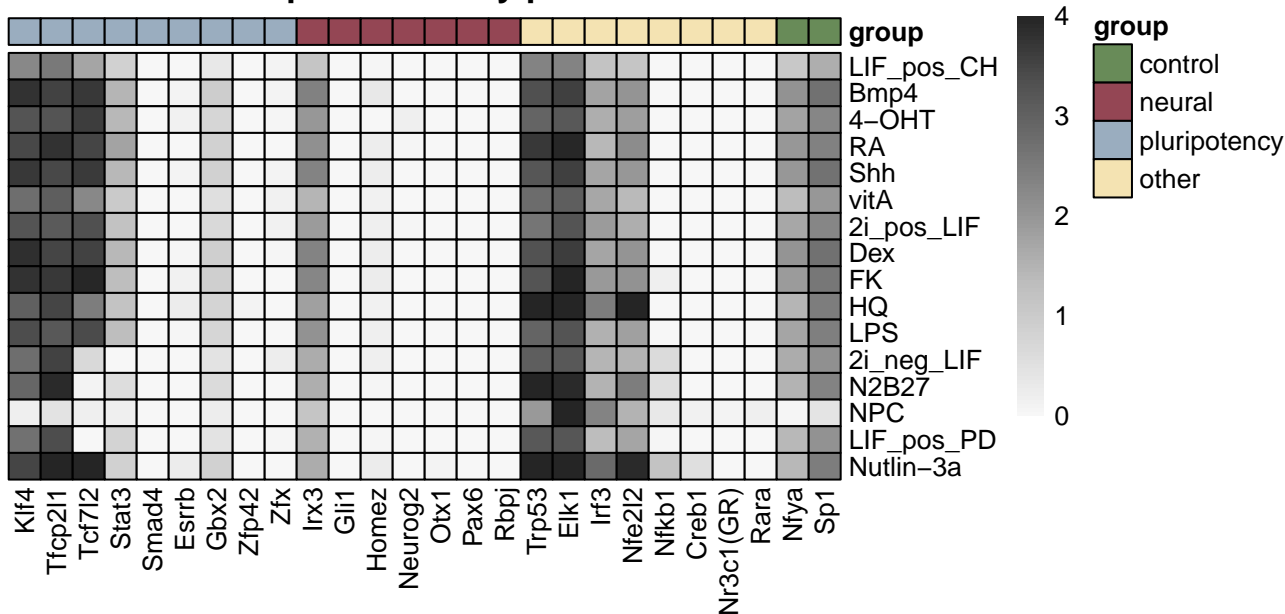
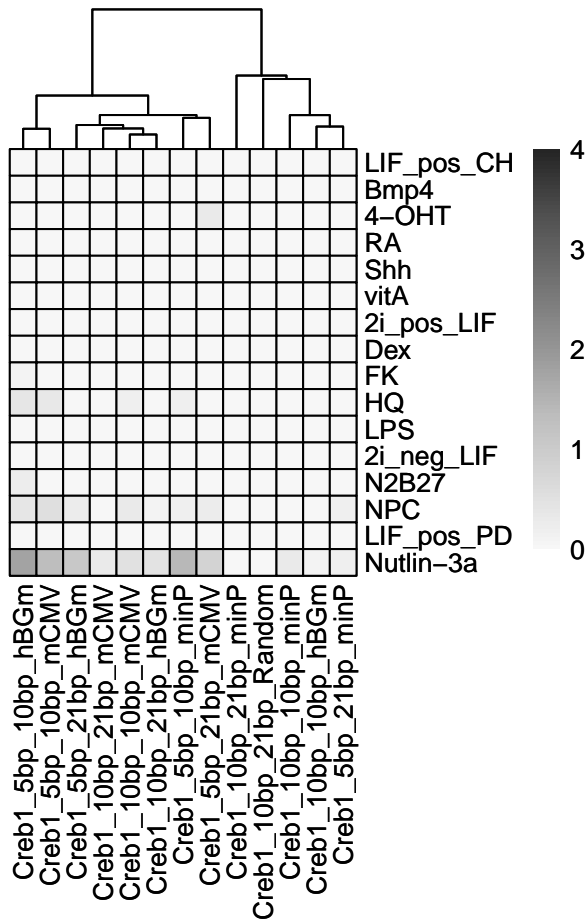


