

*Article*

# Impact of secretion-active osteoblast-specific factor 2 in promoting progression and metastasis of head and neck cancer

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**Table S1 – Clinical and histopathological characteristics of microarray cohort**

Patient	Primary site	Age	Sex	TNM	Grade	HPV
1	hypopharynx	47	f	T2 N2 M0	2	neg
2	hypopharynx	46	m	T4 N1 M0	2	neg
3	hypopharynx	48	m	T3 N1 M0	2	neg
4	hypopharynx	58	m	T3 N2 M0	2	neg
5	hypopharynx	72	m	T3 N2 M0	2	neg
6	hypopharynx	57	m	T1 N2 M0	3	neg
7	hypopharynx	57	m	T2 N3 M0	2	neg
8	hypopharynx	47	m	T2 N2 M0	2	neg
9	hypopharynx	56	m	T2 N3 M0	2	neg
10	larynx	68	m	T4 N2 M0	2	neg
11	larynx	56	m	T4 N3 M0	3	neg
12	oropharynx	58	f	T3 N2 M0	2	neg
13	oropharynx	56	m	T3 N2 M0	2	neg
14	oropharynx	49	m	T3 N1 M0	2	neg
15	oropharynx	53	m	T2 N2 M0	2	neg

**Table S2 – Clinical characteristics of TCGA Head and Neck Cancer (HNSC) cohort (n=612).**

		<i>n</i>
Clinical T	T1	39
	T2	173
	T3	169
	T4a/b	203
Clinical N	N0	288
	N1	99
	N2a/b/c/-	180
	N3a/b/-	9
Clinical M	M0	571
	M1	5
Clinical UICC Stage <sup>1</sup>	Stage I	22
	Stage II	117
	Stage III	134
	Stage IV	311
Sex	Male	440
	Female	164

<sup>1</sup> 8th Edition

**Table S3. Top 30 list of significantly up-regulated genes in PTvs.N ( $\log_2\text{FC} > 1.5$ , p-value < 0.0001)**

No.	Gene	Protein	$\log_2\text{FC}$	p-value
1	<b>MMP1</b>	Matrix Metallopeptidase1	4,86	2,38E-09
2	<b>SPP1</b>	Osteopontin	3,27	2,01E-05
3	<b>POSTN</b>	Periostin, Osteoblast Specific Factor	3,06	9,40E-09
4	<b>COL1A1</b>	Collagen, Type I, Alpha 1	2,79	1,42E-06
5	<b>COL11A1</b>	Collagen, Type XI, Alpha 1	2,78	1,67E-05
6	<b>ASPN</b>	Asporin	2,73	6,72E-06
7	<b>INHBA</b>	Inhibin, Beta A	2,72	4,22E-07
8	<b>MMP3</b>	Matrix Metallopeptidase 3	2,71	2,00E-07
9	<b>COL5A2</b>	Collagen, Type V, Alpha 2	2,57	7,59E-07
10	<b>COL1A2</b>	Collagen, Type I, Alpha 2	2,56	1,22E-07
11	<b>MMP13</b>	Matrix Metallopeptidase 13	2,37	6,55E-07
12	<b>COL5A1</b>	Collagen, Type V, Alpha 1	2,23	5,68E-06
13	<b>CDH11</b>	Cadherin 11, Type 2 (OSF-4)	2,13	7,13E-05
14	<b>COL3A1</b>	Collagen, Type III, Alpha 1	2,11	1,19E-06
15	<b>LOX</b>	Lysyl Oxidase	1,98	2,32E-05
16	<b>LAMC2</b>	Laminin, Gamma 2	1,97	2,06E-05
17	<b>COL4A2</b>	Collagen, Type IV, Alpha 2	1,96	5,42E-08
18	<b>COL4A1</b>	Collagen, Type IV, Alpha 1	1,95	2,58E-08
19	<b>FN1</b>	Fibronectin 1	1,87	7,56E-06
20	<b>LOXL2</b>	Lysyl Oxidase-like 2	1,79	8,94E-08
21	<b>COL6A3</b>	Collagen, Type VI, Alpha 3	1,76	1,03E-06
22	<b>SULF1</b>	Sulfatase1	1,75	4,77E-05
23	<b>MAGEA4</b>	Melanoma Antigen Family A, 4	1,74	5,33E-06
24	<b>COL10A1</b>	Collagen, Type X, Alpha 1	1,71	3,33E-06
25	<b>SERPINE1</b>	Serpin Peptidase Inhibitor	1,70	9,23E-05
26	<b>FAP</b>	Fibroblast Activation Protein	1,67	9,01E-05

