

Figure S5. mRNA and protein expression of FBLN2 after stable transfection with the FBLN2 expression vector.

S6

Cell adhesion molecules	ECM associated genes
Tight junction: JAM1, ZO1/2	Fibronectin, Vimentin, Elastin
Adherent junction: CDH1-3	Keratin
Desmosome: DSP, DSG2, DSC3, PKG	COL1A1, COL3A1, COL4A1
Gap junction: CX26, CX26	MMPs (MMP2, 7, 9, 14)
Hemidesmosome: ITGB4, ITGB1, ITGA6	TIMP1/2 (MMPs inhibitor)

Figure S6. Selected panel of cell adhesion molecules and ECM-associated genes for the expression analysis.

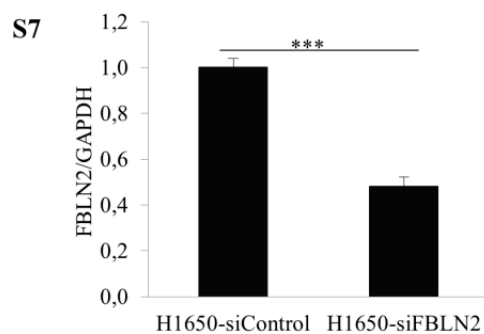


Figure S7. mRNA expression of FBLN2 after siRNA transient transfection in H1650 cells, mRNA expression level compared to GAPDH in cells transfected with siRNA-Control was set to 1.0. *** $p < 0.001$.

S8

	miRNA Name	Cat. No. (Qiagen)
1	miR-29a-3p	YP00204698
2	miR-29b-3p	YP00204679
3	miR-1-3p	YP00204344
4	miR-19a-3p	YP00205862
5	miR-199a-5p	YP00204494
6	miR-9-3p	YP00204620
7	miR-200b-3p	YP00206071
8	miR-155-5p	YP00204308
9	miR-125b-5p	YP00205713
10	U6 snRNA	YP00203907

Figure S8. The list of miRNAs analyzed in this study.

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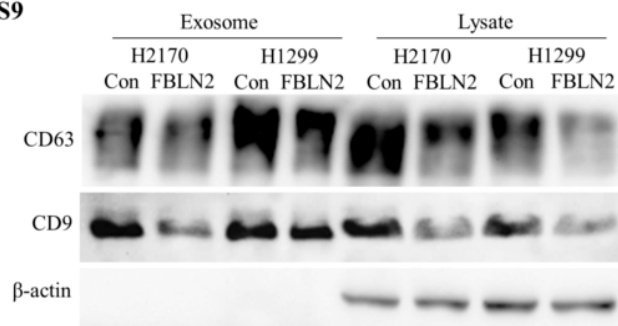


Figure S9. Western blotting of exosome markers CD63 and CD9 in both cell lysates and isolated exosomes of H2170 and H1299 cells. β -actin was used as loading control for proteins isolated from the cell lysates.

S10



Figure S10. (a) Schematic profile of the FBLN2 CpG islands (<http://dbcat.cgm.ntu.edu.tw/>). (b) Primers used for BS (black underlined) and MSP (green underlined) are indicated within the CpG island spanning from -280 to +384 bp (Chr.: 13565409-13566075).