

# Figure 1

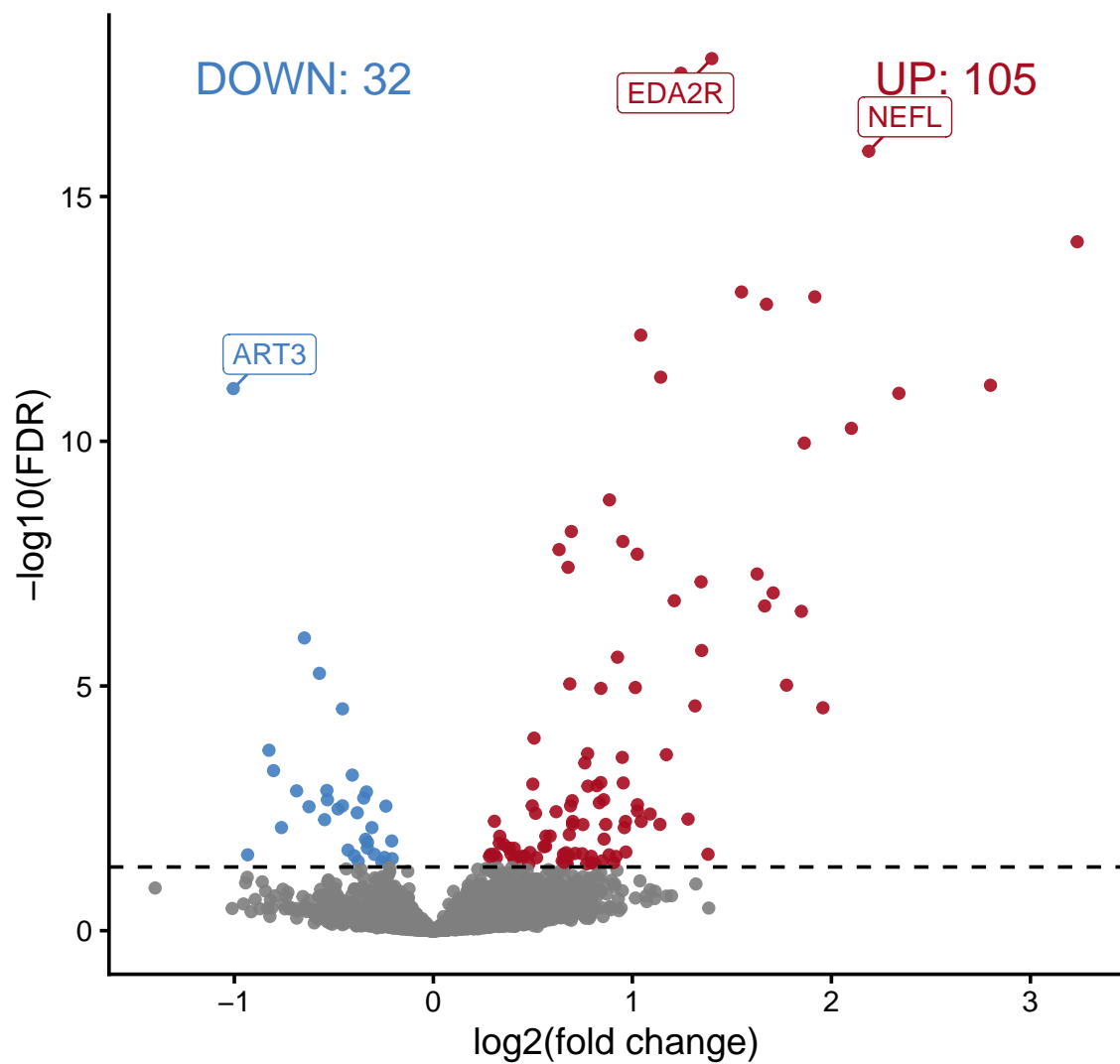
Ximing Ran

2025-10-14

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Fig 1A: Volcano plot of differential expression proteins  
ALS versus Control (Miami cohort)



