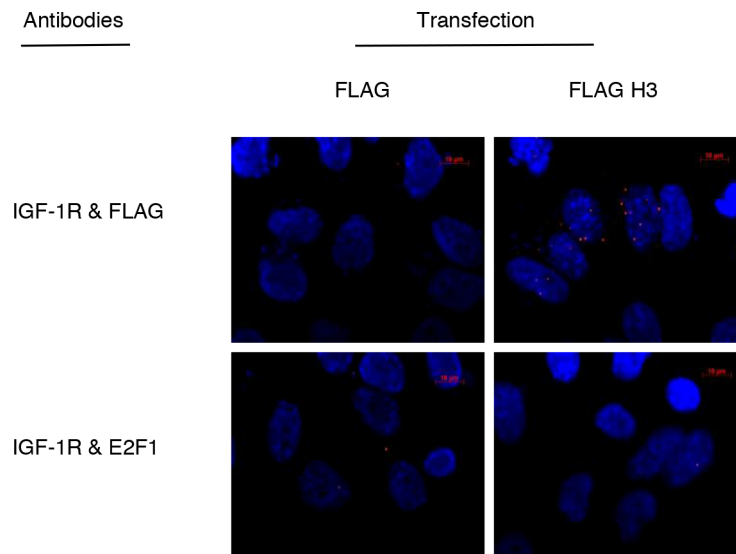
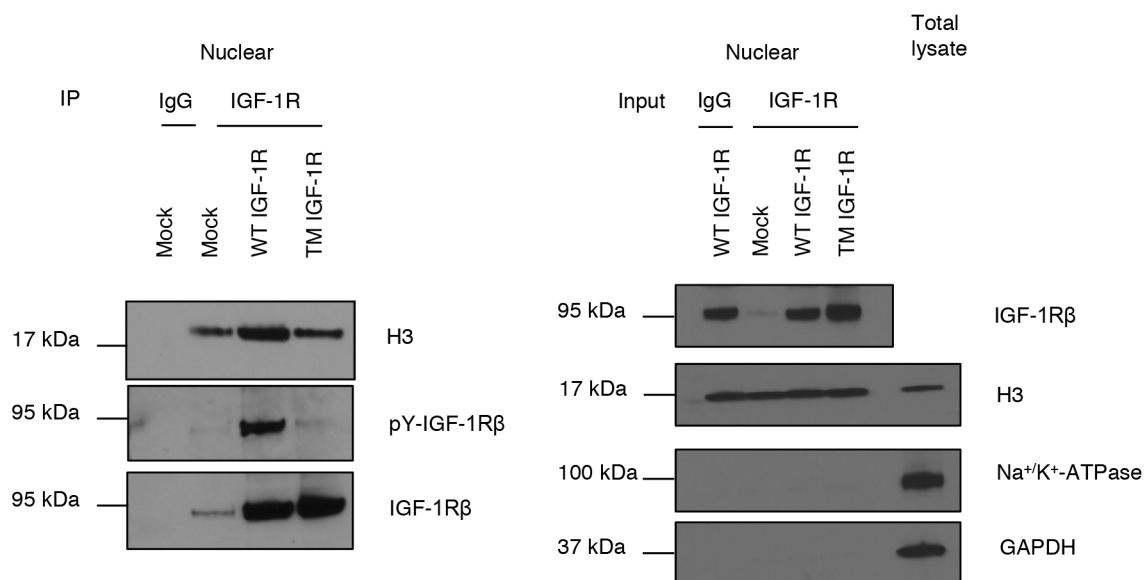


Nuclearly translocated insulin-like growth factor 1 receptor phosphorylates histone H3 at tyrosine 41 and induces *SNAI2* expression via Brg1 chromatin remodeling protein

