

Tesi et al. Appendix

Appendix Legends	2
Appendix Figure S1	3
Appendix Figure S2	4
Appendix Figure S3	5
Appendix Table S1	6

Appendix Figure Legends

Appendix Figure S1 Tat-Cre transiently induces LPS-responsive genes, but does not interfere with the subsequent LPS response, and does not act as a mitogenic stimulus.

A. To control for LPS contamination in bacterially purified Tat-Cre we treated freshly purified primary *c-myc*^{wt/wt} B-cells for 1 hour and monitored the expression of 2 LPS-responsive genes (*junb* and *Ikβa*) by RT-PCR. RNA was isolated at the indicated time-points after treatment with either Tat-Cre or LPS as positive control. While induced by Tat-CRE treatment at short time-points, the *junb* and *Ikβa* mRNAs returned to background levels (i.e. the same as in untreated cells) after 12h, at which time we started our LPS time-course experiments (i.e. time 0 in our LPS time-courses).

B. Pretreatment with Tat-CRE does not affect the response of *junb* and *Ikβa* to LPS. RNA was collected 12h after Tat-Cre treatment (0h LPS) and 1,2 and 4h after LPS stimulation.

C. Growth curves of wt B-cells treated with Tat-Cre alone, LPS alone, Tat-Cre followed by LPS or left untreated (none).

Appendix Figure S2

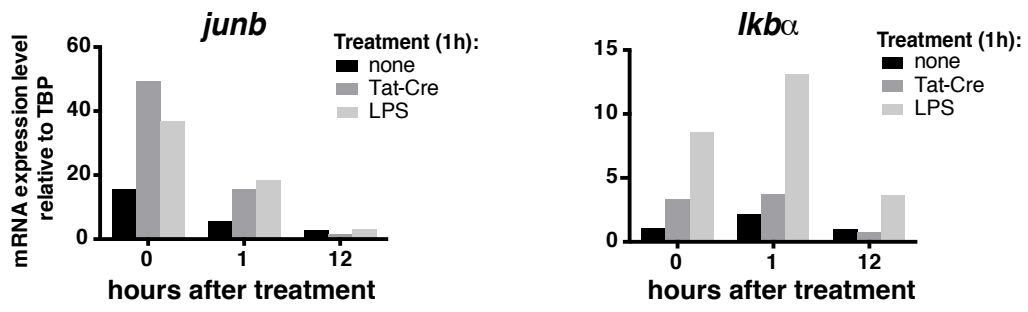
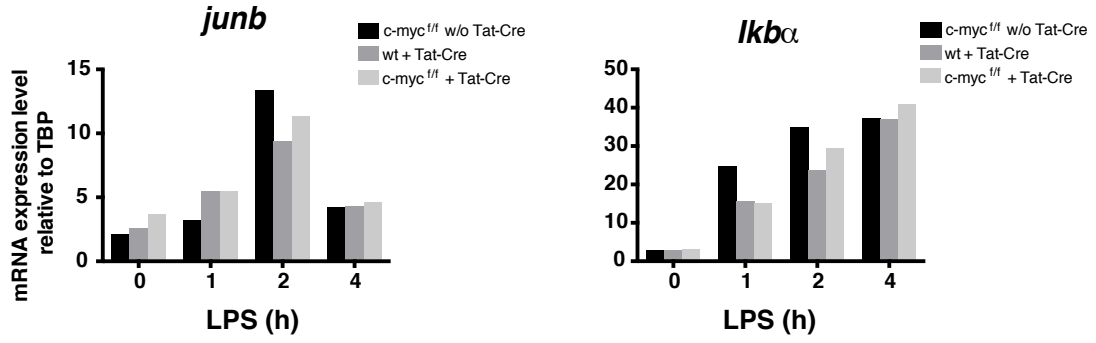
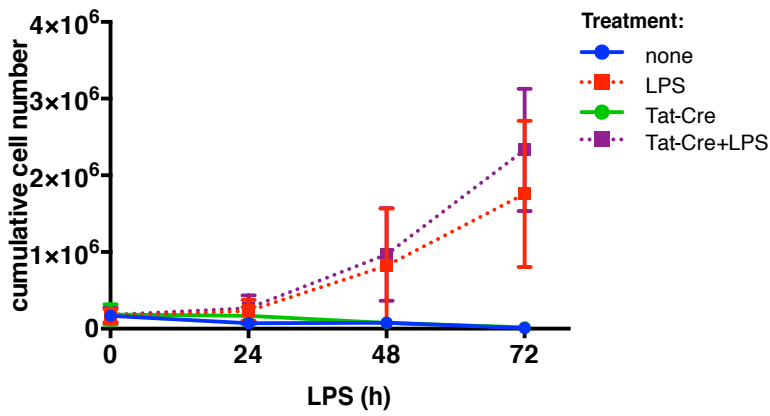
Zoom-in heatmaps and boxplot representations for the clusters represented in Fig. 4D.

