

# PNAS

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Supporting Information for

## **Global epigenomic analysis of KSHV-infected primary effusion lymphoma identifies functional MYC super-enhancers and enhancer RNAs**

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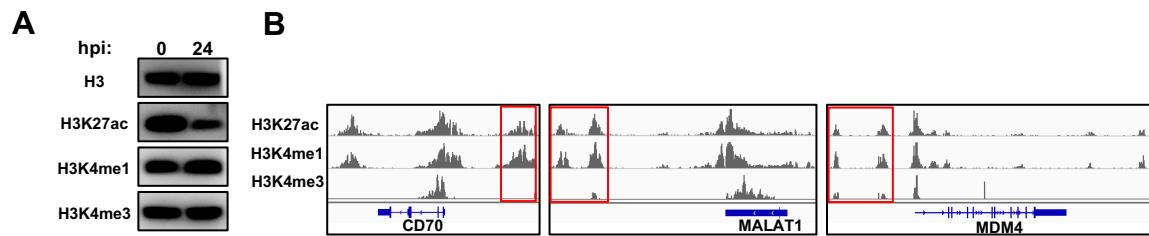
Email: [mrosenfeld@ucsd.edu](mailto:mrosenfeld@ucsd.edu)

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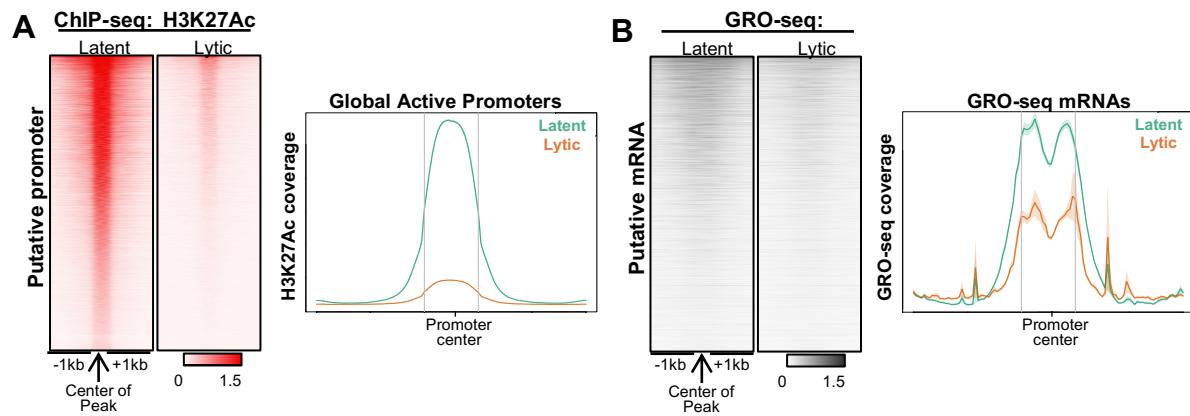
Figures S1 to S3

Tables S1 to S3

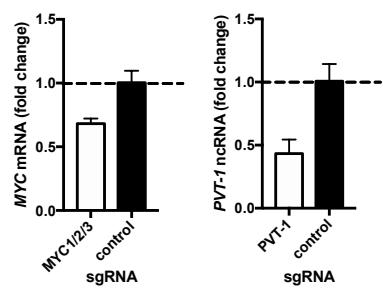
## Supplementary Information



**Fig. S1.** (A) Immunoblot analysis of total H3 and H3 modification levels in TRExBCBL1-RTA cells during latency and lytic replication (induced by doxycycline for 24 h). (B) Three strongest super-enhancers (red box) in TRExBCBL1-RTA cells during latency showing high levels of H3K27ac and H3K4me1 ChIP-seq signals relative to H3K4me3 signals.



**Fig. S2.** (A) Left: Heatmap of H3K27ac ChIP-seq signals at putative active promoters in TRExBCBL1-RTA cells during latency and lytic replication induced by doxycycline for 24 h; each row represents one promoter region. Right: Density plot of average ChIP-seq signals in 10 kb windows around the center of promoter of KSHV latent-infected (green) or lytic-infected (orange) cells. (B) Left: Heatmap of GRO-seq signals at putative active promoters identified in A. Right: Density plot of average GRO-seq signals in 10kb windows around the center of promoter of KSHV latent-infected (green) or lytic-infected (orange) cells.



**Fig. S3.** CRISPRi-mediated depletion. dCas9-KRAB-MYC-sgRNA or dCas9-KRAB-PVT1-sgRNA was used to test system efficacy in TRExBCBL1-RTA cells. 3 different sgRNAs were mixed to repress *MYC* mRNA.

