.Rdata"))
load(file.path("Output R Data", "cca_splithalf_teacher.Rdata"))
```

## Analysis 0 - Hierarchial cluster analysis of variables

To simply aid visualisation, we have cluster the predictor and outcome variables into 4 (5 groups including teacher metrics)

```
distances = stats::dist(t(df0_imputed_BothGroups[,outcomes]), method="manhattan")
hclust_results = stats::hclust(distances)
Outcome_Order = hclust_results$order
Outcome_Ordered = outcomes[Outcome_Order]
Outcome_Labels_Ordered = outcomes_labels[Outcome_Order]
plot(hclust_results, hang=-1, main = "Hierarchial Clustering of Outcome Variables")
```

## Hierarchical Clustering of Outcome Variables



```
Outcome_Cluster = stats::cutree(hclust_results, 4)
# sort(Outcome_Cluster)
Outcome_Cluster_Labels = Convert(Outcome_Cluster,1:4,c("Behavioural Problems","Mental Health", "Drug-")

# For ease of presentation, teacher's responses are not clustered in the usual way, but just tagged on
Outcome_Cluster2 = c(rep(0,6),Outcome_Cluster)
Outcome_Ordered2 = c(outcomes2[1:6],Outcome_Ordered)
Outcome_Labels_Ordered2 = c(outcomes_labels2[1:6],Outcome_Labels_Ordered)
names(Outcome_Cluster2)[1:6] = outcomes2[1:6]
Outcome_Cluster_Labels2 = Convert(Outcome_Cluster2,0:4,c("Teacher Ratings","Behavioural Problems","Mental Health", "Drug-")

# not used in results - but out of curiosity looked at clustering all outcomes with teacher-reports
#
#   hclust_results = hclust(dist(t(df0_imputed_Teach[,outcomes2])), method="manhattan"))
#   plot(hclust_results, hang=-1)

#Cluster Predictors
distances = dist(t(df0_imputed_group1[,envvar]), method="manhattan")
hclust_results = hclust(distances)
envvar_Order = hclust_results$order
envvar_Ordered = envvar[envvar_Order]
envvar_Labels_Ordered = envvar_labels[envvar_Order]
# plot(hclust_results, hang=-1)
```

## Plot All Outcomes

```
pdf(file=file.path("Plots","histograms_variables_combined.pdf"),height=40, width=15)
par(mfrow=c(30,5), mar=c(1,1,3,1))
for (i in c(envvar, outcomes)){
  hist(df0_imputed_BothGroups[,i], main=i, col="grey")
}
dev.off()
```

```
## pdf
## 2
```

## Figure 2

note to giacomio - fix axes!

```
PlotLoadings = function(dat, recode=FALSE, titletext=""){

  if (recode){
    dat = dat*-1
  }

  dat$negative = dat$original<0
  # dat[!dat$negative,1:4]=dat[!dat$negative,1:4]*-1
  dat$FancyLabels = all_var_labels2[match(rownames(dat),all_var2)]
  # dat$Group = factor(as.numeric(grepl("S[5-6]",dat$varlabel)), labels=c("Environmental Predictors","O
  dat = dat[order(abs(dat$original)),]

  dat$errormin = dat$X2.5.
  dat$errormin[dat$negative] = dat$errormin[dat$negative]*-1
  dat$errormax = dat$X97.5
  dat$errormax[dat$negative] = dat$errormax[dat$negative]*-1

  plot=
  ggplot(dat, aes(x=1:nrow(dat), y=abs(original), fill=negative)) + geom_bar(stat="identity") +
    geom_errorbar(aes(ymin=errormin,ymax=errormax)) +
    jtools::theme_apo() +
    coord_flip() + labs(y=NULL,x="") + theme(legend.position = "none") + scale_fill_manual(values=c( "#
    scale_x_continuous(breaks=1:nrow(dat),labels=dat$FancyLabels) +
    theme(text=element_text(family="serif")) +
    geom_hline(aes( alpha=.9, yintercept=.1), linetype="dashed", col="grey") +
    geom_hline(aes( alpha=.9, yintercept=0), linetype="dashed") +
    labs(title=titletext)

  plot
  # facet_wrap(~Group,strip.position = "top", scales = "free", nrow=1, ncol=2) + theme(panel.spacing =
  # scale_x_continuous(breaks=1:nrow(dat),labels=dat$FancyLabels)
  #
  return(plot)
}
```

```

p1 = PlotLoadings(cca_boot$xcoef_Quantiles[[1]], titletext = "ERF CCA Weights Comp1")
p2 = PlotLoadings(cca_boot$xcoef_Quantiles[[2]], titletext = "ERF CCA Weights Comp2")
p3 = PlotLoadings(cca_boot$xcoef_Quantiles[[3]], titletext = "ERF CCA Weights Comp3")

p4 = PlotLoadings(cca_boot$ycoef_Quantiles[[1]], titletext = "Outcome CCA Weights Comp1")
p5 = PlotLoadings(cca_boot$ycoef_Quantiles[[2]], titletext = "Outcome CCA Weights Comp2")
p6 = PlotLoadings(cca_boot$ycoef_Quantiles[[3]], titletext = "Outcome CCA Weights Comp3")

library(patchwork)

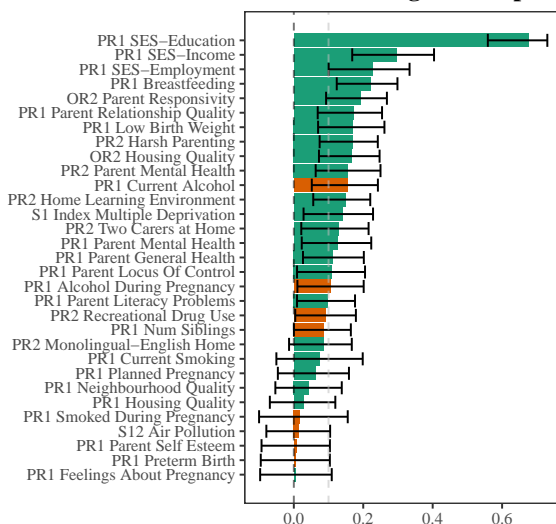
##
## Attaching package: 'patchwork'

