Targeting the Nuclear Protein Export Reverses Epithelial to Mesenchymal Transition

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Supplemental information

Supplementary Figure 1



Supplementary Figure 1. Leptomycin B treatment causes reversal of mesenchymal phenotype to epithelial. HMLE-Snail cells were grown in 100 mm petri plates overnight. After 24 hrs the media was aspirated and the cells were washed with PBS followed by incubation with LMB (300 nM) for 3 hrs.



Supplementary Figure 2. siRNA silencing of CRM1 or snail reverses EMT. 50,000 HMLE-Snail cells grown in six well plates in duplicate and were exposed to either control siRNA, CRM1 siRNA or Snail siRNA for 72 hrs. At the end of the siRNA treatment cells were trypsinized and re-seeded and again treated with siRNA for 72 hrs. The cells were photographed at 40X magnification using an EVOS microscope system. Images representative of two independent experiments.



Supp Fig3. SINE do not induce apoptosis at early time points in HMECs. HMLE-Snail and HMLER-Snail seeded at a density of 50,000 cell per well in six well plate were grown overnight. After 24 hrs the cells were exposed to either DMSO or 1 micro M concentrations of SINE (KPT-185 or Selinexor KPT-330) for 24 hrs. At the end of the reaction period, the cells were processed for Annexin V FITC apoptosis analysis using Biovision apoptosis assay kit according to manufacturers protocol. Note: no significant apoptosis is observed at 24 hrs (early) time point. **[B] SINE do not induce apoptosis in MCF-10A NeoT normal cells.** MCF10A-NeoT cells were grown at a density of 50,000 cells per well and exposed to selinexor at 150 nM for 72 hrs. Annexin V FITC assay was performed using Biovision Apoptosis analysis kit according to manufacturer's protocol.

Supplementary Figure 3



Supplementary Figure 4. Selinexor (KPT-330) suppresses spheroids in HMLE cellular models. HMLE-snail or HMLER-snail spheroids were exposed to selinexor (150 nM) once a week for two weeks. The spheroids were counted under an inverted microscope. See methods section for details on the experimental procedure. **Supplemental Figure 4**



Supplementary Figure 5. SINE suppress colony formation in HMEC derived cells. Clonogenic Assay: 50,000 cells seeded in six well plates and allowed to grow for 24 hrs. Once attached, the cells were exposed to XPO1 inhibitors (0-150 nM) for 72 hrs. At the end of the treatment period, 1,000 cells were taken from each reaction well and re-seeded in 100 mm petri dish and allowed to grow for 2 weeks at 37 °C in a 5% $CO_2/5\% O_2/90\% N_2$ incubator. Colonies were stained with 2% crystal violet, counted, and quantitated. SINE suppress colony formation in HMEC derived cells.



Figure 6. SiRNA silencing of [A] CRM1 or [B] snail enhances the expression of Ecadherin in HMLE-snail cells. HMLE-snail cells were exposed to either CRM1 or snail siRNA and RNA was isolated followed by RT-PCR analysis for E-cadherin expression. Graph representative of two independent experiments.



Supplementary Figure 7. [A] LMB induces nuclear localization of FBXL5 in MCF-7 breast cancer cells. Cells were grown in 4 well chambered slides at a density of 3000 cells per well. The next day the cells were exposed to 150 nM LMB for further 24 hrs and immunofluorescence was performed using FBXL5 antibody. **[B]** Western blotting of HMLE-Snail exposed to either untreated, control siRNA or FBXL5 siRNA in the presence or absence of selinexor 150 nM. Note: lack of snail reduction upon FBXL5 siRNA treatment Note. Nuclear retention of FBXL5 in treated group. Figure representative of three independent experiment.

