

The Wnt/beta-catenin pathway regulates the expression of early embryonic stem cell genes in human parathyroid tumors.

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Introduction

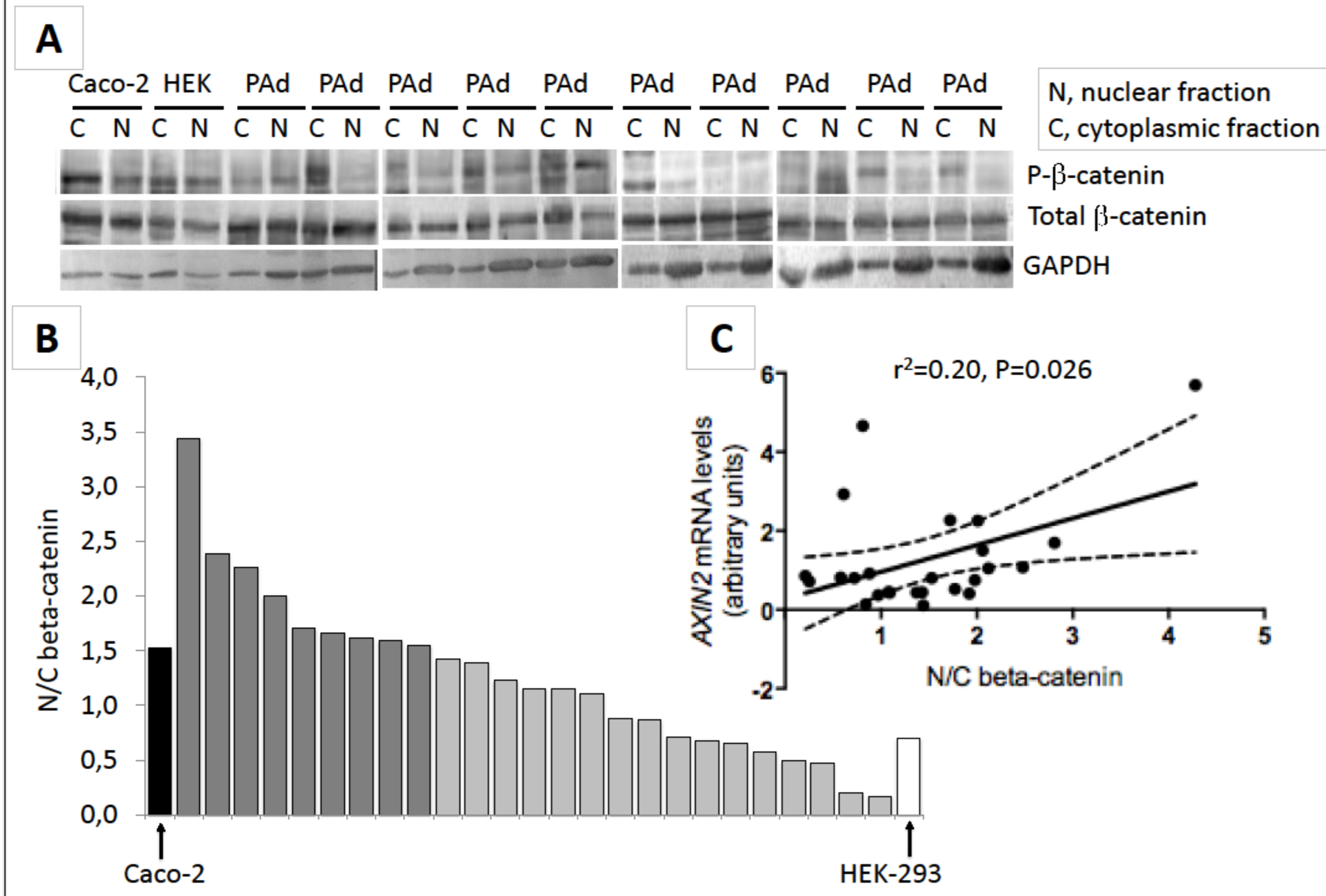
Evidence suggested an embryonic epigenetic signature in parathyroid tumors, with deregulated miRNAs and gene methylation. In embryonic stem cells, the Wnt/ β -catenin signaling regulates the expression of the core stemness genes, namely NANOG, OCT4 and SOX2. Though constitutive nuclear accumulation of β -catenin has not been detected, the Wnt/ β -catenin pathway might be deregulated in parathyroid tumors, as Wnt signaling inhibitors have been found reduced.

Aim of the study To investigate the embryonic signaling Wnt/ β -catenin – core stem cells genes in adult human tumoral parathyroids.

