

MSKIDS: Exploratory Analysis and ComBat Harmonization with Healthy Controls

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18 November, 2020, 15:03

Batch Variables

```
##
## SIEMENSPRISMAFIT    SIEMENSTIMTRIO    SIEMENSVERIO
##                58                1192                36
```

```
##
##  CHP  HSC  PNC
##   36   65 1185
```

Scanners by sites

```
##
##                CHP  HSC  PNC
## SIEMENSPRISMAFIT    0  58    0
## SIEMENSTIMTRIO      0   7 1185
## SIEMENSVERIO       36   0    0
```

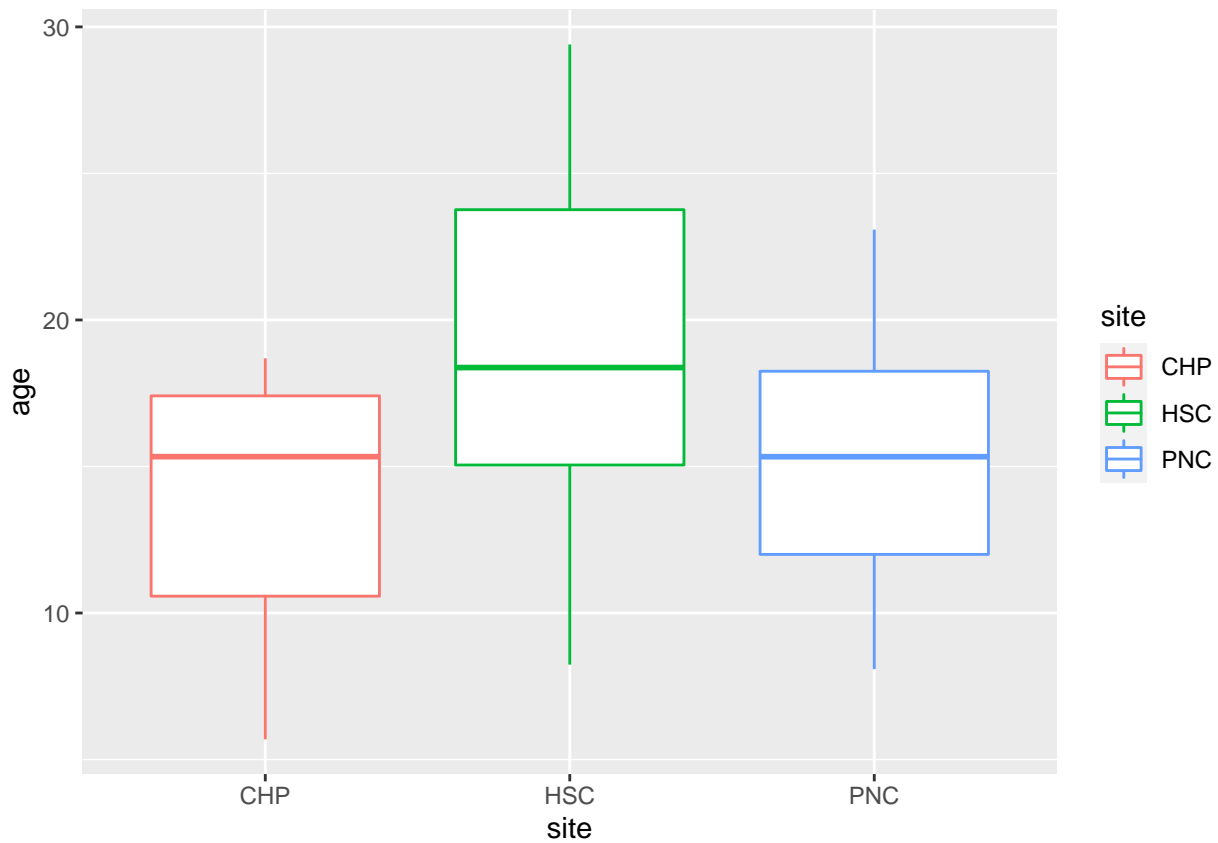
Excluding HSC-SIEMENSTIMTRIO (n=7) for now.

```
## [1] "Batches:"
```

```
## [1] "CHP-SIEMENSVERIO"      "HSC-SIEMENSPRISMAFIT" "PNC-SIEMENSTIMTRIO"
```

Confounding across sites

Age



```
## Analysis of Variance Table
##
## Response: age
##           Df Sum Sq Mean Sq F value    Pr(>F)
## site       2  1012.5   506.27   34.606 2.313e-15 ***
## Residuals 1276 18667.2    14.63
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Age is confounded across sites.

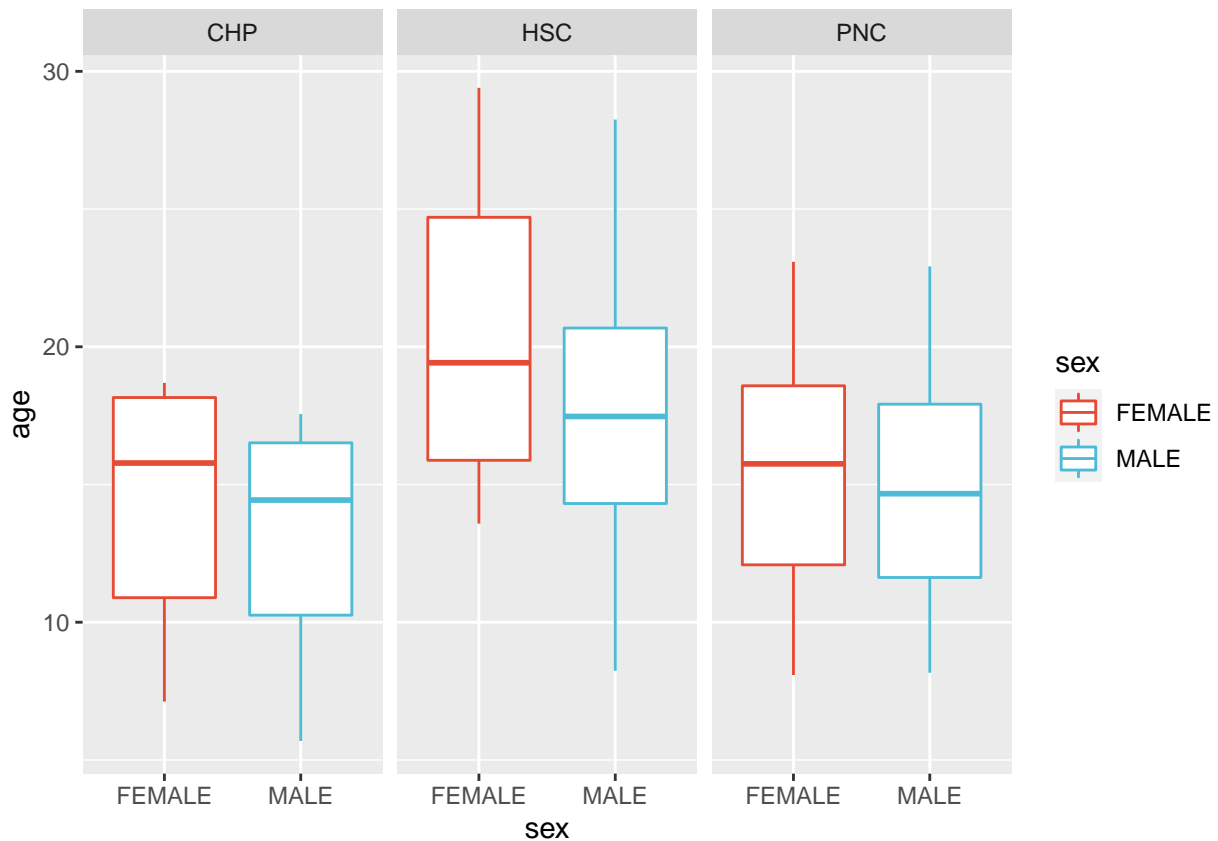
Sex

```
##
##      FEMALE MALE
##  CHP      24  12
##  HSC      37  21
##  PNC     630 555
```

```
##
##      FEMALE      MALE
##  CHP 0.6666667 0.3333333
##  HSC 0.6379310 0.3620690
##  PNC 0.5316456 0.4683544
```

```
##
## Pearson's Chi-squared test
##
## data: df$site and df$sex
## X-squared = 4.8976, df = 2, p-value = 0.0864
```

Sex slightly confounded.



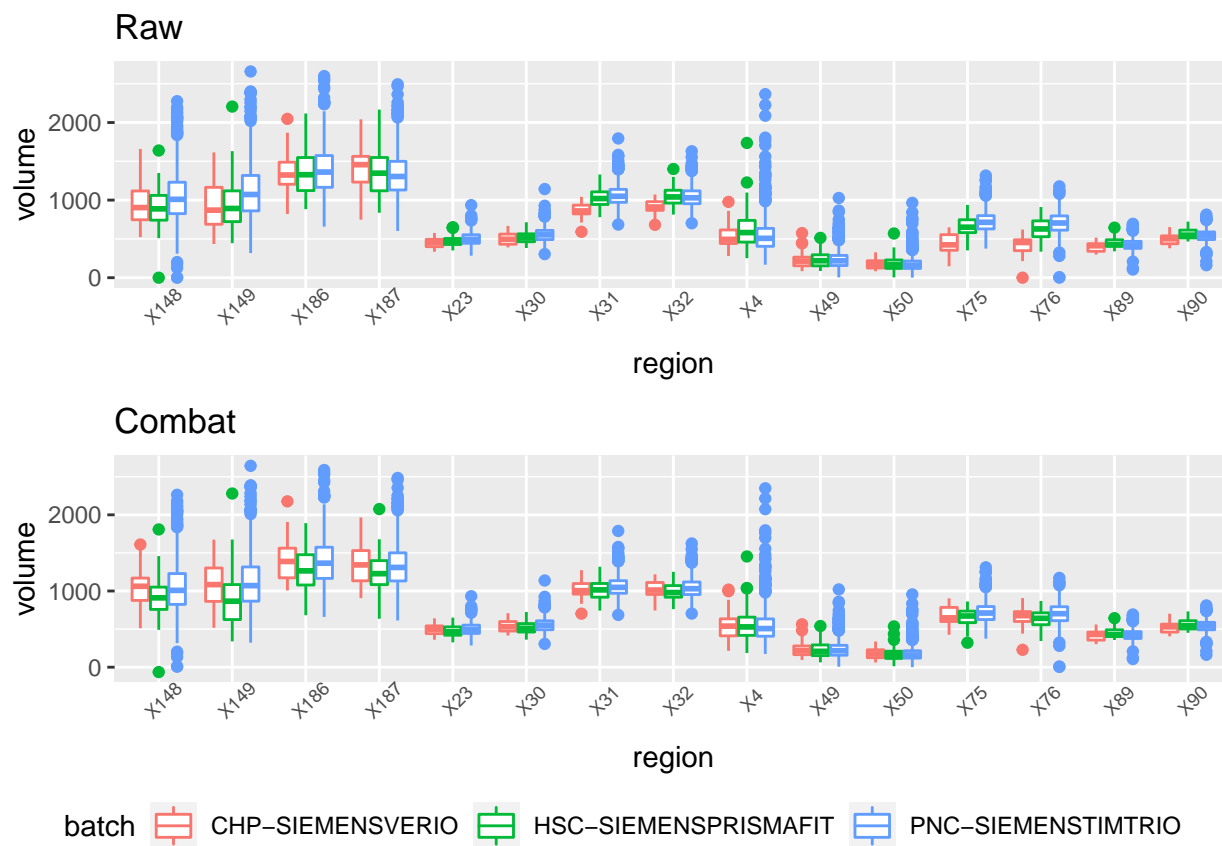
ROI volumes

Number of significant ANOVA after accounting for sites

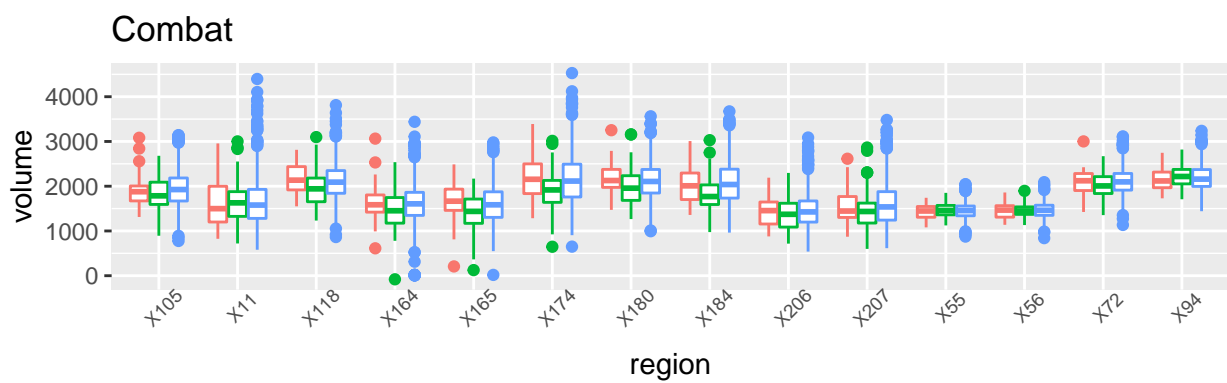
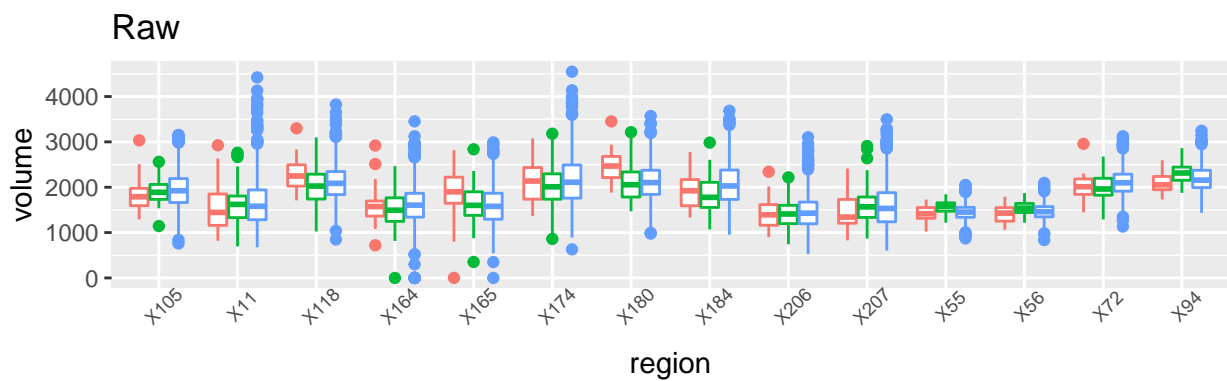
```
##      Raw Combat
##      33      1
```




ROI plots