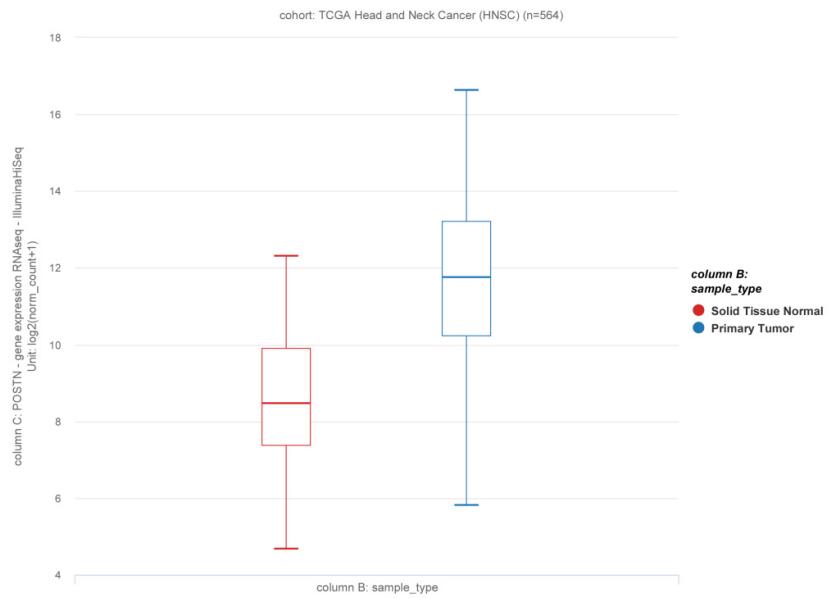
		Alpha		
27	<b>MFAP2</b>	Microfibrillar-Associated Protein 2	1,65	3,23E-07
28	<b>SPARC</b>	Osteonectin	1,64	3,77E-07
29	<b>IGFBP3</b>	Insulin-Like Growth Factor Binding Protein 3	1,64	6,06E-05
30	<b>PXDN</b>	Peroxidasin Homolog (Drosophila)	1,61	7,32E-06

**Table S4.** List of used primers for RT-PCR, quantitative Real-Time PCR, and cloning

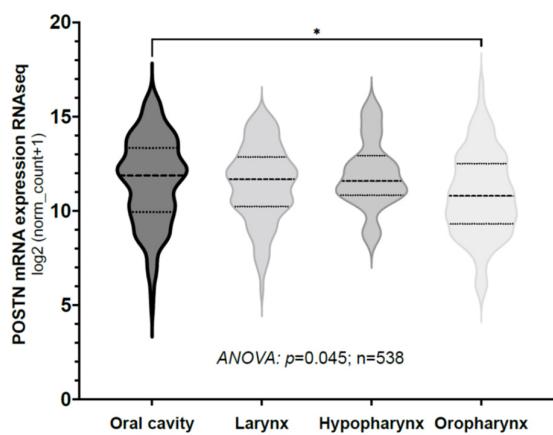
Primer	Sequence
OSF-2 RT PCR for	5`-ATTAGGCTTGGCATCTGCTC-3`
OSF-2 RT PCR rev	5`-CTCGCGGAATATGTGAATCG-3`
GAP-DH RT PCR for	5`-ACCACAGTCCATGCCATCAC-3`
GAP-DH RT PCR rev	5`-TCCACCACCCCTGTTGCTGTA-3`
OSF-2 for	5`-AAACCGCGGATGATTCCCTTTACCCATGTT-3`
OSF-2 rev	5`-AAAGCTAGCCTGAGAACGACCTCCCTTAATC-3`
OSF 2-ΔSec. for	5`-AAACCGCGGATGACAATCATTATGACAAGATCTG-3`
Secretion signal for	5`-CTAGCGCGTTTATAGGGTTAACATAAGCAGCAATAGTAG-3`
Secretion signal rev	5`-GGATGATTCCCTTTACCCATGTTCTACTATTGCTG-3`
RNA POL II for	5`-GCACCACGTCCAATGACAT-3`
RNA POL II rev	5`-GTGCGGCTGCTTCATAA-3`