Appendix Figure S3

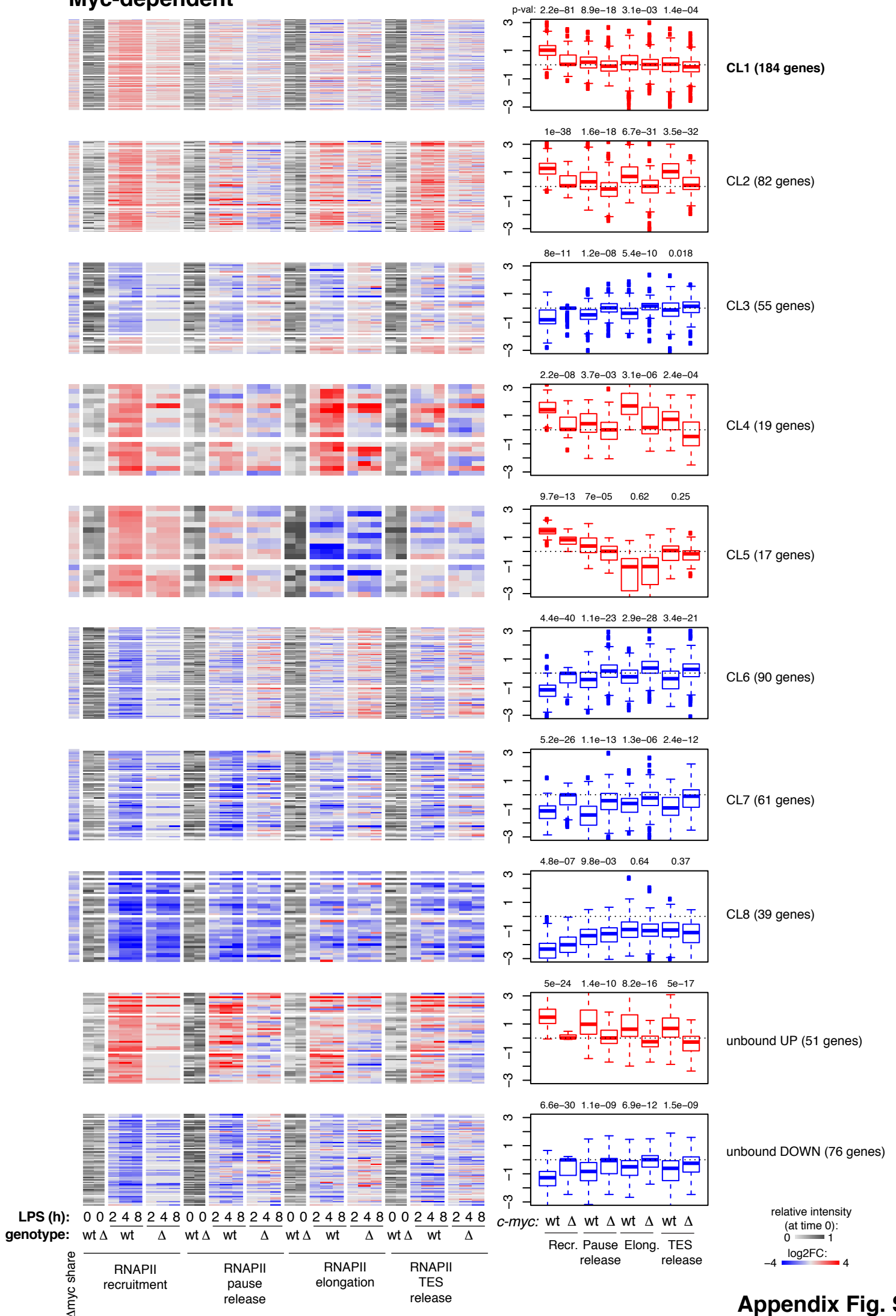
Zoom-in heatmaps and boxplot representations for the clusters represented in Fig. 4E.

