

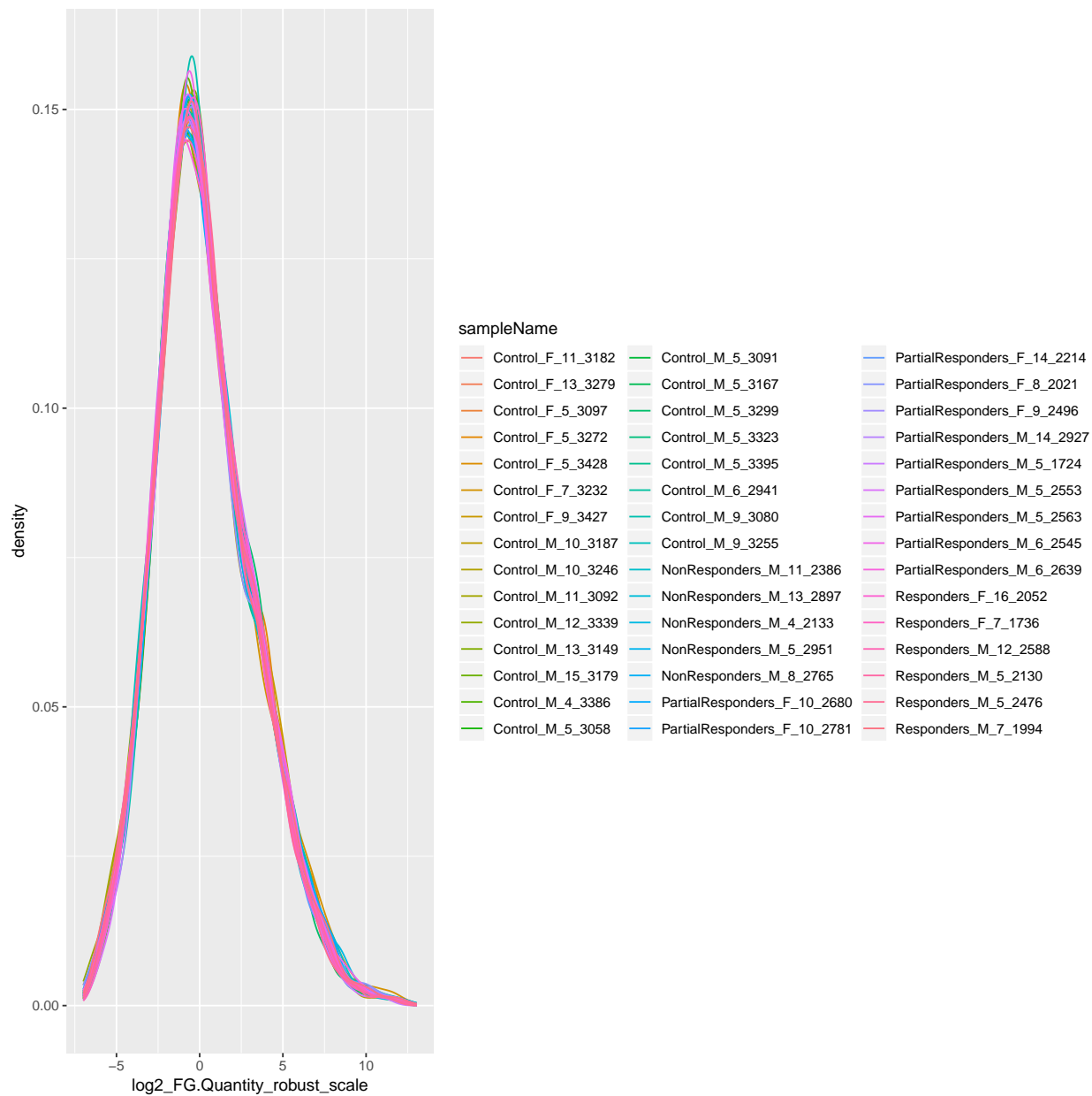
DIA data analysis with MSqROB

Functional Genomics Center Zurich

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```
##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.
## 0.000e+00 0.000e+00 1.894e-05 2.337e-02 1.279e-02 2.848e-01
## [1] 0
```

normalize data —



```
## [1] "PartialResponders" "PartialResponders" "Control"
## [4] "Control"           "Responders"          "Control"
```

Fixed effects

Do you want to save the model?

Set the significance threshold (default: 5% FDR)

```
## [1] "codingPartialResponders - codingResponders"
```

