

Loss of EHF promotes the development of treatment-induced neuroendocrine prostate cancer

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I. Antibody Information

Antibody	Vendor	Catalogue Number	Application	Dilution
AR	Proteintech	22089-1-AP	WB	1:1000
EHF/ESE-3	Proteintech	67125-1-Ig	WB	1:1000
	Abcam	Ab272671	IHC	1:200
ENO2	Proteintech	10149-1-AP	WB	1:1000
			IHC	1:200
CHGA	Proteintech	60135-1-Ig	WB	1:2000
H3K27me3	Omnimabs	OM256819	WB	1:1000
Histone H3	CST	4499	WB	1:1000
EZH2	Proteintech	21800-1-AP	WB	1:1000
SOX2	Abcam	Ab97959	WB	1:1000
			IHC	1:200
Vinculin	Sigma Aldrich	V9131-2ML	WB	1:2000
Ki-67	Genetex	GTX16667	IHC	1:100
AR	CST	5153	ChIP	1:100
H3K9Ac	CST	9649	ChIP	1:50
Rabbit IgG	Invitrogen	10500C	ChIP	1:50/100

II. RNA and Plasmid Information

Reagent	Provider	Catalogue #
ON-TARGETplus Human AR siRNA	Dharmacon	L-003400-00-0005
ON-TARGETplus Human EZH2 siRNA	Dharmacon	L-004218-00-0005
ON-TARGETplus Non-targeting Pool	Dharmacon	D-001810-10-05
TRC Lentiviral Human EHF shRNA	Dharmacon	RHS4533-EG26298
pGL3-Promoter Vector	Promega	E1761
pCMV-hAR	Addgene; pCMV-hAR was a gift from Elizabeth Wilson	89078

III. Primers for qPCR

Primer name	Forward Primer Sequence (5'-3')	Reverse Primer Sequence (5'-3')
CHGA	TAAAGGGGATACCGAGGTGATG	TCGGAGTGTCTCAAAACATTCC
ENO2	CTGTGGTGGAGCAAGAGAAA	ACACCCAGGATGGCATTG
PSA	AGTGCGAGAAGCATTCCCAAC	CCAGCAAGATCACGCTTTTGTT
SLIT2	CGGAGCAGCAAGCTAAAGAA	GCGACAGGGACAGCATCT

DAB2IP	CTGAGCGGGATAAGTGGATGG	AAACATTGTCCGTCTTGAGCT T
ADRB2	TTCTTGCTGGCACCCAATA	GCCAGGACGATGAGAGACAT
GAPDH	GGACCTGACCTGCCGTCTAGAA	GGTGTCGCTGTTGAAGTCAGAG
ARE1 (ChIP-Seq)	GAAGCCTGGCATGTTGTACTTG	GGTAGGAACAATTGGAGGGAA GT
ARE2 (ChIP-Seq)	TGGAGAACTGGCCCATATTGTT	TGAACCAGTGCCAAATCTGTCT
PSA (ChIP-Seq)	CCTAGATGAAGTCTCCATGAGCTA CA	GGGAGGGAGAGCTAGCACTT G

