

The Transcriptional Foundations of Sp110-mediated Macrophage (RAW264.7) Resistance to *Mycobacterium tuberculosis* H37Ra

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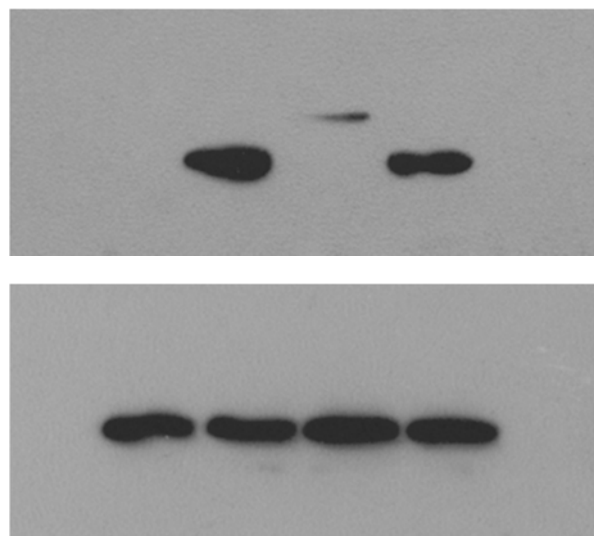
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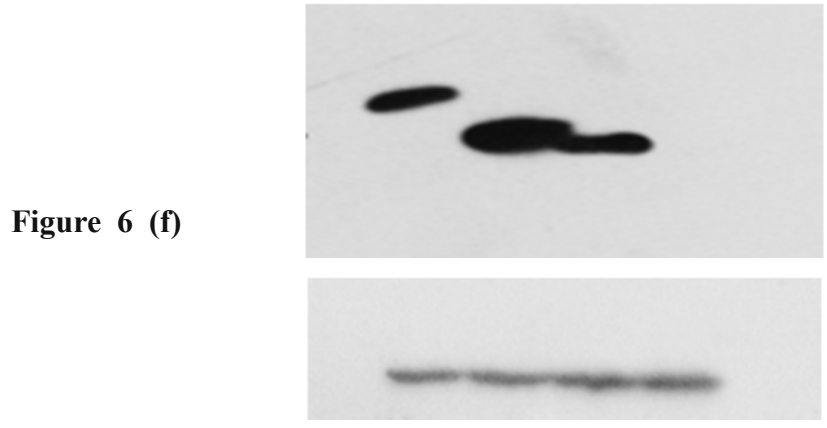
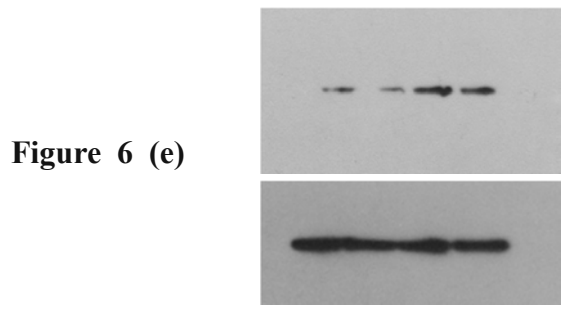
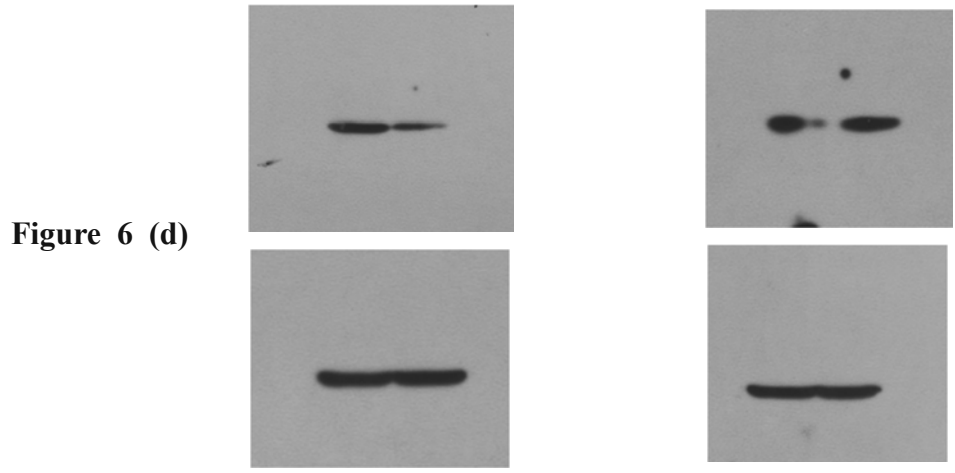
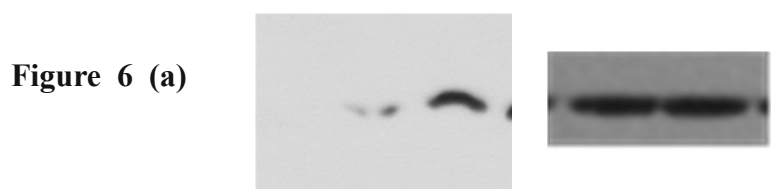
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Supplementary Figure 1. Full-length blots of Figure 1 (b) in the main text.



