

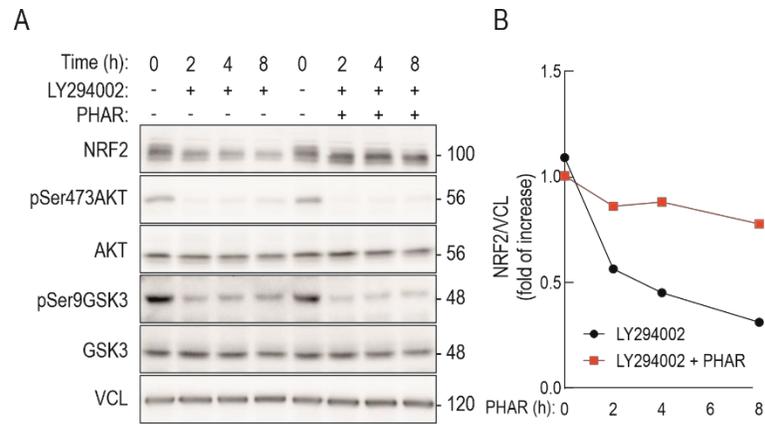
**An inhibitor of interaction between the transcription factor NRF2 and the E3 ubiquitin ligase adapter  $\beta$ -TrCP delivers anti-inflammatory responses in mouse liver**

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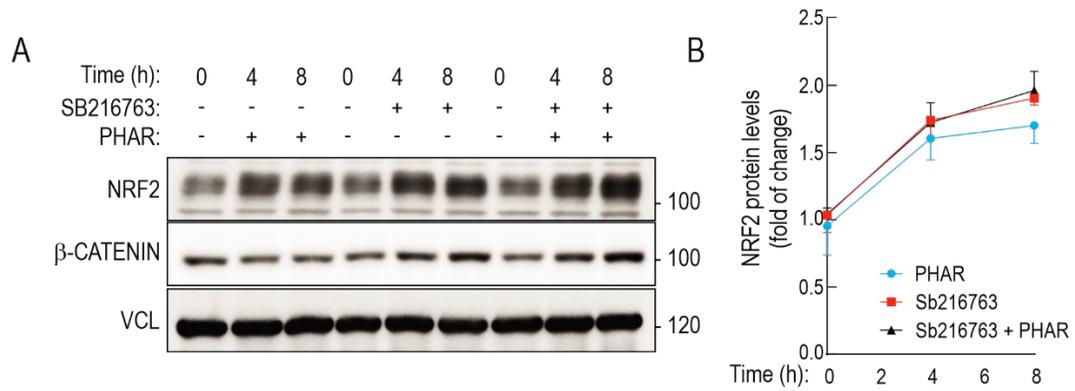
**SUPPLEMENTARY MATERIAL**

Gene product	Forward primer (5'-3')	Reverse primer (5'-3')
<i>Hmox1</i>	CACAGATGGCGTCACTTCGTC	GTGAGGACCCACTGGAGGAG
<i>Aox1</i>	CTTTTGACCAAAGCATCAGTCTC	CCTTTCTCCCAGTCTATATTCGA
<i>Nqo1</i>	GGTAGCGGCTCCATGTACTC	CATCCTCCAGGATCTGCAT
<i>Nrf2</i>	CCCGAAGCACGCTGAAGGCA	CCAGGCGGTGGGTCTCCGTA
<i>Gclc</i>	TTACCGAGGCTACGTGTCAGAC	TATCGATGGTCAGGTTCGATGTC
<i>Gclm</i>	AATCAGCCCCGATTTAGTCAGG	CCAGCGTGCAACTCCAAGGAC
<i>Btrc</i>	AGCGGCTCCTCTGACAACACCAT	AGCACGGGGTCCAAAGCA
<i>Fbxw11</i>	GTCCGCACTCTGAATGGGCACA	GCACCGGACCAATTCTTCG
<i>Ptgs2</i>	TTCGGGAGCACAACAGAGTG	TAACCGCTCAGGTGTTGCAC
<i>Nos2</i>	CCTCCTTTGCCTCTCACTCTTC	AGTATTAGAGCGGTGGCATGGT
<i>Il1b</i>	CTGGTGTGTGACGTTCCATTA	CCGACAGCACGAGGCTTT
<i>Tnf</i>	CATCTTCTCAAATTCGAGTGACAA	TGGGAGTAGACAAGGTACAACCC
<i>Il6</i>	CCTACCCCAATTTCCAATGCT	TATTTTCTGACCACAGTGAGGAATG
<i>Actb</i>	TCCTTCTGGGCATGGAG	AGGAGGAGCAATGATCTTGATCTT
<i>Gapdh</i>	CGACTTCAACAGCAACTCCCACTCTTCC	TGGGTGGTCCAGGGTTTCTTACTCCTT
<i>Tbp</i>	TGCACAGGAGCCAAGAGTGAA	CACATCACAGCTCCCCACCA