```
## [[1]]
```

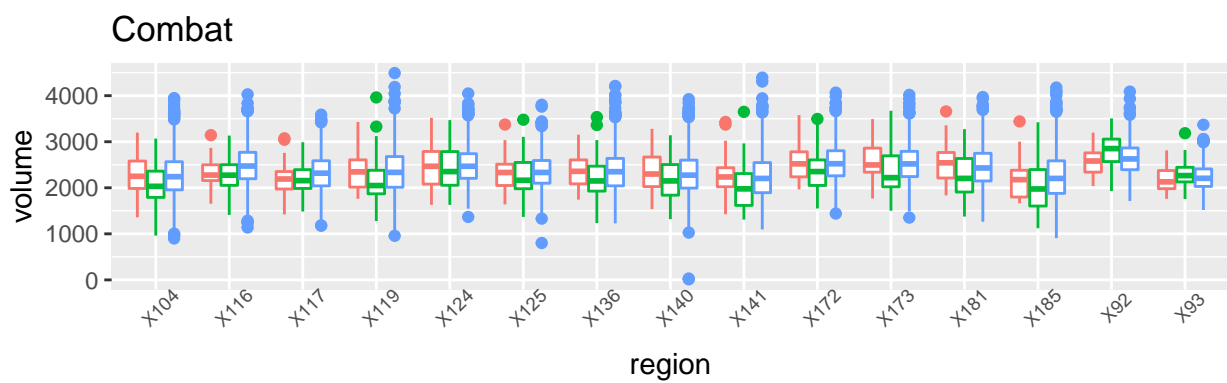
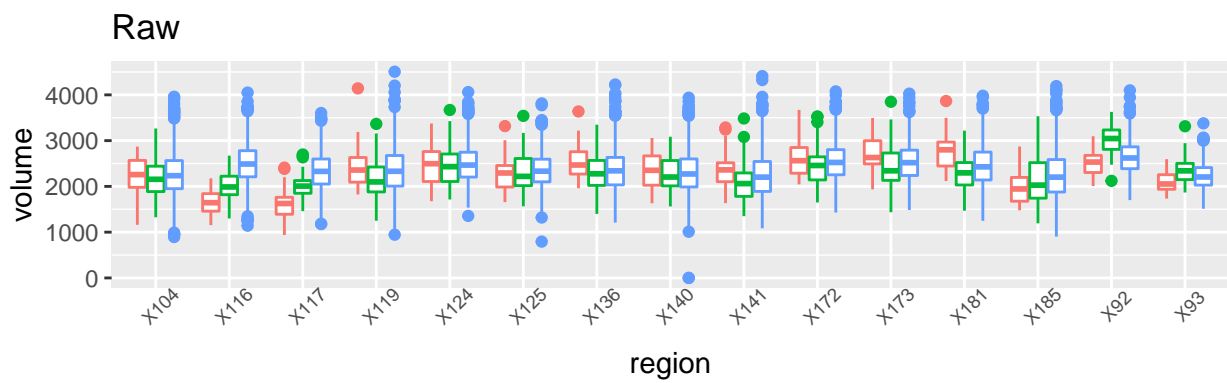





```
##
## [[2]]
```



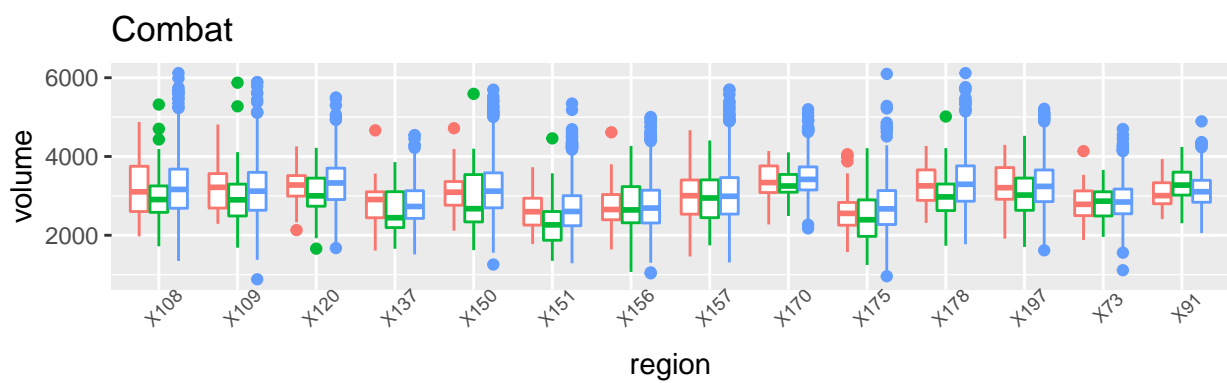
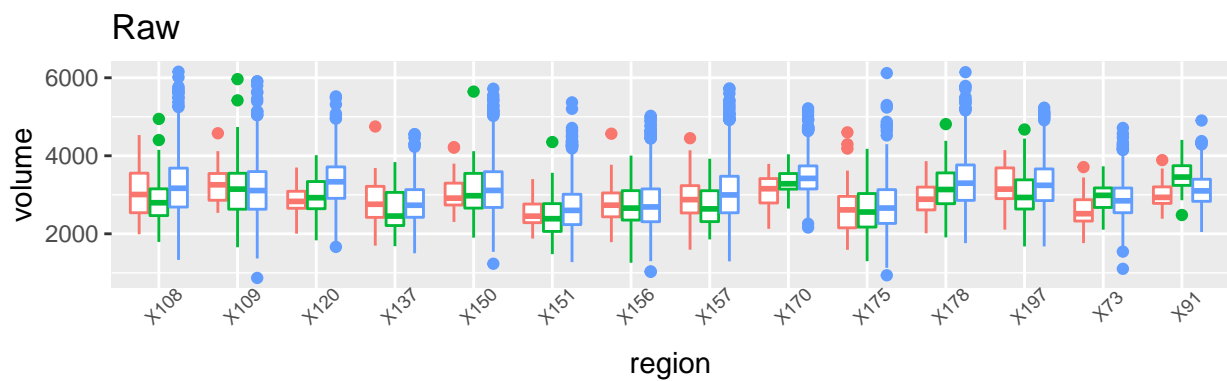
batch  CHP-SIEMENSVERIO  HSC-SIEMENSPRISMAFIT  PNC-SIEMENSTIMTRIO




[[3]]



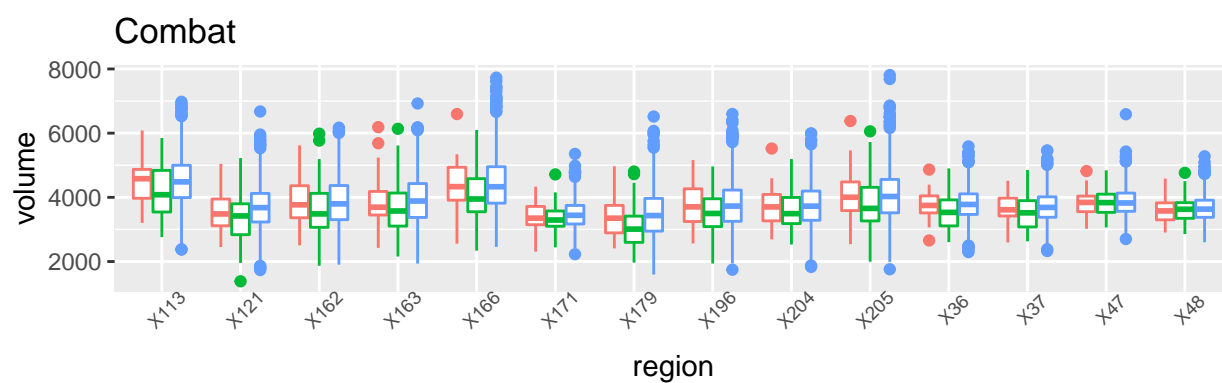
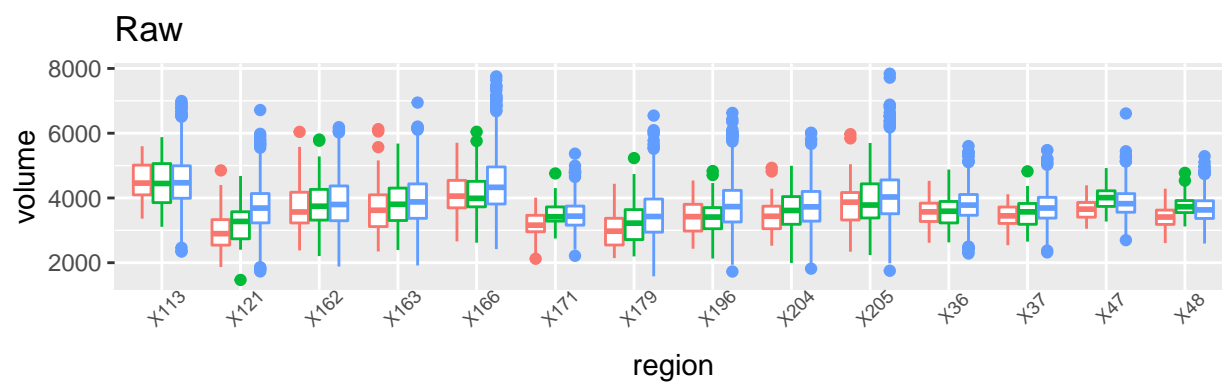
batch  CHP-SIEMENSVERIO  HSC-SIEMENSPRISMAFIT  PNC-SIEMENSTIMTRIO




