

EDA: MS ComBat GAM vs Linear

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Objective

To compare effects of applying ComBat-GAM vs ComBat-Linear in the MS data.

IMPORTANT

MS patients from the *HSC* site that were scanned in a *SIEMENSTIMTRIO* scanner were excluded from the current harmonization.

[1] "In MS:"

| | CHP | HSC |
|------------------|-----|-----|
| SIEMENSPRISMAFIT | 0 | 28 |
| SIEMENSTIMTRIO | 0 | 18 |
| SIEMENSVERIO | 21 | 0 |

IMPORTANT (cont.)

The corresponding group in the Healthy Control data was similarly excluded when modeling site effects with ComBat-GAM.

```
[1] "In HC:"
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| | CHP | HSC | PNC |
|------------------|-----|-----|------|
| SIEMENSPRISMAFIT | 0 | 58 | 0 |
| SIEMENSTIMTRIO | 0 | 7 | 1185 |
| SIEMENSVERIO | 36 | 0 | 0 |

```
|
```

Datasets: MS

MS (dimensions):

[1] 67 161

MS (count by site)

| site | n |
|----------------------|----|
| CHP | 21 |
| HSC-SIEMENSPRISMAFIT | 28 |
| HSC-SIEMENSTIMTRIO | 18 |

MS (count by sex)

| sex | site | n |
|--------|----------------------|----|
| FEMALE | CHP | 18 |
| FEMALE | HSC-SIEMENSPRISMAFIT | 17 |
| FEMALE | HSC-SIEMENSTIMTRIO | 15 |
| MALE | CHP | 3 |
| MALE | HSC-SIEMENSPRISMAFIT | 11 |
| MALE | HSC-SIEMENSTIMTRIO | 3 |

Datasets: HC

Dimensions HC (no PNC):

[1] 101 161

Count per site:

| site | n |
|----------------------|----|
| CHP | 36 |
| HSC-SIEMENSPRISMAFIT | 58 |
| HSC-SIEMENSTIMTRIO | 7 |

Count by sex and site:

| site | sex | n |
|----------------------|--------|----|
| CHP | FEMALE | 24 |
| CHP | MALE | 12 |
| HSC-SIEMENSPRISMAFIT | FEMALE | 37 |
| HSC-SIEMENSPRISMAFIT | MALE | 21 |
| HSC-SIEMENSTIMTRIO | FEMALE | 5 |
| HSC-SIEMENSTIMTRIO | MALE | 2 |

Datasets: HC + MS

Dimensions (no PNC):

[1] 168 161

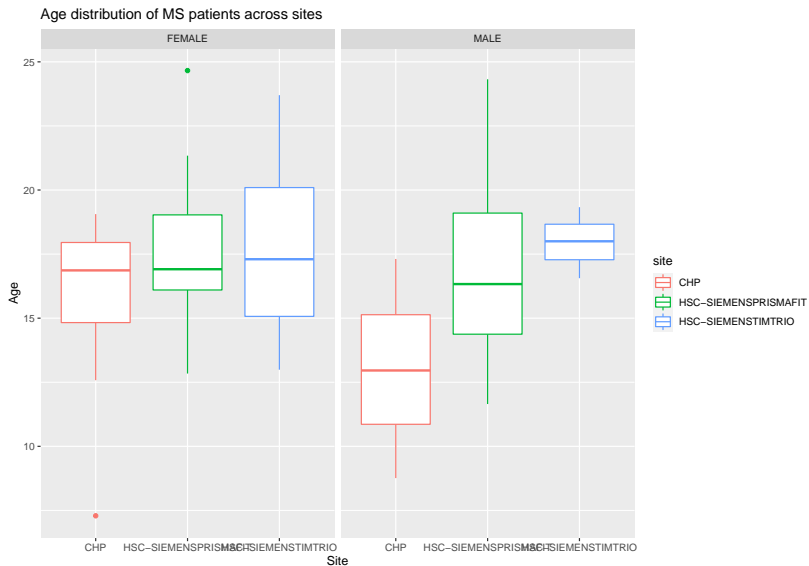
Count per site:

| site | n |
|----------------------|----|
| CHP | 57 |
| HSC-SIEMENSPRISMAFIT | 86 |
| HSC-SIEMENSTIMTRIO | 25 |

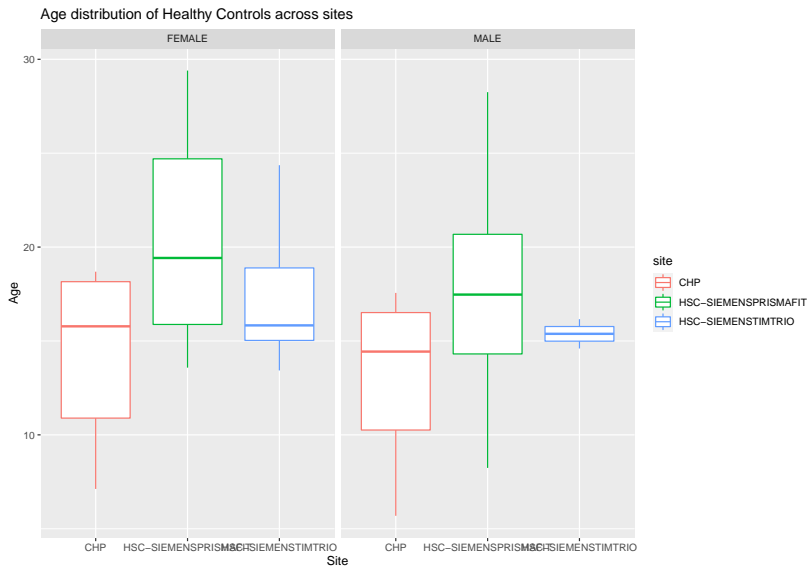
Count by sex and site:

| site | sex | n |
|----------------------|--------|----|
| CHP | FEMALE | 42 |
| CHP | MALE | 15 |
| HSC-SIEMENSPRISMAFIT | FEMALE | 54 |
| HSC-SIEMENSPRISMAFIT | MALE | 32 |
| HSC-SIEMENSTIMTRIO | FEMALE | 20 |
| HSC-SIEMENSTIMTRIO | MALE | 5 |

Age across sites [MS]



Age across sites [HC]



Harmonization Approach

Adjusted data are shown for the following approach:

- ▶ Join HC and MS data into one dataset
- ▶ Split this dataset into males and females
- ▶ Run ComBat (GAM and Linear) on in parallel on males and females.

Models used in harmonization

- ▶ GAM: $s(\text{age}) + \text{MS} + \text{MS} \times \text{age}$
- ▶ Linear: $\text{age} + \text{age}^2 + \text{MS} + \text{MS} \times \text{age} + \text{MS} \times \text{age}^2$

Site effects: MS + HC [GAM]

Number of ROIs showing site effects:

Full covariate model:

[1] “~ ICV + age + age2 + sex + sexage + sexage2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Ignore sex: [1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Site effects by sex: MS + HC [GAM]

Females:

[1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Males:

[1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Site effects: MS + HC [Linear]

Full covariate model:

[1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Sex not considered: [1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Site effects by sex: MS + HC [Linear]

Females:

[1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Males:

[1] “~ ICV + age + age2 + MS + MSage + MSage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Site effects: MS [GAM]

Full model:

[1] “~ ICV + age + age2 + sex + sexage + sexage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Ignore sex:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Site effects: MS [Linear]

Number of ROIs showing site effects:

Full model: [1] “~ ICV + age + age2 + sex + sexage + sexage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 2 |

Sex not considered: [1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 2 |

Site effects by sex: MS [GAM]

Females:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Males:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 8 |

Site effects by sex: MS [Linear]

Females:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Males:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 8 |

Site effects: HC [GAM]

Full model:

[1] “~ ICV + age + age2 + sex + sexage + sexage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Sex not considered:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Site effects by sex: HC [GAM]

Females:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 2 |

Males:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Site effects: HC [Linear]

Number of ROIs showing site effects:

Full model: [1] “~ ICV + age + age2 + sex + sexage + sexage2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Sex not considered: [1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 0 |

Site effects by sex: HC [Linear]

Females:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Males:

[1] “~ ICV + age + age2”

| FDR | Bonferroni | Uncorrected P |
|-----|------------|------------------|
| 0 | 0 | 1 |

Harmonization Conclusions

Harmonization is effective.

Signal check

Full model = $\text{ROI} \sim \text{ICV} + \text{age} + \text{age}^2 + \text{sex} + \text{sexage} + \text{sexage}^2 + \text{MS} + \text{MSage} + \text{MSage}^2$

Raw data

► MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 10 |

► age:MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 10 |

► age2:MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 12 |

Harmonized data (GAM)

► MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 10 |

► age:MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 10 |

► age2:MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 12 |

Raw data (w/ site)

► MS:

| FDR | Bonferroni | Uncorrected P_s |
|-----|------------|----------------------|
| 0 | 0 | 9 |

► age:sexMALE:

| FDR | Bonferroni | Uncorrected P_s |
|-----|------------|----------------------|
| 0 | 0 | 18 |

► age2:sexMALE:

| FDR | Bonferroni | Uncorrected P_s |
|-----|------------|----------------------|
| 0 | 0 | 4 |

Harmonized data (w/ site)

► MS:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 9 |

► age:sexMALE:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 18 |

► age2:sexMALE:

| FDR | Bonferroni | Uncorrected P _s |
|-----|------------|-------------------------------|
| 0 | 0 | 4 |

Signal check conclusion

Harmonization does not change the number of significant p-values when testing the full model. Success!