## The following object is masked from 'package:MASS':
##
##      area

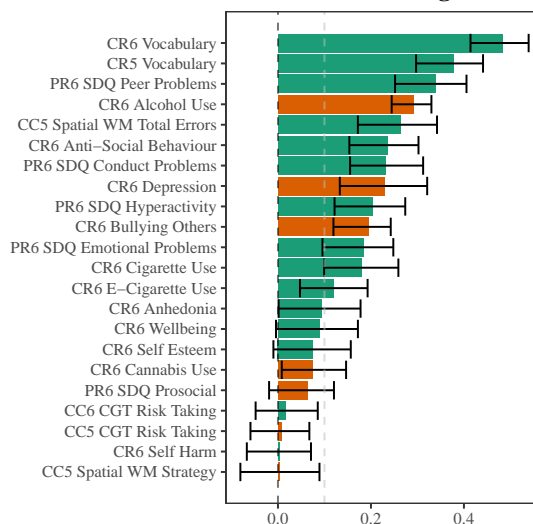
LoadingPlot = p1 + p2 + p3 + p4 + p5 + p6 + patchwork::plot_layout(ncol = 2, byrow = FALSE)
LoadingPlot

```

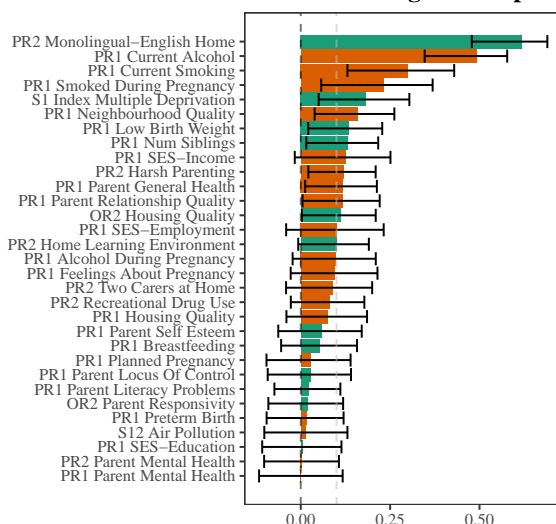
**ERF CCA Weights Comp1**



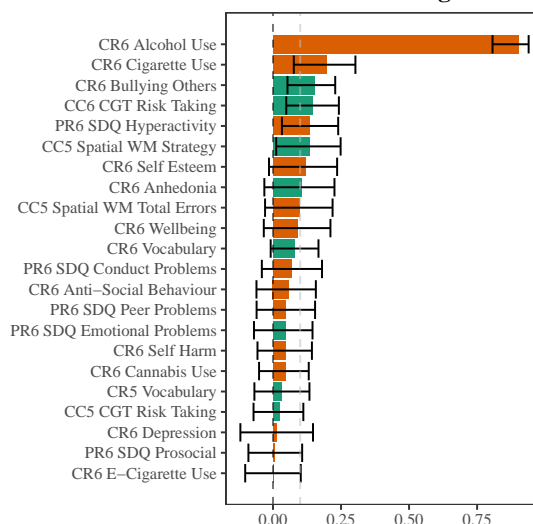
**Outcome CCA Weights Comp1**



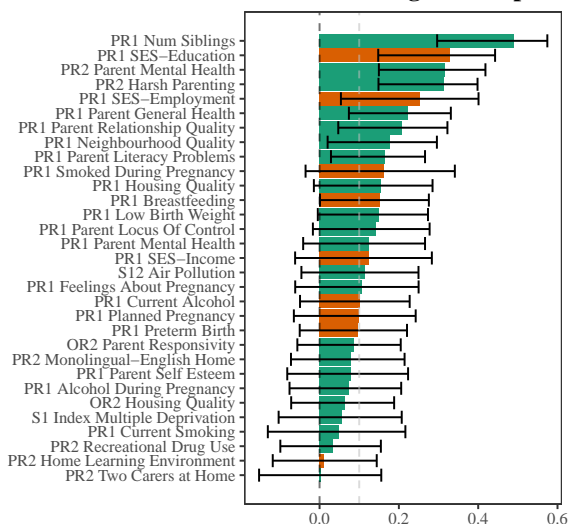
**ERF CCA Weights Comp2**



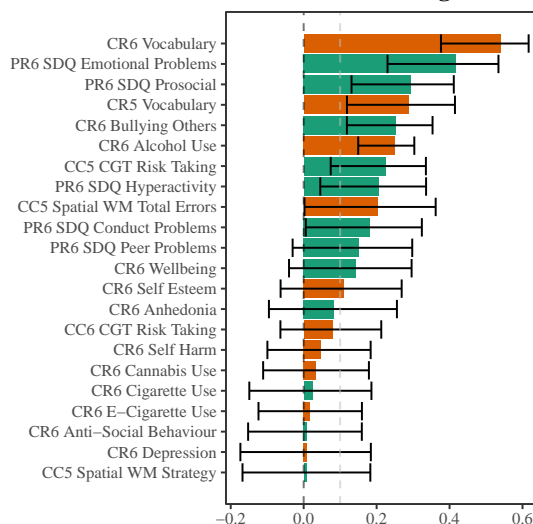
**Outcome CCA Weights Comp2**



**ERF CCA Weights Comp3**



**Outcome CCA Weights Comp3**



```

ggsave(filename=file.path("Plots","LoadingsPlots_group1.pdf"), LoadingPlot, device="pdf", width=9, hei
ggsave(filename=file.path("Plots","LoadingsPlots_group1.png"), LoadingPlot, device="png", width=9, hei

rm(p1,p2,p3,p4,p5,p6,LoadingPlot)

```

### Supplementary Figure 3

```

# cca_splithalf_procrustes$model_results

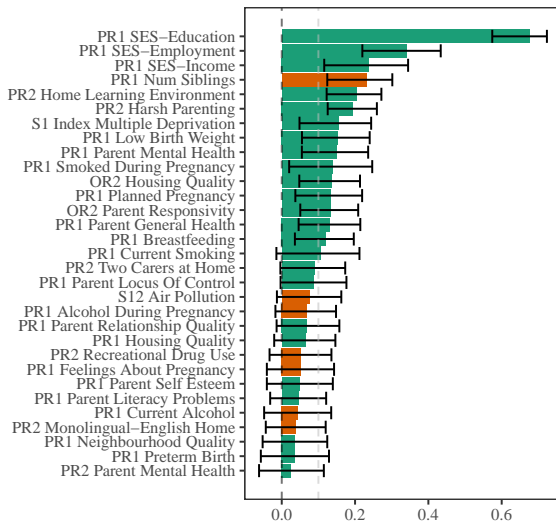
p1 = PlotLoadings(cca_boot_teach$xcoef_Quantiles[[1]], titletext = "ERF CCA Weights Comp1")
p2 = PlotLoadings(cca_boot_teach$xcoef_Quantiles[[2]], titletext = "ERF CCA Weights Comp2")
p3 = PlotLoadings(cca_boot_teach$xcoef_Quantiles[[3]], titletext = "ERF CCA Weights Comp3")

p4 = PlotLoadings(cca_boot_teach$ycoef_Quantiles[[1]], titletext = "Outcome CCA Weights Comp1")
p5 = PlotLoadings(cca_boot_teach$ycoef_Quantiles[[2]], titletext = "Outcome CCA Weights Comp2")
p6 = PlotLoadings(cca_boot_teach$ycoef_Quantiles[[3]], titletext = "Outcome CCA Weights Comp3")