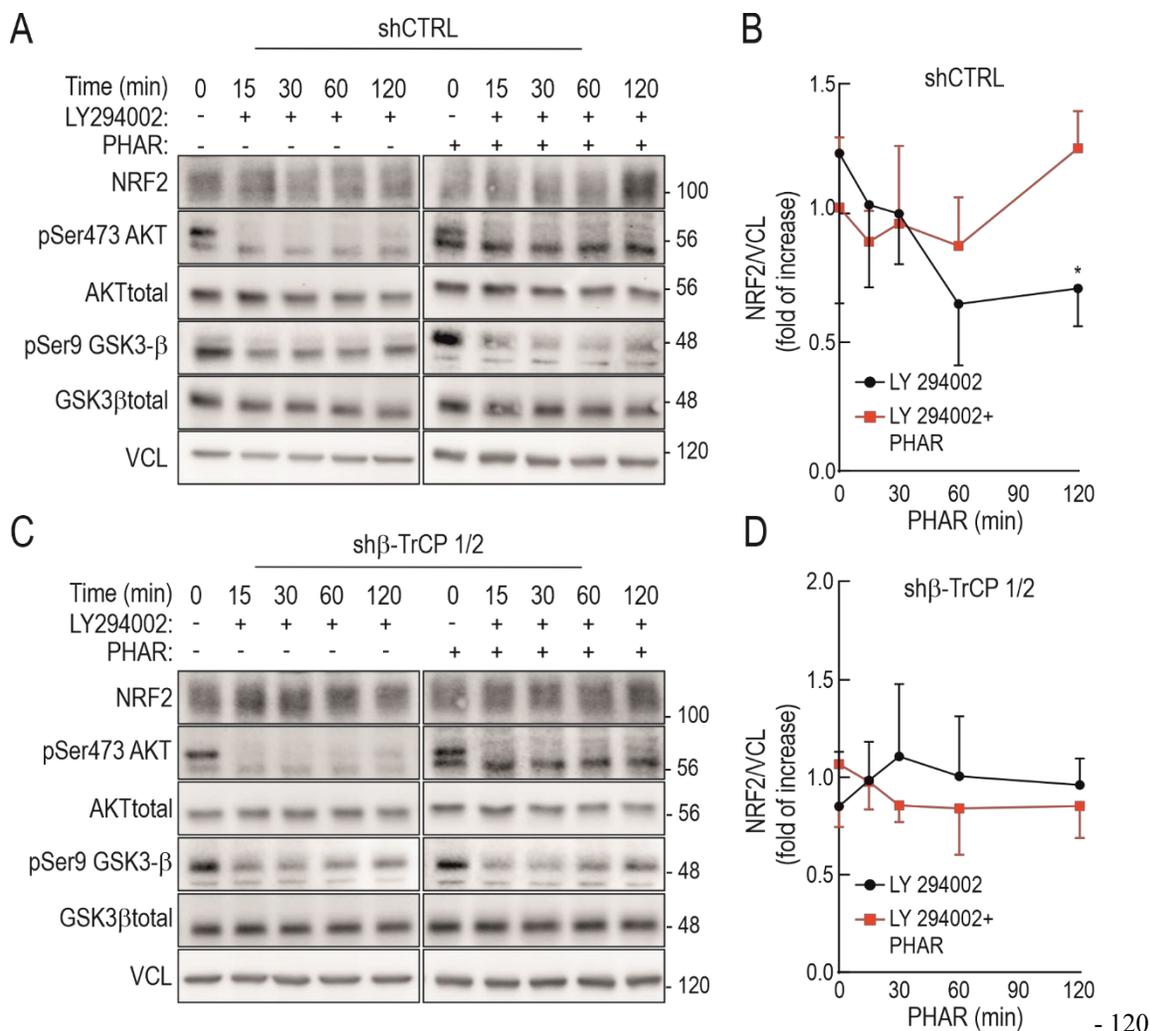
**Supplementary Table 1.** Mouse primers used for RT-PCR.



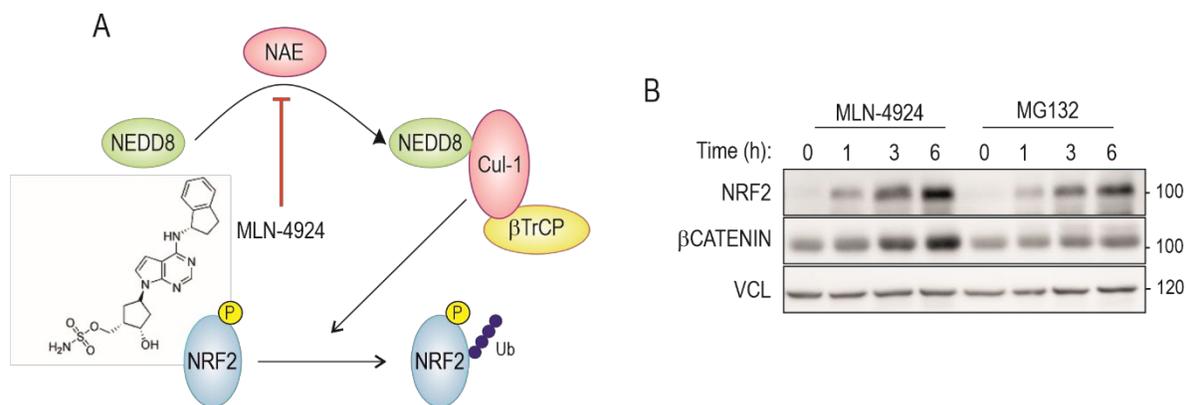
**Figure S1.** PHAR prevents the decrease in NRF2 levels elicited by the phosphoinositide 3-kinase inhibitor LY294002. Serum-depleted *Keap1*<sup>-/-</sup> MEFs were subjected to 20  $\mu$ M LY294002 or 10  $\mu$ M PHAR as indicated. A, representative immunoblots of NRF2, phospho-Ser473AKT (pSer473AKT), AKT, phospho-Ser9GSK3 $\beta$  (pGSK-3), GSK3 and VCL as a loading control. B, densitometric quantification of NRF2 protein levels from representative immunoblots from A, expressed as a ratio of NRF2/VCL.