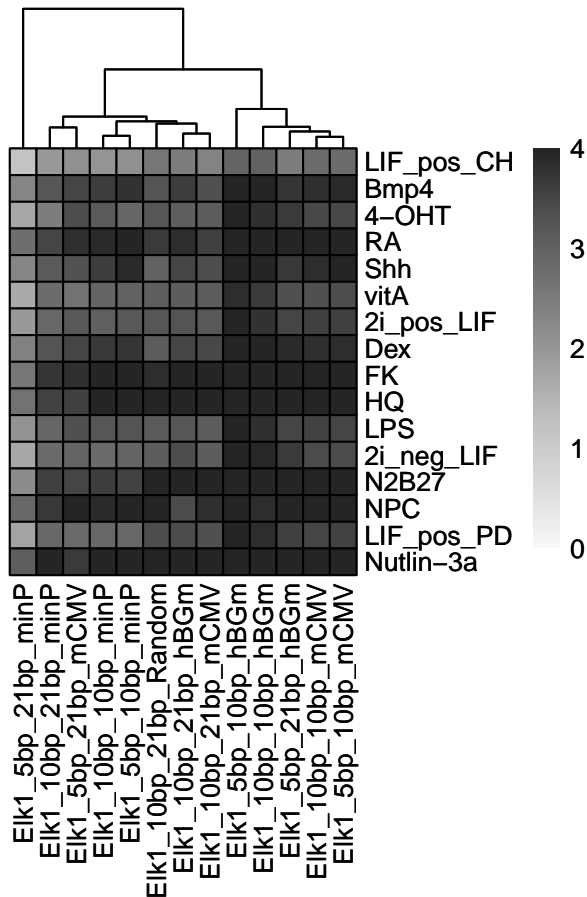
Mean TF reporter activity per condition

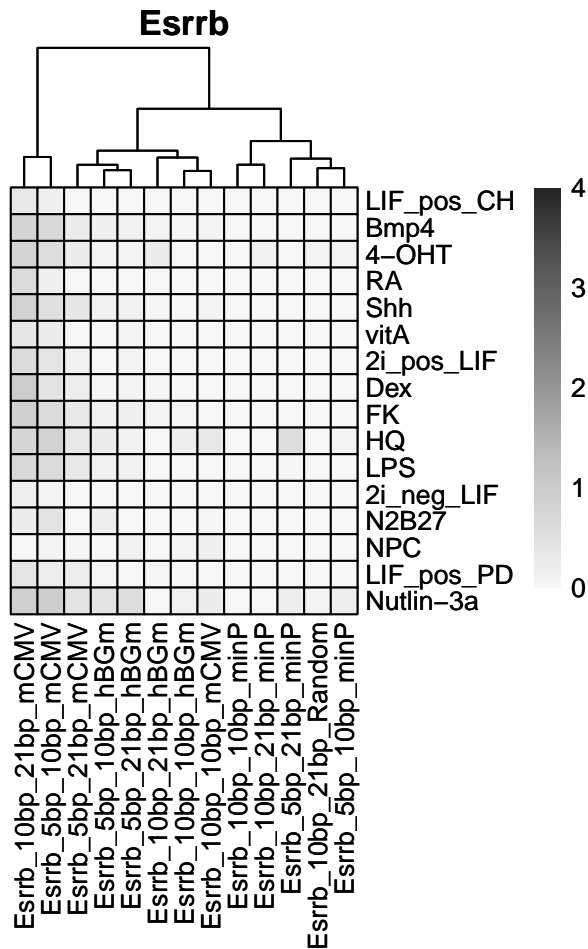


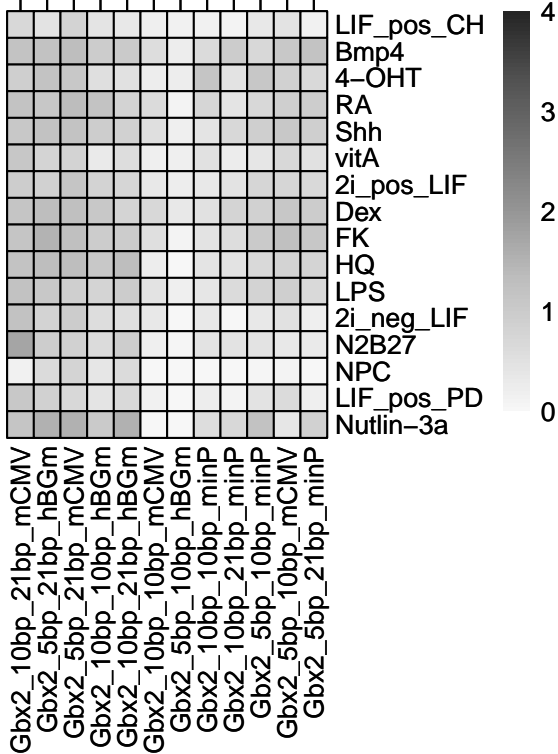
Creb1

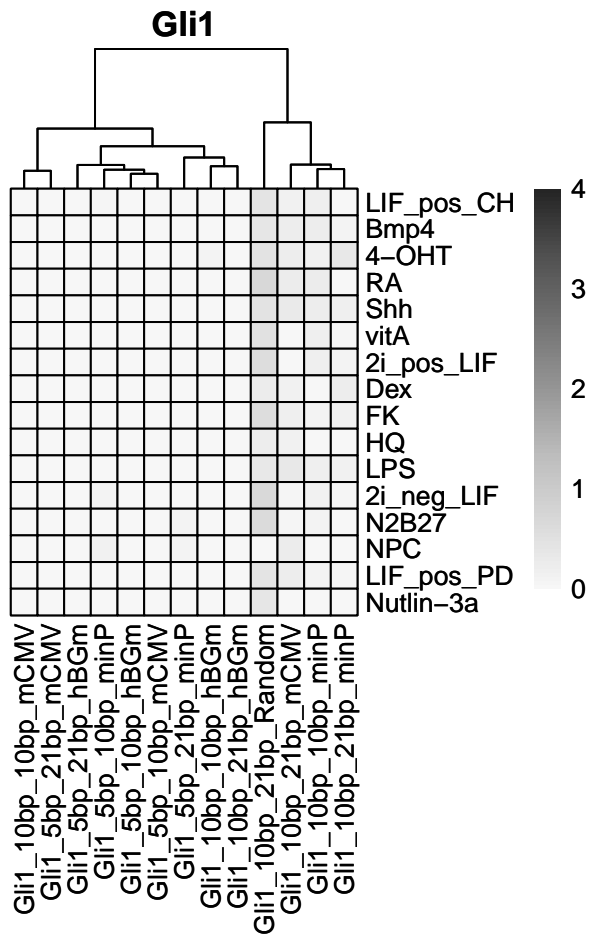


Elk1

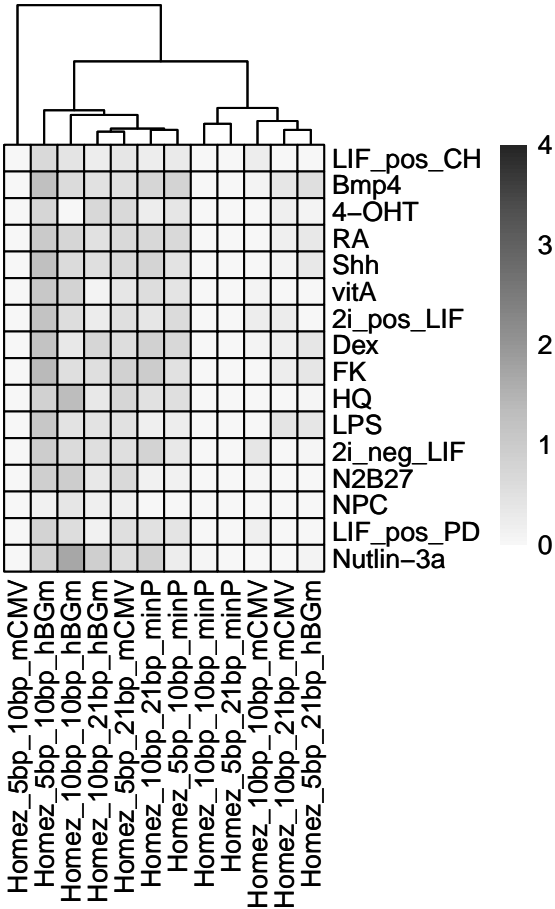