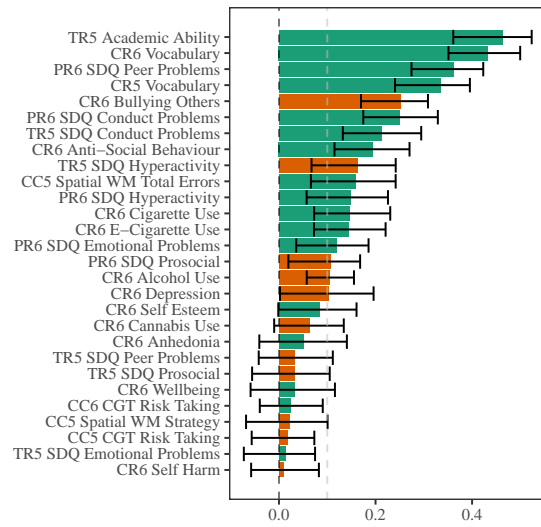
library(patchwork)
LoadingPlot = p1 + p2 + p3 + p4 + p5 + p6 + patchwork::plot_layout(ncol = 2, byrow = FALSE)
LoadingPlot

```

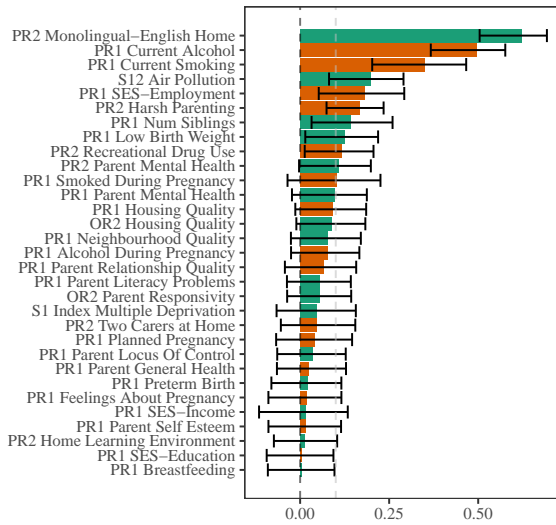
**ERF CCA Weights Comp1**



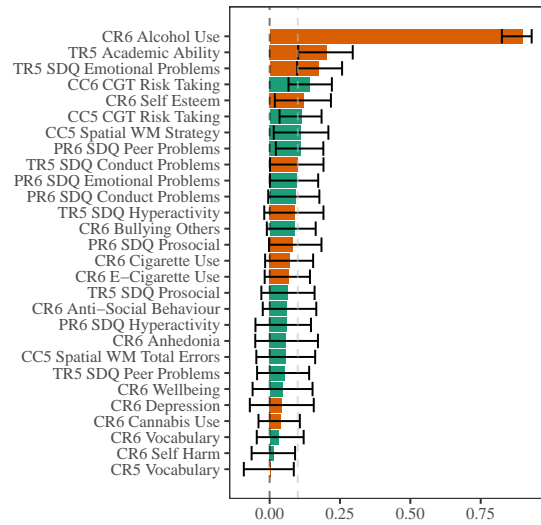
**Outcome CCA Weights Comp1**



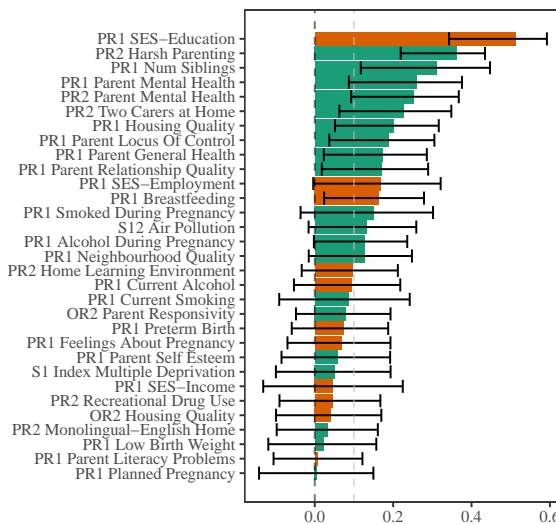
**ERF CCA Weights Comp2**



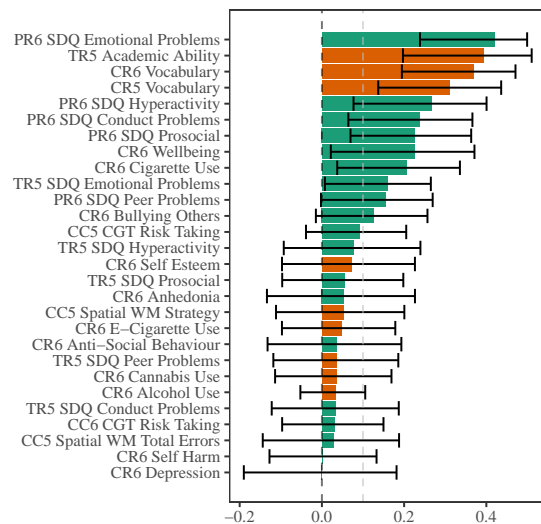
**Outcome CCA Weights Comp2**



**ERF CCA Weights Comp3**



**Outcome CCA Weights Comp3**



```
ggsave(filename=file.path("Plots","LoadingsPlots_teach.pdf"), LoadingPlot, device="pdf", width=9, height=9)
ggsave(filename=file.path("Plots","LoadingsPlots_teach.png"), LoadingPlot, device="png", width=9, height=9)
```

## Stacked bar chart figure

Figure 3

```
PlotData = data.frame(cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined)
PlotData[PlotData<0] = 0
PlotData$Var = rownames(PlotData)
PlotData = melt(PlotData, ID="Var")

## Using Var as id variables

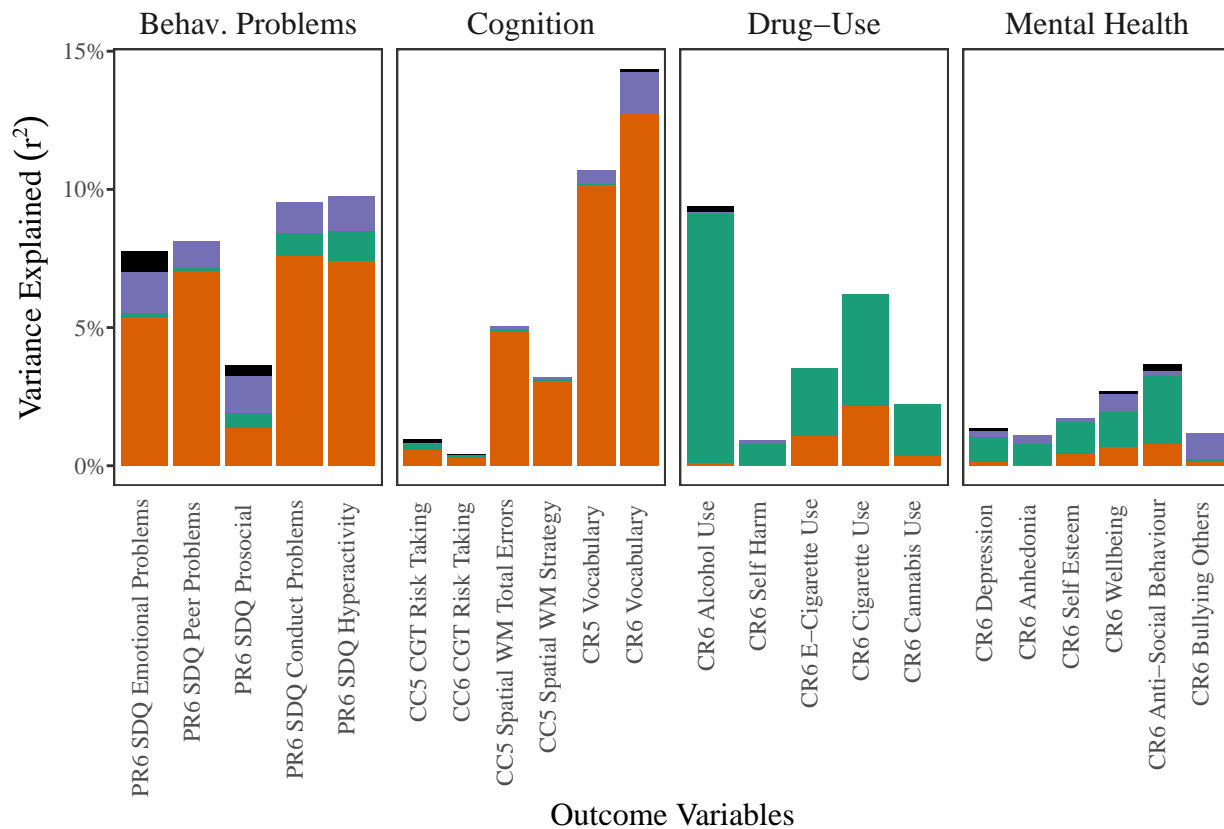
PlotData$variable = factor(PlotData$variable)
PlotData$Cluster = (Outcome_Cluster_Labels2[match(PlotData$Var,outcomes2)])
PlotData$Cluster[PlotData$Cluster=="Behavioural Problems"] = "Behav. Problems"
PlotData$Cluster = factor(PlotData$Cluster)
PlotData$Var = factor(PlotData$Var, levels=Outcome_Ordered2, labels=Outcome_Labels_Ordered2)
#Fill Colour
FillCol = c(as.vector( RColorBrewer::brewer.pal(5,"Dark2")), as.vector(wesanderson::wes_palette("Darjeeling",5)))