**Fig 1B: Up enrichment analysis**

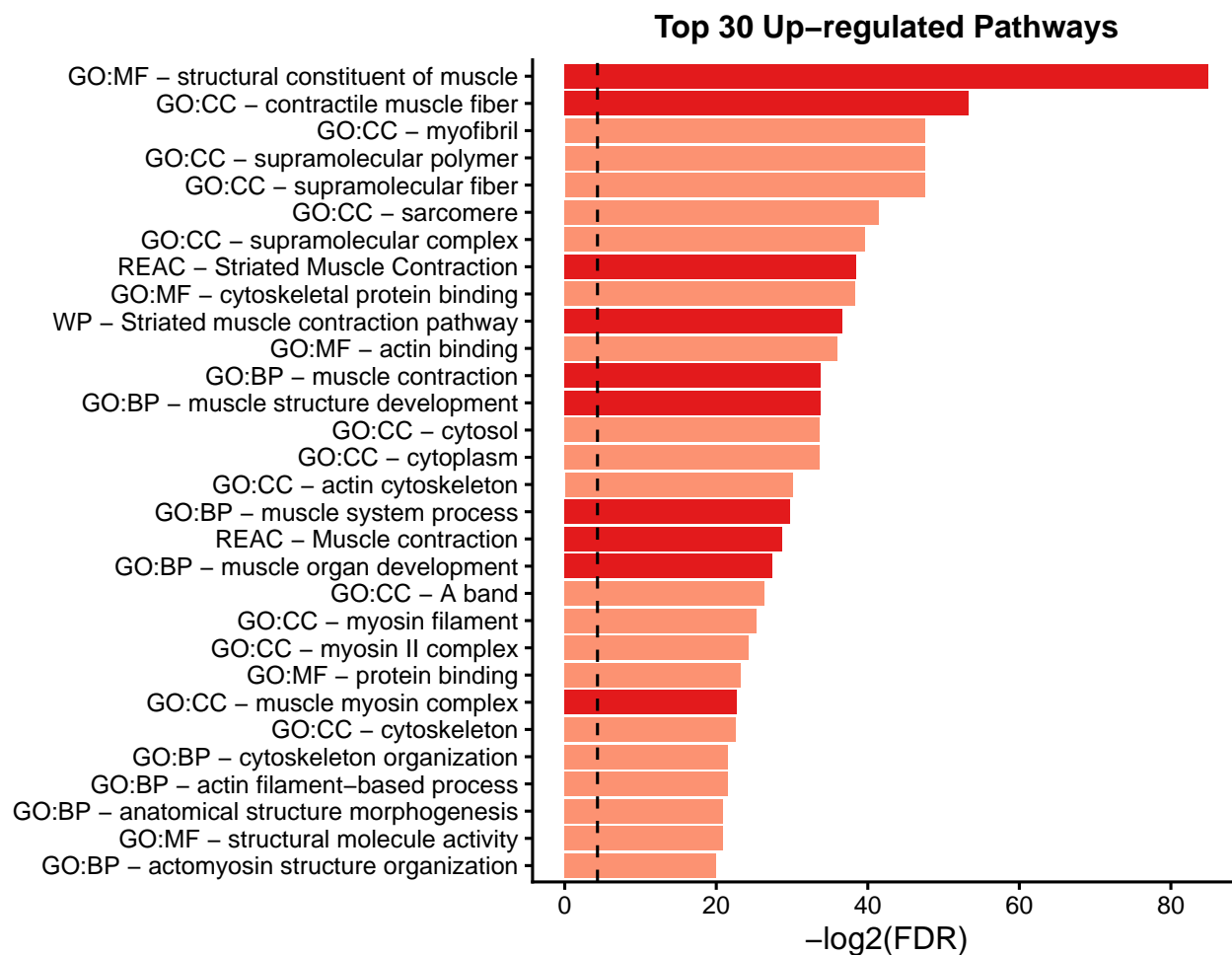


Fig 1C: Down enrichment analysis

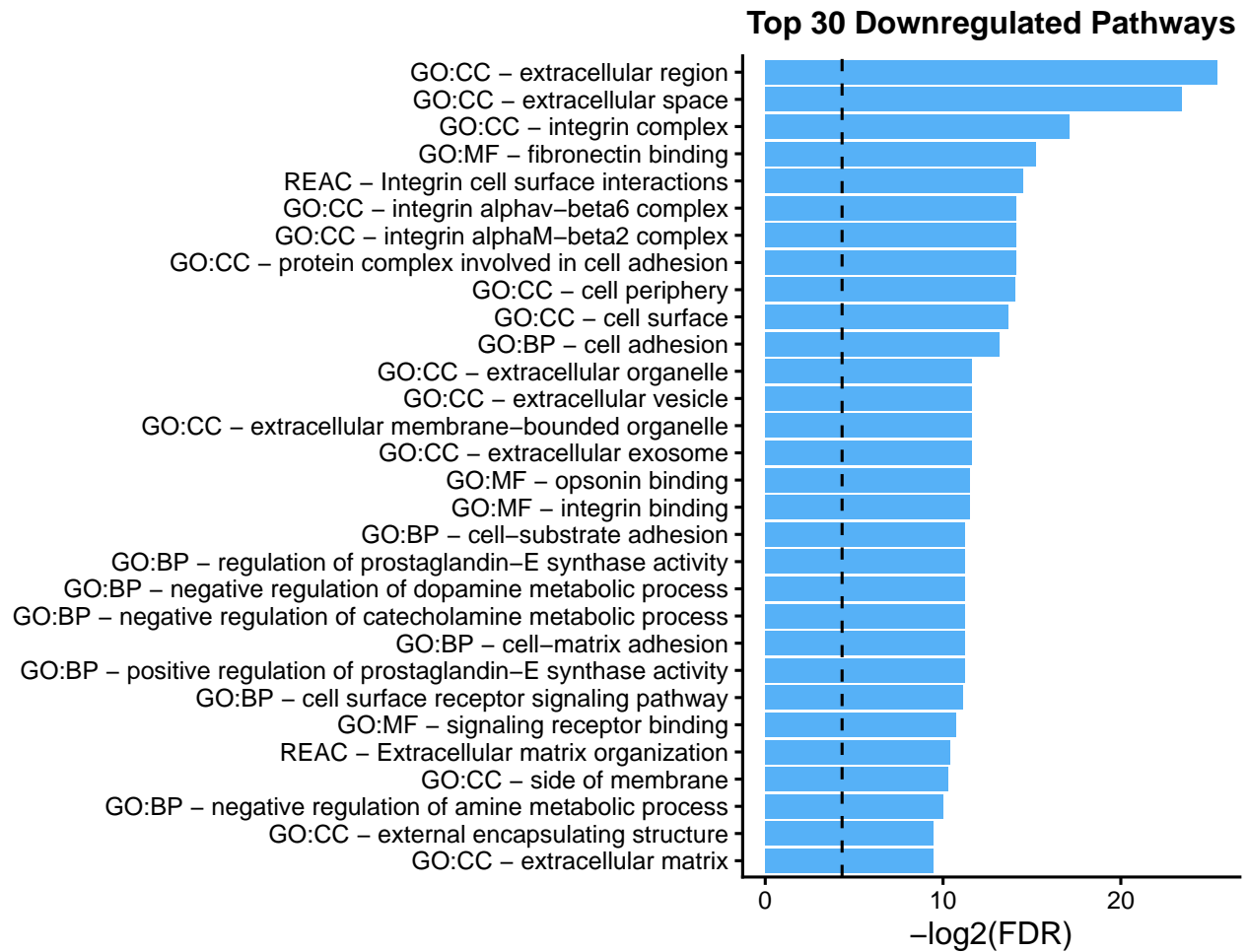


Fig 1D: PPI network of DE proteins

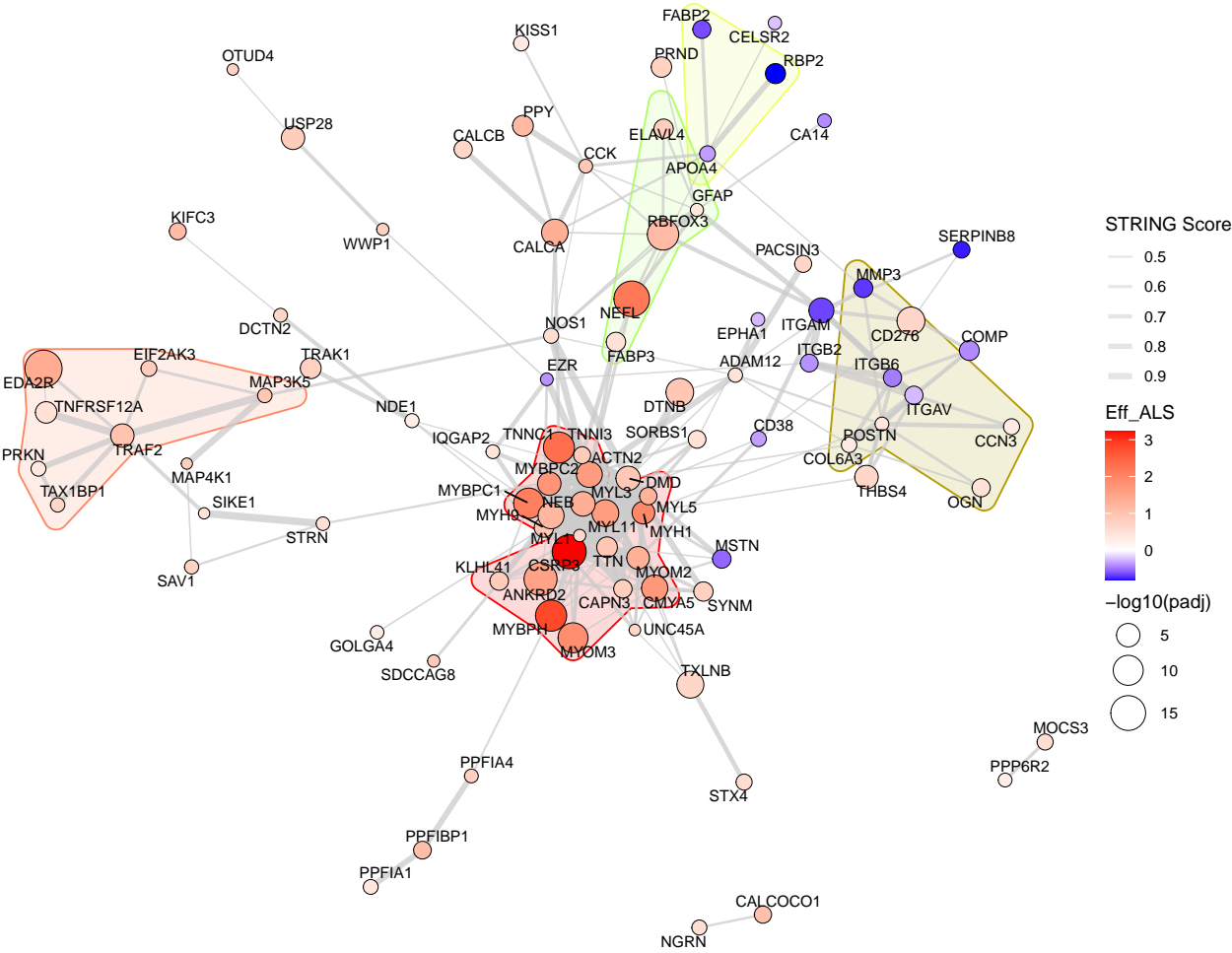
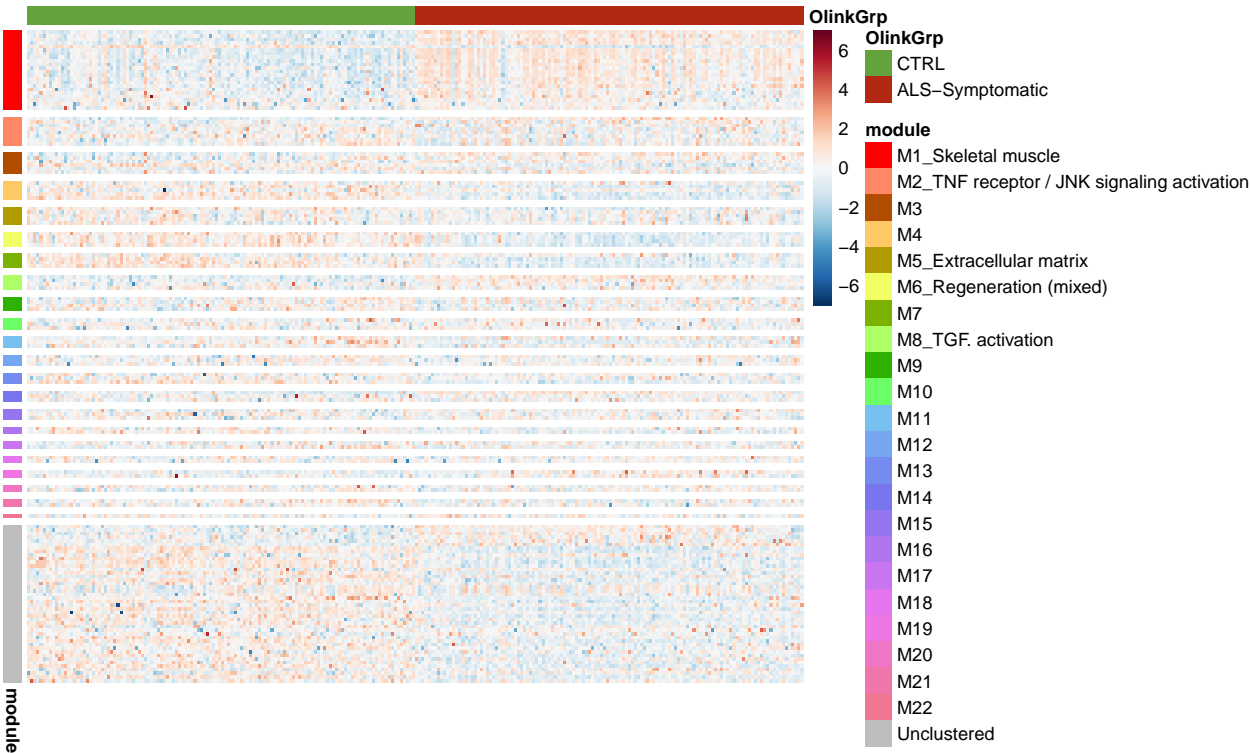