IV. Supplementary Figures

Supplementary Figure 1

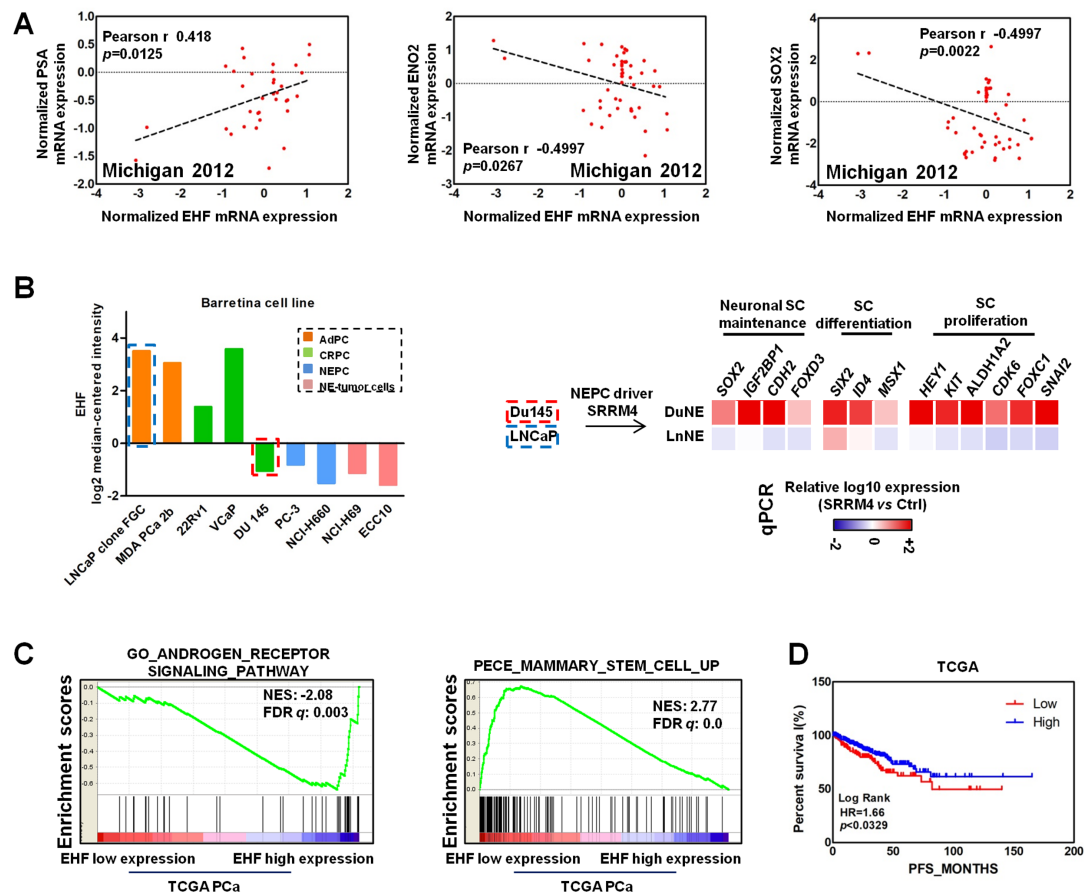


Figure. S1. (A) Pearson correlation between EHF and PSA, EHF and ENO2, and EHF and SOX2 expressions was determined using transcriptomic data of Michigan 2012 cohort¹. **(B)** EHF expression in different cancer cell lines from Barretina *et al.* 2016². Results are presented as mean $\pm$ SD (left). The mRNA expression of the genes with functions on neuronal stem cell maintenance, stem cell differentiation and stem cell proliferation in the SRRM4-overexpressing DU145 and LNCaP cells was compared to that of their respective control cells via qPCR. Heat maps represent the relative fold change in log10. **(C)** GSEA analysis indicated that PCa tumors with low EHF expression in TCGA clinical prostate cancer cohort³ had similar characteristics with t-NEPC in alleviating the dependency on AR signaling and gaining stem-cell features. **(D)** Kaplan-Meier analysis indicated that PCa tumors with low EHF

expression had a poor overall survival. Note: AdPC, prostate adenocarcinoma; CRPC, castration-resistant prostate cancer; NEPC, neuroendocrine prostate cancer; NE, neuroendocrine; SC, stem cell; PCa, prostate cancer; NES, Normalized enrichment score; FDR, false discovery rate; PFS, progression-free survival; HR, hazard rate.