**Table S5.** Sequence of recombinant OSF-2 (BioVendor)

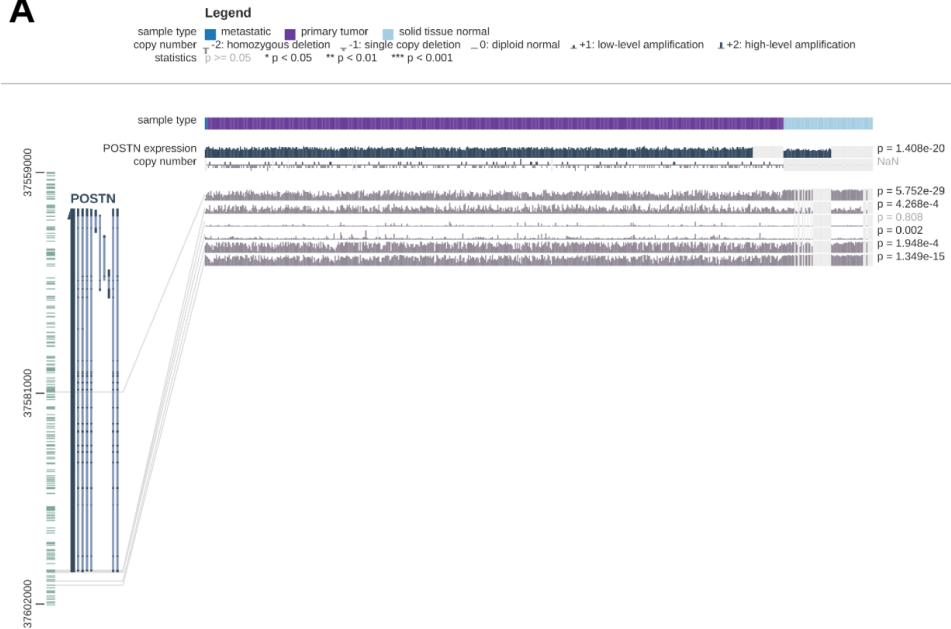
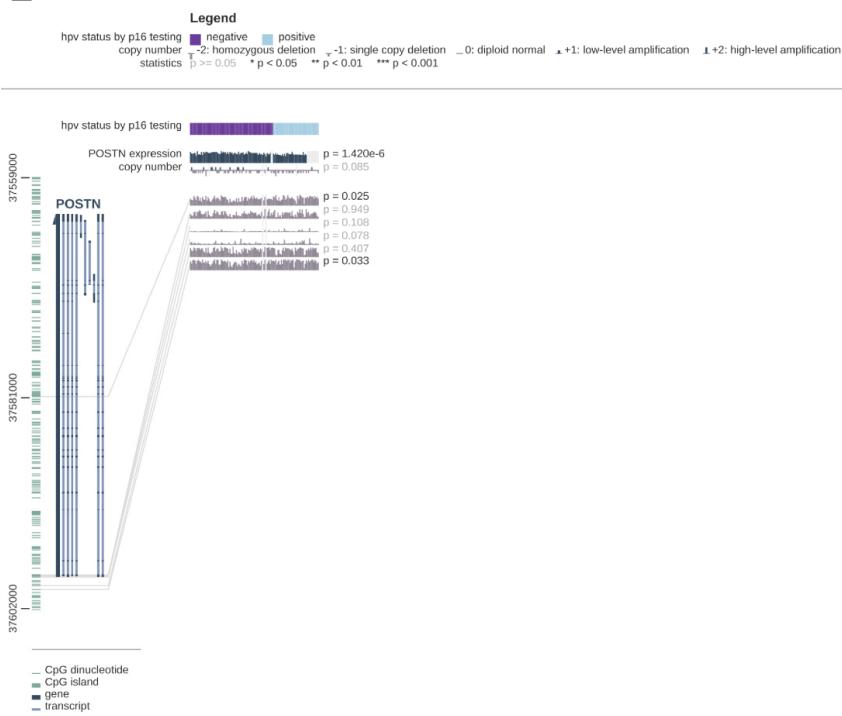
r-OSF-2 (Total 671 AA. MW: 75 kDa. UniProtKB Q15063. <b>N-Terminal HisTag and Xa – cleavage site 23 AA)</b>	MGHHHHHHHHSSGHIEGRHMRNNHYDKILAHSRIRGRDQGPNVCALQQILG TKKKYFSTCKNWYKKSICGQKTTVLYECCPGYMRMEGMKGCPAVLPIDHVYGT LGIVGATTQRYSASKLREEIEGKGSFTYFAPSNEAWNDLSDIRRGLESNVN VELLNALHSHMINKRMLTKDLKNGMIIPSMYNNLGLFINHYPNGVVTNCARIHG NQIATNGVHVIDRVLTQIGTSIQDFIEAEDLSSFRAAAITSDEALGRDGHFTL FAPTNEAFEKLPRGVLERFMGDKVASEALMKYHILNTLQCSESIMGGAVFETLE