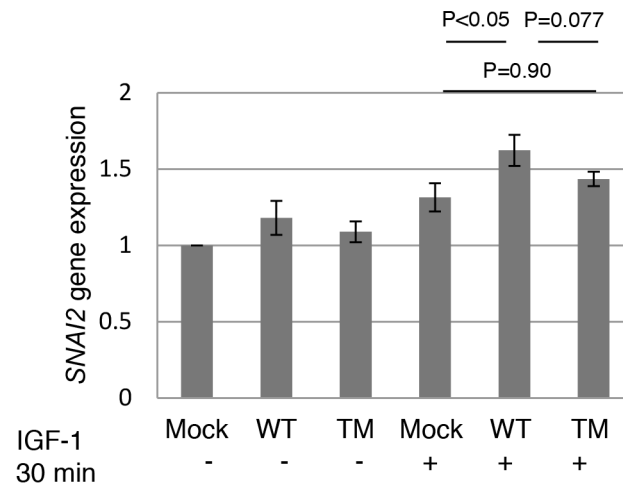
SUPPLEMENTARY FIGURES AND TABLE



Supplementary Figure S1: nIGF-1R binds to histone H3. nIGF-1R and histone H3 binding determined with PLA (red dots) in HeLa cells transfected with an empty Flag vector (top left) or Flag H3 construct (top right). Anti-Flag and anti-IGF-1R antibodies were used. For negative control, PLA between IGF-1R and E2F1 did not show any signal (bottom panels). Cells were growing in basal conditions.



Supplementary Figure S2: IGF-1R phosphorylation is important for its binding to histone H3. HeLa cells cultured under basal conditions were transiently transfected with mock, *wt-IGF1R* or *tm-IGF1R* for 48 hours. Nuclear extracts were subjected to IP with IgG or IGF-1R antibodies and blotted for histone H3, phosphotyrosine-IGF-1R and IGF-1R (left panels). Right panels show purity of nuclear fractions.



Supplementary Figure S3: *SNAI2* gene expression by wt and tm-IGF-1R. HeLa cells cultured under basal conditions were transfected with mock, *wt-IGF-1R* or *tm-IGF-1R* for 24 hours. After 24 hours, the cells were serum starved for an additional 24 hours. Serum starved cells were stimulated with 100 ng/ml IGF-1 for 0-30 minutes and *SNAI2* expression was analyzed with qPCR. One-way ANOVA and Tukey's HSD test were performed. P-values are indicated. Means and s.d. are shown (n=3).

Supplementary Table S1: Identification of nIGF-1R target genes by PCR array

Gene	<i>wt-IGF1R</i>	<i>tsm-IGF1R</i>
	Fold change	Fold change
<i>ARNT</i>	-1.0754	-1.7223
<i>AURKA</i>	-1.1051	-1.3286
<i>BIRC3</i>	1.2182	-1.0754
<i>BMI1</i>	-1.0498	-1.5638
<i>CCND2</i>	3.3681	5.3244
<i>DDB2</i>	-1.4075	-1.8561
<i>DDIT3</i>	2.157	1.3646
<i>DKC1</i>	1.2103	-1.027
<i>E2F4</i>	-1.0373	-1.5605
<i>ERCC3</i>	1.0774	-1.1268
<i>ERCC5</i>	1.2552	-1.2045
<i>ETS2</i>	-1.2203	-2.0247
<i>FOXC2</i>	-1.1939	-1.4475
<i>GSC</i>	1.6681	3.0432
<i>MCM2</i>	1.0559	-1.3938
<i>MKI67</i>	-1.2054	-1.7043
<i>NOL3</i>	1.114	-1.3191
<i>PINX1</i>	1.1587	-1.3956
<i>POLB</i>	1.1331	-1.3281
<i>SKP2</i>	1.0207	-1.3643
<i>SNAI1</i>	-1.0064	-2.3167
<i>SNAI2</i>	1.1915	-1.1338
<i>SNAI3</i>	1.2047	-1.5561
<i>SOX10</i>	-1.279	1.565
<i>TBX2</i>	1.4937	-1.4026
<i>TEP1</i>	1.2796	-1.3499
<i>TERF1</i>	1.6648	-1.2004
<i>TERF2IP</i>	1.4605	1.1349
<i>TINF2</i>	-1.0302	-1.3631
<i>WEE1</i>	2.2683	-1.3681

HeLa cells cultured under basal conditions were transiently transfected with mock, *wt-IGF1R* or *tsm-IGF1R* for 48 hours. RNA was extracted and converted into cDNA (see Materials & Methods). Gene expression was measured using the RT² Profiler PCR Array. The results include genes whose proteins are nuclear/nuclear-associated. Gene expression is shown as fold change compared to mock transfected cells.