```
##
## [[4]]
```



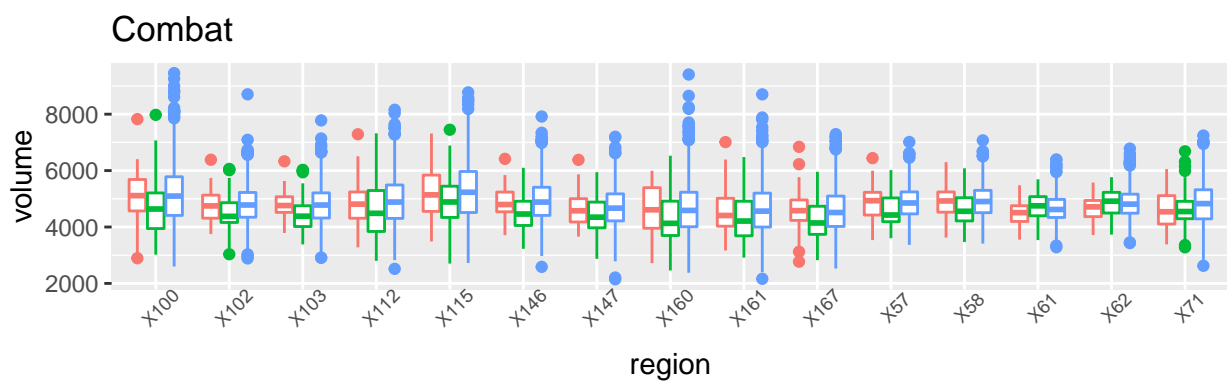
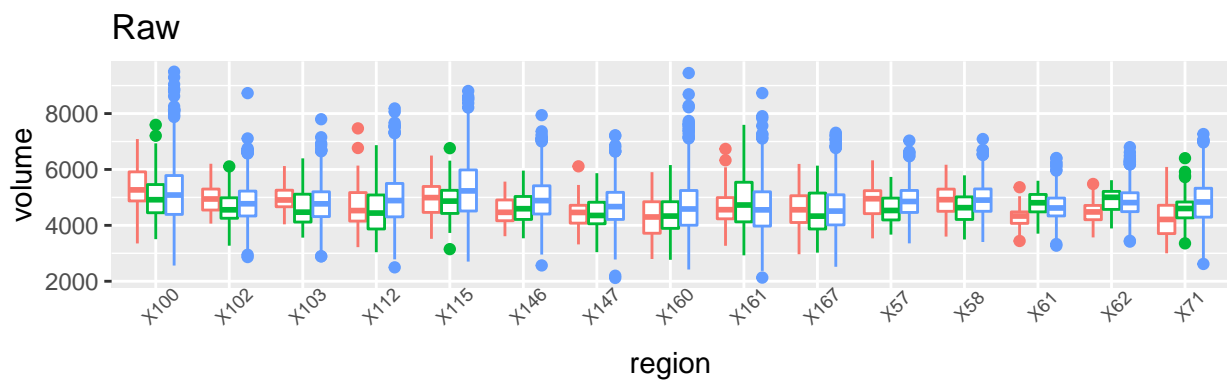
batch  CHP-SIEMENSVERIO  HSC-SIEMENSPRISMAFIT  PNC-SIEMENSTIMTRIO




[[5]]



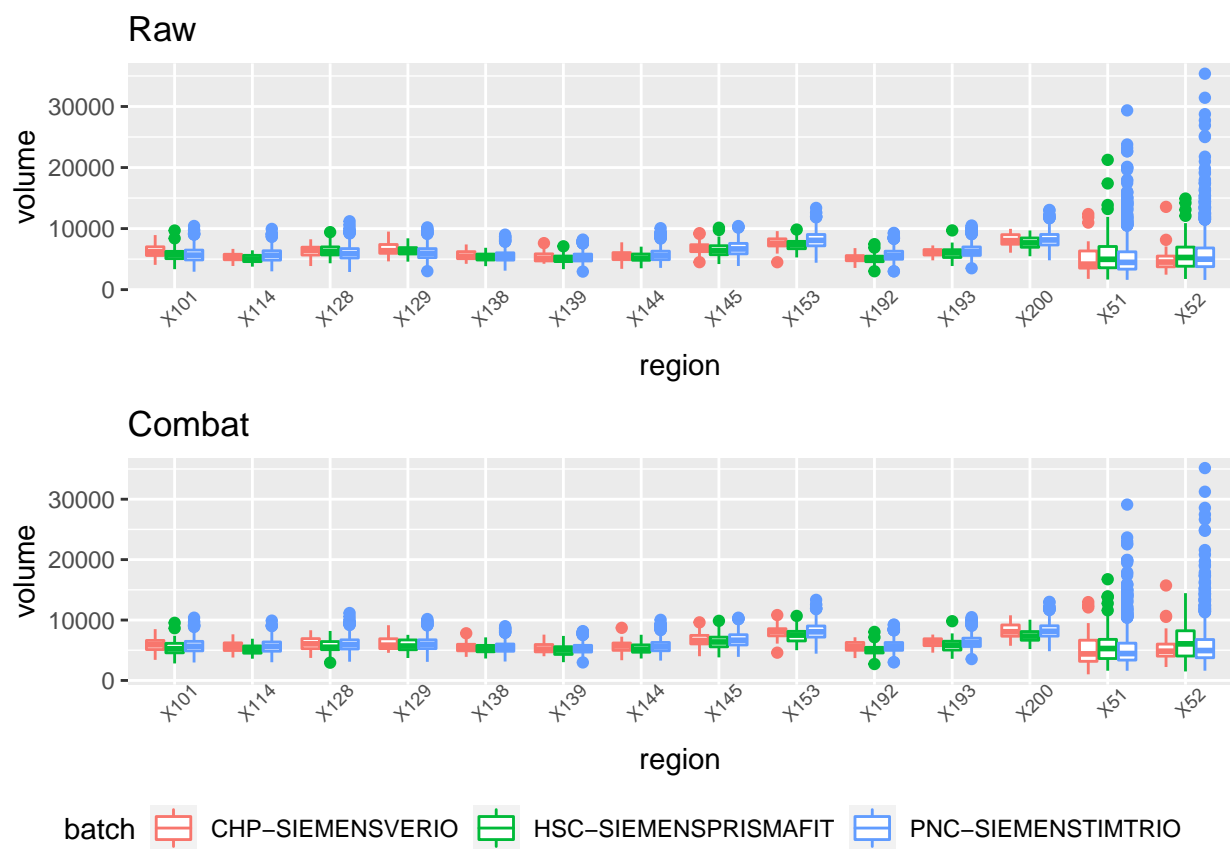
batch  CHP-SIEMENSVERIO  HSC-SIEMENSPRISMAFIT  PNC-SIEMENSTIMTRIO

[[6]]

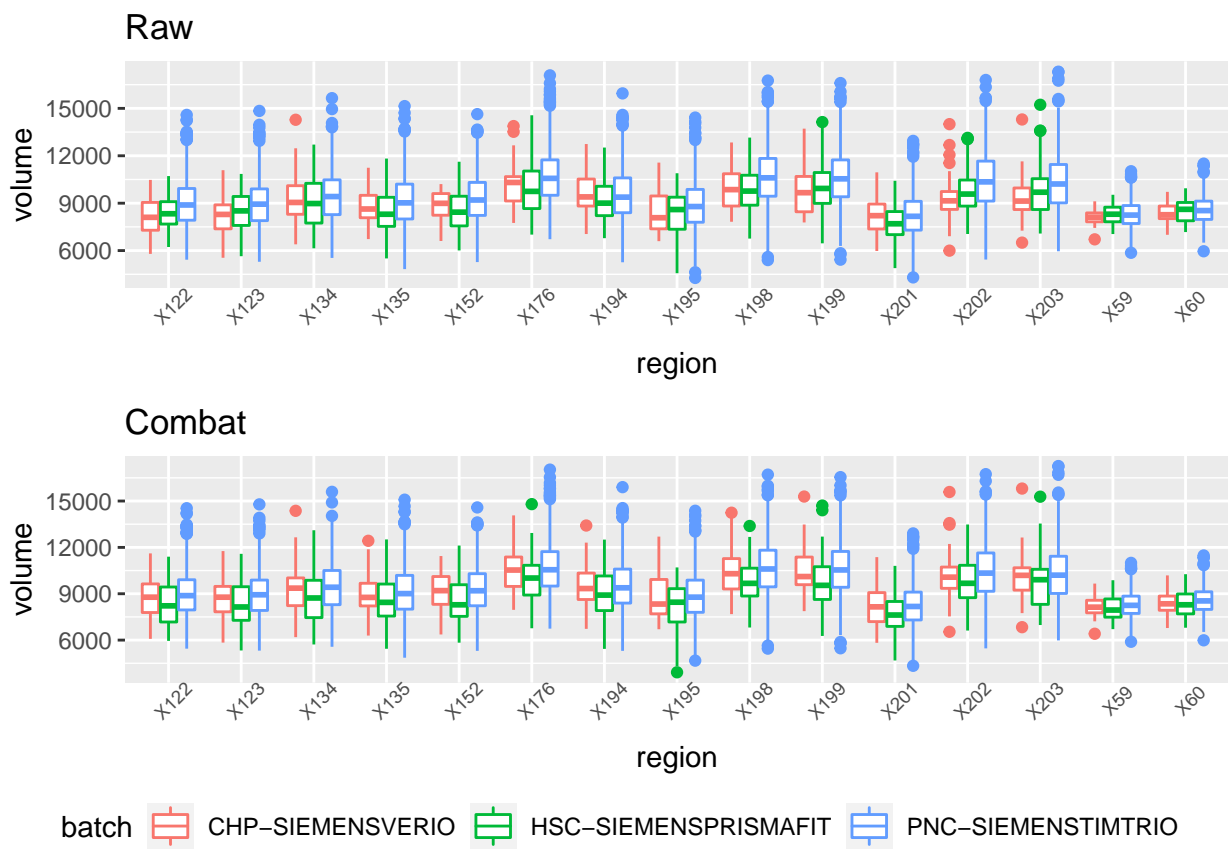


batch  CHP-SIEMENSVERIO  HSC-SIEMENSPRISMAFIT  PNC-SIEMENSTIMTRIO

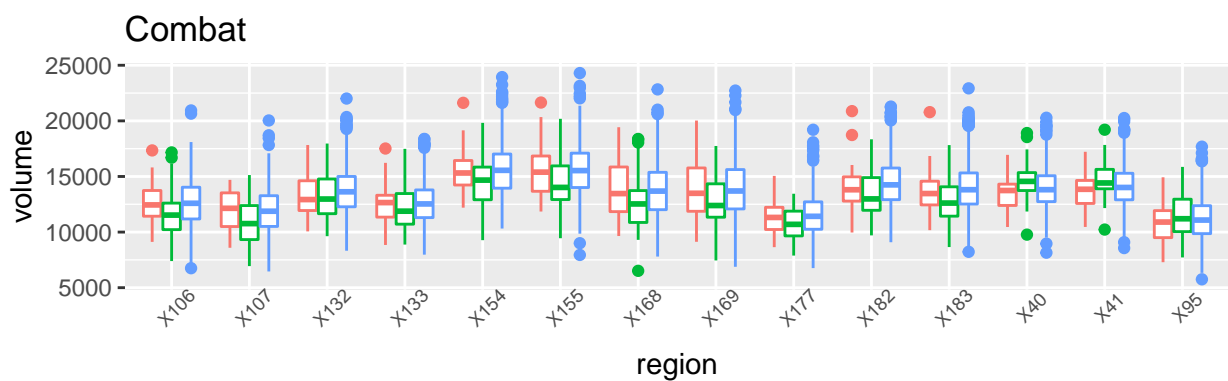
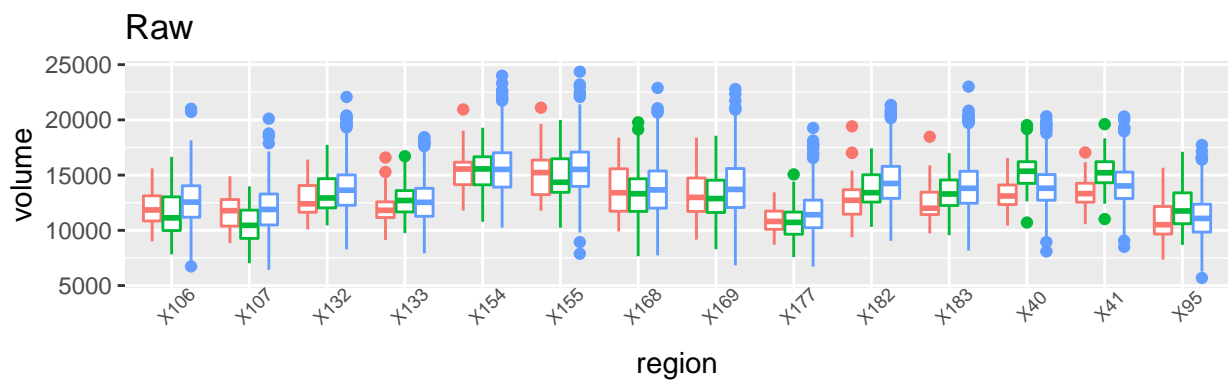
[[7]]






[[8]]

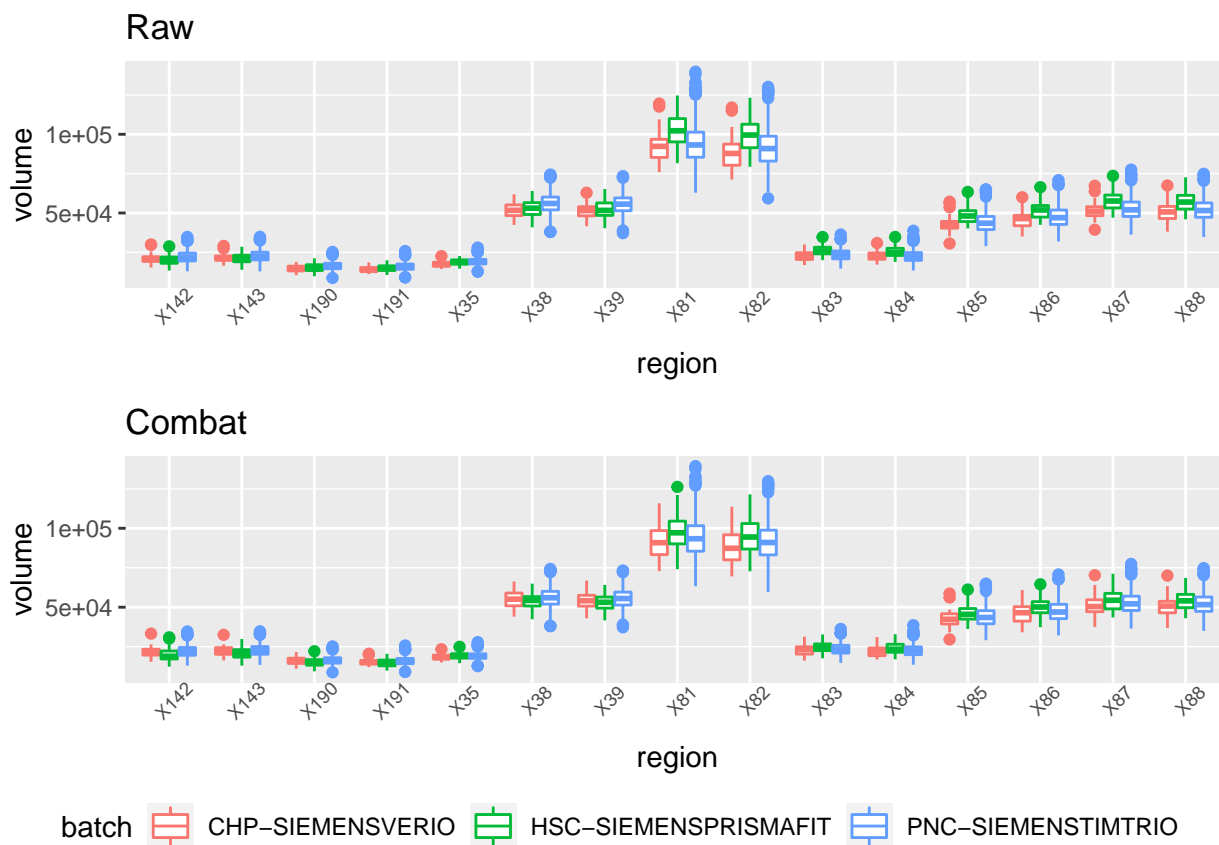


[[9]]



batch  CHP-SIEMENSVERIO  HSC-SIEMENSPRISMAFIT  PNC-SIEMENSTIMTRIO