GNTIEIGCDGDSITVNGIKMVNKDIVTNNGVIHLIDQVLIPDSAKQVIELAGKQQT TFTDLVAQLGLASALRPDGLEYTLLAPVNNAFSDDTLSMVQRLKLILQNHILKV GLNELYNGQILETIGGKQLRWFVYRTAVCIENSCMEKGSKQGRNGAIHFREIIKP AEKSLHEKLQDKRFSTFLSLEAADLKEELLTQPGDWTLFVPTNDAFKGMTSEE KEILIRDKNALQNIILYHLPGVFIGKGFEPGVNLKTTQGSKIFLKEVNDTLLVNE LKSKESEDIMTTNGVIHVVDKLLYPADTPVGNDQLLIELNKLIKYIQIKFVRGSTFKEI PVTVY
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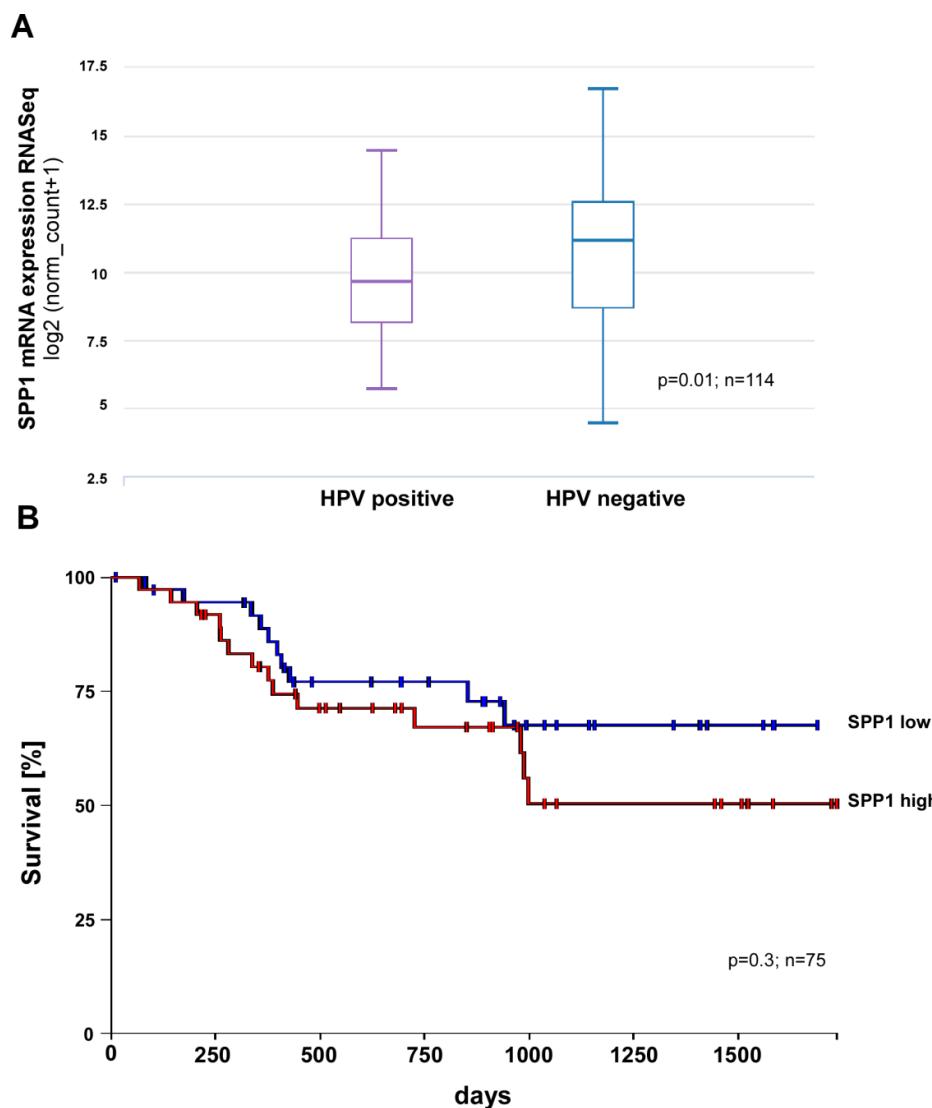
**Figure S1. OSF-2 (*POSTN*) is significantly up-regulated in TCGA HNSCC patient cohort (n=566) comparing N vs PT. Welch's t-test p = 2.341e-16**