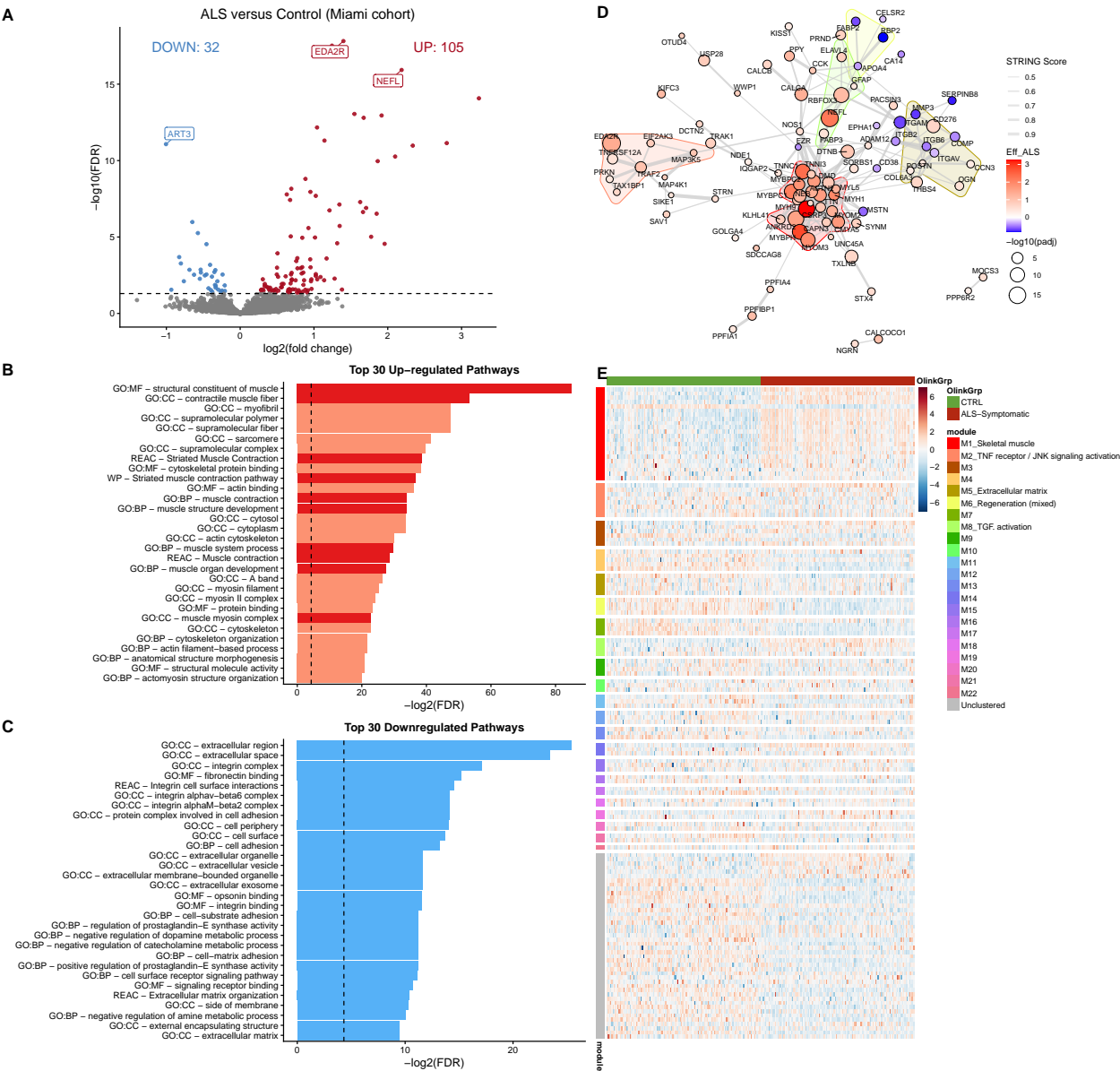


Fig 1E: Heatmap of selected proteins



# Combine all plots into one figure



## Session Info

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## R version 4.4.0 (2024-04-24)
## Platform: aarch64-apple-darwin20
## Running under: macOS 26.0.1
##
## Matrix products: default
## BLAS:   /Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/lib/libRblas.0.dylib
## LAPACK: /Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/lib/libRlapack.dylib; LAPACK v
##
## locale:
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##
## time zone: America/New_York
## tzcode source: internal
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## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods    base
##
## other attached packages:
## [1] here_1.0.1      ggplotify_0.1.2  RColorBrewer_1.1-3  pheatmap_1.0.12
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## [9] tidygraph_1.3.1 ggraph_2.2.1     gprofiler2_0.2.3    ggrepel_0.9.6
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## [4] estimability_1.5.1 farver_2.1.2        nloptr_2.2.1
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