Appendix Table S1

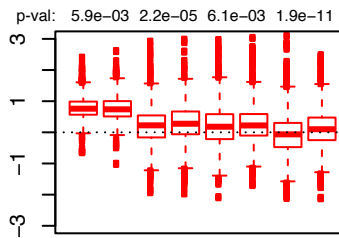
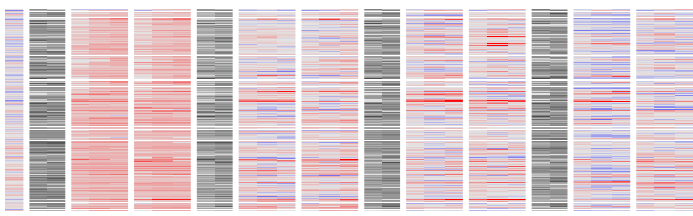
List of the primers used in this work.

A**B****C**

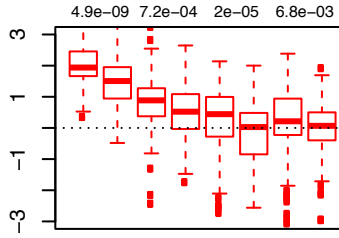
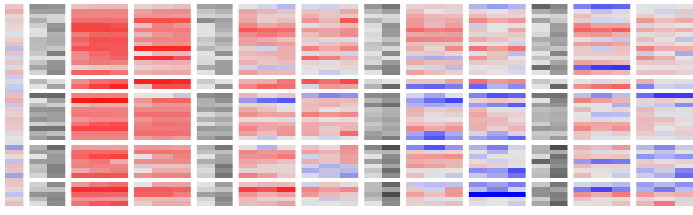
Myc-dependent



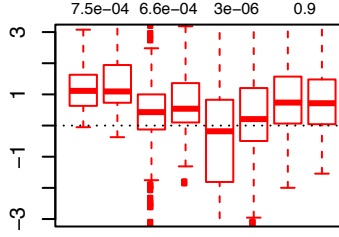
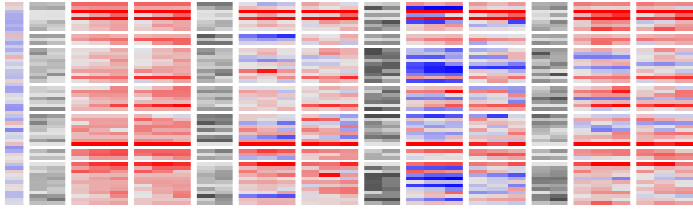
Myc-independent



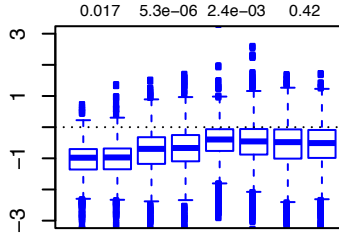
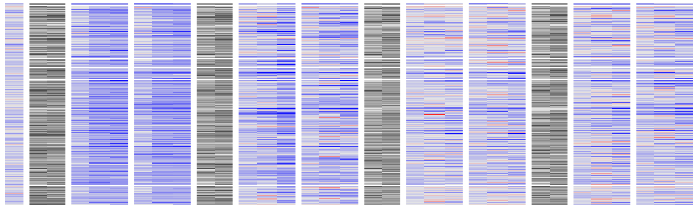
CL9 (260 genes)



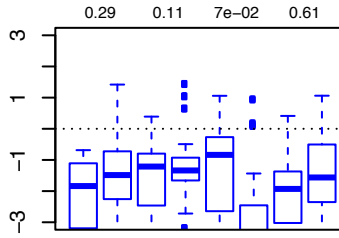
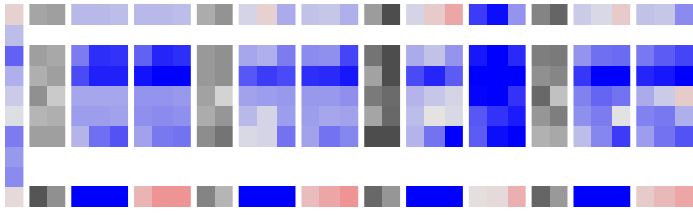
CL10 (43 genes)



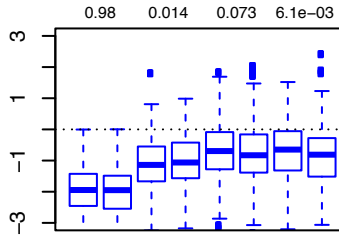
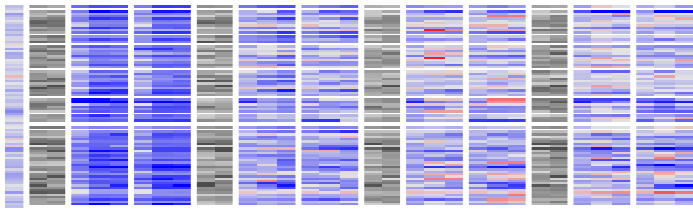
CL11 (58 genes)