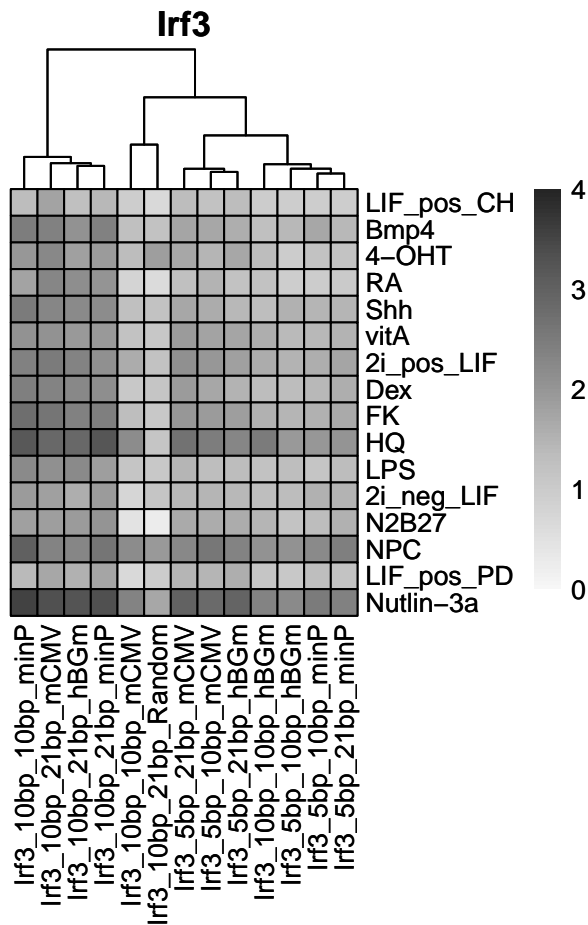


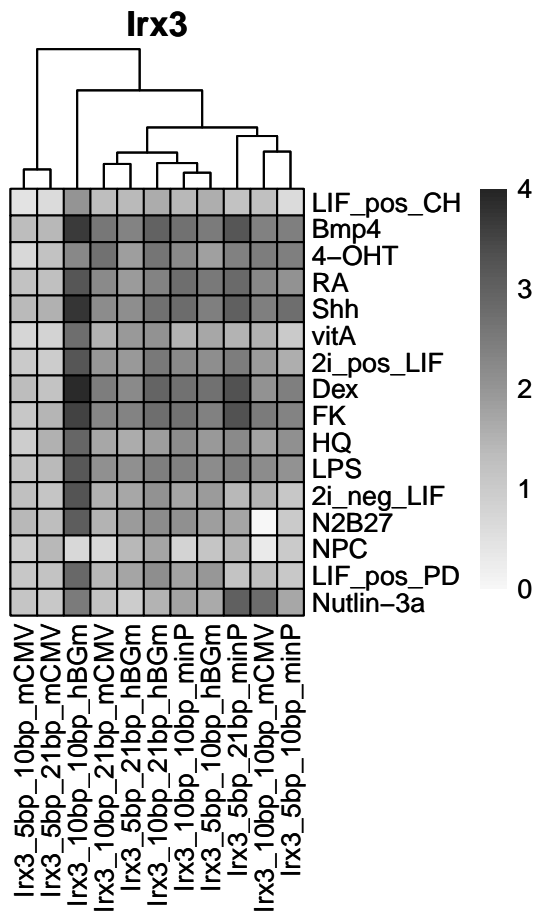


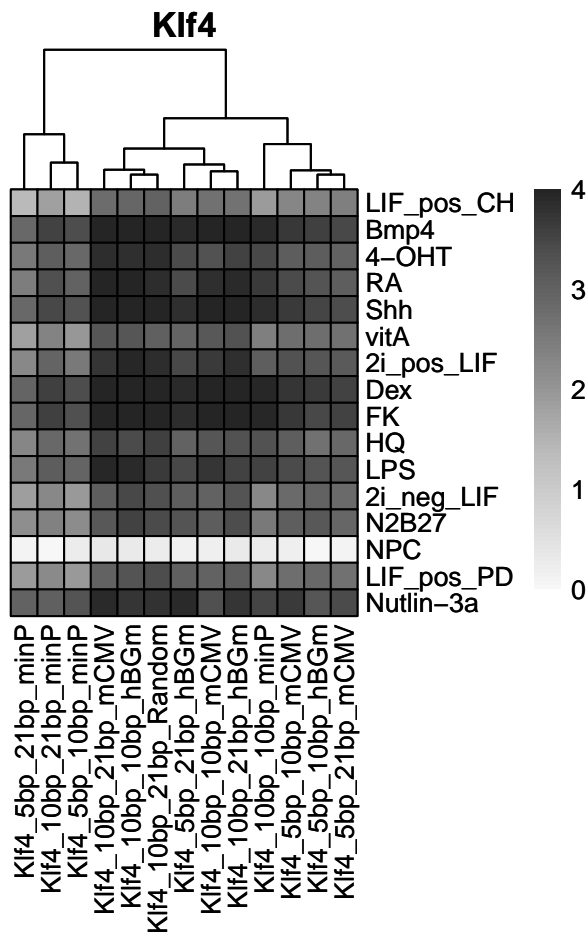


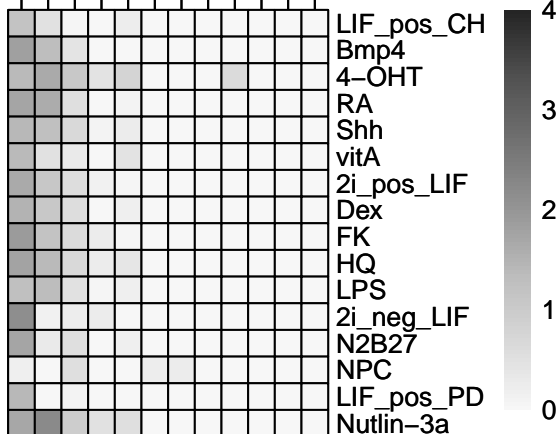
Homez



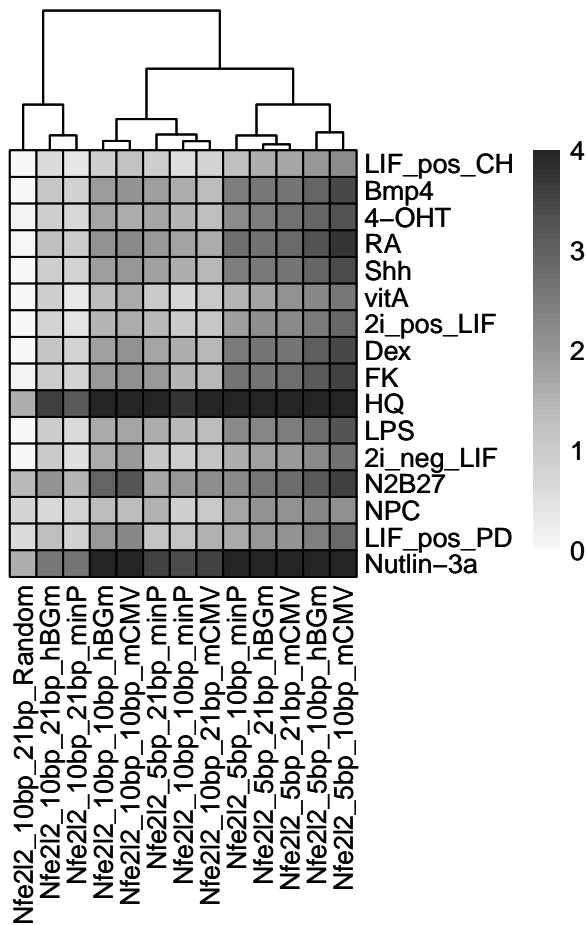








Nfe2l2



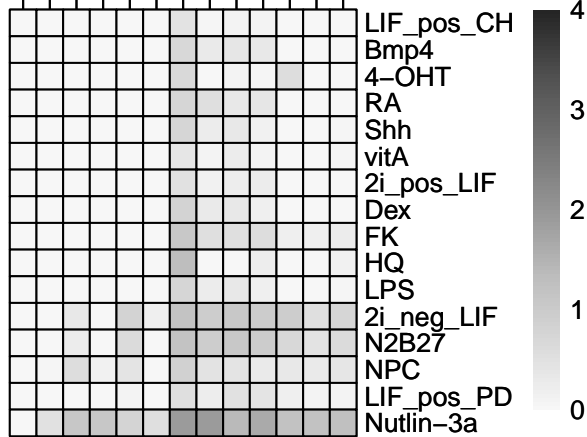