Supplementary Figure 2

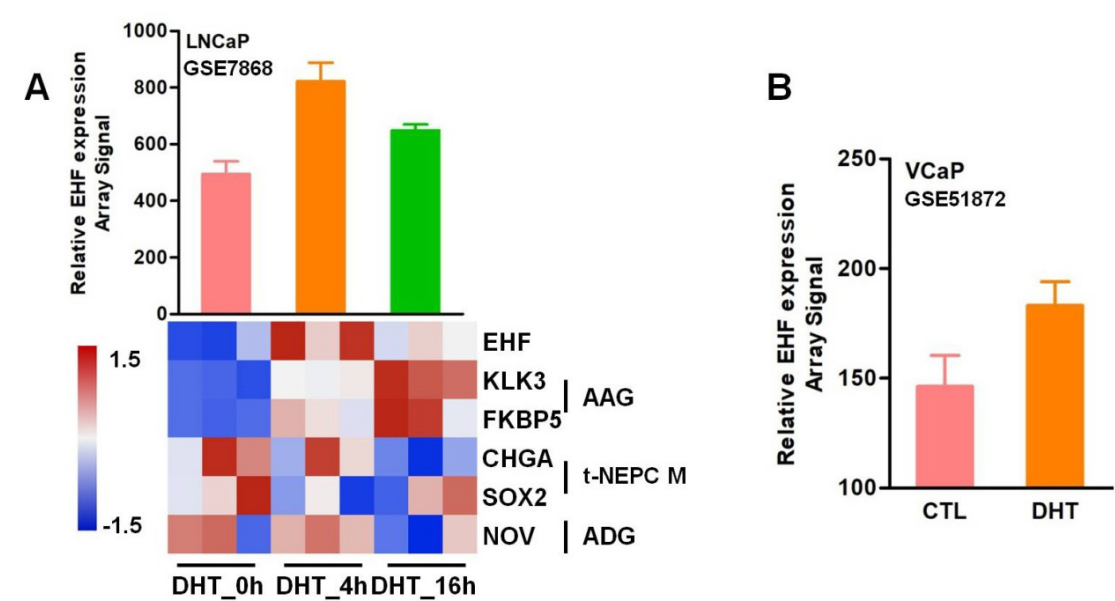


Figure. S2. (A) Bar graph showing EHF expression (Top) and heatmap of AR-associated genes including EHF, t-NEPC markers and an androgen-depressed gene in DHT treated LNCaP cells (GSE7868). (B) Bar graph showing EHF expression in DHT or ENZ treat VCaP cells (GSE51872). Bar graphs show means $\pm$ SD. Note: AAG, AR-associated genes; M, markers; ADG, androgen-depressed gene; AD, androgen deprivation; CTL, control; DHT, double hydrogen testosterone.