**Table S1.** Primers used for qPCR assays.

| Primer name          | Fwd                      | Rev                     | Assay                     |
|----------------------|--------------------------|-------------------------|---------------------------|
| <b>18S</b>           | TTCGAACGTCTGCCCTATCAA    | GATGTGGTAGCCGTTCTCAGG   | RT-qPCR                   |
| <b>MYC</b>           | AGAGTTCATCTCGCACCG       | AAGCCGCTCCACATACAGTC    | RT-qPCR                   |
| <b>MYC_e486</b>      | GCCCTGTGAAACCTAATGACA    | AAGAGGGCATGGAGAGTGATT   | RT-qPCR, ChIP-qPCR, ddPCR |
| <b>MYC_e507</b>      | CCCACCGTGATTCTGAGG       | CACACCCAAGACTGGGAAG     | RT-qPCR, ChIP-qPCR, ddPCR |
| <b>MYC_e530</b>      | GAATCCATTCAGCCTTGCT      | TCTGTCCTCCTTGGGCTCT     | RT-qPCR, ChIP-qPCR, ddPCR |
| <b>LANA</b>          | GAGTCTGGTACGACTTGGAG     | AGGAAGGCCAGACTCTTCAAC   | RT-qPCR                   |
| <b>RTA</b>           | TTGCCAACAGTTGTACAAC TGCT | ACCTTGCAAAGACCATT CAGAT | RT-qPCR                   |
| <b>K2</b>            | TCACTGCGGGTTAATAGGATT    | CATGACGTCCACGTTATCACT   | RT-qPCR                   |
| <b>ORF 25</b>        | ACAGTTATGGCACGCATAGTG    | GGTTCTCTGAATCTCGTCGTG T | RT-qPCR                   |
| <b>cellular IRF4</b> | GGCCAGAGGAAAAACATTGA     | ATCCTGCTCTGGCACAGTCT    | RT-qPCR                   |
| <b>viral IRF4</b>    | GAGCTCCTCAACCAGACAGG     | GCTGACTATCAGGGGGATCA    | RT-qPCR                   |
| <b>PANRNA</b>        | GAATCCATTCAGCCTTGCT      | TCTGTCCTCCTTGGGCTCT     | RT-qPCR, ddPCR            |

**Table S2.** Primers used for 3C & 4C-seq assays.

| Primer name    | Fwd   | Rev  |
|----------------|---|--|
| 3C-e486        | CAGCCGAGCACTCTAGCTCT                                | CCAAAGCACTGACACCTGTG                         |
| 3C-e507        | CAGCCGAGCACTCTAGCTCT                                | AGGTTGCCAGCATAGACAC                          |
| 3C-e530        | CAGCCGAGCACTCTAGCTCT                                | CAGAAAGTCCCTCAAGGTGG                         |
| 4C-mycPromoter | GTTCAAGAGTTCTACAGTCCGACGATC<br>AGCTGCTGGGAGGAGACATG | AGACGTGTGCTTCCGATCT<br>CCACGTCTAACACCTCTAG   |
| 4C-mycEnhancer | GTTCAAGAGTTCTACAGTCCGACGATC<br>CTCATCTGCCGAAGCCTT   | AGACGTGTGCTTCCGATCT<br>CTGCTGCTCATTTGCATAATG |

**Table S3.** shRNA and sgRNA sequence.

| Primer name | Fwd                   | Rev                    |
|-------------|-----------------------|------------------------|
| shRNA-e486  | GGAGGACATGACAGCAGAAGT | ACTTCTGCTGTCATGTCCCTCC |
| shRNA-e507  | GCCTGCCACATAACATCAATC | GATTGATGTTATGTGGCAGGC  |
| shRNA-e530  | GCTCTGCTTGCTAGTTATCT  | AGATAACTAGCAAAGCAGAGC  |
| sgRNA-e486  | GCTTACAAGCCACAGTGTTC  | ACCCCCGCAAGTTCATAGA    |
| sgRNA-e507  | GGAGTCACAATACATTGGAG  | CAGCCTATGGATGCAAGCTA   |
| sgRNA-e530  | CACCACGTAACCTTCCACCT  | ATCCACTGGCAACCATTCAC   |
| sgRNA-MYC1  | GCGAAGCCCCCTATTGCTC   | CGAAAACCGGCTTTATACT    |
| sgRNA-MYC2  | AGGACGCGACTCTCCGACG   | TCGCATTATAAAGGGCCGGT   |
| sgRNA-MYC3  | AGCTATCCCCCTAAAGCGGCT | GCTATCTGGAGACGCACCTT   |
| sgRNA-PVT1  | TCCTCCGGGCAGAGCGCGTG  | CCACACGCGCTTGCCCCGGA   |