Heatmap visualization showing enrichment scores (Y-axis) for various transcription factors (X-axis) across different conditions. The color scale ranges from 0 (white) to 4 (black).

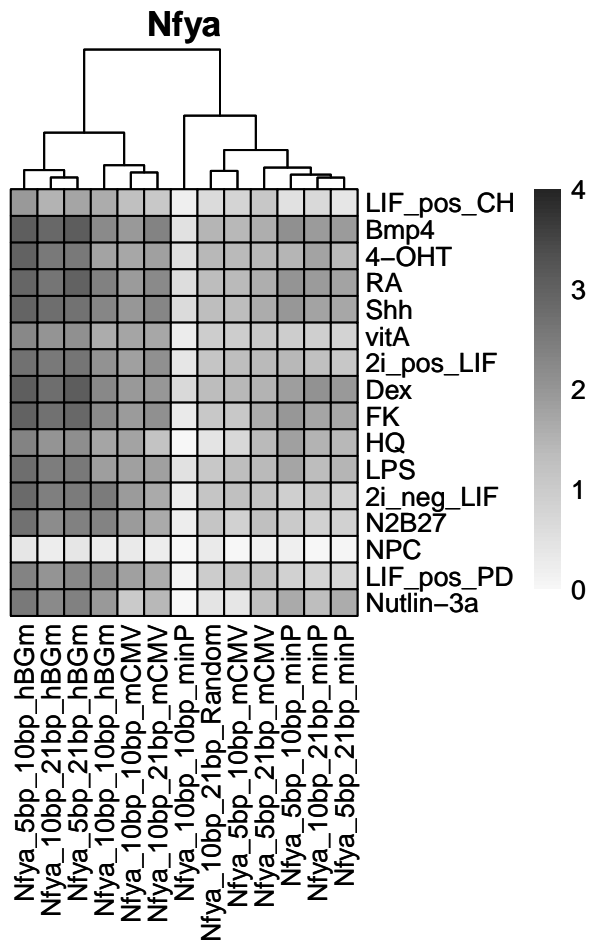
Transcription Factors (X-axis labels):