Supplementary Table 1. FBOX family genes are differentially expressed in SINE treated cells and fall under the canonical biologically meaningful pathways

		Focus Molecule	
Network	Molecules in Network	S	Top Functions
	ABAT,ATP2B2,BBS9,CASK,CTH,Cytokeratin,DDIT3,DDX56,ERF FBXO33, IES4,IPO5,KIF5C,KRT7,KRT19,KRT6A,Laminin1,LARGE,MT1B,NELF,PCM1,P		
1	mca,POLR1B,PSAT1,Ribosomal 40s subunit,Rnr,RPL13,RPL37, RPS24,RFS15A,RPS3A,RPS6KA,RPS6KA,RPS1,TRIB3	28	Gene Expression, Protein Synthesis, Cancer
	ABCC1,ABCD3,Actin,AIMP2,ANXA3,ATPase,BCCIP,Cbp/p300,CCNK,E2F5,EIF6,EPAS1,FAM60A,GRN,GTF2A1,HAUS6,HIRIP3,HIST1H2BJ/HIST1H2BK,H		DNA Replication, Recombination, and Repair, Energy Production, Nucleic Acid
2	olo RNA polymerase II, HSPA4L, KIFC2, NHP2L1, NMI, NUDT5, P-TEFb, Rac, RNF213, RUVBL1, SNRNP48, snRNP, Sos, TAF9, TUFT1, tyrosine kinase, ZCRB1	26	Metabolism
0	ANGP1L4, BMP10, BMP, BNC2, C80rd33, CLGN, EDA, FS1, FUCA1, GF12, GLA, GLHX, IgG2a, IgG2b, IL13HA2, IL18BP, IL18H1, IL23A, INF2, IHAK2, IHAK, IHF5, ITM DD Mae Jacobi NEV (A complexity DPC/FS DD JAA DI)// C4 CAA SECO A C1 C1140, Cared Seco A DD JAA SECO A DI C114	00	Inflammatory Response, Organismal Injury and Abnormalities, Cellular Function
3	2B, MIRC Class II, NYKK (COMPLEX, FUSKS), FYY4H4, HIFK4, SAA, SEHCA, SUC TIAZ, SMBA, SYYLZA, TIKAF, ZHANDS PDNE Collemativilia CASPA conserved CECCU CLUCA CLUB CONJUNA DNA ICS DNA ICCA DNA ICC DNA ICC DNA ICCA STRATCH	26	and Maintenance
	DDINF, Calmodulini, CAST-3, Caspase, CLEC40, CLIC4, CLIC4, CUNNINH, DINAUC6, DINAUC9, DINAUC 15, DINAUC, DTINC 112, ESITOgeni recentre FEMI B. COXMIL HINBRD II Lengy Tengy Hengy Hengy Hengy Hengy Hengy (Hengy Hengy Hengy Hengy Hengy Hengy		Cell Death and Survival Nervous System Development and Eurotion. Cellular
4	Sk/ Prinsulin RNF103 S100A4 SEZ61 Sod TIMUTA UNCES VEGEA	23	Movement
	Alp,ALT,BRD2,CCDC71,CDKN1A,Ck2,COG1,COG5,DDB2,DKK1,DPAGT1,DZIP1,FCRLA,GOT,Histone		
	h3,lgG1,lgG,lgm,lmmunoglobulin,lnsulin,KAT2B,MTOR,NFE2,PBX3,PFKP,PHEX,RAB17,RNA polymerase II,SAP130,SIGLEC5,SIX1,STAT5a/b,Tnf		
5	(family),TYMS,UBQLN1	22	Developmental Disorder, Hereditary Disorder, Metabolic Disease
	26s Proteasome,Alpha tubulin,ANKRD28,Beta Tubulin,CALCOCO2,CCDC28A,CCT3,CCT5,CLDN3,ELMOD3,F		
	Actin, FAM90A1, HDAC6, HIPK2, HISTONE, lkb, LZTS1, MGEA5, NEXN, NFKBIB, NFKBIE, PCDH18, Pka, PP2A, PRKAR1B, RASGRP2, RUNX2, Smad1/5/8, SYNE2, Next,		Cardiovascular Disease, Cell-To-Cell Signaling and Interaction, Cellular
6	TAGLN, Thf receptor, TPPP, Troponin t, Tubulin, Ubiquitin	22	Movement
	APOBEC38, BCL2A1, Caveolin, CES1, CLU, CLUAP1, endothelin receptor, Ferritin, Fibrinogen, FIP1L1, FOS, FOSB, FIH1, FIL, glutathione		Oslindar Davidsana at Oslindar Ossuth and Davidsantian Ossuration Tissue
7	transferase, us 1, us 1A4, us 1M4, nemoglobin, HMOX 1, IFT1 M1, IL 11, Integrin alpha V beta 3, Lon, LDL, MLF1, NAB1, NAB2, NADPH	01	Development, Cellular Growth and Proliferation, Connective Tissue