Supplementary Figure 3

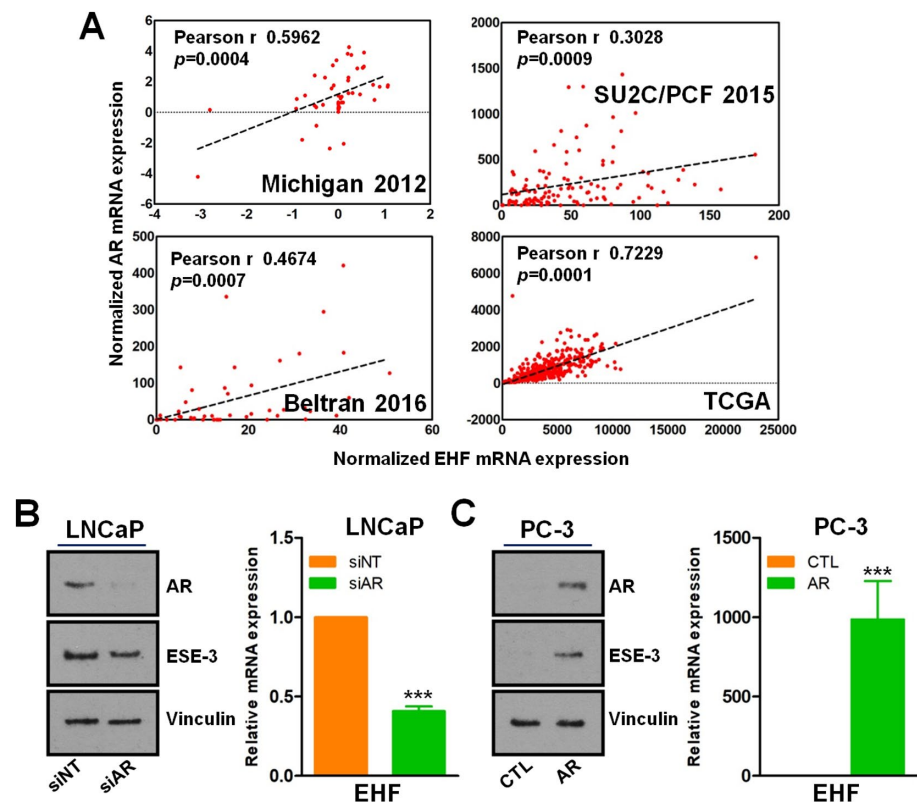


Figure. S3. (A) EHF and AR are positive correlated in Michigan 2012¹, SU2C/PCF 2015⁴, Beltran 2016⁵ and TCGA³ clinical prostate cancer cohort. **(B)** AR and EHF levels in LNCaP cells with/without AR knockdown were measured by qPCR and immunoblotting. **(C)** AR and EHF levels in PC-3 cells with/without AR overexpression were measured by qPCR and immunoblotting. The two-tailed Student's t-test was used to compare results between two groups with *** denoting $p < 0.001$. Bar graphs show means $\pm$ SD. Note: TCGA, the cancer genome atlas; CTL, control; NT, non-target.

Supplementary Figure 4

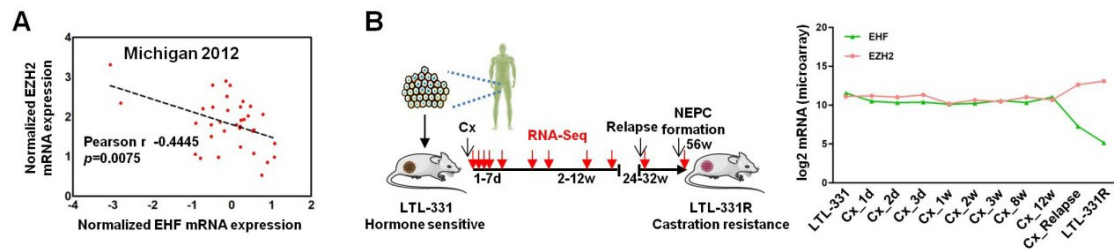


Figure. S4. (A) Pearson correlation between EHF and EZH2 was determined using transcriptomic data of Michigan 2012 cohort¹. (B) EHF and EZH2 mRNA expressions during progression of AdPC (LTL-331) to t-NEPC (LTL-331R) by castration surgery to the host mice⁶ were plotted. Note: Cx, castration.

Supplementary Figure 5

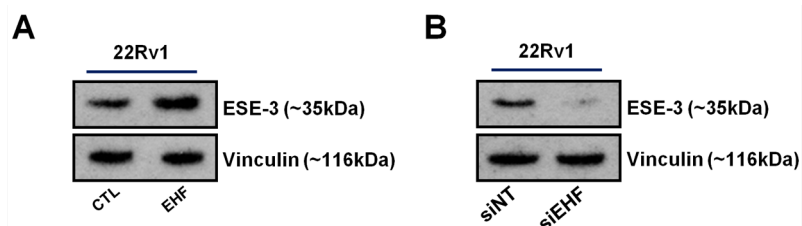


Figure. S5. (A) ESE-3 expression level in 22Rv1 cells with/without EHF overexpression was measured by immunoblotting. (B) ESE-3 expression level in 22Rv1 cells with/without EHF knockdown was measured by immunoblotting.