```
##
## [[10]]
```

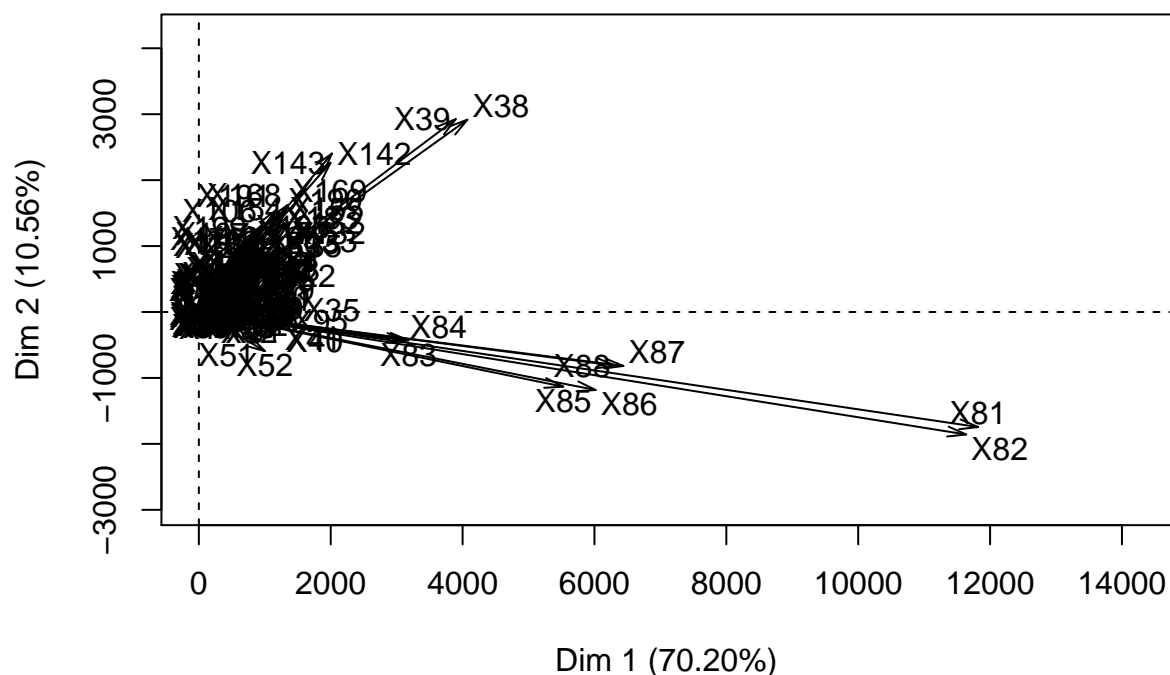


(Covariance) PCA

- PC1: total volume (w/ higher contribution from WM areas)
- PC2: GM/WM
- PC3: Cerebellum
- PC4: L. Ventricles vs Frontal Lobes

##	eigenvalue	percentage of variance	cumulative percentage of variance
## comp 1	529441070	70.202724	70.20272
## comp 2	79664832	10.563382	80.76611
## comp 3	33906223	4.495891	85.26200
## comp 4	17117761	2.269778	87.53177

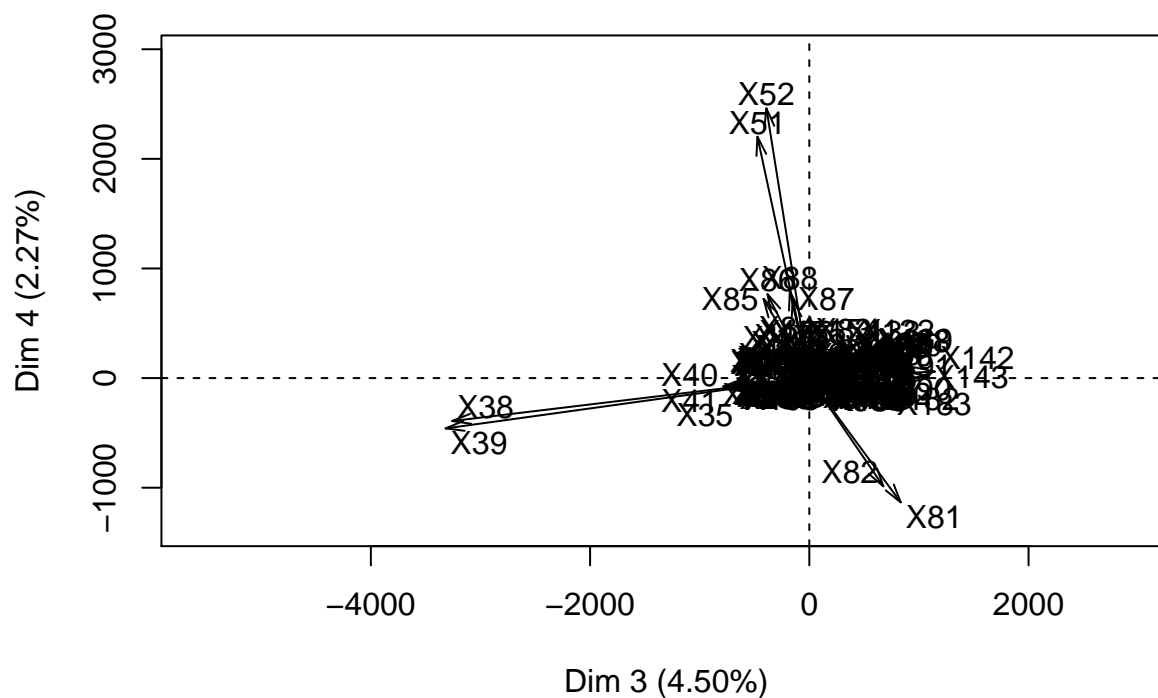
Variables factor map (PCA)



##	contr	ROI_INDEX	ROI_NAME	HEMISPHERE	TISSUE_SEG
## 1	26.382477	81	frontal lobe WM right	Right	WM
## 2	25.582651	82	frontal lobe WM left	Left	WM
## 3	7.830356	87	temporal lobe WM right	Right	WM
## 4	7.542814	88	temporal lobe WM left	Left	WM
## 5	6.840359	86	parietal lobe WM left	Left	WM
## 6	5.770535	85	parietal lobe WM right	Right	WM
## 7	3.132028	38	Right Cerebellum Exterior	Right	GM
## 8	2.868396	39	Left Cerebellum Exterior	Left	GM
## 9	1.909531	83	occipital lobe WM right	Right	WM
## 10	1.848759	84	occipital lobe WM left	Left	WM

##	contr	ROI_INDEX	ROI_NAME	HEMISPHERE	TISSUE_SEG
## 1	10.737997	39	Left Cerebellum Exterior	Left	GM
## 2	10.677509	38	Right Cerebellum Exterior	Right	GM
## 3	7.251971	142	Right MFG middle frontal gyrus	Right	GM
## 4	6.432769	143	Left MFG middle frontal gyrus	Left	GM
## 5	4.332382	82	frontal lobe WM left	Left	WM
## 6	3.820361	81	frontal lobe WM right	Right	WM
## 7	3.296048	169	Left PCu precuneus	Left	GM
## 8	3.053991	168	Right PCu precuneus	Right	GM
## 9	3.012387	191	Left SFG superior frontal gyrus	Left	GM
## 10	2.804837	190	Right SFG superior frontal gyrus	Right	GM

Variables factor map (PCA)

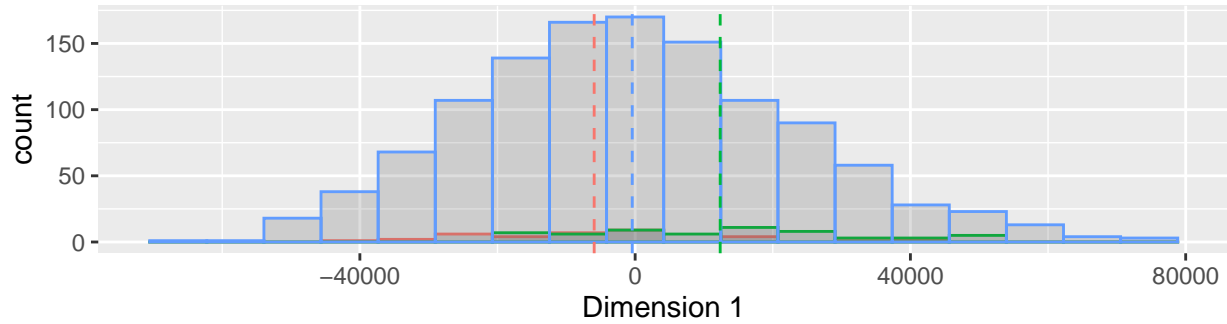


##	contr	ROI_INDEX	ROI_NAME	HEMISPHERE	TISSUE_SEG
## 1	32.469723	39	Left Cerebellum Exterior	Left	GM
## 2	31.319681	38	Right Cerebellum Exterior	Right	GM
## 3	3.867695	142	Right MFG middle frontal gyrus	Right	GM
## 4	3.510285	143	Left MFG middle frontal gyrus	Left	GM
## 5	2.052446	81	frontal lobe WM right	Right	WM
## 6	1.821346	40	Right Cerebellum White Matter	Right	WM
## 7	1.817530	41	Left Cerebellum White Matter	Left	WM
## 8	1.671595	183	Left PrG precentral gyrus	Left	GM
## 9	1.345486	82	frontal lobe WM left	Left	WM
## 10	1.231723	35	Brain Stem	Both	NONE

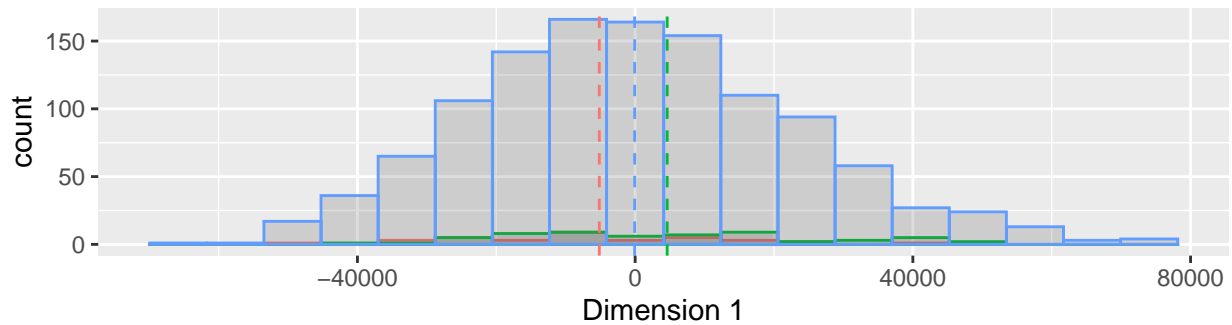
##	contr	ROI_INDEX	ROI_NAME	HEMISPHERE	TISSUE_SEG
## 1	35.3848911	52	Left Lateral Ventricle	Left	VN
## 2	28.3104971	51	Right Lateral Ventricle	Right	VN
## 3	7.5112725	81	frontal lobe WM right	Right	WM
## 4	5.6938487	82	frontal lobe WM left	Left	WM
## 5	3.6520987	88	temporal lobe WM left	Left	WM
## 6	3.4310753	86	parietal lobe WM left	Left	WM
## 7	3.0537511	85	parietal lobe WM right	Right	WM
## 8	3.0201610	87	temporal lobe WM right	Right	WM
## 9	1.2318525	39	Left Cerebellum Exterior	Left	GM
## 10	0.8973908	38	Right Cerebellum Exterior	Right	GM

PC1

Raw Dimension 1 Scores



Harmonized Dimension 1 Scores



batch ■ CHP-SIEMENSVERIO ■ HSC-SIEMENSPRISMAFIT ■ PNC-SIEMENSTIMTRIO

```
## [1] "RAW"
```

```
##
## Call:
## glm(formula = scale(pc1) ~ batch + age + sex, data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.4608  -0.5675  -0.0262   0.5276   2.8222
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -1.592861   0.163874  -9.720  <2e-16 ***
## batchHSC-SIEMENSPRISMAFIT  0.373088   0.179210   2.082  0.0376 *
## batchPNC-SIEMENSTIMTRIO    0.015492   0.140852   0.110  0.9124
## age              0.072177   0.006104  11.824  <2e-16 ***
## sexMALE          0.998936   0.046883  21.307  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.689549)
##
##      Null deviance: 1278.00  on 1278  degrees of freedom
## Residual deviance:  878.49  on 1274  degrees of freedom
```



```

## AIC: 3161.2
##
## Number of Fisher Scoring iterations: 2

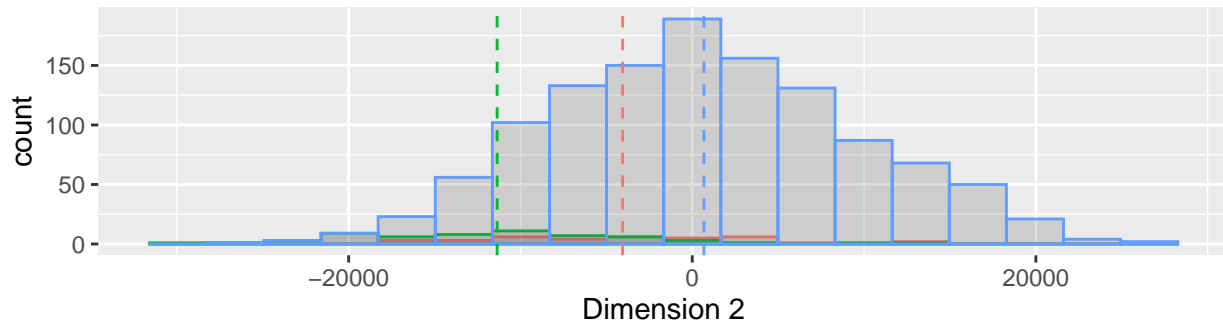
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc1) ~ batch + age + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7952  -0.7223  -0.0701   0.6473   3.2489
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.542306   0.196485  -2.760  0.00586 **
## batchHSC-SIEMENSPRISMAFIT  0.299705   0.214872   1.395  0.16332
## batchPNC-SIEMENSTIMTRIO   0.196690   0.168882   1.165  0.24437
## age              0.023558   0.007319   3.219  0.00132 **
## sexMALE         -0.028981   0.056213  -0.516  0.60625
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9912978)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1262.9  on 1274  degrees of freedom
## AIC: 3625.5
##
## Number of Fisher Scoring iterations: 2

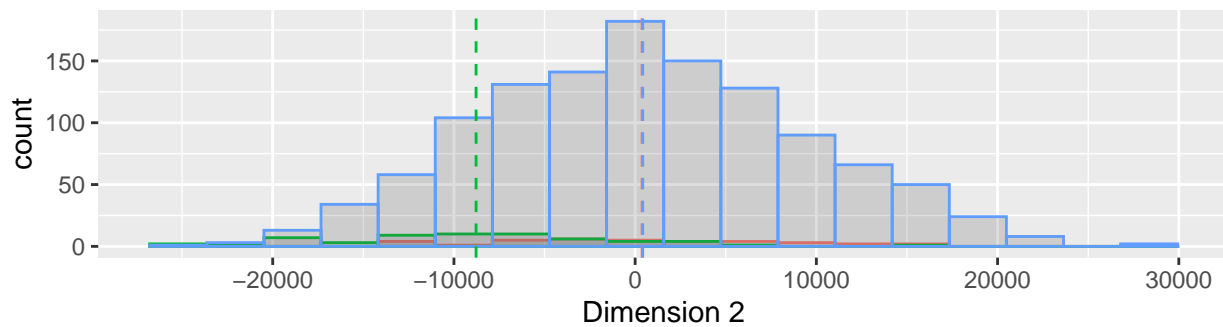
```

PC2

Raw Dimension 2 Scores



Harmonized Dimension 2 Scores



batch ■ CHP-SIEMENSVERIO ■ HSC-SIEMENSPRISMAFIT ■ PNC-SIEMENSTIMTRIO

```
## [1] "RAW"
```

```
##
## Call:
## glm(formula = scale(pc2) ~ batch + age + sex, data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.29198  -0.48643  -0.03406   0.48108   2.05144
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    1.553387   0.135982  11.424 < 2e-16 ***
## batchHSC-SIEMENSPRISMAFIT 0.017259   0.148707   0.116  0.908
## batchPNC-SIEMENSTIMTRIO  0.638097   0.116879   5.459 5.73e-08 ***
## age            -0.155728   0.005065 -30.744 < 2e-16 ***
## sexMALE         0.506897   0.038903  13.030 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.474796)
##
##      Null deviance: 1278.00  on 1278  degrees of freedom
## Residual deviance:  604.89  on 1274  degrees of freedom
```

```

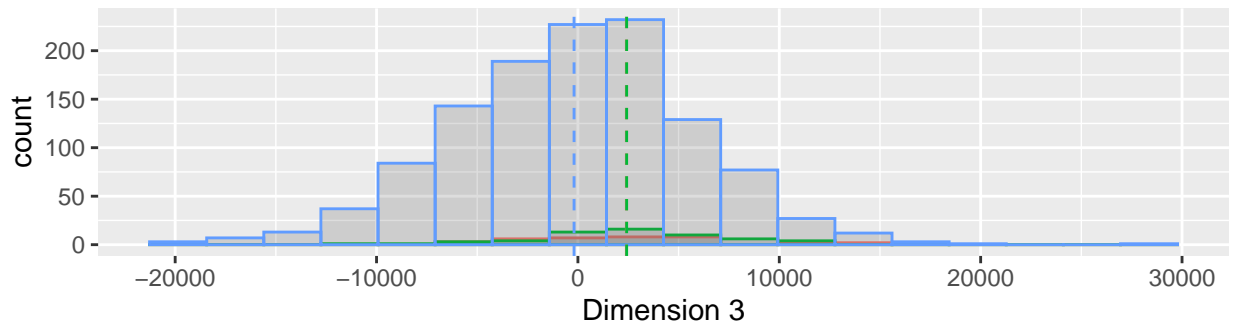
## AIC: 2683.9
##
## Number of Fisher Scoring iterations: 2

