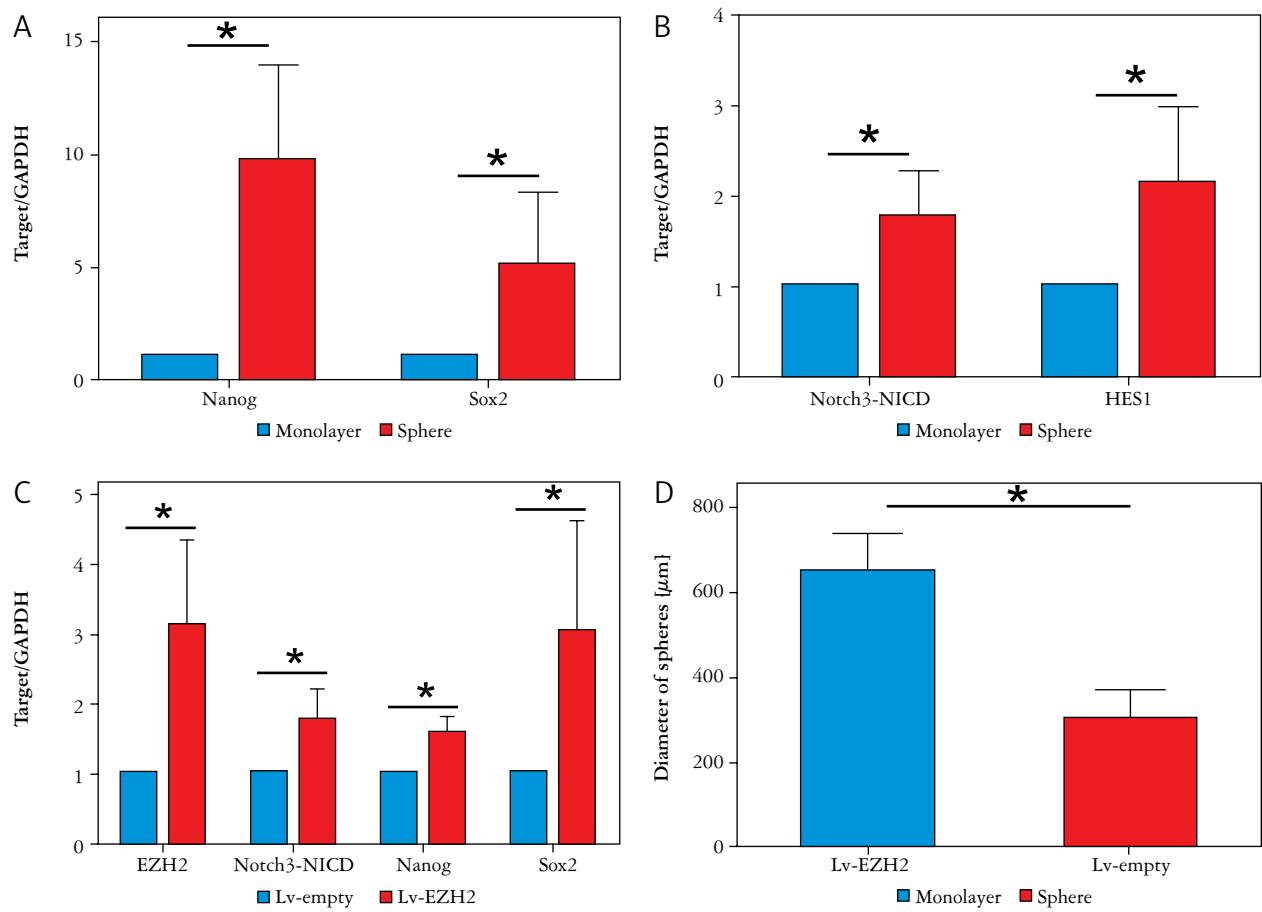


**Suppl. Table 1.** Sequences of primer pairs used in this study

GENE	PRIMER SEQUENCE (5'→3')
GAPDH	F: GGAGCGAGATCCCTCCAAAAT R: GGCTGTTGTCATACTTCTCATGG
Nanog	F: TTTGTGGGCCTGAAGAAAAT R: AGGGCTGTCCTGAATAAGCAG
Sox2	F: GCCGAGTGGAAACTTTGTCG R: GGCAGCGTGTACTTATCCTTCT
Oct4	F: GTGTTCAGCCAAAGACCAC R: GGCCTGCATGAGGGTTCT
Notch1	F: GAGGCGTGGCAGACTATGC R: CTTGTA CCGTCAGCGTGA
Notch2	F: CCTTCCACTGTGAGTGTCTGA R: AGGTAGCATCATTCTGGCAGG
Notch3	F: CGTGGCTTCTTCTACTGTGC R: CGTCACCGGATTGTGTAC
Notch4	F: GATGGGCTGGACACCTACAC R: CACACGCAGTGAAAGCTACCA
Hes1	F: CCTGTCATCCCCGTCTACAC R: CACATGGAGTCCGCCGTAA
Hey1	F: GTTCGGCTCTAGGTTCCATGT R: CGTCGGCGTTCTCAATTATTC

**Suppl. Table 2.** Characteristics of osteosarcoma patients

MALE/FEMALE	37/21				
Age (years)	Range	Mean	Median	≤ 24	> 24
	7–74	24.31	18	38 (65.5%)	20 (34.5%)
Tumour diameter [cm]	Range	Mean	Median	≤ 8	> 8
	1.4–18	8.4	8	32 (55.2%)	26 (44.8%)
Metastatic location	Lung	Pelvis	Scapula	Skull	
	29 (90.6%)	1 (3.1%)	1 (3.1%)	1 (3.1%)	
Enneking staging	IA	IB	IIA	IIB	III
	1 (1.7%)	1 (1.7%)	12 (20.7%)	39 (67.2%)	5 (8.6%)
Locations	Femur	Tibia	Fibula	Humerus	Pelvis
	35 (60.3%)	8 (13.8%)	5 (8.6%)	4 (6.9%)	4 (6.9%)
					Clavicle
					1 (1.7%)
					1 (1.7%)
Surgery method	Excision	Amputation			
	49 (84.5%)	9 (15.6%)			



**Suppl. Fig. 1.** Densitometry analysis of protein expressions in osteosarcoma stem cell (OSC) spheres. A) Densitometry analysis of protein expressions demonstrated that Nanog and Sox2 were upregulated in OSC spheres compared to monolayer cells (\* $p < 0.05$ ); B) Densitometry analysis of protein expressions demonstrated that Notch3 intracellular domain (Notch3-NICD) and HES1 were upregulated in OSC spheres compared to monolayer cells (\* $p < 0.05$ ); C) Densitometry analysis of protein expressions showed that after overexpression of enhancer of zeste homologue 2, Notch3-NICD, Nanog, and Sox2 were upregulated in osteosarcoma cells (\* $p < 0.05$ )