Results

Nuclear active β -catenin levels in parathyroid adenomas:

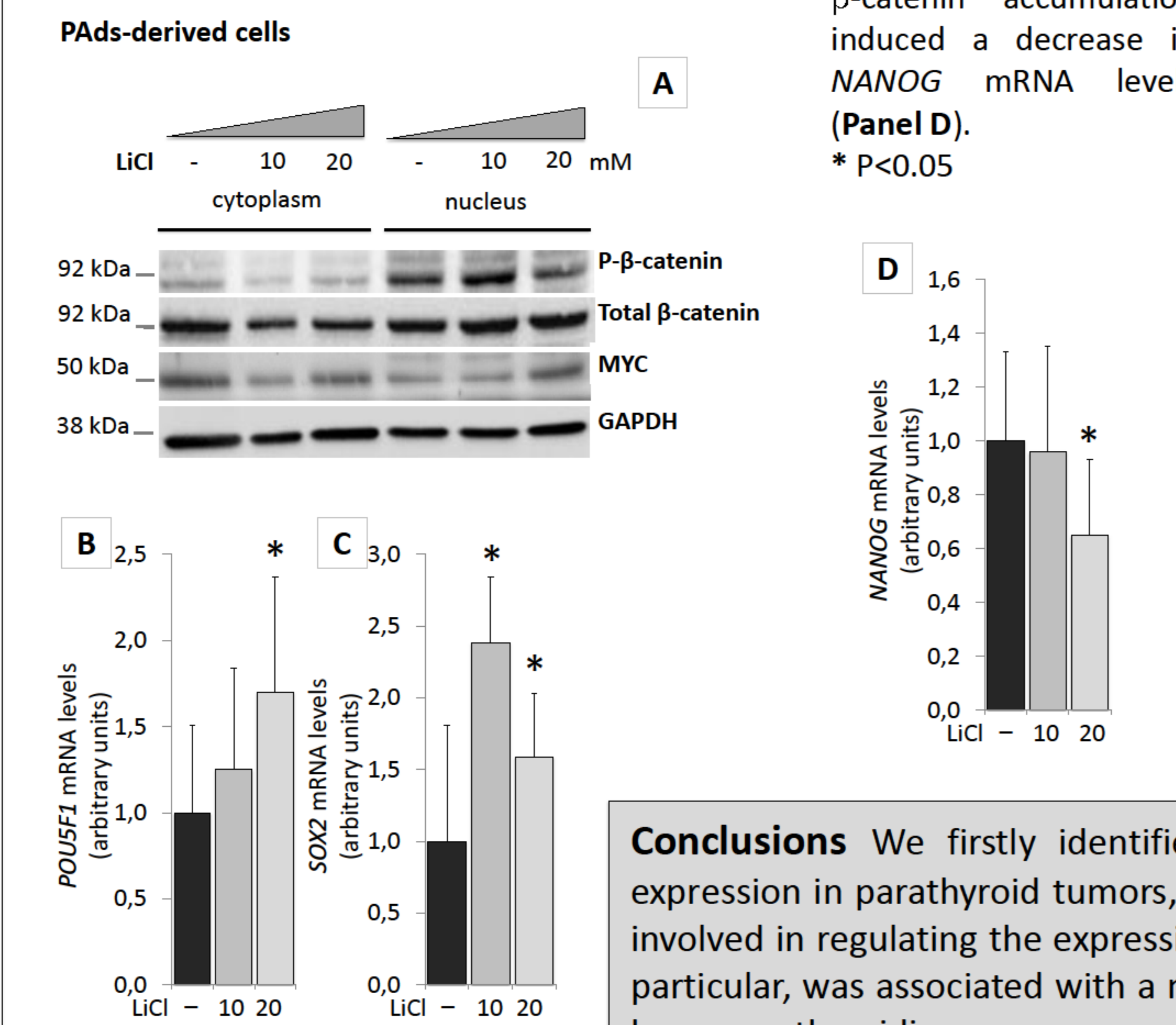
We investigated unphosphorylated active β -catenin distribution by western blot in 25 typical parathyroid adenomas (PADs) (Panel A): β -catenin accumulation in the nuclear protein fractions varied from the levels detected in Caco-2 cells with constitutively active Wnt signaling (9 PADs) to the levels measured in HEK293 cells with intact Wnt signaling (6 PADs) (Panel B) and positively correlated with *AXIN2* mRNA levels ($r=0.445$, $P=0.026$) (Panel C).