## [1] "COMBAT"

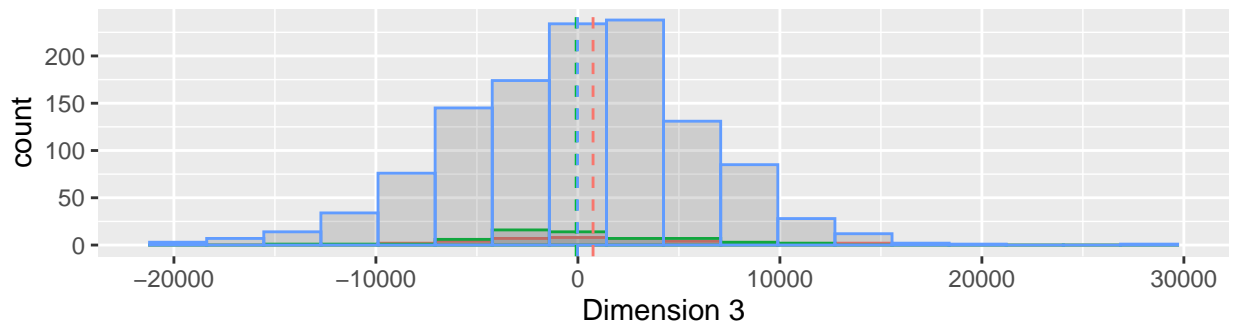
##
## Call:
## glm(formula = scale(pc2) ~ batch + age + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.0459  -0.6557  -0.0314   0.6341   3.1826
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.94783    0.18658   5.080 4.34e-07 ***
## batchHSC-SIEMENSPRISMAFIT -0.67681    0.20404  -3.317 0.000935 ***
## batchPNC-SIEMENSTIMTRIO   0.08105    0.16037   0.505 0.613371
## age            -0.06544    0.00695  -9.416 < 2e-16 ***
## sexMALE         0.01582    0.05338   0.296 0.766950
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.8938674)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1138.8  on 1274  degrees of freedom
## AIC: 3493.1
##
## Number of Fisher Scoring iterations: 2

```

Raw Dimension 3 Scores



Harmonized Dimension 3 Scores



###PC3 batch CHP-SIEMENSVERIO HSC-SIEMENSPRISMAFIT PNC-SIEMENSTIMTRIO

[1] "RAW"

##

Call:

glm(formula = scale(pc3) ~ batch + age + sex, data = dat)

##

Deviance Residuals:

	Min	1Q	Median	3Q	Max
##	-3.2145	-0.6257	0.0497	0.6084	4.3267

##

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	1.547371	0.186955	8.277	3.16e-16 ***
## batchHSC-SIEMENSPRISMAFIT	0.430204	0.204451	2.104	0.0356 *
## batchPNC-SIEMENSTIMTRIO	-0.332042	0.160691	-2.066	0.0390 *
## age	-0.079515	0.006964	-11.418	< 2e-16 ***
## sexMALE	-0.097504	0.053486	-1.823	0.0685 .

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

##

(Dispersion parameter for gaussian family taken to be 0.8974707)

##

Null deviance: 1278.0 on 1278 degrees of freedom

Residual deviance: 1143.4 on 1274 degrees of freedom

AIC: 3498.3

##

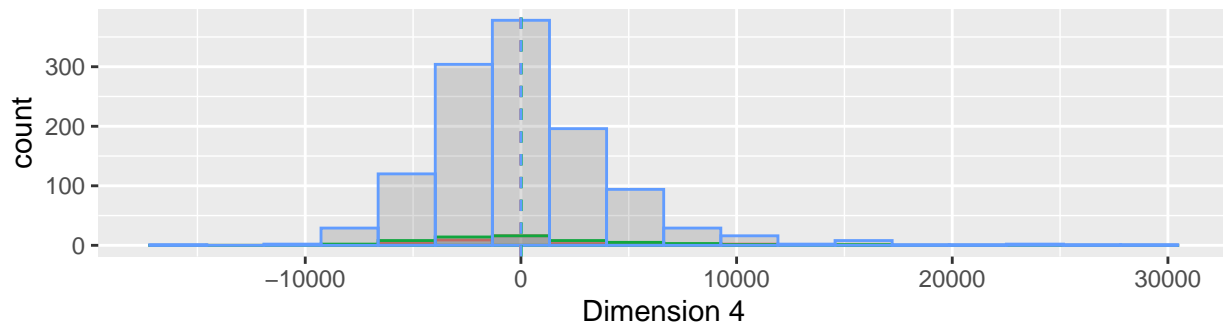
Number of Fisher Scoring iterations: 2

```
## [1] "COMBAT"

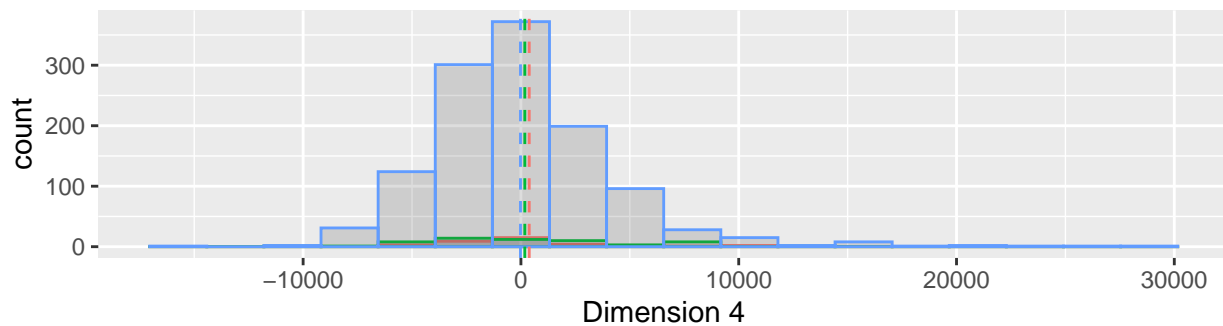
##
## Call:
## glm(formula = scale(pc3) ~ batch + age + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.5735  -0.6406   0.0321   0.6290   4.5153
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      0.60183    0.19597   3.071  0.00218 **
## batchHSC-SIEMENSPRISMAFIT  0.03676    0.21431   0.172  0.86382
## batchPNC-SIEMENSTIMTRIO -0.08939    0.16844  -0.531  0.59571
## age              -0.03369    0.00730 -4.615 4.33e-06 ***
## sexMALE          -0.01337    0.05607  -0.238  0.81156
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9861376)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1256.3  on 1274  degrees of freedom
## AIC: 3618.8
##
## Number of Fisher Scoring iterations: 2
```

PC4

Raw Dimension 4 Scores



Harmonized Dimension 4 Scores



batch ■ CHP-SIEMENSVERIO ■ HSC-SIEMENSPRISMAFIT ■ PNC-SIEMENSTIMTRIO

```
## [1] "RAW"

##
## Call:
## glm(formula = scale(pc4) ~ batch + age + sex, data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.8037  -0.6140  -0.0957   0.4637   7.0235
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.005306   0.197461  -0.027   0.979
## batchHSC-SIEMENSPRISMAFIT  0.016352   0.215941   0.076   0.940
## batchPNC-SIEMENSTIMTRIO  -0.005198   0.169722  -0.031   0.976
## age            -0.001978   0.007355  -0.269   0.788
## sexMALE         0.086108   0.056492   1.524   0.128
##
## (Dispersion parameter for gaussian family taken to be 1.001179)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1275.5  on 1274  degrees of freedom
## AIC: 3638.1
##
## Number of Fisher Scoring iterations: 2
```

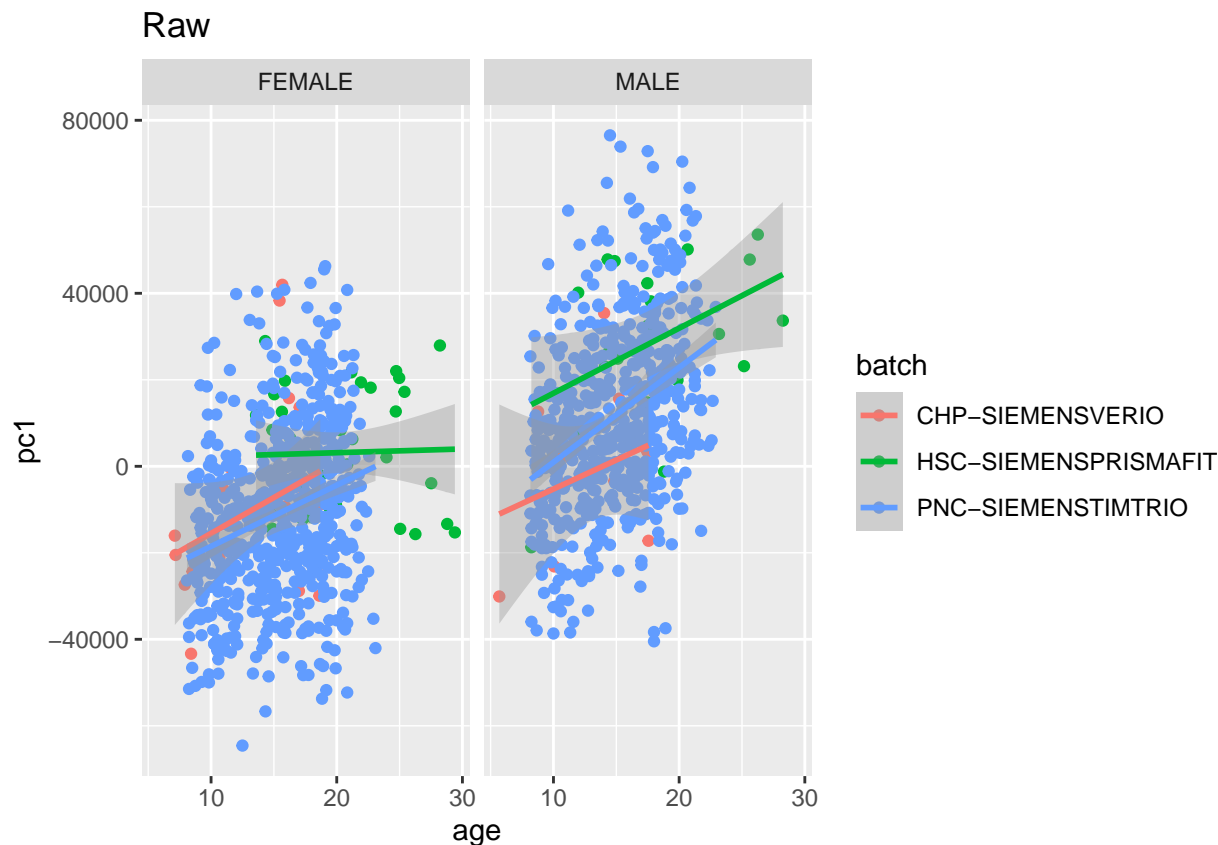
```
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc4) ~ batch + age + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.7491  -0.6170  -0.0837   0.4642   7.0588
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      0.224264   0.197458   1.136   0.256
## batchHSC-SIEMENSPRISMAFIT -0.003739   0.215937  -0.017   0.986
## batchPNC-SIEMENSTIMTRIO  -0.079104   0.169719  -0.466   0.641
## age               -0.008108   0.007355  -1.102   0.271
## sexMALE           -0.058662   0.056491  -1.038   0.299
##
## (Dispersion parameter for gaussian family taken to be 1.001145)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1275.5  on 1274  degrees of freedom
## AIC: 3638.1
##
## Number of Fisher Scoring iterations: 2
```

PCs ~ Age

PC1

```
## `geom_smooth()` using formula 'y ~ x'
```



```
## [1] "RAW"
```

```
##
```

```
## Call:
```

```
## glm(formula = scale(pc1) ~ batch + scale(age) + sex, data = dat)
```

```
##
```

```
## Deviance Residuals:
```

```
##      Min       1Q   Median       3Q      Max
## -2.4608  -0.5675  -0.0262   0.5276   2.8222
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.49052    0.13944  -3.518  0.00045 ***
## batchHSC-SIEMENSPRISMAFIT  0.37309    0.17921   2.082  0.03756 *
## batchPNC-SIEMENSTIMTRIO   0.01549    0.14085   0.110  0.91244
## scale(age)       0.28323    0.02395  11.824 < 2e-16 ***
## sexMALE         0.99894    0.04688  21.307 < 2e-16 ***
```

```
## ---
```

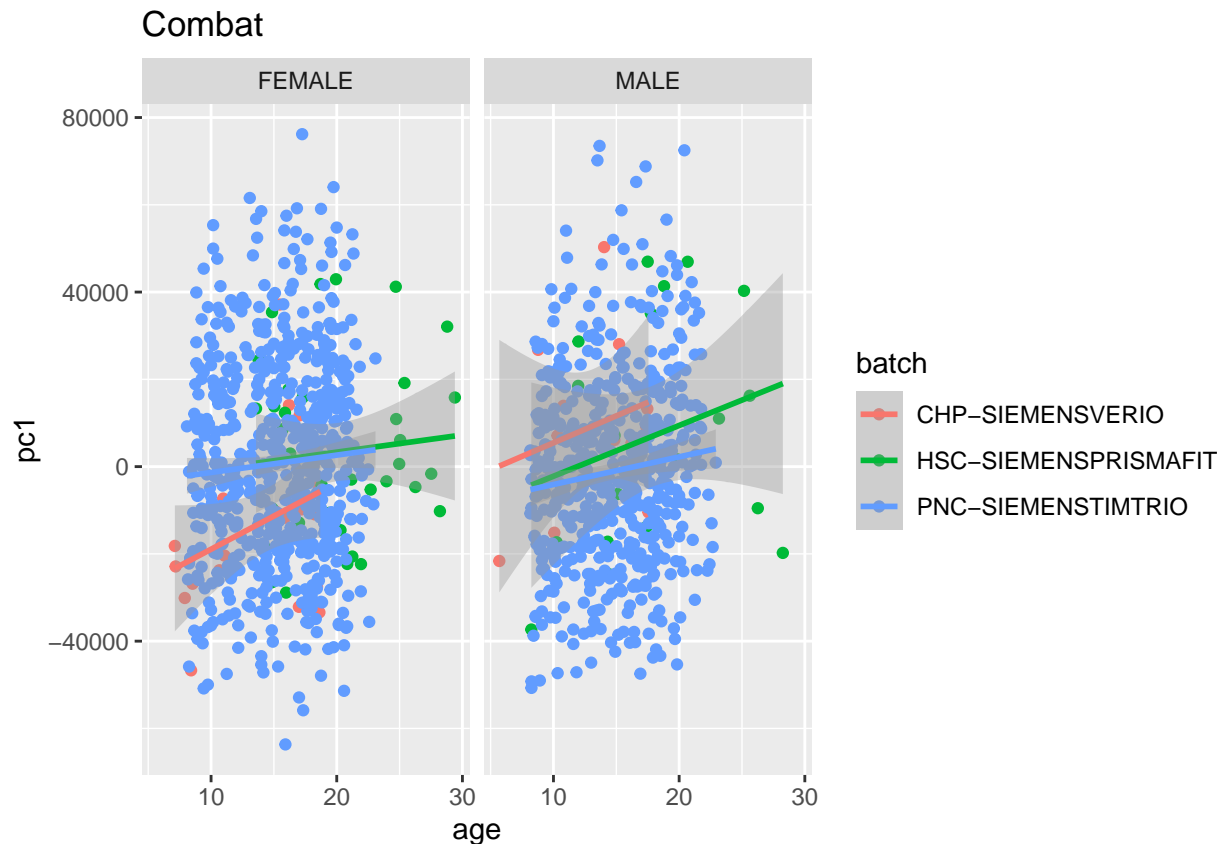
```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
```