- Nfkb1_10bp_21bp_Random
- Nfkb1_5bp_21bp_minP
- Nfkb1_5bp_10bp_mCMV
- Nfkb1_5bp_10bp_minP
- Nfkb1_10bp_10bp_mCMV
- Nfkb1_10bp_21bp_minP
- Nfkb1_5bp_21bp_mCMV
- Nfkb1_5bp_10bp_hBGM
- Nfkb1_10bp_21bp_mCMV
- Nfkb1_5bp_21bp_hBGM
- Nfkb1_10bp_10bp_hBGM
- Nfkb1_10bp_10bp_minP
- Nfkb1_10bp_21bp_hBGM

Conditions (Y-axis labels):

- LIF_pos_CH
- Bmp4
- 4-OHT
- RA
- Shh
- vita
- 2i_pos_LIF
- Dex
- FK
- HQ
- LPS
- 2i_neg_LIF
- N2B27
- NPC
- LIF_pos_PD
- Nutlin-3a





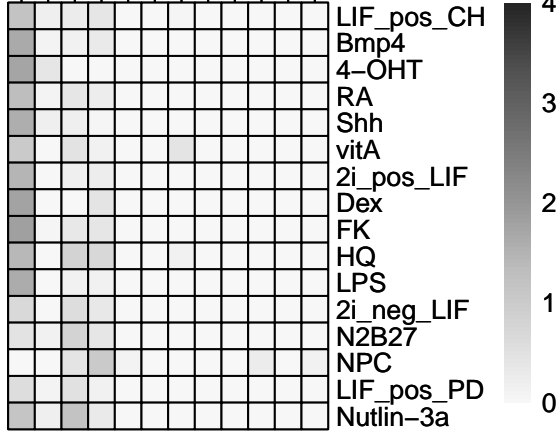
Heatmap showing the effect of various treatments on gene expression across different Nr3c1(GR) constructs. The color scale ranges from 0 (white) to 4 (black).

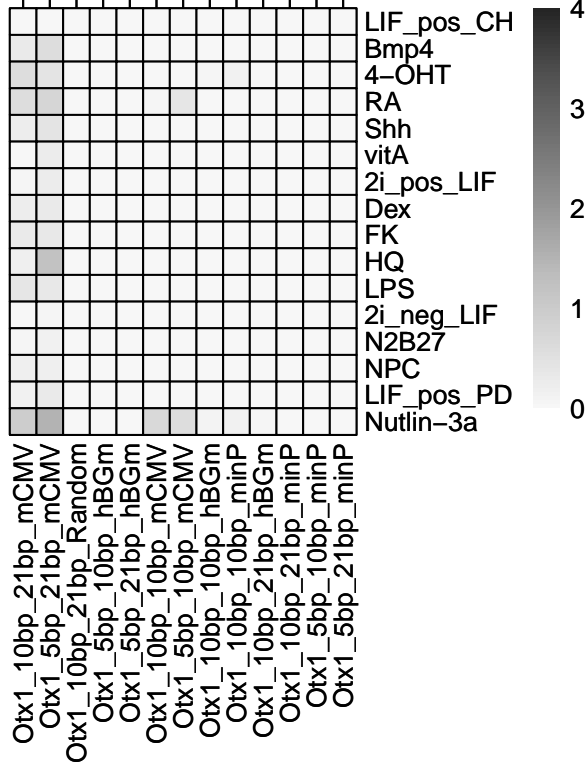
Treatments (Rows):

- LIF_pos_CH
- Bmp4
- 4-OHT
- RA
- Shh
- vitA
- 2i_pos_LIF
- Dex
- FK
- HQ
- LPS
- 2i_neg_LIF
- N2B27
- NPC
- LIF_pos_PD
- Nutlin-3a

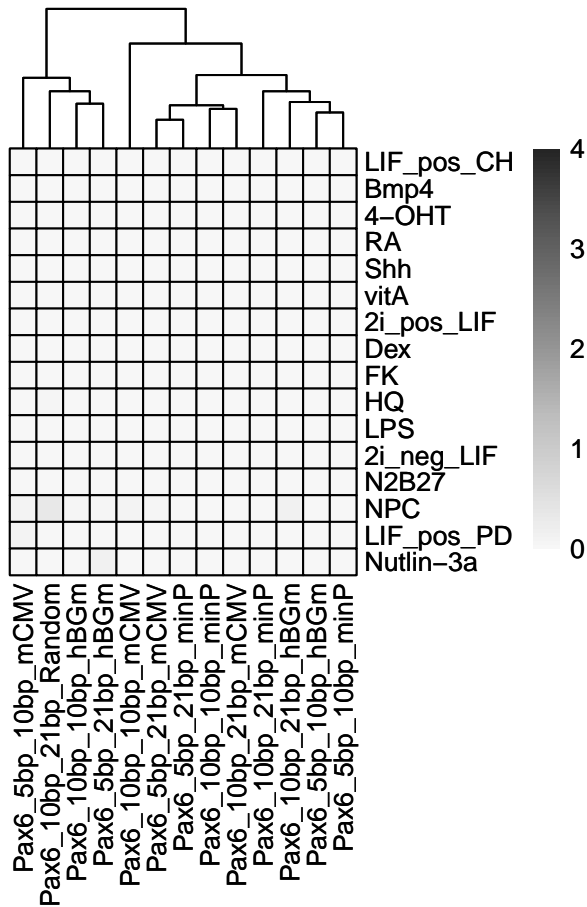
Constructs (Columns):