/	20-alpha-bydroxyorogesterone AES APP ATP282 BCAS1 beta-estradial CC2D1A CCDC88B CGB (includes	21	
	others). CLDN3.DNAH14.DNAH15.ECH1.GJB2.HRK.HSD17B2.ITGBL1.NCAPG.NUDT9.OSBPL5.PPM1K,PPP1R14A.progesterone.PTGFR.RASGRP2.RER1.		Reproductive System Disease, Cell Death and Survival, Nervous System
8	RYR3,SERPINA6,SH3GL3,SIX4,SLC30A3,SLC7A2,SPAG9,SULT1E1,TCF15	22	Development and Function
	ABCA7,Akt,atrial natriuretic peptide,CAV3,creatine kinase,CREG1,CTC1,cyclooxygenase,Foxo,FXR ligand-FXR-Retinoic acid-RXRq,Growth		
	hormone,HDL,HDL-cholesterol,Kallikrein,LDL-		
0	Cholesterol,LPXN,MYL9,Nos,NH4A3,OBFC1,PCSK9,PCYOX1,PDK4,Ppp2c,PHKAR2B,PVHL2,SCAHB1,SEC14L2,SGMS1,SH2B3,SLC9A3H1,SOA12,SYN	01	Linid Matchaliam, Malagular Transport, Small Malagula Bioghamiatry
9	NI, VEDE-GIOISSIEIO, VEDE-GIOISSIEIO, VEDE-GIONA DE CONTRA CARACTERIZZA CONTRA CO	21	Lipid Metabolishi, Molecular Transport, Shian Molecule Biochemistry
	recentor, GTPase I, APTM5, I PAR1, MRPI 20, MTORC2, NPR3, OAZ1, Pkc(s), Pko, PLC, Plc		
10	beta,PLCB2,RGS4,RHEB,SLC30A3,UBR2,ULK2,USP13,USP47,USP48	20	Carbohydrate Metabolism, Molecular Transport, Small Molecule Biochemistry
	Ap1,ATXN10,BAMBI,BTG2,calpain,CaMKII,CCNE1,CENPI,Cg,Creb,EDN1,FJX1,FSH,GLS,Gsk3,Histone		
	h4,HMBS,HSD3B7,IL1,JPH4,Lh,LRP,MCM7,NOS1AP,Notch,NRARP,PDGF BB,PDLIM3,PPARG,Pro-inflammatory		Organismal Functions, Cell Morphology, Renal and Urological System
11	Cytokine,PTX3,SLC7A7,SOCS2,SRH1,Igt beta	20	Development and Function
	AMIRA, ASNS, AT POVICI, C-STC, C022, ODH18, C04K, Cyclin A, Cyclin D, Cyclin E, Cytochrome C, E21, FAHD I, FANG, HMG C0A		
12		19	Cancer, Reproductive System Disease, Carbohydrate Metabolism
	AFAP112,ARHGDI,ATF3,0HRNA7,Collagen		
	Alpha1, Eotaxin, Ern FBX02 Fcer1, Fgf, FGFR3, Fgfr, FRMD3, Gap, GJB2, Hedgehog, HES6, HHIP, Hspg, IGSF8, IL4R, JAK, MGAT5, MMP1, MT2A, Nicotinic		Cellular Growth and Proliferation, Cell Signaling, Connective Tissue Development
13	acetylcholine recepter, Datailed, PI3K (complex), PLC gamma, PRRX1, STAT1/3/5 dimer, SYK/ZAP, Tenascin, TIMP3, TWIST1	18	and Function
	ARHGEF39,C2orf43,COMMD1,DPY19L1,DVL2,ELAVL1,EMC7,FAM136A,GINM1,HKDC1,KCTD14,KDM5B,KIAA0922,LMNA,LRRC37B,LSM1,MIOS,PKD1,		Cellular Assembly and Organization, Organ Morphology, Reproductive System
14	RAELSCONDJSLC22A23,SMARCA4,SUCO,SYVN1,IMEM38B,UBC,VSIG10	16	Development and Function
	14-3-3, ABLIMIN, ADAMITST, ARO, DOLZETT, BOR (COMPLEX), BEGIN, Celonieum Protein(S), OD302, ODA, DERSS, ECH, EGRT,		Cellular Compromise, Organismal Injury and Abnormalities, Cell-mediated
15	(family) OSB1. RABIP. Baf. Bas. RUIX111: Sapt. TCR. TXNRD2	20	Immune Response
	Adaptor protein 1,Caspase 3/7,CFL1,CLCN5,Cofilin,FGD4,GCNT1,GKAP1,Gm-		
	csf,GPS2,HNRNPA3,IRS,Jnk,JUN/JUNB/JUND,KCNJ15,LAMP3,LHX1,Mek,Mlc,MSI2,Myosin,NRG (family),PAK6,Pak,PP1 protein complex		