PLOT3=
ggplot(data=PlotData, aes(x=Var,y=value, fill=variable, group=Cluster)) +
  geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="LM_R2",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc10",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc9",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc8",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc7",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc6",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc5",])
  # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc4",])
  geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc3",],)
  geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc2",],)
  geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc1",],)
  labs(y=bquote('Variance Explained'~(r^2)), x="Outcome Variables") +
  facet_wrap(~Cluster,strip.position = "top", scales = "free_x", nrow=1) + theme(panel.spacing = unit(10,"mm"),
  jtools::theme_apa()+
  theme(axis.title.y.right = element_blank(), # hide right axis title
        axis.text.y.right = element_blank(), # hide right axis labels
        axis.ticks.x = element_blank(), # hide left/right axis ticks
        axis.text.y = element_text(margin = margin(r = 0)), # move left axis labels closer to axis
        #panel.spacing = unit(10, "mm"), # remove spacing between facets
        strip.background = element_blank(), # match default line size of theme_class
        legend.position = "none",
        axis.text.x = element_text(angle = 90,hjust=0.95,vjust=0.2),
        text=element_text(family="serif")
  ) +
  scale_fill_manual(labels = unique(PlotData$variable),values=c( wesanderson::wes_palette("BottleRock",5)))
  scale_y_continuous(breaks=c(0,5,10,15)/100, labels=c("0%","5%", "10%", "15%"))
```



```
ggsave(filename=file.path("Plots","VarianceExplained1.pdf"), PLOT3, device="pdf", width=6, height=4)
ggsave(filename=file.path("Plots","VarianceExplained1.png"), PLOT3, device="png", width=6, height=4, dpi=300)
```

```
PLOT3 ## ylim(c(0,.19))
```



Supplementary Figure 4

```
PlotData = data.frame(cca_splithalf_teacher$R2_matrix_BinaryOutcomes_Combined)
PlotData[PlotData<0] = 0
PlotData$Var = rownames(PlotData)
PlotData = melt(PlotData, ID="Var")
```

```
## Using Var as id variables
```

```
PlotData$variable = factor(PlotData$variable)
PlotData$Cluster = (Outcome_Cluster_Labels2[match(PlotData$Var,outcomes2)])
PlotData$Cluster[PlotData$Cluster=="Behavioural Problems"] = "Behav. Problems"
PlotData$Cluster = factor(PlotData$Cluster)
PlotData$Var = factor(PlotData$Var, levels=Outcome_Ordered2, labels=Outcome_Labels_Ordered2)

#Fill Colour
FillCol = c(as.vector( RColorBrewer::brewer.pal(5,"Dark2")), as.vector(wesanderson::wes_palette("Darjeeling")))
```

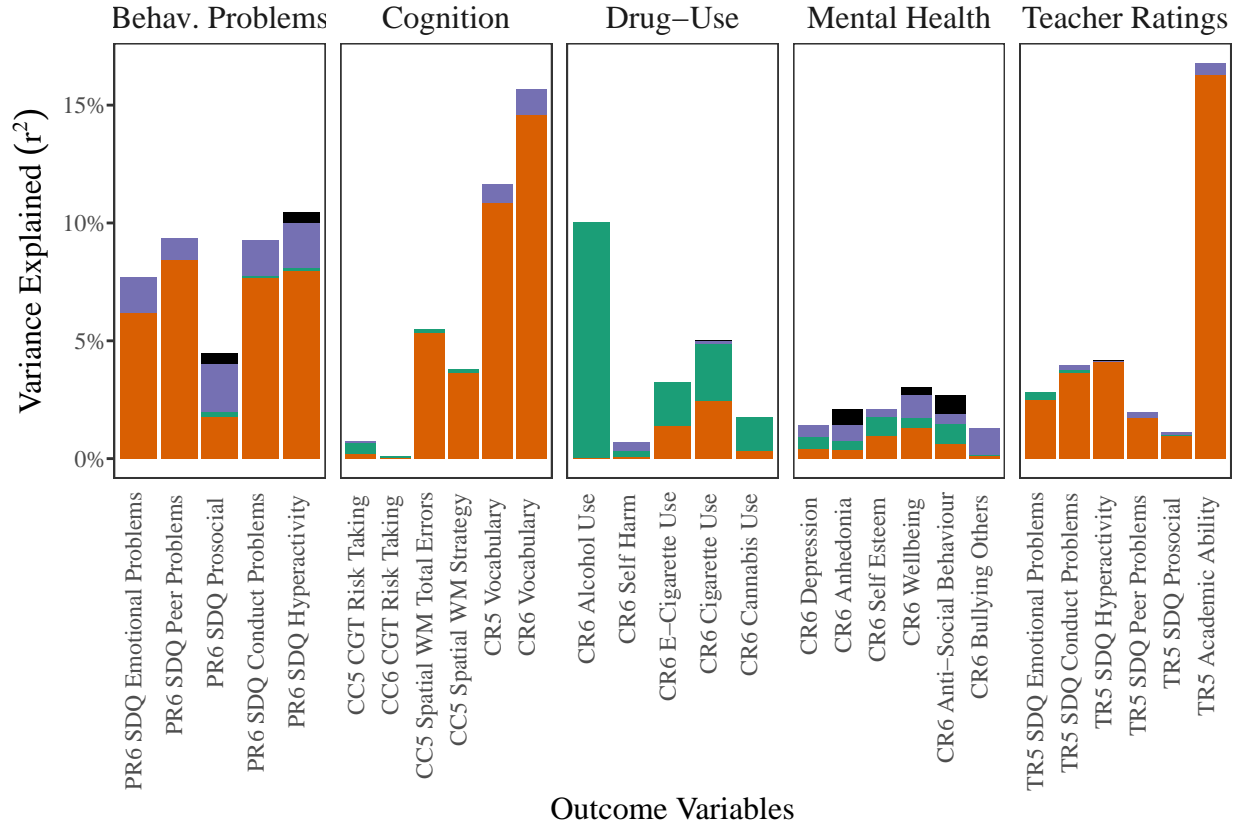
```