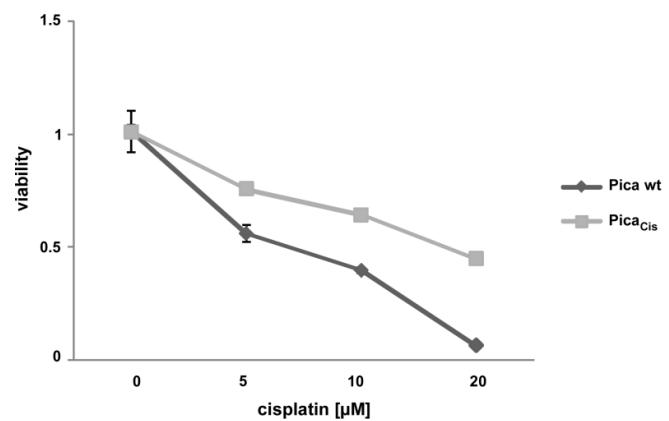
**Figure S2.** OSF-2 expression did not correlate with tumor localization in TCGA HNSCC patient cohort ( $n=538$ ).

**A****B**

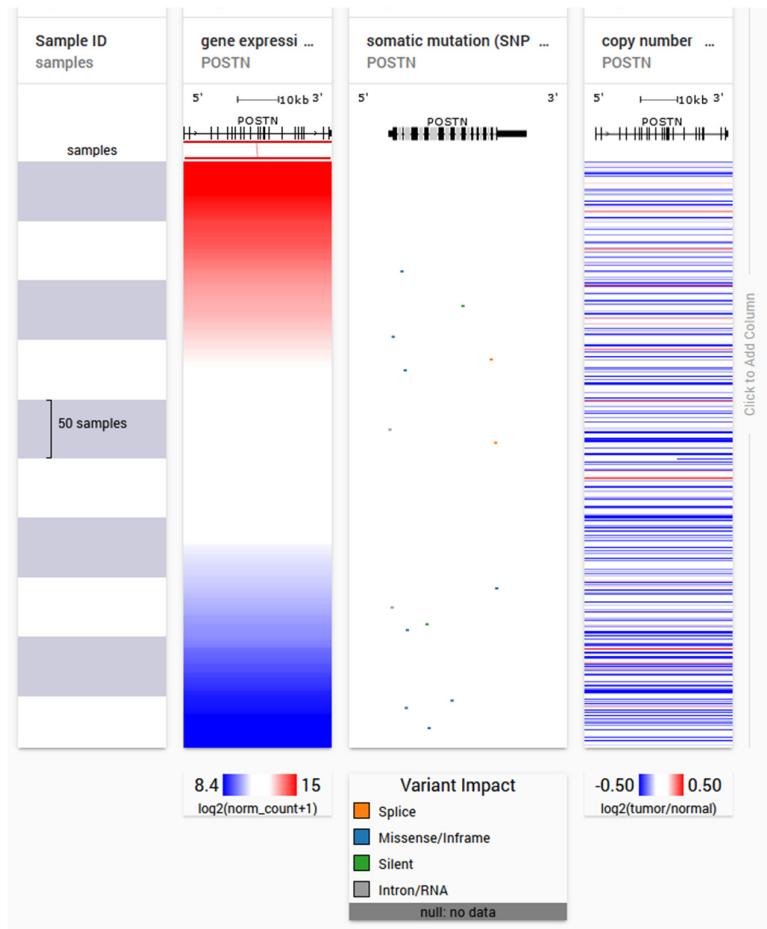
**Figure S3. OSF-2/POSTN DNA methylation of six CpG sites in n=528 HNSCC tumors and n=82 normal tissue samples from the TCGA cohort** (Infinium HumanMethylation450 BeadChip; cg23202139, cg04922971, cg24719107, cg05604486, cg13634560, cg18341491). Analysis was performed with MEXPRESS online tool (<https://www.mexpress.be>) (A) Methylation in 5 of 6 CpG sites is significantly decreased in PT vs. N correlating with increased POSTN expression levels. (B) POSTN methylation is reduced in HPV negative tumors compared to HPV positive samples. p values as indicated.



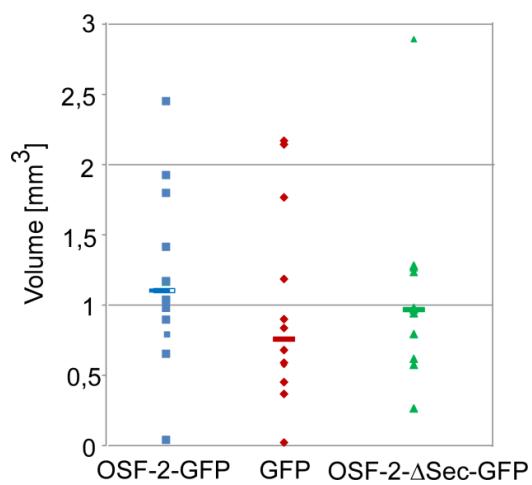
**Figure S4.** (A) Osteopontin (*SPP1*) is significantly upregulated in HPV negative vs. positive tumors of the TCGA HNSCC patient cohort (n=114). (B) High *SPP1* expression levels did not correlate with overall survival of HPV negative HNSCC patients. n=75; p=0.3.



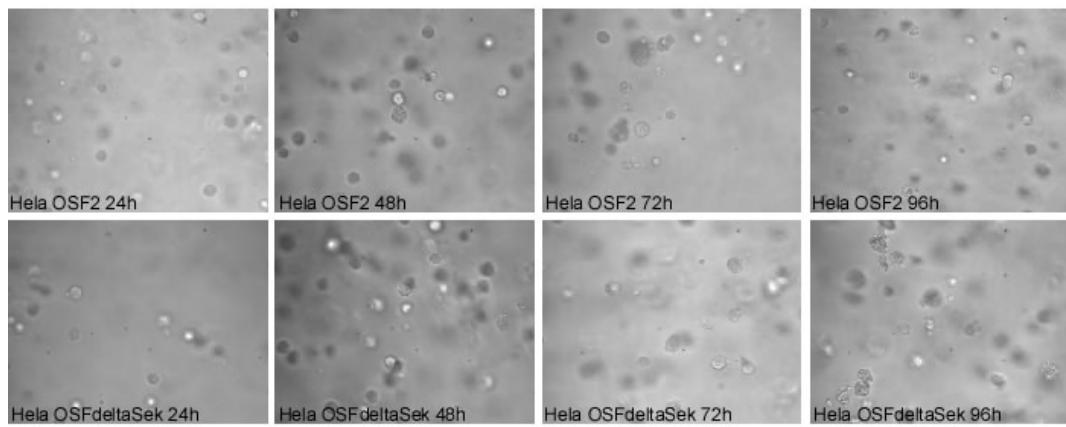
**Figure S5.** Spheroids of cisplatin-resistant pica cell line (PicaCis) exhibit significant chemoresistance towards different concentrations of cisplatin compared to wt cell line.



**Figure S6. Mutational analysis of OSF-2 gene using TCGA database (n=493).** Different types of mutations (middle column) were found in 15 HNC patients. There seems to be no correlation between high OSF-2 expression (left, red) and the occurrence of somatic mutations and/or copy number variations (right).