Effects of Lithium Chloride (LiCl) treatment on stem cell markers in parathyroid adenomas (PADs)-derived cells:

The Wnt/ β -catenin pathway is intimately connected to the embryonic pluripotent core circuitry. Treatment of PADs-derived cells ($n=5$) with 10-20 mM lithium chloride (LiCl) for 8 hours induced nuclear accumulation of β -catenin (Panel A) and concomitant increases in mRNA levels of *POU5F1* (Panel B and C) and *SOX2* genes, as reported in embryonic stem cells. Nonetheless, at variance with embryonic stem cells,

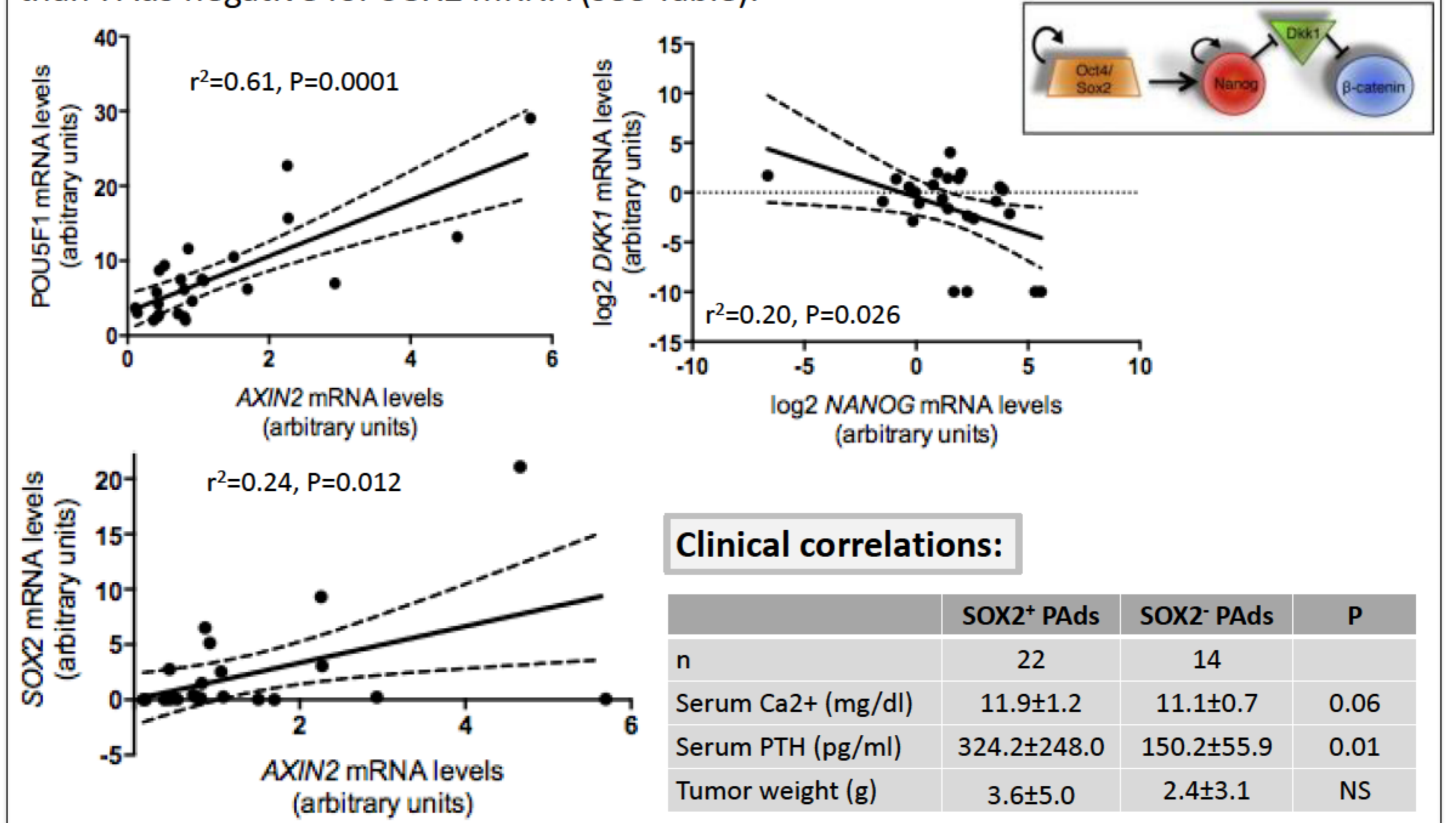
β -catenin accumulation induced a decrease in *NANOG* mRNA levels (Panel D). * $P<0.05$



Conclusions We firstly identified an embryonic pattern of gene expression in parathyroid tumors, where β -catenin signaling might be involved in regulating the expression of the core stem genes. *SOX2*, in particular, was associated with a more severe presentation of primary hyperparathyroidism.