PLOT3=
  ggplot(data=PlotData, aes(x=Var,y=value, fill=variable, group=Cluster)) +
    geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="LM_R2",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc10",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc9",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc8",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc7",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc6",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc5",])
    # geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc4",])
    geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc3",],)
    geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc2",],)
    geom_bar(position="identity",stat="identity", alpha=1, data=PlotData[PlotData$variable=="cc1",],)
  labs(y=bquote('Variance Explained'~(r^2)), x="Outcome Variables") +
  facet_wrap(~Cluster,strip.position = "top", scales = "free_x", nrow=1) + theme(panel.spacing = unit(
  jtools::theme_apo()+
  theme(axis.title.y.right = element_blank(),           # hide right axis title
        axis.text.y.right = element_blank(),           # hide right axis labels
        axis.ticks.x = element_blank(),                 # hide left/right axis ticks
        axis.text.y = element_text(margin = margin(r = 0)), # move left axis labels closer to axis
        #panel.spacing = unit(10, "mm"),                 # remove spacing between facets
        strip.background = element_blank(),             # match default line size of theme_classic
        legend.position = "none",
        axis.text.x = element_text(angle = 90,hjust=0.95,vjust=0.2),
        text=element_text(family="serif")
  ) +
    scale_fill_manual(labels = unique(PlotData$variable),values=c( wesanderson::wes_palette("BottleRock",10)
    scale_y_continuous(breaks=c(0,5,10,15)/100, labels=c("0%","5%", "10%", "15%"))

  ggsave(filename=file.path("Plots","VarianceExplained_teach.pdf"), PLOT3, device="pdf", width=8, height=8)
  ggsave(filename=file.path("Plots","VarianceExplained_teach.png"), PLOT3, device="png", width=8, height=8)

PLOT3 #+ ylim(c(0,.19))

```



## Supplementary Figure 2

```
# apply(A3_teach$R2_matrix,2,mean)
# apply(A3_teach$R2_matrix,2,mean)
```

### Variance Explained in each outcome tables

Correlation Confidence Interval Fucntion

```
.CorrelationCIEstimator = function(r, n, alpha=0.05){
  Fr = base::atanh(r) # Fisher Z Transform
  SE = 1/((n-3)^.5) # Standard Error
  CI = stats::qnorm(c(alpha/2,1-alpha/2), mean=Fr, sd=SE)
  CI = base::tanh(CI)
  # p = (1-pnorm(abs(Fr), mean=0, sd=SE))*2 # Fisher Z P value
  t = r*base::sqrt((n-2)/(1-r^2)) # P-value estimated from t-distribution
  p = (1-stats::pt(base::abs(t), df=n-2))*2
  return(base::list(CI=CI,p=p))
}
```

Table 2

Variance Explained for each outcome variables

```

absSquare = function(x){
  neg = x<0
  out = x^2
  out[neg] = out[neg]*-1
  return(out)
}

cc1 = cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined$cc1
# First CCA variate
# cc1_CI_LB = sapply(cc1, function(x)
#                   .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_group2), alpha=0.05)$CI[1]) #Need to
# cc1_CI_UB = sapply(cc1, function(x)
#                   .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_group2), alpha=0.05)$CI[2])

# Second CCA variate
cc2 = cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined$cc2
# cc2_CI_LB = sapply(cc2, function(x)
#                   .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_group2), alpha=0.05)$CI[1])
# cc2_CI_UB = sapply(cc2, function(x)
#                   .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_group2), alpha=0.05)$CI[2])

# Third CCA variate
cc3 = cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined$cc3
# cc3_CI_LB = sapply(cc3, function(x)
#                   .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_group2), alpha=0.05)$CI[1])
# cc3_CI_UB = sapply(cc3, function(x)
#                   .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_group2), alpha=0.05)$CI[2])

# Variance Explained by Linear Model
LM_VE = cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined$LM_R2

Table2 =
cbind.data.frame(cc1, #absSquare(cc1_CI_LB), absSquare(cc1_CI_UB),
                 cc2, #absSquare(cc2_CI_LB), absSquare(cc2_CI_UB),
                 cc3, #absSquare(cc3_CI_LB), absSquare(cc3_CI_UB),
                 LM_VE
                 )

AverageVE = apply(Table2, 2, mean)
Table2 = rbind.data.frame(Table2, AverageVE)

Table2 %>%
  # format(.,nsmall=3, digits=0) %>%
  # mutate_all(.,str_replace_all, pattern = "0\\.\"", replacement="\\.") %>%
  dplyr::slice(c(Outcome_Order,23)) %>%
  'rownames<-'(c(Outcome_Labels_Ordered, "Mean")) %>%
  'colnames<-'(c("cc1","cc2","cc3","LM")) %>%
  knitr::kable(digits=3)

```

	cc1	cc2	cc3	LM
PR6 SDQ Emotional Problems	0.054	0.055	0.070	0.078
PR6 SDQ Peer Problems	0.070	0.071	0.081	0.080
PR6 SDQ Prosocial	0.014	0.019	0.032	0.036
PR6 SDQ Conduct Problems	0.076	0.084	0.095	0.093
PR6 SDQ Hyperactivity	0.074	0.085	0.098	0.098
CR6 Depression	0.001	0.010	0.012	0.014
CR6 Anhedonia	-0.001	0.008	0.011	0.010
CR6 Self Esteem	0.004	0.016	0.017	0.017
CR6 Wellbeing	0.007	0.019	0.026	0.027
CR6 Anti-Social Behaviour	0.008	0.033	0.034	0.037
CR6 Bullying Others	0.001	0.002	0.012	0.009
CR6 Alcohol Use	0.001	0.091	0.092	0.094
CR6 Self Harm	-0.001	0.008	0.009	0.007
CR6 E-Cigarette Use	0.011	0.035	0.035	0.033
CR6 Cigarette Use	0.022	0.062	0.062	0.059
CR6 Cannabis Use	0.003	0.022	0.022	0.018
CC5 CGT Risk Taking	0.005	0.008	0.008	0.010
CC6 CGT Risk Taking	0.003	0.004	0.004	0.004
CC5 Spatial WM Total Errors	0.049	0.049	0.050	0.044
CC5 Spatial WM Strategy	0.031	0.031	0.032	0.029
CR5 Vocabulary	0.102	0.102	0.107	0.106
CR6 Vocabulary	0.127	0.127	0.142	0.144
Mean	0.030	0.043	0.048	0.047

Table2 %>%

```
dplyr::slice(c(Outcome_Order,23)) %>%
  'rownames<-'(c(Outcome_Labels_Ordered, "Mean")) %>%
  'colnames<-'(c("cc1","cc2","cc3","LM")) %>%
  format(., digits=0, nsmall=3) %>%
  apply(., 2, function(x) sub("0\\.","\\.",x, perl=TRUE)) %>%
  write.csv(., file=file.path("Output R Data", "Table2.csv"), row.names = TRUE)
```

```
rm(cc1,#cc1_CI_LB,cc1_CI_UB,
   cc2,#cc2_CI_LB,cc2_CI_UB,
   cc3,#cc3_CI_LB,cc3_CI_UB,
   LM_VE
)
```

```
format(cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined, digits=0, nsmall=4)[,c(1:4,22,23)]
```