```
## (Dispersion parameter for gaussian family taken to be 0.689549)
##
## Null deviance: 1278.00 on 1278 degrees of freedom
## Residual deviance: 878.49 on 1274 degrees of freedom
## AIC: 3161.2
##
## Number of Fisher Scoring iterations: 2

## `geom_smooth()` using formula 'y ~ x'
```



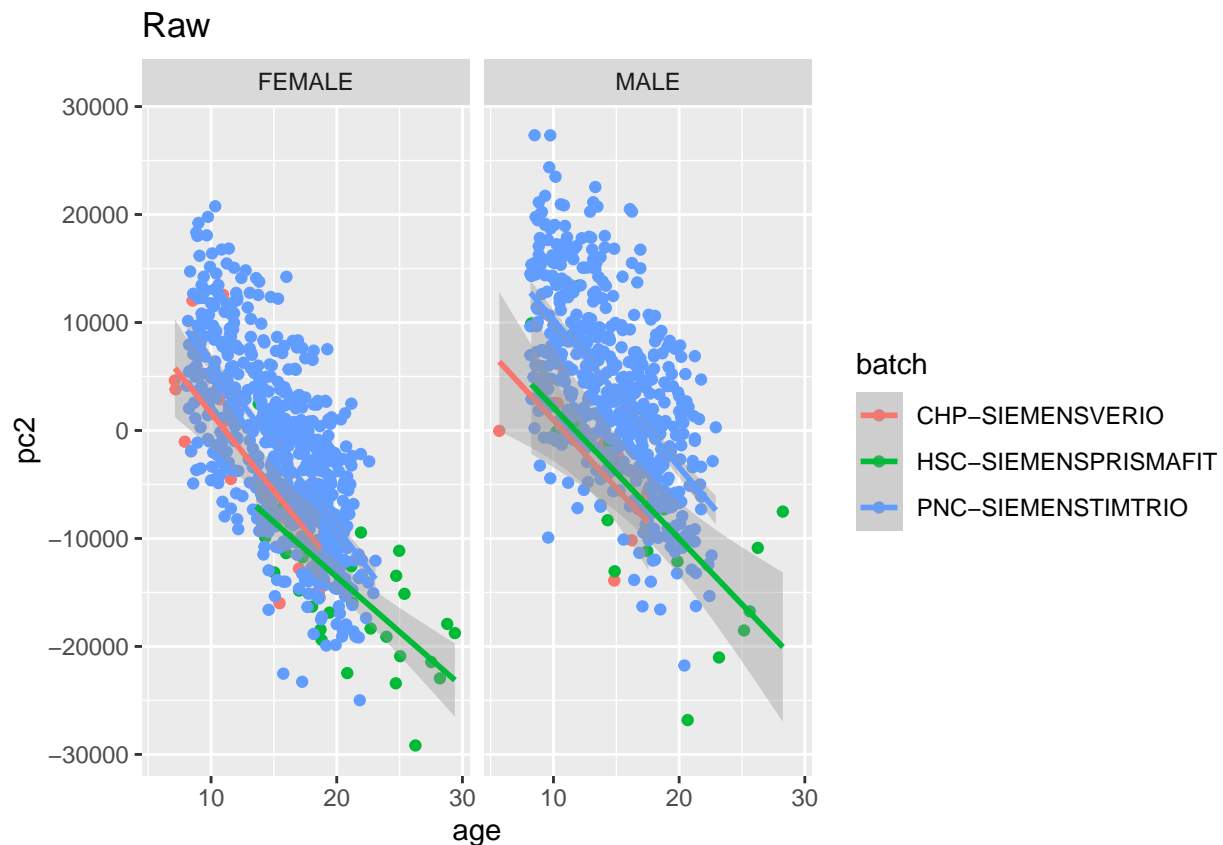
```
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc1) ~ batch + scale(age) + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7952  -0.7223  -0.0701   0.6473   3.2489
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.18250    0.16719  -1.092  0.27522
## batchHSC-SIEMENSPRISMAFIT  0.29970    0.21487   1.395  0.16332
## batchPNC-SIEMENSTIMTRIO   0.19669    0.16888   1.165  0.24437
## scale(age)       0.09245    0.02872   3.219  0.00132 **
```

```
## sexMALE          -0.02898    0.05621   -0.516   0.60625
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9912978)
##
## Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1262.9  on 1274  degrees of freedom
## AIC: 3625.5
##
## Number of Fisher Scoring iterations: 2
```

PC2

```
## `geom_smooth()` using formula 'y ~ x'
```

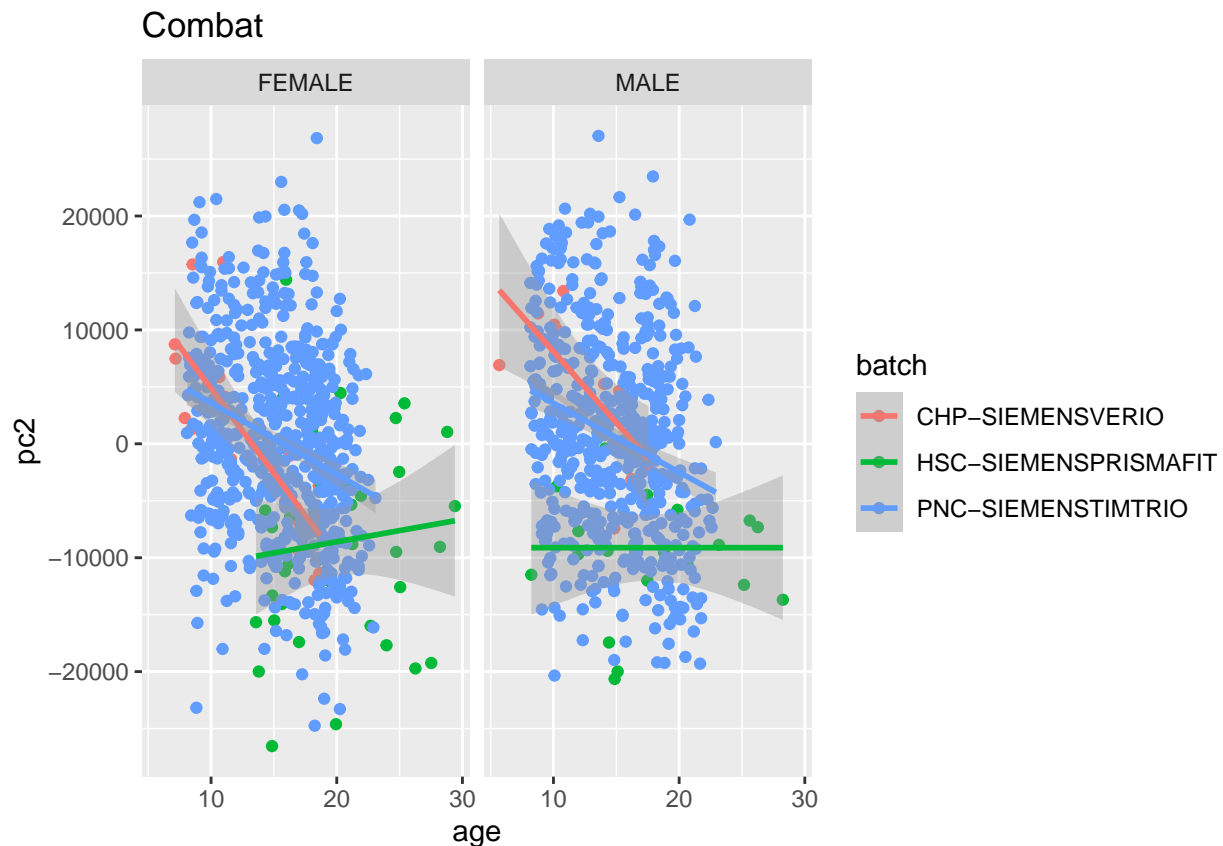


```
## [1] "RAW"

##
## Call:
## glm(formula = scale(pc2) ~ batch + scale(age) + sex, data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.29198  -0.48643  -0.03406   0.48108   2.05144
```

```
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.82502   0.11571  -7.130 1.67e-12 ***
## batchHSC-SIEMENSPRISMAFIT  0.01726   0.14871   0.116   0.908
## batchPNC-SIEMENSTIMTRIO    0.63810   0.11688   5.459 5.73e-08 ***
## scale(age)      -0.61110   0.01988 -30.744 < 2e-16 ***
## sexMALE         0.50690   0.03890  13.030 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.474796)
##
## Null deviance: 1278.00 on 1278 degrees of freedom
## Residual deviance: 604.89 on 1274 degrees of freedom
## AIC: 2683.9
##
## Number of Fisher Scoring iterations: 2

## `geom_smooth()` using formula 'y ~ x'
```



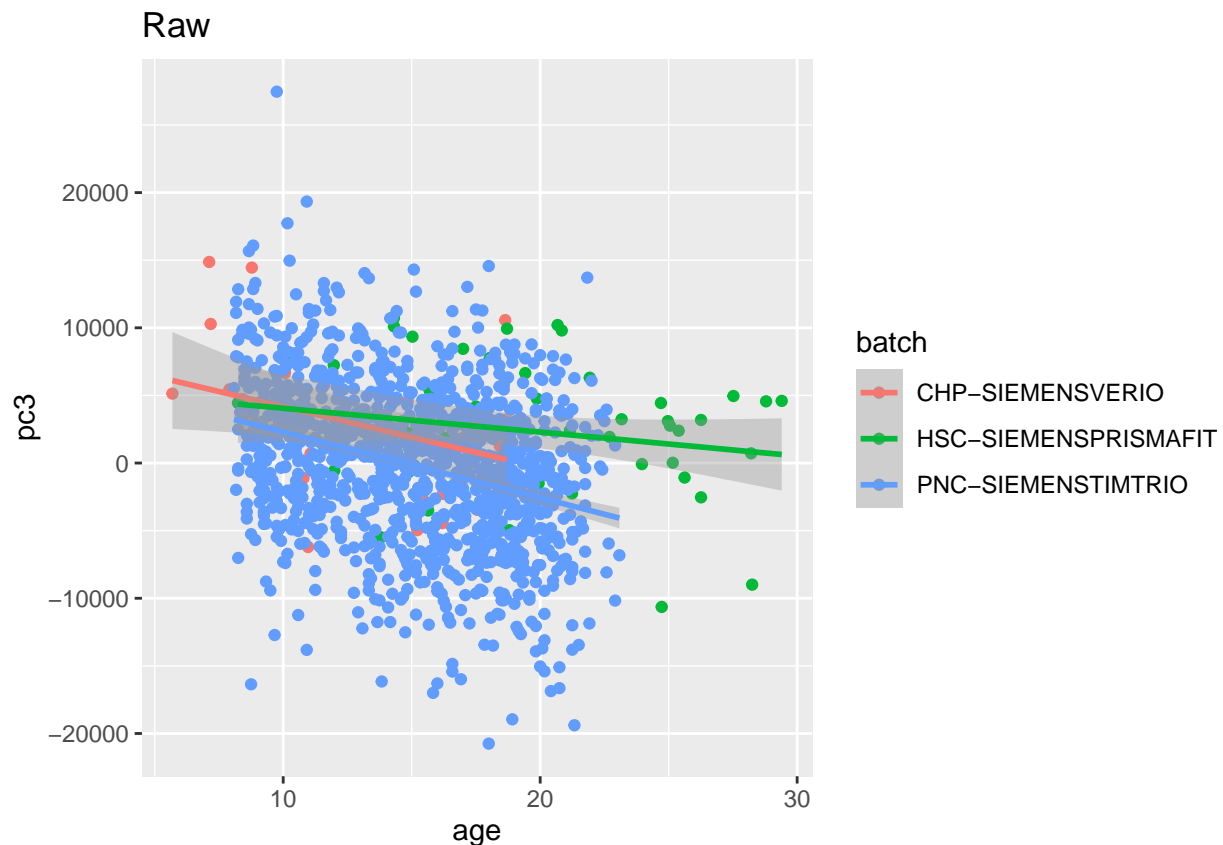
```
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc2) ~ batch + scale(age) + sex, data = dat_combat)
```

