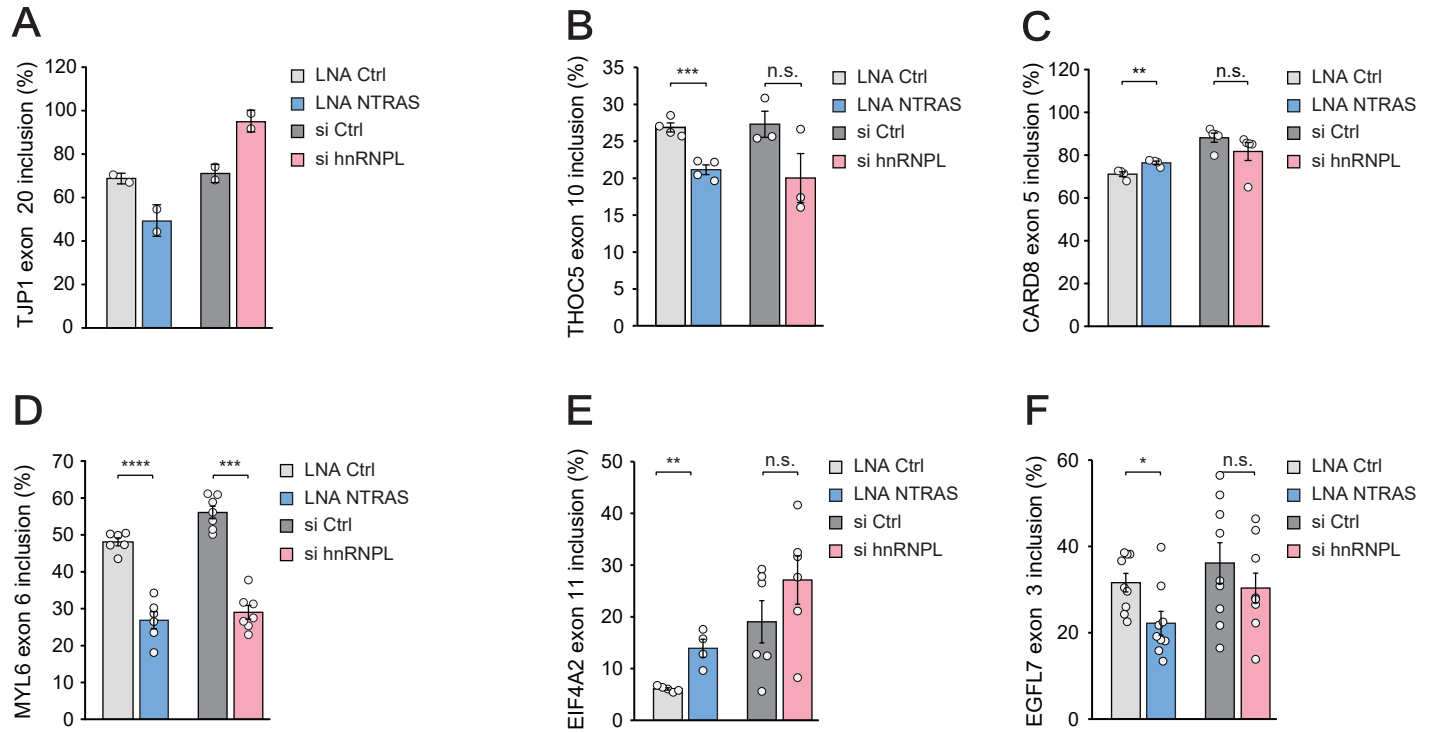


# Appendix

- Appendix Figure S1

# Appendix Figure S1



## Appendix Figure S1 | NTRAS and hnRNPL co-regulated alternative splicing (AS) events.

(A) RNA deep sequencing-based analysis of TJP1 exon 20 inclusion upon silencing of NTRAS or hnRNPL in HUVECs (n = 2 independent biological replicates). Data refers to Fig 2H (B) RT-PCR-based analysis of THOC5 exon 10 (n = 3-4 independent biological replicates), (C) CARD8 exon 5 (n = 5-6 independent biological replicates), (D) MYL6 exon 6 (n = 6-7 independent biological replicates), (E) EIF4A2 exon 11 (n = 4-6 independent biological replicates), and (F) EGFL7 exon 3 (n = 9 independent biological replicates) inclusion in NTRAS- or hnRNPL-silenced HUVECs.

Data information: In (A-E) data are represented as mean  $\pm$  S.E.M. n.s.: non-significant, \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001, \*\*\*\*p < 0.0001.