```
##          cc1    cc2    cc3    cc4    cc22  LM_R2
## S6_Parent_SDQ_Emotion    0.0537 0.0550 0.0700 0.0763 0.0777 0.0777
## S6_Parent_SDQ_Conduct    0.0760 0.0840 0.0952 0.0934 0.0932 0.0932
## S6_Parent_SDQ_Hyper      0.0741 0.0848 0.0975 0.0969 0.0976 0.0976
## S6_Parent_SDQ_Peer       0.0702 0.0713 0.0814 0.0836 0.0799 0.0799
## S6_Parent_SDQ_Prosocial   0.0137 0.0189 0.0322 0.0376 0.0362 0.0362
## S6_MentalHealth_HateSelf   0.0012 0.0103 0.0124 0.0150 0.0136 0.0136
## S6_MentalHealth_Tired     -0.0014 0.0076 0.0109 0.0131 0.0103 0.0103
```

```
## S6_SelfEsteem          0.0040 0.0159 0.0173 0.0182 0.0167 0.0167
## S6_Wellbeing           0.0068 0.0194 0.0260 0.0287 0.0271 0.0271
## S6_RiskyBehaviours_Cig 0.0217 0.0620 0.0618 0.0610 0.0594 0.0589
## S6_RiskyBehaviours_ECig 0.0107 0.0353 0.0354 0.0344 0.0328 0.0325
## S6_RiskyBehaviours_Alcohol 0.0010 0.0910 0.0917 0.0927 0.0943 0.0941
## S6_RiskyBehaviours_Cannabis 0.0035 0.0222 0.0222 0.0218 0.0164 0.0176
## S6_ChildQ_SelfHarm     -0.0013 0.0076 0.0093 0.0111 0.0070 0.0069
## S5_Cog_VS_Ability      0.1017 0.1020 0.1071 0.1069 0.1056 0.1056
## S6_Cog_WordScoreIRT    0.1272 0.1270 0.1424 0.1460 0.1436 0.1436
## S5_Cog_SWM_TotalErrors48 0.0487 0.0489 0.0504 0.0505 0.0441 0.0441
## S5_Cog_SWM_Strategy    0.0307 0.0312 0.0320 0.0323 0.0288 0.0288
## S5_Cog_CGT_RiskTaking  0.0055 0.0076 0.0082 0.0087 0.0097 0.0097
## S6_Cog_CGT_RiskTaking  0.0028 0.0039 0.0036 0.0032 0.0041 0.0041
## S6_AntiSocialBehaviour 0.0079 0.0325 0.0344 0.0342 0.0368 0.0368
## S6_Bullying            0.0012 0.0023 0.0117 0.0140 0.0090 0.0090
```

```
# format(cca_splithalf_teacher$R2_matrix_unbiased, digits=0, nsmall=4)[,c(1:4,23)]
```

```
apply(cca_splithalf_procrustes$R2_matrix_BinaryOutcomes_Combined,1,which.max)
```

```
##      S6_Parent_SDQ_Emotion      S6_Parent_SDQ_Conduct
##                10                3
##      S6_Parent_SDQ_Hyper      S6_Parent_SDQ_Peer
##                16                4
##      S6_Parent_SDQ_Prosocial  S6_MentalHealth_HateSelf
##                8                8
##      S6_MentalHealth_Tired      S6_SelfEsteem
##                4                7
##                S6_Wellbeing      S6_RiskyBehaviours_Cig
##                6                2
##      S6_RiskyBehaviours_ECig  S6_RiskyBehaviours_Alcohol
##                3                5
## S6_RiskyBehaviours_Cannabis      S6_ChildQ_SelfHarm
##                8                4
##                S5_Cog_VS_Ability      S6_Cog_WordScoreIRT
##                8                4
##      S5_Cog_SWM_TotalErrors48      S5_Cog_SWM_Strategy
##                4                4
##                S5_Cog_CGT_RiskTaking      S6_Cog_CGT_RiskTaking
##                10                6
##      S6_AntiSocialBehaviour      S6_Bullying
##                5                8
```

**Table 2 - Including Teacher Data**

Variance Explained for each outcome variables - TEACHER DATA INCLUDED

```
cc1 = cca_splithalf_teacher$R2_matrix_BinaryOutcomes_Combined$cc1
# First CCA variate
# cc1_CI_LB = sapply(cc1, function(x)
#               .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_Teach_group2), alpha=0.05)$CI[1])
# cc1_CI_UB = sapply(cc1, function(x)
```

```

#                               .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_Teach_group2), alpha=0.05)$CI[2])

# Second CCA variate
cc2 = cca_splithalf_teacher$R2_matrix_BinaryOutcomes_Combined$cc2
# cc2_CI_LB = sapply(cc2, function(x)
#                               .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_Teach_group2), alpha=0.05)$CI[1])
# cc2_CI_UB = sapply(cc2, function(x)
#                               .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_Teach_group2), alpha=0.05)$CI[2])

# Third CCA variate
cc3 = cca_splithalf_teacher$R2_matrix_BinaryOutcomes_Combined$cc3
# cc3_CI_LB = sapply(cc3, function(x)
#                               .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_Teach_group2), alpha=0.05)$CI[1])
# cc3_CI_UB = sapply(cc3, function(x)
#                               .CorrelationCIEstimator(x^.5, n=nrow(df0_imputed_Teach_group2), alpha=0.05)$CI[2])

# Variance Explained by Linear MOdel
LM_VE = cca_splithalf_teacher$R2_matrix_BinaryOutcomes_Combined$LM_R2

Table2 =
cbind.data.frame(cc1,
                 cc2,
                 cc3,
                 LM_VE
                 )

AverageVE = apply(Table2, 2, mean)
Table2 = rbind.data.frame(Table2, AverageVE)

Table2 %>%
  # format(.,nsmall=3, digits=0) %>%
  # mutate_all(.,str_replace_all, pattern = "0\\.\\.", replacement="\\.\\.") %>%
  dplyr::slice(c(match(Outcome_Ordered2, outcomes2),29)) %>%
  'rownames<-'(c(Outcome_Labels_Ordered2, "Mean")) %>%
  'colnames<-'(c("cc1","cc2","cc3","LM")) %>%
  knitr::kable(digits=3)

```

	cc1	cc2	cc3	LM
TR5 SDQ Emotional Problems	0.025	0.028	0.028	0.022
TR5 SDQ Conduct Problems	0.036	0.038	0.040	0.040
TR5 SDQ Hyperactivity	0.041	0.041	0.042	0.042
TR5 SDQ Peer Problems	0.017	0.017	0.020	0.010
TR5 SDQ Prosocial	0.010	0.010	0.011	0.008
TR5 Academic Ability	0.163	0.162	0.168	0.166
PR6 SDQ Emotional Problems	0.062	0.062	0.077	0.077
PR6 SDQ Peer Problems	0.084	0.083	0.094	0.075
PR6 SDQ Prosocial	0.018	0.020	0.040	0.045
PR6 SDQ Conduct Problems	0.077	0.077	0.093	0.088
PR6 SDQ Hyperactivity	0.080	0.081	0.100	0.104
CR6 Depression	0.004	0.009	0.014	0.009
CR6 Anhedonia	0.004	0.008	0.014	0.021
CR6 Self Esteem	0.009	0.018	0.021	0.020

	cc1	cc2	cc3	LM
CR6 Wellbeing	0.013	0.017	0.027	0.030
CR6 Anti-Social Behaviour	0.006	0.015	0.019	0.027
CR6 Bullying Others	0.001	0.002	0.013	0.004
CR6 Alcohol Use	-0.001	0.100	0.100	0.100
CR6 Self Harm	0.000	0.003	0.007	-0.002
CR6 E-Cigarette Use	0.014	0.032	0.032	0.031
CR6 Cigarette Use	0.024	0.049	0.050	0.050
CR6 Cannabis Use	0.003	0.018	0.018	0.000
CC5 CGT Risk Taking	0.002	0.007	0.007	0.001
CC6 CGT Risk Taking	-0.001	0.001	0.001	-0.008
CC5 Spatial WM Total Errors	0.053	0.055	0.055	0.049
CC5 Spatial WM Strategy	0.036	0.038	0.035	0.026
CR5 Vocabulary	0.108	0.108	0.116	0.108
CR6 Vocabulary	0.146	0.146	0.157	0.157
Mean	0.037	0.044	0.050	0.046