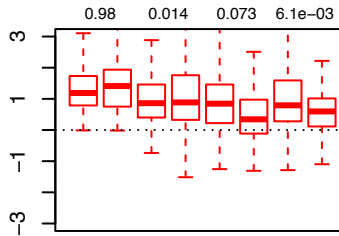
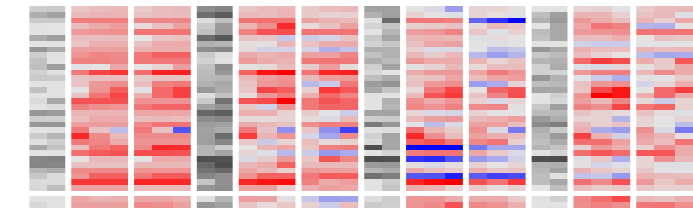
CL12 (398 genes)



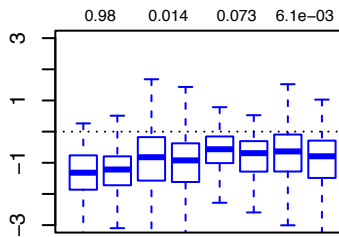
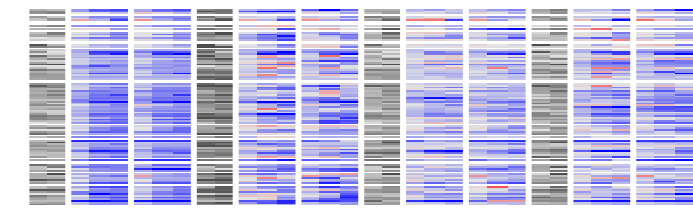
CL13 (10 genes)



CL14 (87 genes)



unbound UP (35 genes)



unbound DOWN (114 genes)

LPS (h): 0 0 2 4 8 2 4 8 0 0 2 4 8 2 4 8 0 0 2 4 8 2 4 8 0 0 2 4 8 2 4 8
 genotype: wt Δ wt Δ wt Δ wt Δ wt Δ wt Δ wt Δ wt Δ wt Δ

Δmyc share

RNAPII recruitment

RNAPII pause release

RNAPII elongation

RNAPII TES release

c-myc: wt Δ wt Δ wt Δ wt Δ
 Recr. Pause Elong. TES
 release release release release

relative intensity (at time 0):
 0 1
 log2FC:
 -4 4

<u>List of primers for gene expression</u>			
Primers number	Gene Symbol	Forward	Reverse
AP10011-12	Tbp	TAATCCCAAGCGATTGCTG	CAGTTGTCCGTGGCTCTCTT
AP6540-41	Myc	TTTTTGTCTATTTGGGACAGTG	CATCGTCGTGGCTGTCTG
AP1985-86	Ncl	GTCTGAGGATACCACTGAAG	GCCCAGTCCAAGGTAAC
AP8859-60	Pus7	GCAGAAGAATTGAGGAGTACGG	GGATGGAGTAGTTGTCAACGTCA
<u>List of primers for CHIP</u>			
Primers number	Gene Symbol	Forward	Reverse
AP4231-32	AchR	AGTGCCCCCTGCTGTCAGT	CCCTTTCCTGGTGCCAAGA
AP2829-30	Ncl	GGCGTGGTGACTCCACGT	CGAAATCACCTCTAAAGCAGA
<u>List of primers for gDNA</u>			
Primers number	Gene Symbol	Forward	Reverse
AP2829-30	Ncl	GGCGTGGTGACTCCACGT	CGAAATCACCTCTAAAGCAGA
AP1167-68	myc fl/fl	TCTAGACTTGCTTCCCTTGCTGT	TTCCTGTTGGTGAAGTTCACGT
AP1169-70	myc Δ	AAATAGTGATCGTAGTAAAATTTAGCCTG	ACCGTTCTCCTTAGCTCTCACG
<u>List of primers for genotyping</u>			
Primers number	Gene Symbol	Forward	Reverse
AP6509-10	myc	CACCGCTACATCCTGTCCATTC	TACAGTCCCAAGCCCCAGCCAAG

Appendix Table S1