- Nr3c1(GR)_10bp_21bp_hBGm
- Nr3c1(GR)_10bp_10bp_hBGm
- Nr3c1(GR)_10bp_21bp_mCMV
- Nr3c1(GR)_5bp_21bp_mCMV
- Nr3c1(GR)_10bp_21bp_minP
- Nr3c1(GR)_5bp_10bp_mCMV
- Nr3c1(GR)_5bp_10bp_hBGm
- Nr3c1(GR)_5bp_21bp_minP
- Nr3c1(GR)_10bp_10bp_mCMV
- Nr3c1(GR)_5bp_10bp_minP
- Nr3c1(GR)_10bp_10bp_minP
- Nr3c1(GR)_5bp_21bp_hBGm

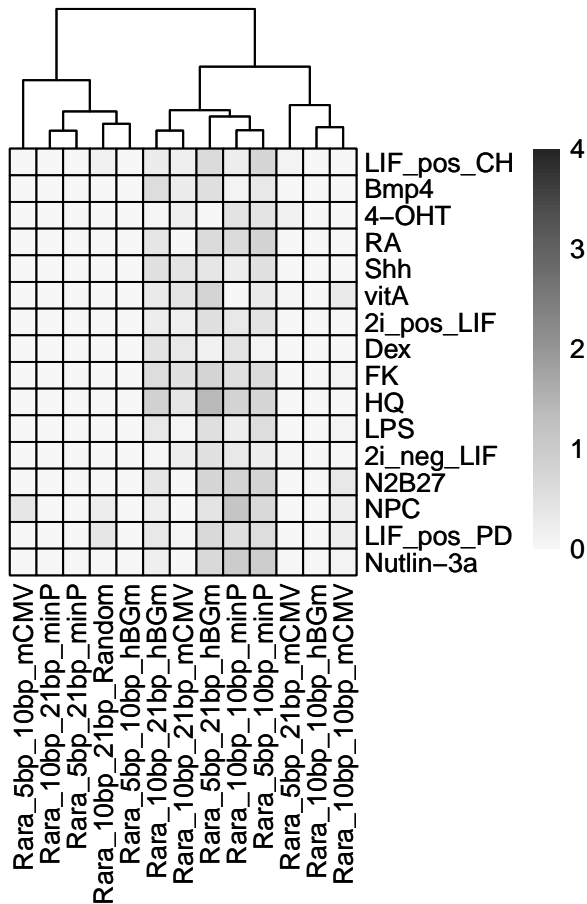


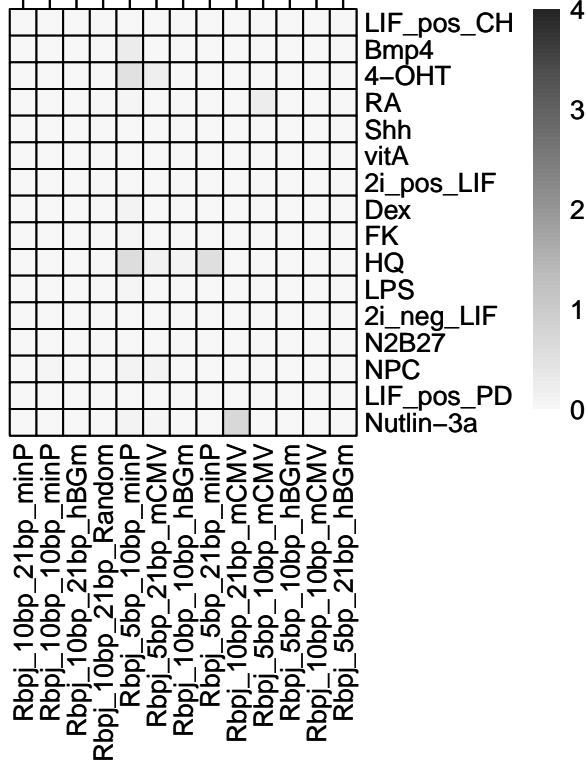
[illegible]