Supplementary Figure 2. Full-length blots of Figure 6 (a), (d), (e) and (f) in the main text.



Supplementary Table 10. **Primer sequences for qPCR analysis.**

Gene name	Forward primer (5'- 3')	Reverse primer (5'-3')
Bmf	GGAGCGGGCGTATTTTGGAA	ACACTCGATTGGGAAGAAGGG
Dock1	AGTACGGCGTGGCCTTTTAC	GACGGTTTCATGTTGCCCTTT
Il10	GCTCTTACTGACTGGCATGAG	CGCAGCTCTAGGAGCATGTG
Ccl2	TTAAAAACCTGGATCGGAACCAA	GCATTAGCTTCAGATTTACGGGT
Aatk	GGGAGGTATGAGTGTGGTGG	TAGATCCCCATTCGTGGTGTG
Adrbk1	GCGCCAGCAAGAAGATCCT	GCAGAAGTCCCGGAAAAGCA
Atp1a3	TCTCAGATGTGTCCGTTCTTCT	TGGAAAGAGAGTGAAAGGCAAG
Dusp14	TTGCTCAGATCACCTCCTCTC	AGTACAGTCTAATGGGGGCAT
Chek1	GTTAAGCCACGAGAATGTAGTGA	GATACTGGATATGGCCTTCCCT
Csf3	ATGGCTCAACTTTCTGCCAG	CTGACAGTGACCAGGGGAAC
Cxcl2	CCAACCACCAGGCTACAGG	GCGTCACACTCAAGCTCTG
Irak3	AGGCCAAGTTAAGACCCACG	ACCTCAGACTGGCTGCATTC
Pdcd4	CGAGGGGATCCTCAAGGAAGA	TTTCCGCAGTCGTCTTTTGG
Pdcd1	TCATGAGTGCCCTAGTGGGT	AGGGAGCTCTGGTGTCTTCT
Cend2	AGTCCCAGACTCCTAAGACCC	GAAGTCGTGAGGGGTGACTG
Pmp22	CATCGCGGTGCTAGTGTTG	AAGGCGGATGTGGTACAGTTC
Ccl4	TTCCTGCTGTTTCTCTTACACCT	CTGTCTGCCTCTTTTGGTCAG
Il6	GCCTTCTTGGGACTGATGCT	TGCCATTGCACAACCTTTTTCT
Tnf	CCCTCACACTCACAACCAC	ATAGCAAATCGGCTGACGGT
Nos2	GGTGAAGGGACTGAGCTGTT	ACGTTCTCCGTTCTCTTGCAG
Casp2	CACAGACTCCTGCGTAGTGG	TGTCTCATCTCCACGACATGC
Casp3	GAGCTTGAACGGTACGCTA	GCGAGATGACATTCCAGTGC
Casp6	GAGGTGTCGACTTCAAGCCA	CCGACAGGCCTGGATGATAA
Casp7	CCTCTGGGACTTTTGCTTTCA	CTGATCATCGGTCATCGTTCC
Casp8	CGGAGGCATTTCTGTCCCCTA	AGAGCTGTAACCTGTGGCCG
Casp9	ACCTTCCCAGGTTGCCAATG	AGCCTGCCCCAGTTCAAAT
Gapdh	GTGTTCTACCCCAATGTGT	ATTGTCATACCAGGAAATGAGCTT
miR-125a	TCCCTGAGACCCTTTAACC	The reverse primer for qPCR experiments of miRNAs is a universal primer provided by the miScript II RT Kit (QIAGEN).
miR-21a	GGGTAGCTTATCAGACTGATGTTGA	
miR-27b	AGAGCTTAGCTGATTGGTGAAC	
miR-29a	ACTGATTTCTTTTGGTGTTCAG	

miR-146a	GGTGAGAACTGAATTCCATG	
miR-155	TTAATGCTAATTGTGATAGGGGT	
miR-99b	CACCCGTAGAACCGACCTTGC	
Rnu6	TGCGCAAGGATGACACGC	