16	group,RBM47,Rock,RPS6K42,Rsk,SESN2,SHANK1,SMPD4,sphingomyelinase,ST6GAL1,TESK2	19	Cellular Assembly and Organization, Hereditary Disorder, Neurological Disease
	BHLHE40,C/ebp,CIITA,CXCL14,CXCH3,EIF3,EIF3,EIF4G3,elastase,HEXB,HLA-DR,HLA-DRB5,HSP,IFI44L,IFIT1,Ifn,IFN alpha/beta,IFN Beta,IFN gamma,IFN type		Outline Exaction and Maintenance Hamatala sized Outland Development and
17	1, IFINAC1, IQE, LL2 (COMPLEX), IL22FA1, IMBHERON AIPTA, MINO CIASS II (COMPLEX), INIKD1-REIA, P36 MAPA, PCDF1, FISA (familu) D1 SCF113 SCFDINBA TIr THESEA TEIPIO	17	Function, Humoral Immune Response
17	ABCA6 AOX1 BATF2.BATF3.BATF3.CL2A1.CACNB3.Cbo/p300-Maf-	17	r unction, numeral minune nesponse
	Nfe2l2,CEBPD,CYB561,EIF2AK4,FGF1,GCLM,GSTA4,GSTA5,GSTP1,HCP5,IFNG,IKZF2,IL11,KRTAP2-3 (includes others),neuroprotectin		
18	D1,NFE2L2,NFE2L3,PRICKLE1,RAB34,SHISA5,SLC14A , 3LS54H9 ,SLC6A1,SLC7A11,SMAD4,TBXAS1,TCTA,TRIM33,UGT2B15	17	Drug Metabolism, Protein Synthesis, Cellular Assembly and Organization
10	BABAM1,BRD4,CLN3,COMMD1,COMMD7,COPS6,DIS3 2,FBXL17 EZ1,G3BP1,HERC4,HERC6,KBTBD7,MLH1,MRPL45,MYADM,NME2,PDX1,PRRC1,		
19	RADI8, KTN4, SELTU3, SELRU1, SFAN4, SLU25A23, SPRIM, TAGENSILUT, IRMI16, IRMI161A, IRMI11, IUBB3, IUBB4A, UBC, ZNF14	16	Cancer, Hereditary Disorder, Renai and Urological Disease
20	ANNA, ANTIGO, ALL, ATIWO, OLTIN,	16	Carbohydrate Metabolism, Small Molecule Biochemistry, Hereditary Disorder
20	ACADSB,AP4B1,AP4M1,AP4S1,CHST12,CLASP1,CYP4F3,CYP4F11,DDX52,GCLM,GFM1,H2AFV,LRBA,LSR,MCTS1,MEF2B,PIGH,PIGQ,PIGY,PPHLN1,	-	
21	PSMF1,PYCR1,RAB10,RPL17,SEPT8,SLC35B2,SRSF7,STOX1,TBC1D7,TGOLN2,UBC,UQCR10,UQCR11,UQCRB,WDR41	16	Hereditary Disorder, Neurological Disease, Metabolic Disease
	ADCK1,ANXA8L2 (includes		
00	omers), ARHIGETZ6, ASFM, OTTOM96, CHMIL, CYBS61, DARG, ESMT, FLT3LG, GAS1, GCN11, GLT1, GPEH, HSPB2, hydroxyproline, ICAM1, ILT7F, Integrin alpha V bata 2, MANIG1, NEED (VIGL 4) DI IM2 REV. DI OD1 PARTA PAREA DI GES OLDO COS CUI CADA TOCTO TU DE STUTUIO DI OD1	16	Cell-To-Cell Signaling and Interaction, Cellular Movement, Hematological System
22	Ueta SJWANTO LIJANE F, O VOL J, F ULIWI, FNIG, FLUD I, HAD IA, HADOA, HIV 192, SI FNS, SOUS, SLU I SAS, I GFDI, I LHS, I MEMIUU, I HIVST I 38 ADAM32 CI JIAFZA CEPIA CI DNI DHX8 DNA IC30 F XOSC2 FAMIATR GATAA GCK GOTI GPR3 GENPR HNFIA HNFAA HPV HSD1782 HSDAAL I EBS	10	Development and Function
23	ITH4, KIAA0101, LAS1L, MLF1IP, NUF2, REPIN1, RPS25, RRP8, SLC26A1, SLC37A4, SLC38A4, SPCS3, TBX18, UG72815, ZNF443, ZNF576	16	Carbohydrate Metabolism, Molecular Transport, Small Molecule Biochemistry
	ATM,CCND1,CCR6,CD3,CDX1,DEF6,DICER1,eIF2B,EPSTI1,FAM129A,FAM65B,ICOS,IL2,IL-2R,IL12RB1,IL18R1,IL23R,IL7R,KRT80,LINC00667,miR-19b-		- ,
	3p (and other miRNAs w/seed		Cell-mediated Immune Response, Cellular Development, Cellular Function and
24	GUGGAAA), MSLN, NANOSI, NEATI, PIK3R3, PRKAR2B, PTPN22, RNF139, SATB1, SIPA1L2, SYAP1, Terd, TMEM117, TMSB15A, TNFRSF4	16	Maintenance