Pax6

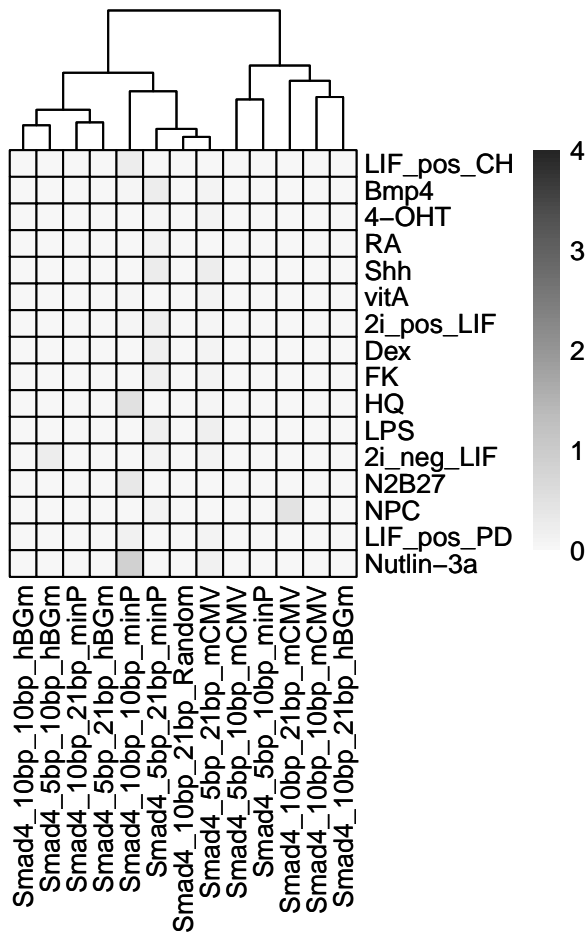


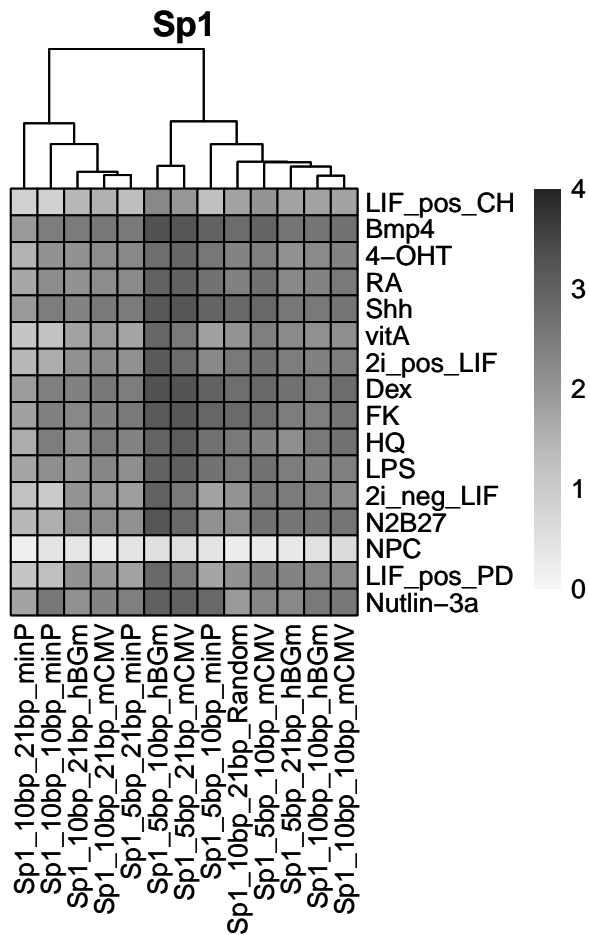
Rara

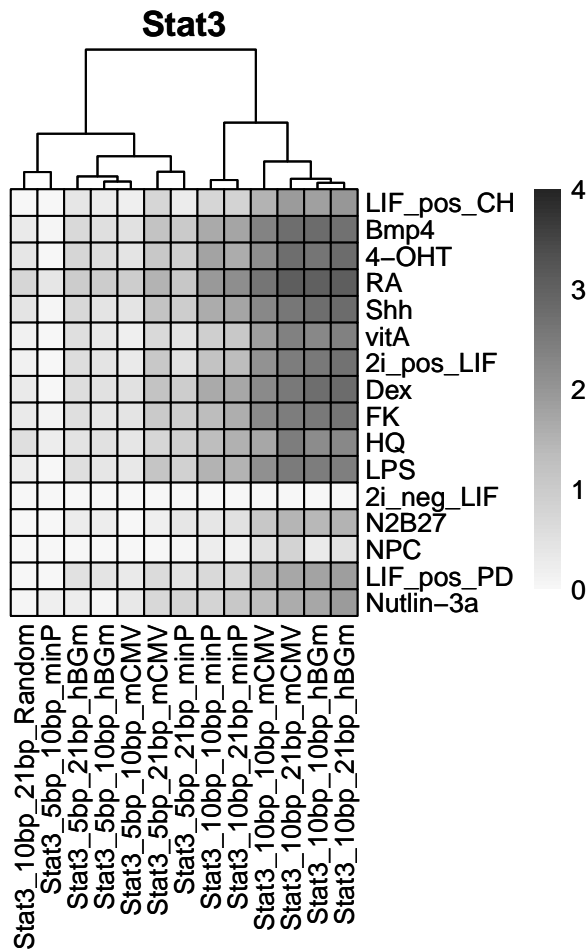




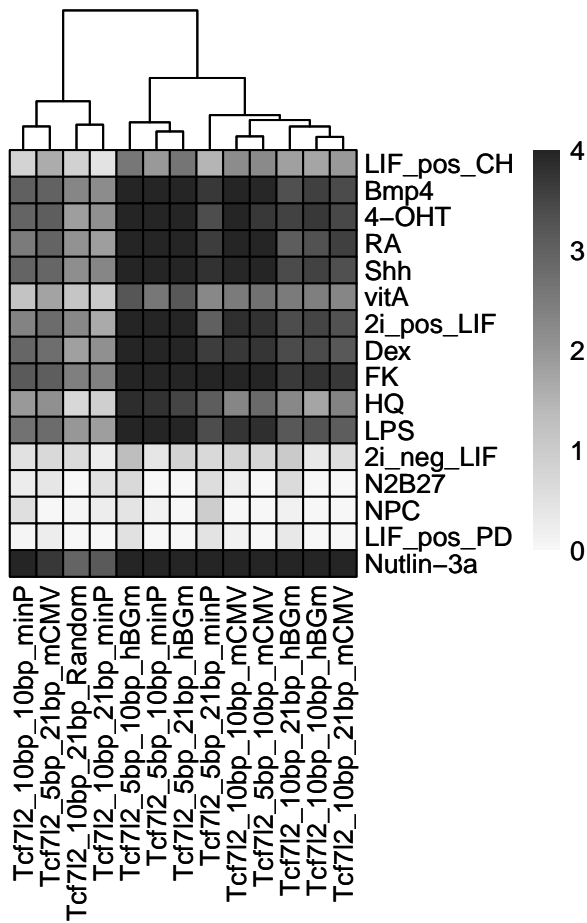