**Figure S7. Overexpression of OSF-2 in head and neck cancer cells did not alter tumor growth after subcutaneous injection into nude mice.** The average ratio of the tumor volume between tumors derived from periostin-overexpressing cells and control cells is less than 2-fold.



**Figure S8. OSF-2 secretion has no effect on cancer cell invasion.** Matrigel invasion assay of transfected HeLa cells expressing secretion active *versus* secretion deficient OSF-2 (deltaSek). 50 - 100  $\mu$ l diluted matrigel (BD Bioscience) was pipetted into the upper chamber of the 24-well transwell (Falcon BD, 8  $\mu$ m pore size). 24h after transfection,  $1 \times 10^5$  cells were seeded in 100  $\mu$ l of cell suspension in medium with 1% FCS into upper chamber of transwell. After gelling at 37° C for at least 4-5 hours, lower chamber was filled with 10 % FCS medium, and incubated at 37° C. Cells were observed for 4 days.



## Raw Data

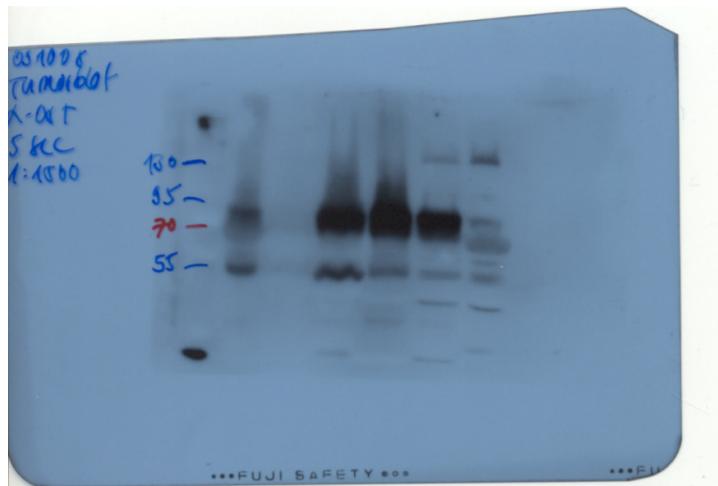


Fig1D

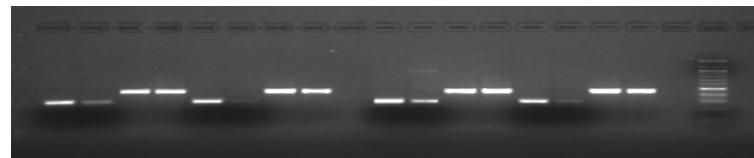
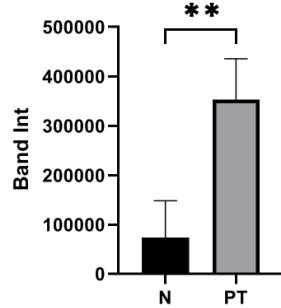


Fig1C

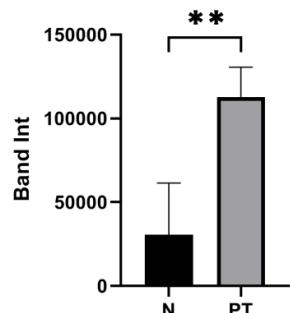


Figure S9. Western blot (upper) and agarose gel (lower) referring to validation of microarray data (Figure 1C, D) including densitometric analysis of bands ( $n=2$ ) (right).

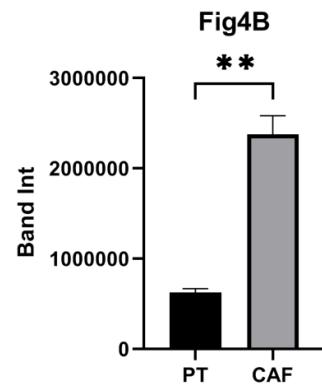
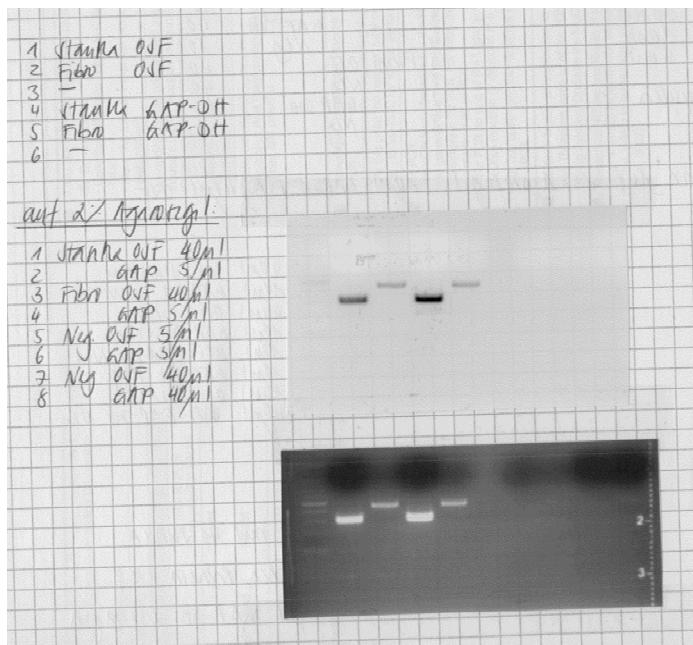


Figure S10. Agarose gel referring to RT-PCR analysis of POSTN expression in tumor cells and CAFs (Figure 4B) including densitometric analysis of bands ( $n=2$ ) (right).