```
rm(cc1,#cc1_CI_LB,cc1_CI_UB,
    cc2,#cc2_CI_LB,cc2_CI_UB,
    cc3,#cc3_CI_LB,cc3_CI_UB,
    LM_VE
)
```

## Supplementary Alcohol Plots

```
load(file.path("Data","MCS Data","Processed Data","DataWithExclusionsNoImputations","df0_BothGroups_F"))
```

```
##SES and alcohol use - SUPPLEMENTARY MATERIALS
```

```
#Recode variables so that higher alcohol use is greater, and normalise vars
```

```
df0$S1_Parent_AlcoholCurrent_REVERSED = (Convert(df0$S1_Parent_AlcoholCurrent,1:7,7:1))
```

```
## before
```

```
##      1      2      3      4      5      6      7
## 404  377 1419 4468 3819 3562 4480
```

```
## after
```

```
##      1      2      3      4      5      6      7
## 4480 3562 3819 4468 1419  377  404
```

```
df0$S1_Parent_AlcoholPregnant_REVERSED = (Convert(df0$S1_Parent_AlcoholPregnant,1:7,7:1))
```

```
## before
```

```
##      1      2      3      4      5      6      7
##   68   48  201 1270 1268 2461 13180
```

```
## after
```

```
##      1      2      3      4      5      6      7
## 13180 2461 1268 1270  201   48   68
```



```
df0$S1_Parent_AlcoholCurrent_REVERSED_NORM = (Convert(df0$S1_Parent_AlcoholCurrent,1:7,7:1))
```

```
## before
##      1      2      3      4      5      6      7
## 404 377 1419 4468 3819 3562 4480
## after
##      1      2      3      4      5      6      7
## 4480 3562 3819 4468 1419 377 404
```

```
df0$S1_Parent_AlcoholPregnant_REVERSED_NORM = (Convert(df0$S1_Parent_AlcoholPregnant,1:7,7:1))
```

```
## before
##      1      2      3      4      5      6      7
##    68    48   201  1270  1268  2461 13180
## after
##      1      2      3      4      5      6      7
## 13180 2461  1268  1270   201    48    68
```

```
# brewer.pal(n = 4, name = "Dark2")
```

```
# Parents and children's alcohol consumption (without using normed data)
```

```
# ggplot(data=df0, aes(y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholCurrent_REVERSED)) + geom_hline(y=
# jtools::theme_apa() + scale_x_continuous(breaks=1:7, labels = c("Never", "<1 Month", "1-2 Times a
# geom_boxplot(aes(group=S1_Parent_AlcoholCurrent_REVERSED),col="black", fill="#1B9E77", outlier.sh
# coord_flip() + labs(y="Equivilised Income (ADOEDE00) Normalised",x="Alcohol Consumption After Pre
#
#
```

```
# ggplot(data=df0, aes(y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholPregnant_REVERSED)) + geom_hline(
# jtools::theme_apa() + scale_x_continuous(breaks=1:7, labels = c("Never", "<1 Month", "1-2 Times a
# geom_boxplot(aes(group=S1_Parent_AlcoholPregnant_REVERSED),col="black", fill="#D95F02", outlier.s
# coord_flip() + labs(y="Equivilised Income (ADOEDE00) Normalised",x="Alcohol Consumption During Pr
```

```
# Alcohol Consumption after pregnancy and family income
```

```
p1=
```

```
ggplot(data=df0, aes(y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholCurrent_REVERSED)) + geom_hline(yin
geom_smooth(aes( y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholCurrent_REVERSED_NORM), method="lm",
geom_jitter(aes(y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholCurrent_REVERSED_NORM-.35),width=.095,
jtools::theme_apa() +
scale_x_continuous(breaks=sort(na.omit(unique(df0$S1_Parent_AlcoholCurrent_REVERSED_NORM))), labels
geom_boxplot(aes(group=S1_Parent_AlcoholCurrent_REVERSED_NORM, y=S1_SES_EquivIncome_Norm,x=S1_Paren
coord_flip() + labs(y="Equivilised Income (ADOEDE00) Normalised",x="Alcohol Consumption After Pregna
ggtitle("A")
```

```
# Alcohol Consumption DURING pregnancy and family income
```

```
p2=
```

```
ggplot(data=df0, aes(y=S1_Parent_AlcoholCurrent_REVERSED_NORM,x=S1_Parent_AlcoholPregnant_REVERSED_NORM
geom_smooth(aes( y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholPregnant_REVERSED_NORM), method="lm",
geom_jitter(aes(y=S1_SES_EquivIncome_Norm,x=S1_Parent_AlcoholPregnant_REVERSED_NORM-.35),width=.095,
jtools::theme_apa() +
scale_x_continuous(breaks=sort(na.omit(unique(df0$S1_Parent_AlcoholPregnant_REVERSED_NORM))), label
geom_boxplot(aes(group=S1_Parent_AlcoholPregnant_REVERSED_NORM, y=S1_SES_EquivIncome_Norm,x=S1_Paren
coord_flip() + labs(y="Equivilised Income (ADOEDE00) Normalised",x="Alcohol Consumption During Preg
```

```

ggtitle("B")

#Parent and children's alcohol
df0$S6_RiskyBehaviours_Alcohol_factor = factor(df0$S6_RiskyBehaviours_Alcohol, levels=c(0,1),labels=c(
"0","1"))

#Tables used to add count information to plots
Table_1 = table(df0$S6_RiskyBehaviours_Alcohol_factor, df0$S1_Parent_AlcoholCurrent_REVERSED_NORM)
Table_1 = melt(Table_1)
Table_1$y = rep(c(.85,.15),7)
Table_2 = table(df0$S6_RiskyBehaviours_Alcohol_factor, df0$S1_Parent_AlcoholPregnant_REVERSED_NORM)
Table_2 = melt(Table_2)
Table_2$y = rep(c(.85,.15),7)

#Parent alcohol consumption after pregnancy and CHILDREN'S consumption
p3=
ggplot(data=df0[!is.na(df0$S6_RiskyBehaviours_Alcohol),], aes(x=(S1_Parent_AlcoholCurrent_REVERSED_NORM),
y=(S6_RiskyBehaviours_Alcohol_factor), fill="fill", stat="count", width=.4) +
scale_x_continuous(breaks=sort(na.omit(unique(df0$S1_Parent_AlcoholCurrent_REVERSED_NORM))), labels=
c("0","25","50","75","100")), theme(jtools::theme_apo() + theme(axis.text.x = element_text(angle = 90)) +
labs(x="Alcohol Consumption After Pregnancy (amldr00)", y = "Children's Alcohol Consumption (Pink :
S6_RiskyBehaviours_Alcohol_factor", angle=90) + theme(legend.position = "right")) +
scale_fill_manual(values = c("#1B9E77", "#E7298A"))+
scale_y_continuous(breaks=seq(0,1,by=.25), labels=c("0%", "25%", "50%", "75%", "100%")) +
ggtitle("C"))