Supplementary video 1. HMLE-snail cells were grown at a density of 1x10⁶ in 100 mm petri dish overnight. After 24 hrs the media was aspirated and the cells were washed with PBS followed by incubation with DMSO for 3 hrs with simultaneous live video recording at 40 X magnification under a EVOS FL microscope system.

Supplementary video 2. HMLE-snail cells were grown at a density of 1x10⁶ in 100 mm petri dish overnight. After 24 hrs the media was aspirated and the cells were washed with PBS followed by incubation with selinexor (KPT-330) for 3 hrs with simultaneous live video recording at 40 X magnification under a EVOS FL microscope system. **Supplementary video 3.** HMLE-snail cells were grown at a density of 1x10⁶ in 100 mm petri dish overnight. After 24 hrs the media was aspirated and the cells were washed with PBS followed by incubation with KPT-185 for 3 hrs with simultaneous live video recording at 40 X magnification under a EVOS FL microscope system. **Supplementary video 4.** HMLE-snail cells were grown at a density of 1x10⁶ in 100 mm petri dish overnight. After 24 hrs the media was aspirated and the cells were washed with PBS followed by incubation with KPT-185 for 3 hrs with simultaneous live video recording at 40 X magnification under a EVOS FL microscope system. **Supplementary video 4.** HMLE-snail cells were grown at a density of 1x10⁶ in 100 mm petri dish overnight. After 24 hrs the media was aspirated and the cells were washed with PBS followed by incubation with LMB (500 nM) for 3 hrs with simultaneous live video recording at 40 X magnification under a EVOS FL microscope system.