Supplementary Figure 6

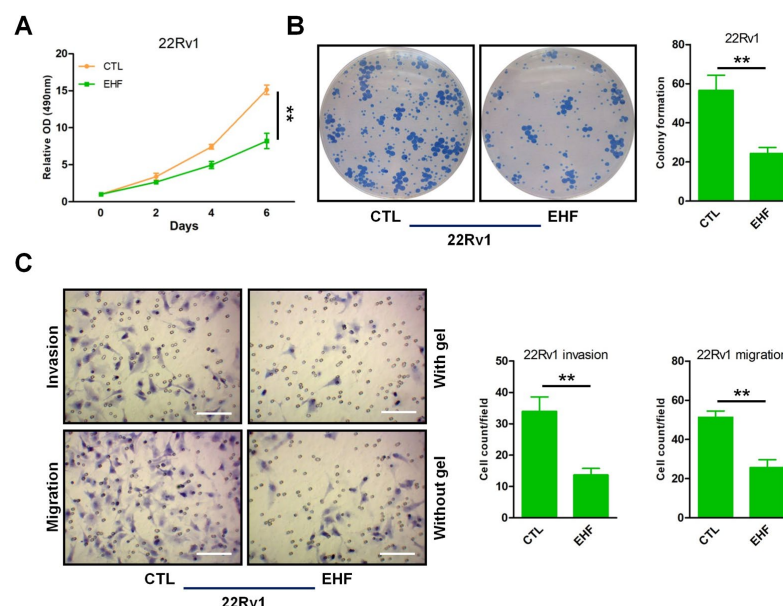


Figure. S6. (A) MTS assays measured cell proliferation of 22Rv1 cells with/without EHF

overexpression. **(B)** Same cells as in **A** were used to perform colony formation assays. Representative images are shown. **(C)** Same cells as in **A** were used to perform cell invasion and migration assays in transwell chambers with/without matrigel coated. Representative images are shown. Scale bar 100 μ m. Two-tailed Student's t-test was used to compare results between two groups with ** denoting $p < 0.01$. Experiments were performed with three biologically independent samples. Bar graphs show means $\pm$ SD. Note: OD, optical density.

Supplementary Figure 7

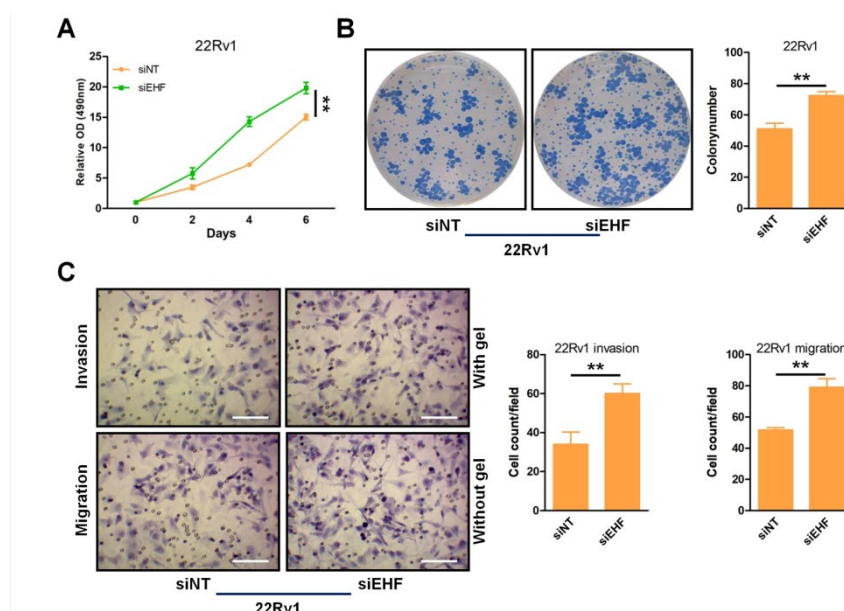


Figure. S7. (A) MTS assays measured cell proliferation of 22Rv1 cells with/without EHF knockdown. **(B)** Same cells as in **A** were used to perform colony formation assays. Representative images are shown. **(C)** Same cells as in **A** were used to perform cell invasion and migration assays in transwell chambers with/without matrigel coated. Representative images are shown. Scale bar 100 μ m. Two-tailed Student's t-test was used to compare results between two groups with ** denoting $p < 0.01$. Experiments were performed with three biologically independent samples. Bar graphs show means $\pm$ SD. Note: OD, optical density.

V. References

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