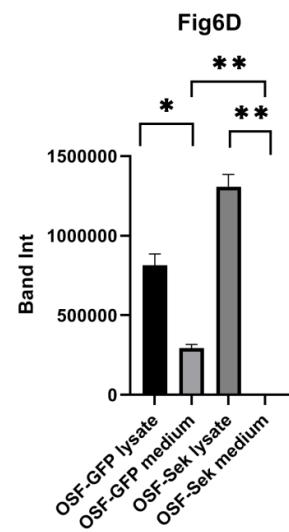
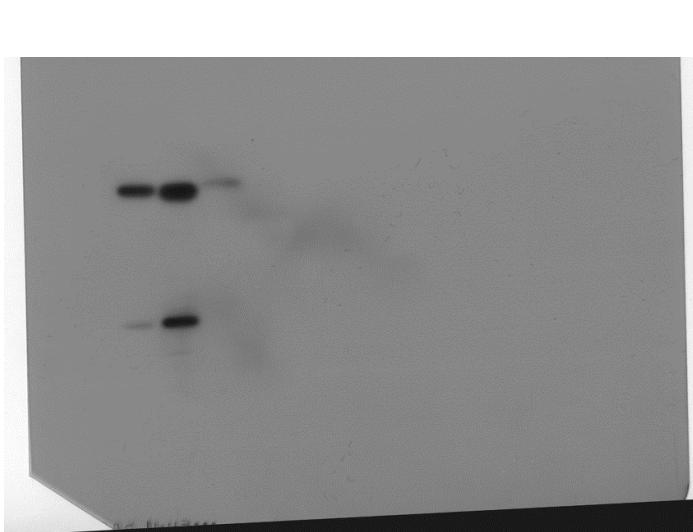


Figure S11. Western blot referring to analysis of OSF-2 secretion signal (Figure 6D) including densitometric analysis of bands ( $n=2$ ) (right).

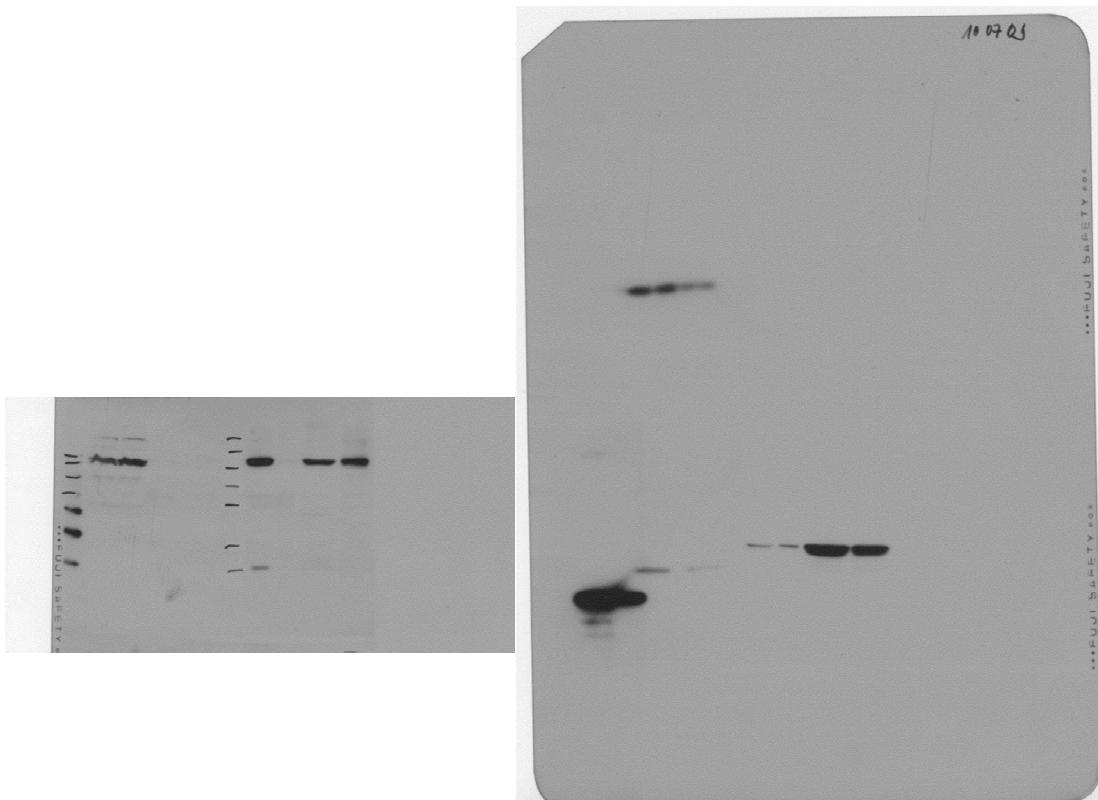
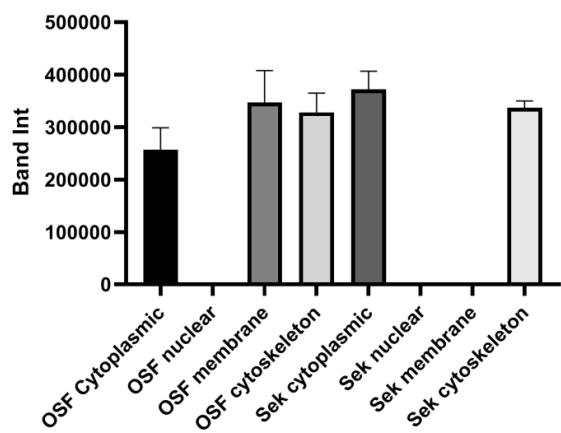