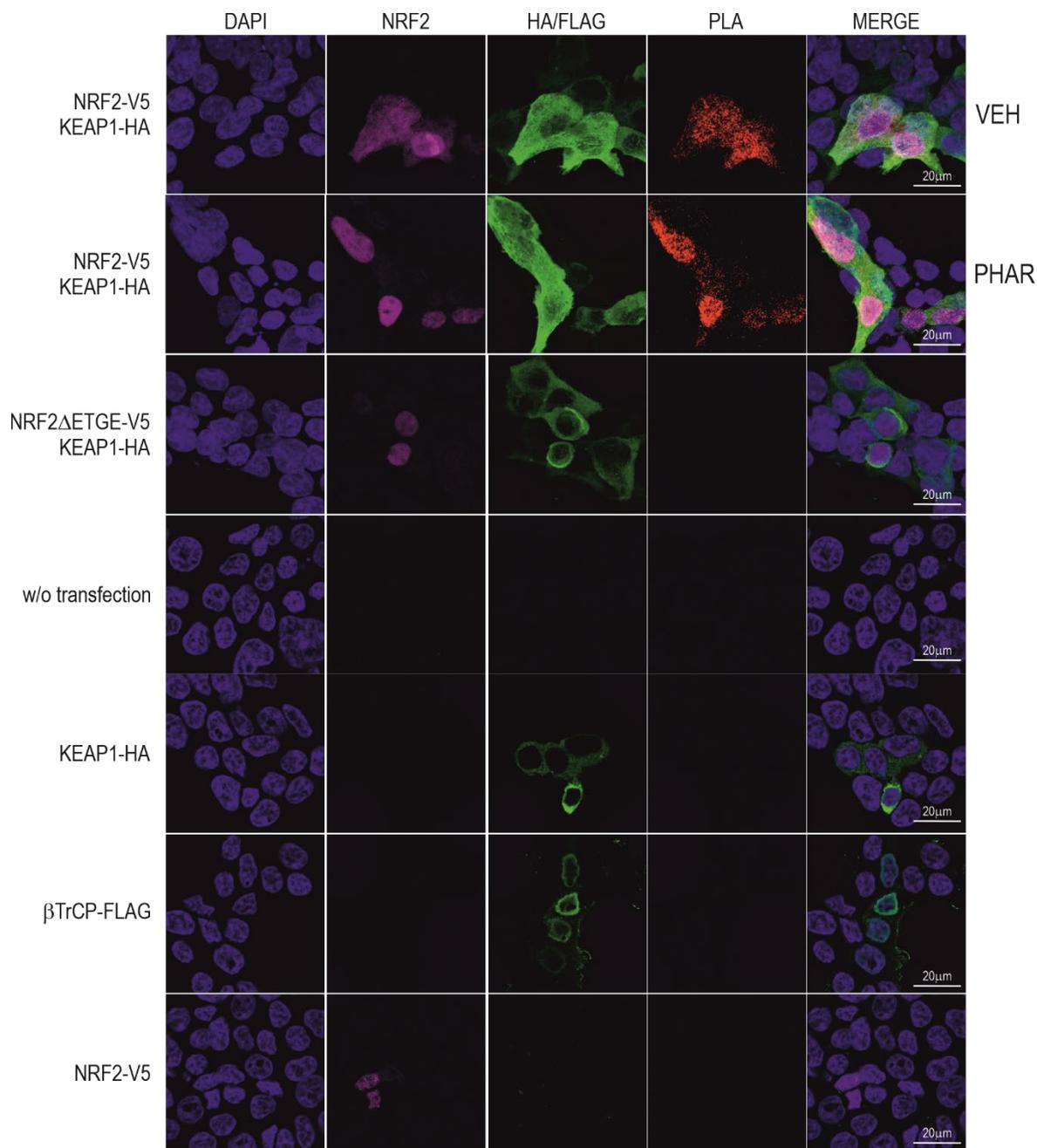
**Figure S2.** GSK-3 inhibition abolishes further NRF2 accumulation in response to PHAR. *Keap1*<sup>-/-</sup> MEFs were serum-depleted for 16 h and then subjected to 10  $\mu$ M PHAR, 10  $\mu$ M SB216763 (GSK-3 inhibitor) or to both treatments for the indicated times. A, representative immunoblots of NRF2,  $\beta$ -Catenin and VCL as loading control. B, densitometric quantification of NRF2 protein levels from representative immunoblots from A, expressed as a ratio of NRF2/VCL. Data are mean  $\pm$  S.D. ( $n=3$ ). A Student t test did not detect statistically significant differences.