```
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.0459  -0.6557  -0.0314   0.6341   3.1826
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.05167    0.15876  -0.325  0.744864
## batchHSC-SIEMENSPRISMAFIT -0.67681    0.20404  -3.317  0.000935 ***
## batchPNC-SIEMENSTIMTRIO   0.08105    0.16037   0.505  0.613371
## scale(age)      -0.25681    0.02727  -9.416 < 2e-16 ***
## sexMALE         0.01582    0.05338   0.296  0.766950
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.8938674)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1138.8  on 1274  degrees of freedom
## AIC: 3493.1
##
## Number of Fisher Scoring iterations: 2
```

PC3

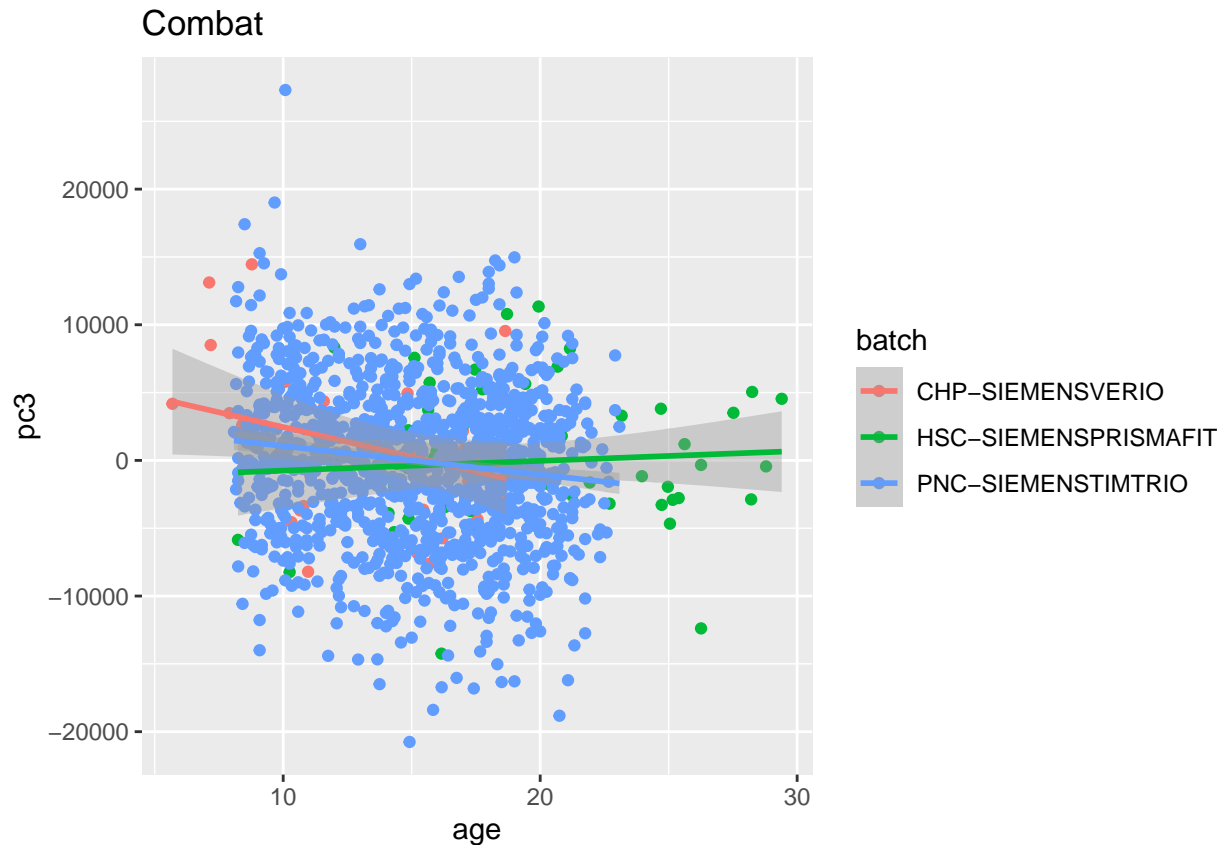
```
## `geom_smooth()` using formula 'y ~ x'
```



```
## [1] "RAW"

##
## Call:
## glm(formula = scale(pc3) ~ batch + scale(age) + sex, data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.2145  -0.6257   0.0497   0.6084   4.3267
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      0.33296    0.15908   2.093  0.0365 *
## batchHSC-SIEMENSPRISMAFIT 0.43020    0.20445   2.104  0.0356 *
## batchPNC-SIEMENSTIMTRIO -0.33204    0.16069  -2.066  0.0390 *
## scale(age)      -0.31203    0.02733 -11.418 <2e-16 ***
## sexMALE         -0.09750    0.05349  -1.823  0.0685 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.8974707)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1143.4  on 1274  degrees of freedom
## AIC: 3498.3
##
## Number of Fisher Scoring iterations: 2

## `geom_smooth()` using formula 'y ~ x'
```

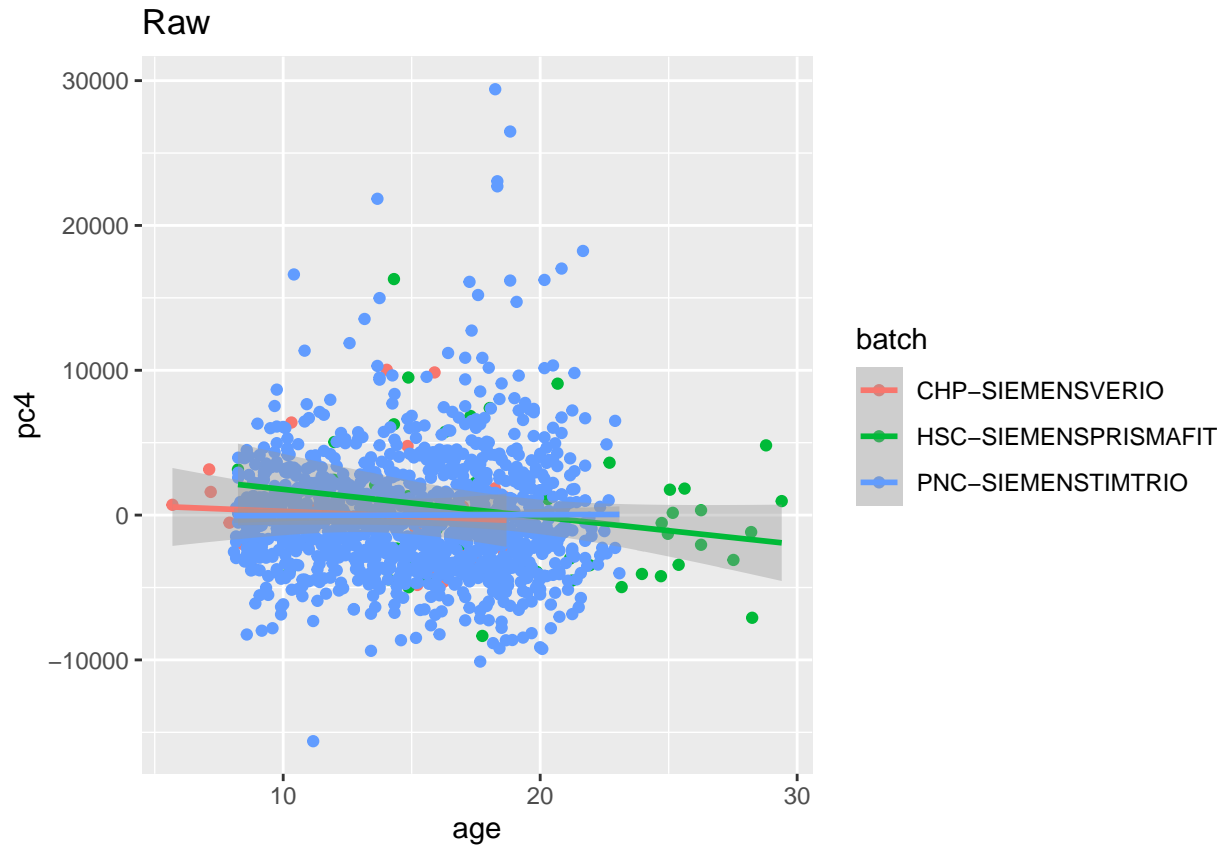


```
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc3) ~ batch + scale(age) + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.5735  -0.6406   0.0321   0.6290   4.5153
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      0.08730    0.16675   0.524   0.601
## batchHSC-SIEMENSPRISMAFIT 0.03676    0.21431   0.172   0.864
## batchPNC-SIEMENSTIMTRIO -0.08939    0.16844  -0.531   0.596
## scale(age)       -0.13220    0.02865  -4.615 4.33e-06 ***
## sexMALE          -0.01337    0.05607  -0.238   0.812
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9861376)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1256.3  on 1274  degrees of freedom
## AIC: 3618.8
##
## Number of Fisher Scoring iterations: 2
```

PC4

```
## `geom_smooth()` using formula 'y ~ x'
```

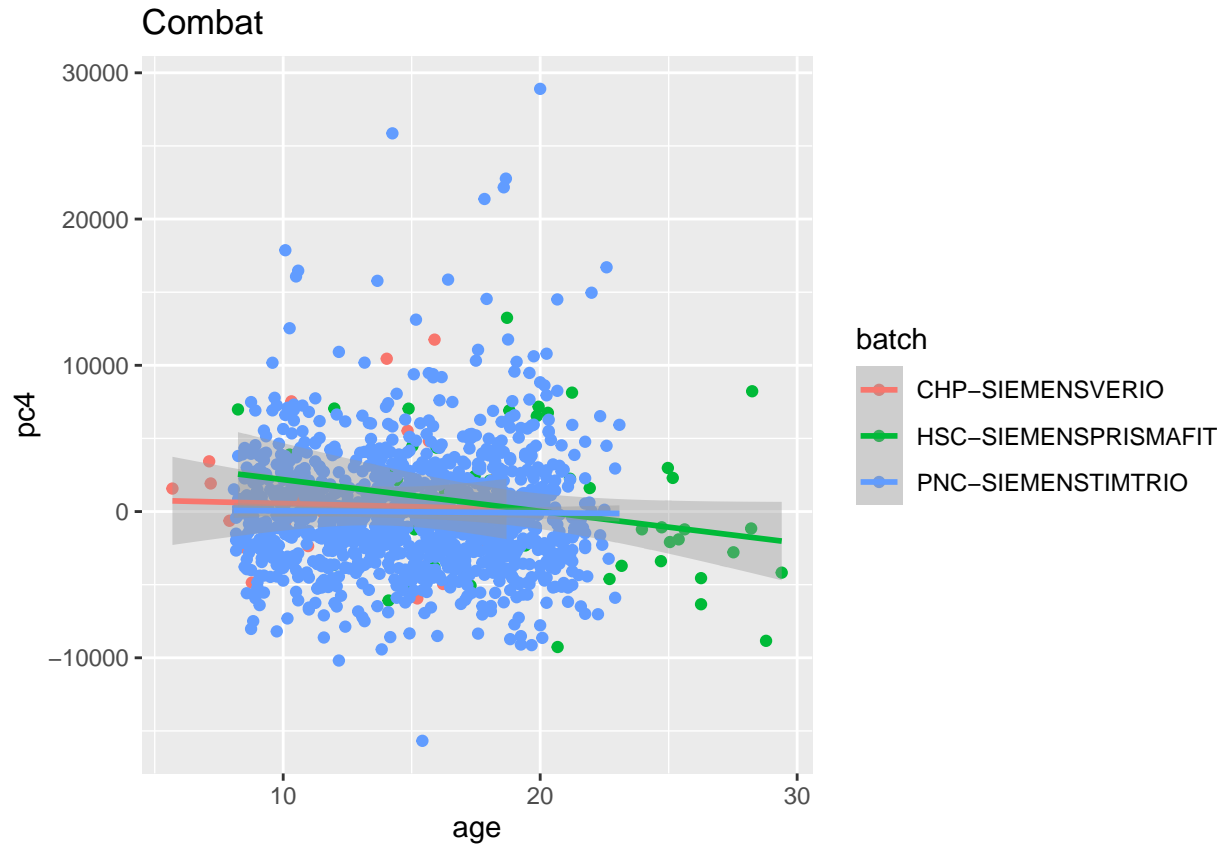


```
## [1] "RAW"
```

```
##
## Call:
## glm(formula = scale(pc4) ~ batch + scale(age) + sex, data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.8037  -0.6140  -0.0957   0.4637   7.0235
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.035512   0.168019  -0.211   0.833
## batchHSC-SIEMENSPRISMAFIT  0.016352   0.215941   0.076   0.940
## batchPNC-SIEMENSTIMTRIO  -0.005198   0.169722  -0.031   0.976
## scale(age)     -0.007761   0.028864  -0.269   0.788
## sexMALE         0.086108   0.056492   1.524   0.128
##
## (Dispersion parameter for gaussian family taken to be 1.001179)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1275.5  on 1274  degrees of freedom
```

```
## AIC: 3638.1
##
## Number of Fisher Scoring iterations: 2

## `geom_smooth()` using formula 'y ~ x'
```



```
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc4) ~ batch + scale(age) + sex, data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.7491  -0.6170  -0.0837   0.4642   7.0588
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.100429   0.168017   0.598   0.550
## batchHSC-SIEMENSPRISMAFIT -0.003739   0.215937  -0.017   0.986
## batchPNC-SIEMENSTIMTRIO  -0.079104   0.169719  -0.466   0.641
## scale(age)      -0.031818   0.028863  -1.102   0.271
## sexMALE         -0.058662   0.056491  -1.038   0.299
##
## (Dispersion parameter for gaussian family taken to be 1.001145)
##
```

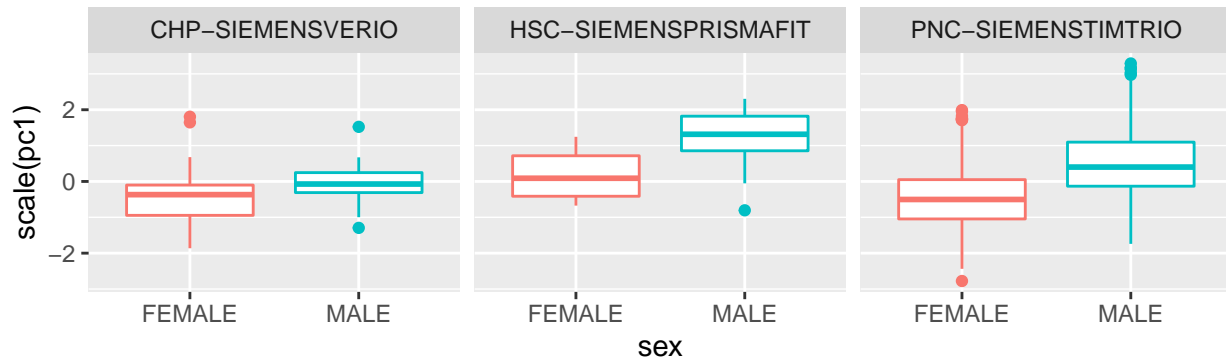


```
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1275.5  on 1274  degrees of freedom
## AIC: 3638.1
##
## Number of Fisher Scoring iterations: 2
```

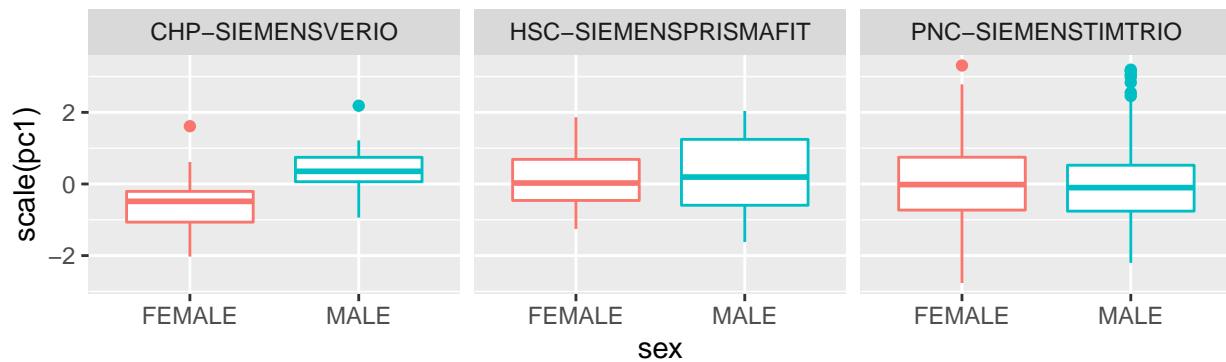
PCs ~ Sex

PC1

Raw



Combat