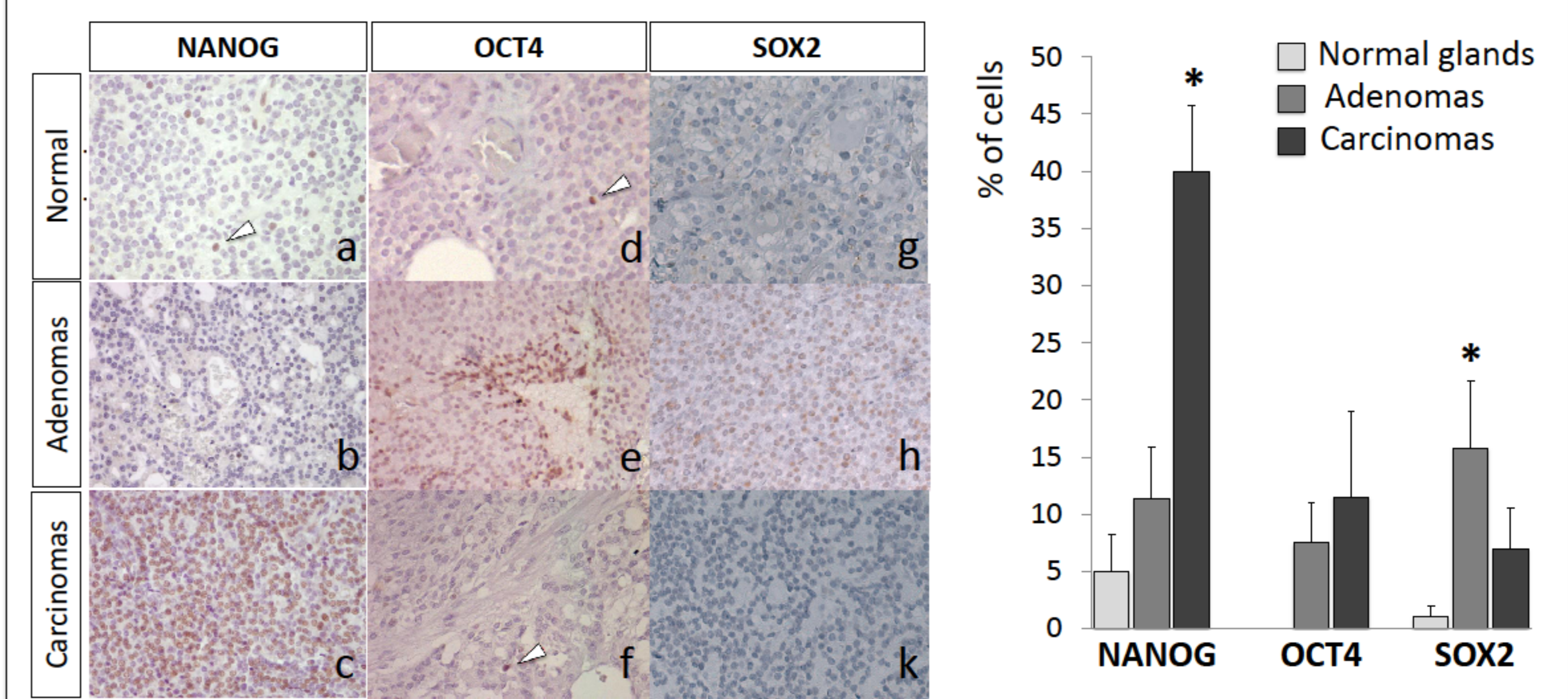
AXIN2 mRNA expression correlated with *POU5F1* and *SOX2* mRNA expression levels:

the mRNA levels of *POU5F1* and *SOX2* genes in 25 PADs positively correlated with *AXIN2* mRNA levels, in agreement with the effect of LiCl in dispersed cells. *NANOG* mRNA levels negatively correlated with *DKK1* mRNA levels. PADs expressing *SOX2* transcripts were associated with more severe hyperparathyroidism than PADs negative for *SOX2* mRNA (see Table).



Core stem genes analysis by immunohistochemistry in parathyroid tumors:

Immunohistochemistry of tumor sections [11 PADs, 8 carcinomas (PCas)] identified cells expressing at nuclear level the core stem cell genes:



Immunofluorescence analysis of core stem genes in PADs-derived cells:

Immunofluorescence detected few cells coexpressing *SOX2* and *NANOG* or *OCT4* (Panel A), in agreement with the positive correlation between *NANOG* and *SOX2* mRNA levels in Pads.

Parathyroid tumor cells expressing PTH were negative for all the stem cell genes (Panel B).