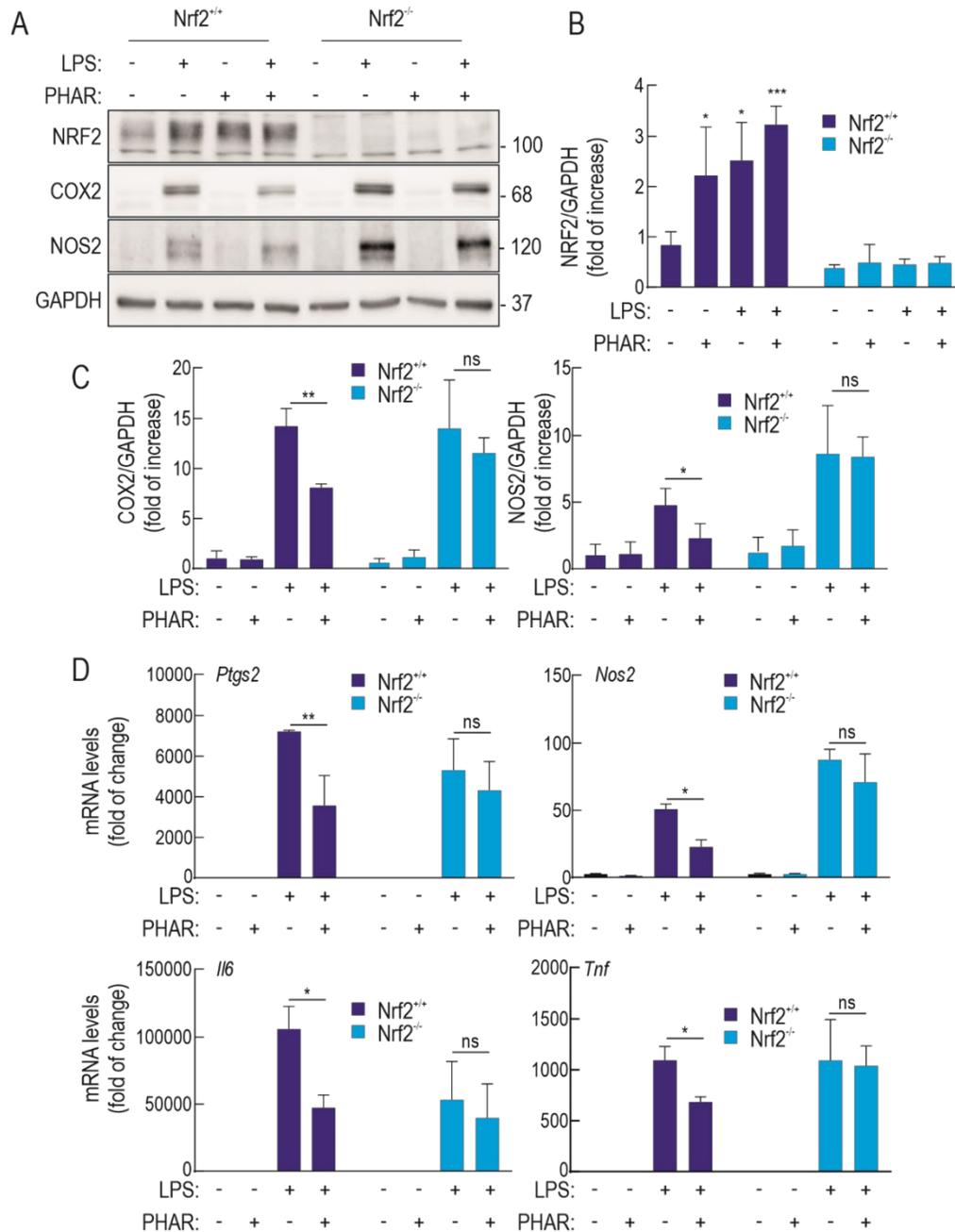
**Figure S3.** PHAR increases NRF2 protein levels in a  $\beta$ -TrCP dependent-manner. Keap1<sup>-/-</sup> MEFs were transduced with lentivirus encoding shCTRL or sh against mouse  $\beta$ -TrCP1/2. After 5 days, cells were serum-depleted for 16 h and then subjected to 10  $\mu$ M PHAR for 30 min. 0.1% DMSO was used as vehicle. Then, cells were treated with 20  $\mu$ M LY294002 for the indicated times. A and C, representative immunoblots of NRF2, pSer473-AKT, AKT, pSer9-GSK3 $\beta$ , GSK3, and VCL as loading control from shCTRL and sh  $\beta$ -TrCP1/2. B-D, densitometric quantification of NRF2 protein levels from representative immunoblots from A and C, expressed as a ratio of NRF2/GAPDH. Data are mean  $\pm$  S.D. ( $n=4$ ). \* $p<0.01$  vs LY294002 according to a two-way ANOVA test.