```
## [1] "RAW"
```

```
##
## Call:
## glm(formula = scale(pc1) ~ batch + sex + scale(age), data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.4608  -0.5675  -0.0262   0.5276   2.8222
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.49052    0.13944  -3.518  0.00045 ***
## batchHSC-SIEMENSPRISMAFIT  0.37309    0.17921   2.082  0.03756 *
## batchPNC-SIEMENSTIMTRIO   0.01549    0.14085   0.110  0.91244
## sexMALE          0.99894    0.04688  21.307 < 2e-16 ***
## scale(age)       0.28323    0.02395  11.824 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.689549)
##
```

```

##      Null deviance: 1278.00  on 1278  degrees of freedom
## Residual deviance:  878.49  on 1274  degrees of freedom
## AIC: 3161.2
##
## Number of Fisher Scoring iterations: 2

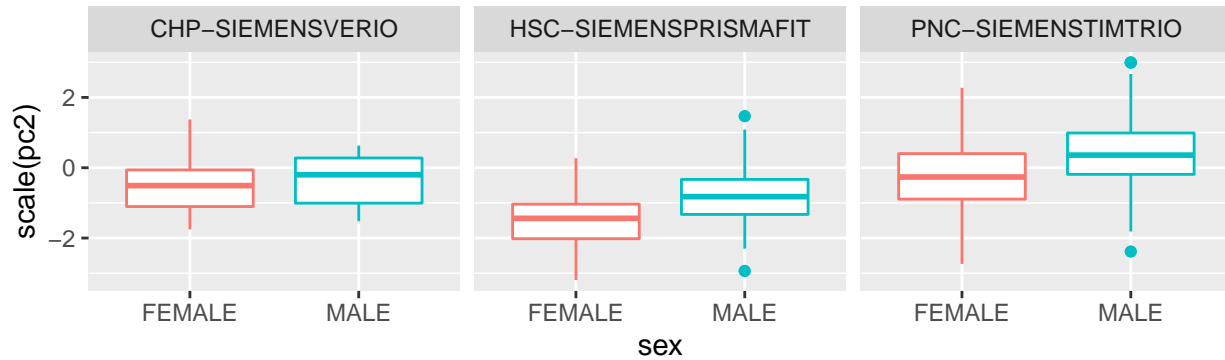
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc1) ~ batch + sex + scale(age), data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7952  -0.7223  -0.0701   0.6473   3.2489
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.18250    0.16719  -1.092  0.27522
## batchHSC-SIEMENSPRISMAFIT  0.29970    0.21487   1.395  0.16332
## batchPNC-SIEMENSTIMTRIO   0.19669    0.16888   1.165  0.24437
## sexMALE        -0.02898    0.05621  -0.516  0.60625
## scale(age)       0.09245    0.02872   3.219  0.00132 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9912978)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1262.9  on 1274  degrees of freedom
## AIC: 3625.5
##
## Number of Fisher Scoring iterations: 2

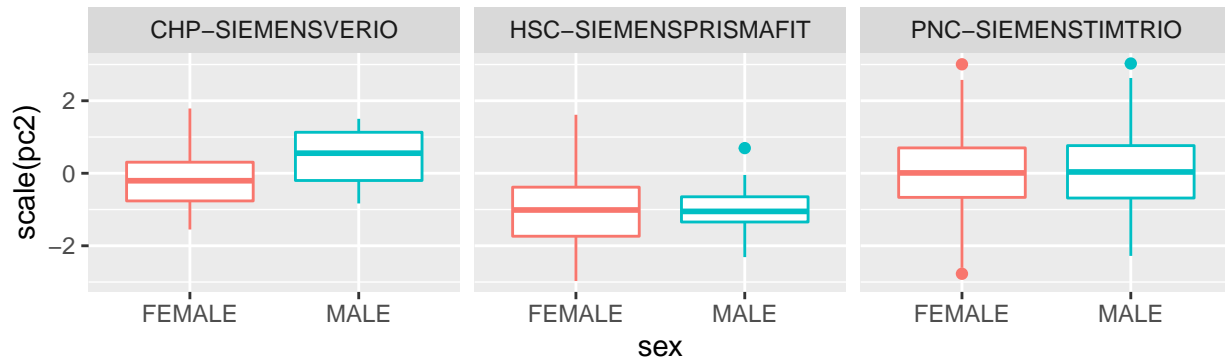
```

PC2

Raw



Combat



```
## [1] "RAW"
```

```
##
## Call:
## glm(formula = scale(pc2) ~ batch + sex + scale(age), data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.29198  -0.48643  -0.03406   0.48108   2.05144
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -0.82502    0.11571  -7.130 1.67e-12 ***
## batchHSC-SIEMENSPRISMAFIT  0.01726    0.14871   0.116  0.908
## batchPNC-SIEMENSTIMTRIO   0.63810    0.11688   5.459 5.73e-08 ***
## sexMALE          0.50690    0.03890  13.030 < 2e-16 ***
## scale(age)       -0.61110    0.01988 -30.744 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.474796)
##
##      Null deviance: 1278.00  on 1278  degrees of freedom
## Residual deviance:  604.89  on 1274  degrees of freedom
```

```

## AIC: 2683.9
##
## Number of Fisher Scoring iterations: 2

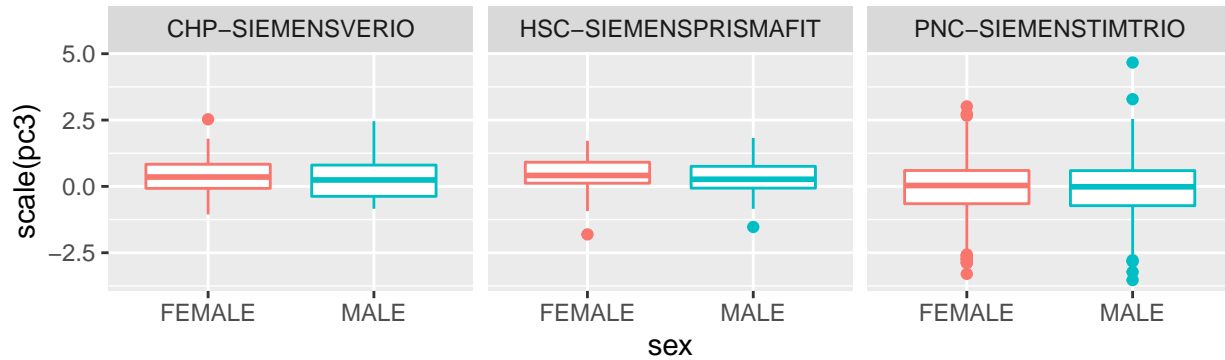
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc2) ~ batch + sex + scale(age), data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.0459  -0.6557  -0.0314   0.6341   3.1826
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   -0.05167    0.15876  -0.325  0.744864
## batchHSC-SIEMENSPRISMAFIT -0.67681    0.20404  -3.317  0.000935 ***
## batchPNC-SIEMENSTIMTRIO   0.08105    0.16037   0.505  0.613371
## sexMALE         0.01582    0.05338   0.296  0.766950
## scale(age)     -0.25681    0.02727  -9.416 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.8938674)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1138.8  on 1274  degrees of freedom
## AIC: 3493.1
##
## Number of Fisher Scoring iterations: 2

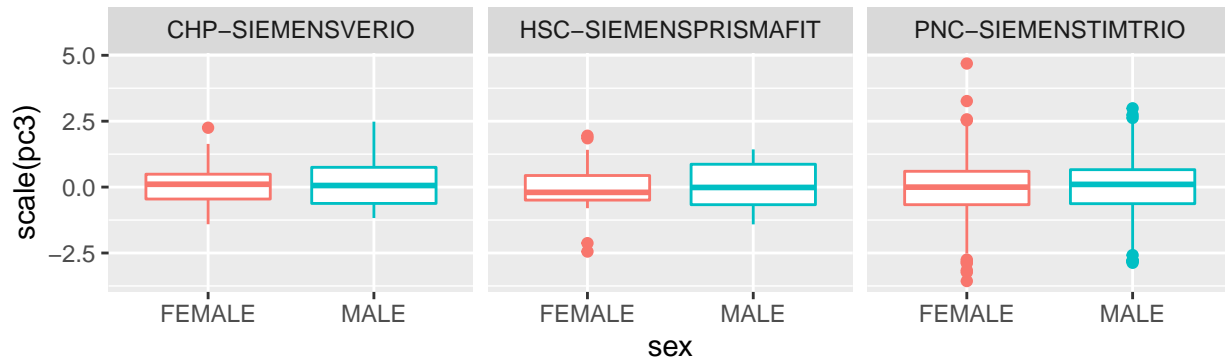
```

PC3

Raw



Combat



```
## [1] "RAW"
```

```
##
## Call:
## glm(formula = scale(pc3) ~ batch + sex + scale(age), data = dat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.2145  -0.6257   0.0497   0.6084   4.3267
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.33296    0.15908   2.093  0.0365 *
## batchHSC-SIEMENSPRISMAFIT 0.43020    0.20445   2.104  0.0356 *
## batchPNC-SIEMENSTIMTRIO -0.33204    0.16069  -2.066  0.0390 *
## sexMALE        -0.09750    0.05349  -1.823  0.0685 .
## scale(age)     -0.31203    0.02733 -11.418 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.8974707)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1143.4  on 1274  degrees of freedom
```

```

## AIC: 3498.3
##
## Number of Fisher Scoring iterations: 2

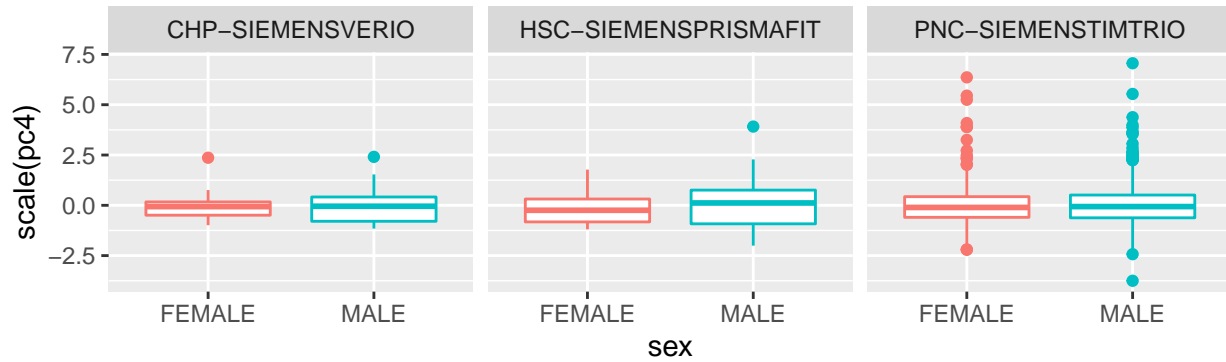
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc3) ~ batch + sex + scale(age), data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.5735  -0.6406   0.0321   0.6290   4.5153
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      0.08730    0.16675   0.524   0.601
## batchHSC-SIEMENSPRISMAFIT  0.03676    0.21431   0.172   0.864
## batchPNC-SIEMENSTIMTRIO -0.08939    0.16844  -0.531   0.596
## sexMALE           -0.01337    0.05607  -0.238   0.812
## scale(age)        -0.13220    0.02865  -4.615 4.33e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9861376)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1256.3  on 1274  degrees of freedom
## AIC: 3618.8
##
## Number of Fisher Scoring iterations: 2

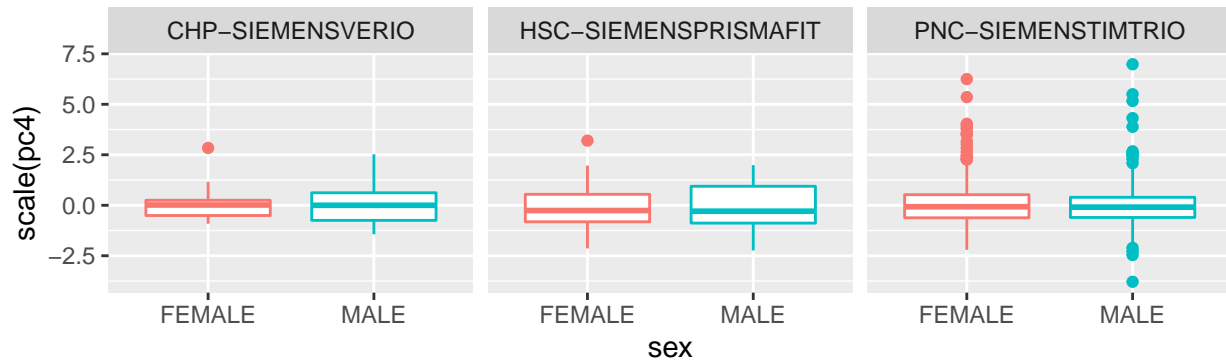
```

PC4

Raw



Combat



```
## [1] "RAW"
```

```
##
```

```
## Call:
```

```
## glm(formula = scale(pc4) ~ batch + sex + scale(age), data = dat)
```

```
##
```

```
## Deviance Residuals:
```

```
##      Min       1Q   Median       3Q      Max
```

```
## -3.8037  -0.6140  -0.0957   0.4637   7.0235
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error t value Pr(>|t|)
```

```
## (Intercept)      -0.035512   0.168019  -0.211   0.833
```

```
## batchHSC-SIEMENSPRISMAFIT  0.016352   0.215941   0.076   0.940
```

```
## batchPNC-SIEMENSTIMTRIO -0.005198   0.169722  -0.031   0.976
```

```
## sexMALE           0.086108   0.056492   1.524   0.128
```

```
## scale(age)        -0.007761   0.028864  -0.269   0.788
```

```
##
```

```
## (Dispersion parameter for gaussian family taken to be 1.001179)
```

```
##
```

```
##      Null deviance: 1278.0  on 1278  degrees of freedom
```

```
## Residual deviance: 1275.5  on 1274  degrees of freedom
```

```
## AIC: 3638.1
```

```
##
```

```
## Number of Fisher Scoring iterations: 2
```



```
## [1] "COMBAT"

##
## Call:
## glm(formula = scale(pc4) ~ batch + sex + scale(age), data = dat_combat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.7491  -0.6170  -0.0837   0.4642   7.0588
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      0.100429   0.168017   0.598   0.550
## batchHSC-SIEMENSPRISMAFIT -0.003739   0.215937  -0.017   0.986
## batchPNC-SIEMENSTIMTRIO  -0.079104   0.169719  -0.466   0.641
## sexMALE           -0.058662   0.056491  -1.038   0.299
## scale(age)        -0.031818   0.028863  -1.102   0.271
##
## (Dispersion parameter for gaussian family taken to be 1.001145)
##
##      Null deviance: 1278.0  on 1278  degrees of freedom
## Residual deviance: 1275.5  on 1274  degrees of freedom
## AIC: 3638.1
##
## Number of Fisher Scoring iterations: 2
```

““