**Fig6E**


Figure S12. Western blots referring to cell fractionation analysis of secreted/ non-secreted OSF-2 (Figure 6E). a-OSF Ab (left), loading controls (right), and densitometric analysis of bands (n=2) (below).

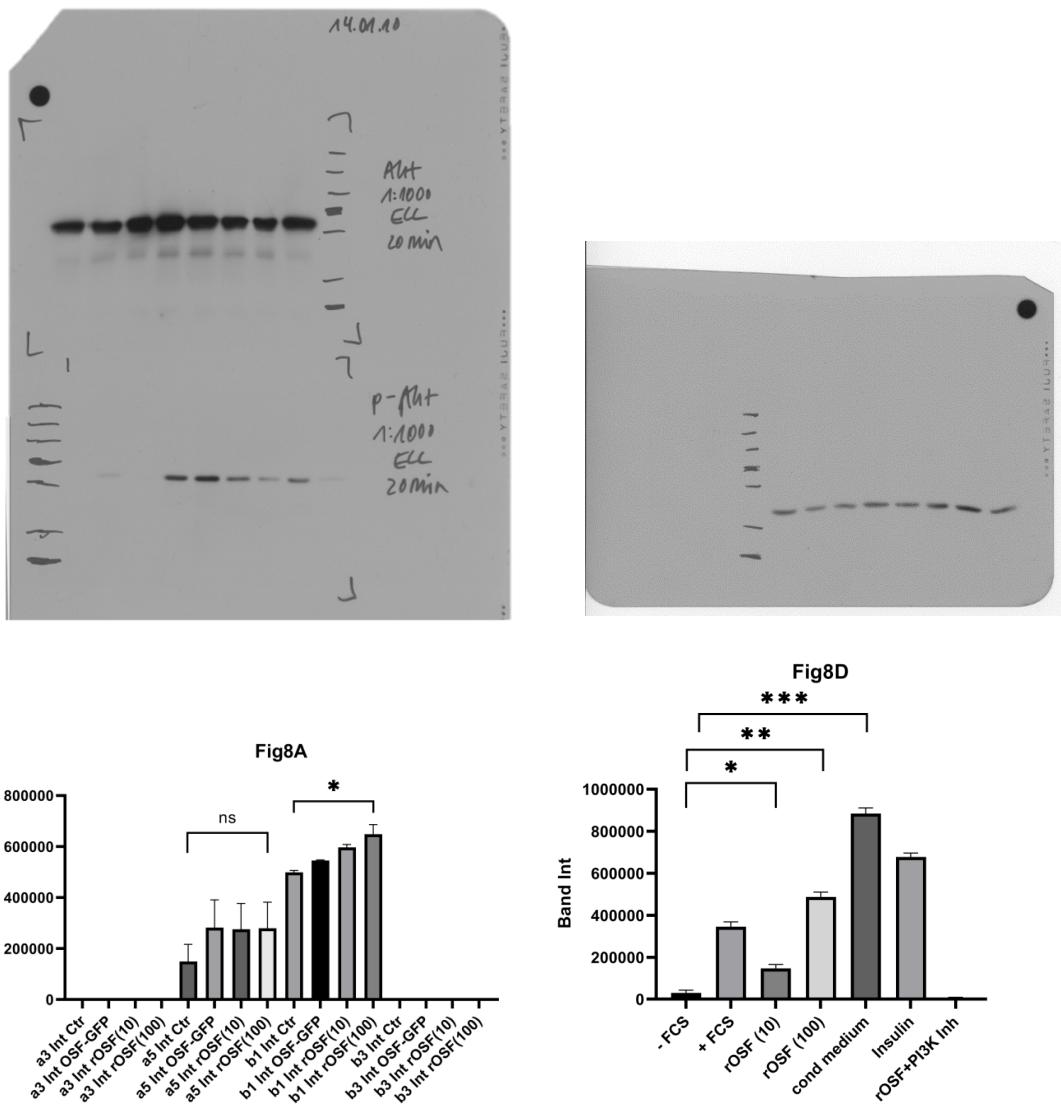


Figure S13. Western blots and densitometric analyses referring to Figure 8 A and D. a-pAkt Ab (left, lower), a-Actin Ab (right). Densitometric analysis of bands ( $n=2$ ) below.