**Figure S4.** Pevonedistat (MLN-4924) promotes the accumulation of NRF2 through NEDD8-activating enzyme (NAE) inhibition. **A**, Mechanism of action of pevonedistat. Pevonedistat (MLN-4924) is a NAE inhibitor that selectively prevents the activation of cullin-based ring-ubiquitin ligases (CRLs), including CUL1 (connected with  $\beta$ -TrCP) and CUL3 (connected with KEAP1), and alters the ubiquitination and proteasomal degradation of cellular proteins, NRF2 in our case, causing its accumulation in the non-ubiquitinated form. **B**, HEK293T were serum-depleted for 16h and then subjected to 1  $\mu$ M MLN-4924 or 20  $\mu$ M MG132 for the indicated time. Representative immunoblots of NRF2,  $\beta$ -CATENIN, and VCL as a loading control.



**Figure S5.** Proximity ligation assay (PLA) for KEAP1 and NRF2 in HEK293T cells. HEK293T cells were transfected with expression vectors for NRF2-V5, NRF2 $\Delta$ ETGE-V5, HA-KEAP1 and  $\beta$ -TrCP-FLAG for 24h. Then, cells were incubated in the presence vehicle (VEH) or PHAR (10 $\mu$ M) for 6 h. Cells were subjected to the PLA assay using rabbit anti-FLAG/anti-HA and mouse anti-V5. Fluorescent images were captured, and red puncta represent co-localization of KEAP1 and NRF2. To confirm the validity of the assay, we used as negative control ectopic expression of NRF2 $\Delta$ ETGE-V5 which lacks the high affinity binding site for KEAP1. Calibration bar = 20  $\mu$ m. When no antibody was present or when only one plasmid was present (KEAP1-HA only,  $\beta$ -TrCP-FLAG only or NRF2-V5 only), no red fluorescent puncta were detected.



**Figure S6.** PHAR decreases the inflammatory response in peritoneal macrophages derived from wild type but not from Nrf2-knockout mice. Serum-depleted peritoneal macrophages were pre-treated with 10  $\mu$ M PHAR for 8h. Then, cells were treated with 100 ng/ml LPS for the indicated times. A, representative immunoblots of NRF2, COX2, NOS2, and GAPDH as a loading control. B-C, densitometric analysis of NRF2, COX2, and NOS2 protein levels from representative immunoblots from (A), expressed as a ratio of protein levels/GAPDH. Data are mean  $\pm$  S.D. ( $n=3$ ). \* $p<0.05$ ; \*\* $p<0.01$ ; \*\*\* $p<0.001$  vs. vehicle or LPS treatment, as indicated, according to a one-way ANOVA test. C, transcript levels of *Il1b*, *Ptg2*, *Nos2*, *Il6*, and *Tnf* were determined by qRT-PCR and normalized by the average of *Gapdh*, *Tbp*, and *Actb*. Data are mean  $\pm$  S.D. ( $n=3$ ). \* $p<0.05$ ; \*\* $p<0.01$ ; \*\*\* $p<0.001$  vs. LPS according to a one-way ANOVA test.