

Figure S1. Oz cells exposed to hSC conditioned medium displayed a trend (yet statistically unsignificant) of increased Matrigel invasiveness. Micrographs and related diagrams show the increase of Matrigel invasion in Oz cells treated with hSCs CM respect to control CM. Student's t-test: p= 0,13. N=3

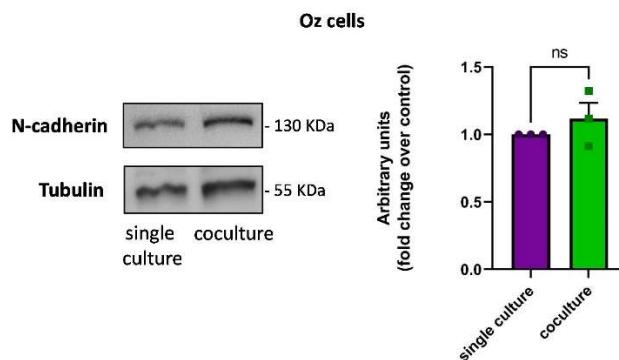


Figure S2. Western blot analyses of N-cadherin expression in Oz cells cocultured with hSCs. The figure shows an upregulation trend (yet statistically unsignificant) of mesenchymal marker N-cadherin in the coculture sample. The graph depicts the result of densitometric analyses. Student's t-test: p = 0.19; N=3.

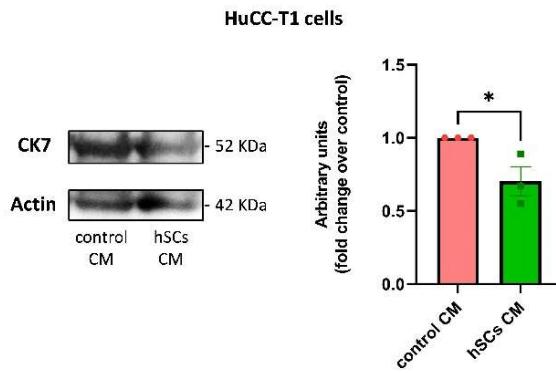


Figure S3. SCs CM-induced downregulation of cytokeratin 7 (CK7) in HuCC-T1 cells. Western blot analysis of CK7 expression in HuCC-T1 cells exposed to hSCs CM. HuCC-T1 cells treated with hSCs CM exhibited lower levels of cytokeratin 7 (CK7), compared to the control group. The graph depicts the result of densitometric analysis. Student's t-test: * $p \leq 0.05$. N=3.

	Accession	Protein description	Abundance Ratio: coculture / single culture	Abundance Ratio Adj. P-Value: coculture / single culture	number of detections in HuCC-T1 single culture	number of detections in HuCC-T1/hSCs cocultures
1	O14576-1	cytoplasmic dynein 1 intermediate chain 1	100	6,50E-16	0	4
2	O15239	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 1	100	6,50E-16	0	4
3	P07093-1	Glia-derived nexin	100	6,50E-16	0	3
4	Q02388-1	Collagen alpha-1(VII) chain	100	6,50E-16	0	3
5	Q68CP9-1	AT-rich interactive domain-containing protein 2	2,40	4,32E-02	4	4
6	P51991-2	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A3	2,19	1,81E-09	4	4
7	P11171-1	protein 4.1	2,16	3,55E-06	4	4
8	P62306	Small nuclear ribonucleoprotein F	2,05	2,18E-04	4	4
9	Q6IQ49-1	Protein SDE2 homolog	2,63	4,70E-02	1	3
10	Q96KC8	DnaJ homolog subfamily C member 1	2,34	1,10E-02	3	3
1	Q15828	Cystatin-M	0,01	6,50E-16	3	0
2	Q9UKQ2-1	Disintegrin and metalloproteinase domain-containing protein 28	0,01	6,50E-16	3	0
3	P67775	Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform	0,05	6,50E-16	4	3
4	P13929-1	Beta-enolase	0,39	8,50E-14	4	4
5	P43308	Translocon-associated protein subunit beta	0,41	3,58E-05	4	4
6	P53794	sodium/myo-inositol cotransporter	0,42	1,11E-09	4	4
7	P69905	Hemoglobin subunit alpha	0,43	6,50E-16	4	4
8	Q9UGUO	Transcription factor 20	0,45	1,03E-02	4	4
9	P32456	Guanylate-binding protein 2	0,45	1,33E-02	4	4
10	Q9UM47	Neurogenic locus notch homolog protein 3	0,50	7,93E-08	4	4

Table S1. Proteins differentially regulated in HuCC-T1 cells cocultured with hSCs identified by proteomic analysis. The table shows the complete list of modulated proteins in HuCC-T1/hSCs coculture respect to controls, with the respective accession numbers and p values. Red: upregulated proteins; Blue: downregulated proteins. T-test, Benjamini-corrected. N=4.