Smad4

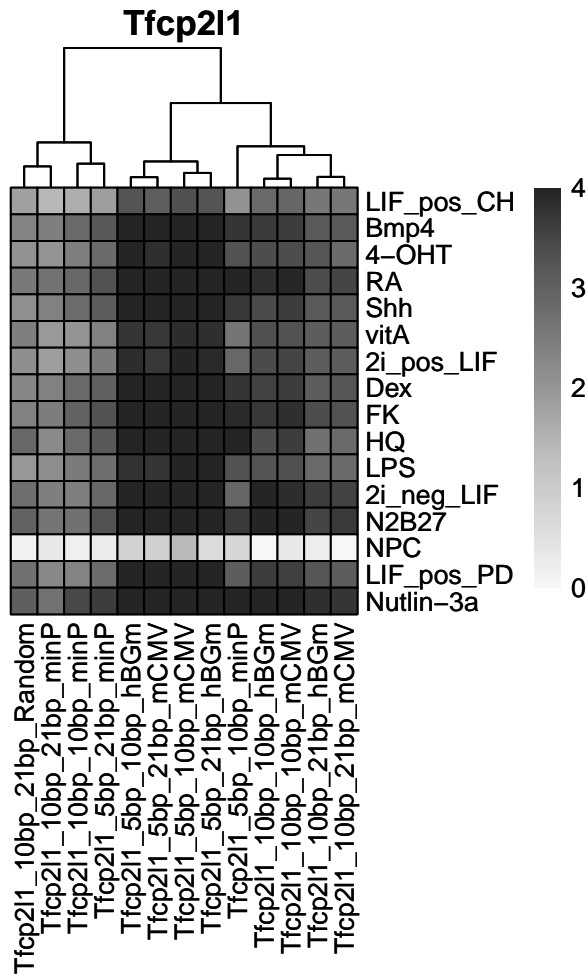




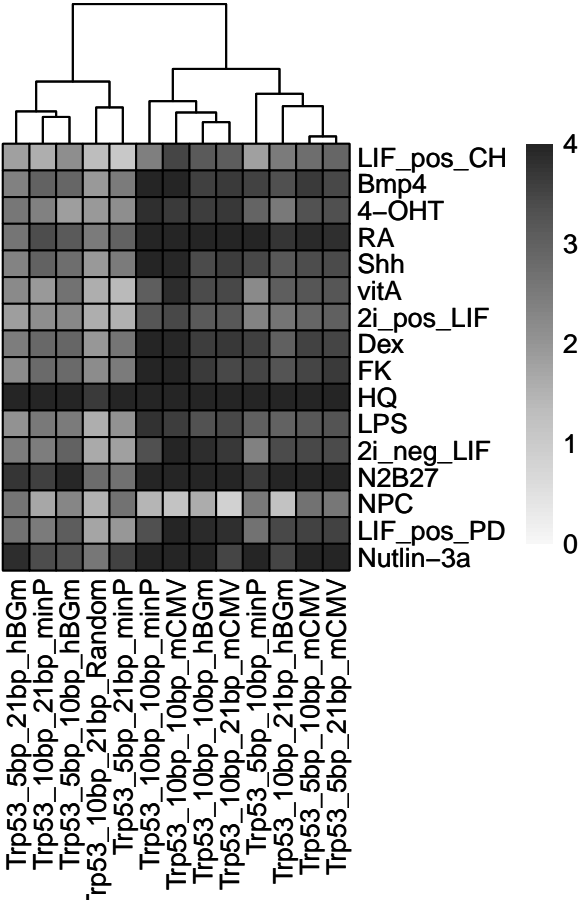


Tcf7l2

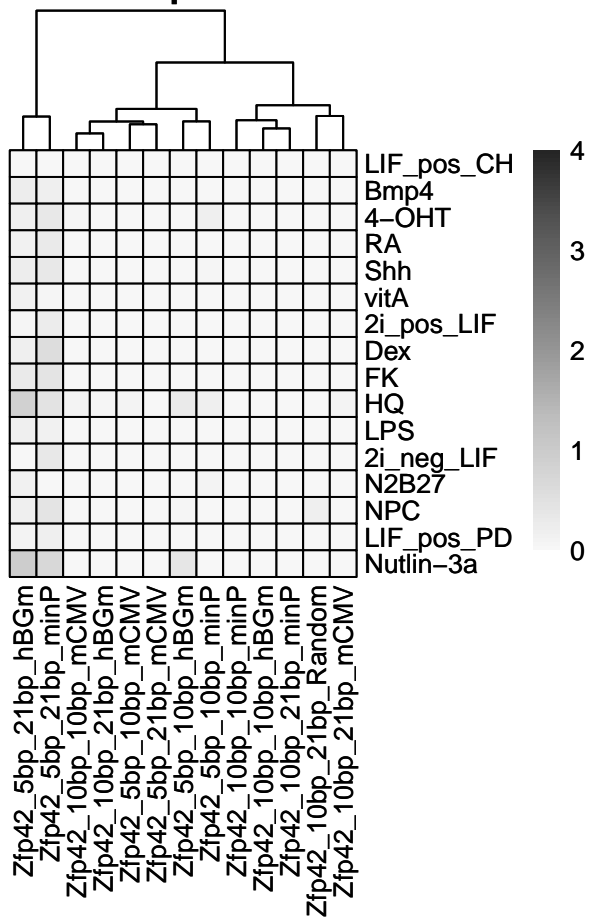




Trp53



Zfp42



Zfx