#Parent alcohol consumption during pregnancy and CHILDREN'S consumption
p4=
ggplot(data=df0[!is.na(df0$S6_RiskyBehaviours_Alcohol),], aes(x=(S1_Parent_AlcoholPregnant_REVERSED_NORM),
y=(S6_RiskyBehaviours_Alcohol_factor), fill="fill", stat="count", width=.4) +
scale_x_continuous(breaks=sort(na.omit(unique(df0$S1_Parent_AlcoholPregnant_REVERSED_NORM))), labels=
c("0","25","50","75","100")), theme(jtools::theme_apo() + theme(axis.text.x = element_text(angle = 90)) +
labs(x="Alcohol Consumption During Pregnancy (amdpr00)", y = "Children's Alcohol Consumption (Pink :
S6_RiskyBehaviours_Alcohol_factor", angle=90) + theme(legend.position = "right")) +
scale_fill_manual(values = c("#1B9E77", "#E7298A"))+
scale_y_continuous(breaks=seq(0,1,by=.25), labels=c("0%", "25%", "50%", "75%", "100%")) +
ggtitle("D"))

# Combine plots
p1 + p2 + p3 + p4

```

```
## Warning: Removed 896 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 714 rows containing missing values (stat_boxplot).
```

```
## Warning: Removed 182 rows containing non-finite values (stat_boxplot).
```

```
## Warning: Removed 896 rows containing missing values (geom_point).
```

```
## Warning: Removed 929 rows containing non-finite values (stat_smooth).
```

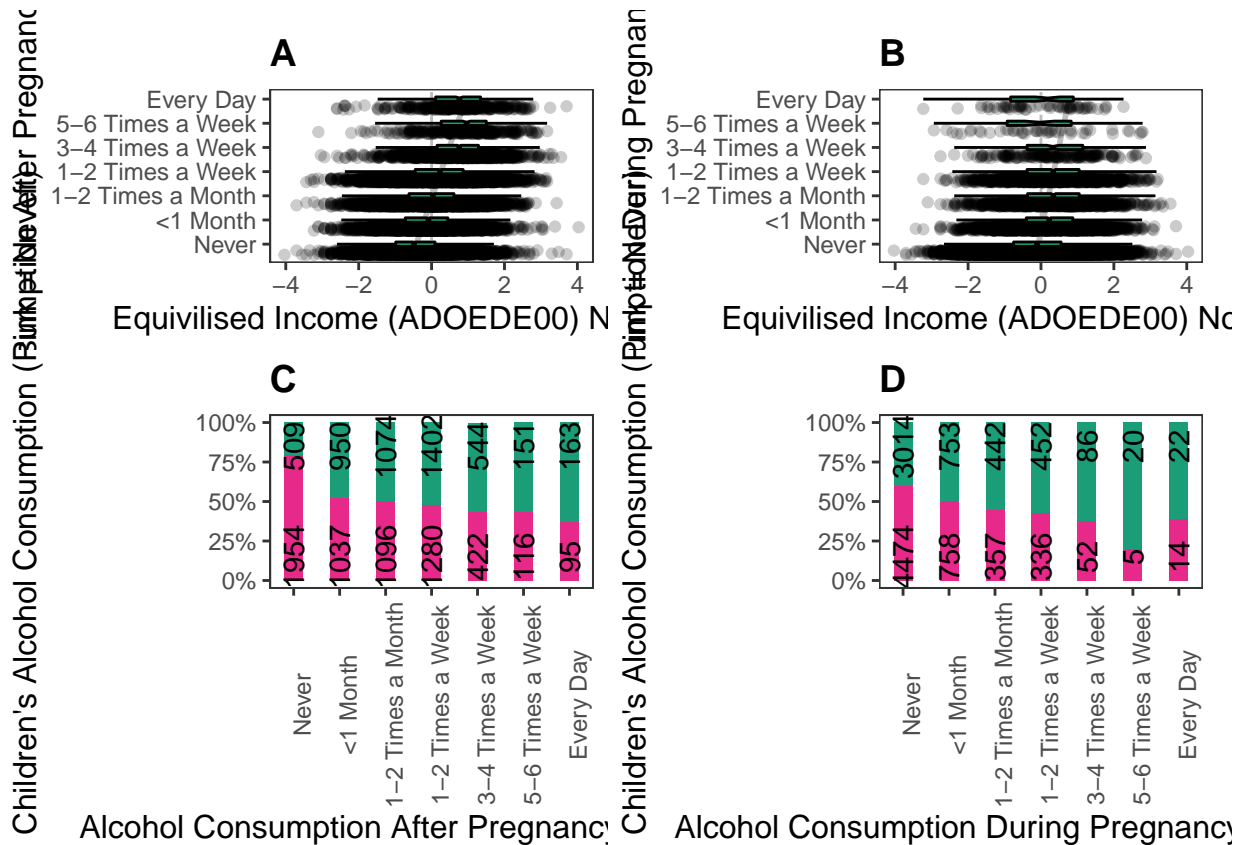
```
## Warning: Removed 747 rows containing missing values (stat_boxplot).

## Warning: Removed 182 rows containing non-finite values (stat_boxplot).

## Warning: Removed 929 rows containing missing values (geom_point).

## Warning: Removed 400 rows containing non-finite values (stat_count).

## Warning: Removed 408 rows containing non-finite values (stat_count).
```



```
ggsave(file="Plots/Alcohol_SupplementaryFigure.png", device="png", width = 15, height = 10)
```

```
## Warning: Removed 896 rows containing non-finite values (stat_smooth).

## Warning: Removed 714 rows containing missing values (stat_boxplot).

## Warning: Removed 182 rows containing non-finite values (stat_boxplot).

## Warning: Removed 896 rows containing missing values (geom_point).

## Warning: Removed 929 rows containing non-finite values (stat_smooth).

## Warning: Removed 747 rows containing missing values (stat_boxplot).
```

```
## Warning: Removed 182 rows containing non-finite values (stat_boxplot).
```

```
## Warning: Removed 929 rows containing missing values (geom_point).
```

```
## Warning: Removed 400 rows containing non-finite values (stat_count).
```

```
## Warning: Removed 408 rows containing non-finite values (stat_count).
```

```
pdf(file="Plots/Alcohol_SupplementaryFigure.pdf", width=15, height=10)
p1 + p3 + p2 + p4
```

```
## Warning: Removed 896 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 714 rows containing missing values (stat_boxplot).
```

```
## Warning: Removed 182 rows containing non-finite values (stat_boxplot).
```

```
## Warning: Removed 896 rows containing missing values (geom_point).
```

```
## Warning: Removed 400 rows containing non-finite values (stat_count).
```

```
## Warning: Removed 929 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 747 rows containing missing values (stat_boxplot).
```

```
## Warning: Removed 182 rows containing non-finite values (stat_boxplot).
```

```
## Warning: Removed 929 rows containing missing values (geom_point).
```

```
## Warning: Removed 408 rows containing non-finite values (stat_count).
```

```
dev.off()
```

```
## pdf
```

```
## 2
```