The role of transcriptional factor brachyury in the development and repair of nucleus pulposus

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Abstract: Transcription factor Brachyury, a protein containing 435 amino acids, has been widely investigated and reported in notochord differentiation and nucleus pulposus development. The crucial functions and underlying mechanisms by Brachyury are discussed in this paper, which suggests Brachyury can be developed into a potential novel target for the therapy of intervertebral disc degeneration.

Abbreviations

IVD:	Intervertebral disc
NP:	Nucleus pulposus
NC:	Notochordal
iPSC:	Induced pluripotent stem cells
PSM:	Primitive streak mesoderm

Introduction

Current therapeutic modalities for low back pain arising from intervertebral disc (IVD) degeneration mainly target the symptoms alleviation while failing to treat the underlying disease pathology (Tang *et al.*, 2021).

Various studies have attempted to develop early-stage intervention to retard and reverse the IVD degeneration process. Recapitulation of the embryonic patterning process of the IVD may help develop novel biological repair approaches for disc degeneration (Cornejo *et al.*, 2015). The central nucleus pulposus (NP) tissue plays a pivotal role in maintaining the function of IVD. It is the mainly affected structure in IVD degeneration (Rodrigues-Pinto *et al.*, 2014), which demonstrates decreased cellularity, water content and proteoglycans.

Various studies have evidenced that the entire cell subpopulations of the NP are descended from the embryonic notochord (Shapiro and Risbud, 2010). The transcriptional factors playing vital roles in the notochord

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development have been attempted to rescue the function of degenerated nucleus pulposus cells.

Main Text

Transcription factor Brachyury, a protein of 435 amino acids, binds with the canonical T-domain DNA sequence: TCACACCT. Brachyury regulates posterior mesoderm formation and notochord differentiation by directly activating downstream mesoderm-specific genes (Chen et al., 2020a; Chen et al., 2020b; Xu et al., 2020). The embryonic notochord is considered to be the developmental origin of mature adult NP tissue (Richardson et al., 2017). Brachyury has widely been recognized as the specific marker for the notochordal (NC) cells (Tang et al., 2016). The human notochord and juvenile NP are populated by larger and vacuolated NC cells, which gradually disappear and are replaced by a population of small and round "chondrocytelike" NP cells after puberty (Risbud et al., 2015). There is a long-held controversy on the ontogeny and heterogeneity of morphologically distinct NC and NP cells. Minogue et al. (2010) found NP cells isolated from adult bovine and human discs also expressed Brachyury, the mRNA level expression remained unchanged in the degenerated human nucleus pulposus, which is suggestive of a common ontogeny of the NP and NC cells. Further study by Richardson et al. (2017) showed Brachyury protein was expressed in the cells of adult human nucleus pulposus tissue, but significantly lower positivity was demonstrated in mature adult compared to young and significantly lower positivity in severely degenerated tissues compared to nondegenerated. Various convincing evidence suggest that most cells in adult nucleus pulposus tissue differentiate along the





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notochordal lineage, morphological differences of NC and NP cells represent different physiological or pathological stages of aging and degeneration (Risbud *et al.*, 2015). Brachyury, a traditional marker of NC cells, has been considered as a critical phenotypic marker of healthy NP (Risbud *et al.*, 2015) and helps to evidence the homology of NC and NP cells.

Several studies have reported that Brachyury was transfected into degenerated NP cells or human induced pluripotent stem cells (iPSC), aiming to promote these cells to transmit or differentiate into healthy NP cells, which can produce more extracellular matrix and restore the homeostasis of disc microenvironment. Brachyury lent virus transfection promoted and enhanced the differentiation of human iPSC toward NP-like cells in vitro (Tang et al., 2018). Tang et al. (2019) demonstrated nonviral Brachyury transfection into mildly degenerated human NP cells promoted the phenotype transition into a pro-anabolic and anti-catabolic/inflammatory one, and significantly increased glycosaminoglycan accumulation in vitro, which suggests Brachyury has the potential to be developed into an useful target for the treatment of IVD degeneration. Sheyn et al. (2019) reported stepwise differentiation strategy of human iPSCs into NC cells. Brachyury-encoding plasmids were nucleofected into primitive streak mesoderm (PSM) cells, which were induced and differentiated from iPSCs, to acquire NC cells. Further testing in vivo in the porcine model showed that the acquired NC cells had a protective role on discs from degeneration.

We have found that Brachyury can promote proteoglycan synthesis by nucleus pulposus cells. Whether Brachyury transfection into remaining degenerated nucleus pulposus cells or exogenously sourced cells can facilitate the transition toward a healthy nucleus pulposus phenotype in the harsh degenerated disc microenvironment requires more in-depth investigations.

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References

- Cornejo M, Cho S, Giannarelli C, Iatridis J, Purmessur D (2015). Soluble factors from the notochordal-rich intervertebral disc inhibit endothelial cell invasion and vessel formation in the presence and absence of pro-inflammatory cytokines. *Osteoarthritis and Cartilage* 23: 487–496. DOI 10.1016/j.joca.2014.12.010.
- Chen M, Wu Y, Zhang H, Li S, Zhou J, Shen J (2020a). The roles of embryonic transcription factor BRACHYURY in tumorigenesis

and progression. *Frontiers in Oncology* **10**: 961. DOI 10.3389/ fonc.2020.00961.

- Chen M, Zou S, He C, Zhou J, Li S et al. (2020b). Transactivation of SOX5 by Brachyury promotes breast cancer bone metastasis. *Carcinogenesis* **41**: 551–560. DOI 10.1093/carcin/bgz142.
- Minogue BM, Richardson SM, Zeef LA, Freemont AJ, Hoyland JA (2010). Transcriptional profiling of bovine intervertebral disc cells: Implications for identification of normal and degenerate human intervertebral disc cell phenotypes. *Arthritis Research & Therapy* **12**: R22. DOI 10.1186/ar2929.
- Richardson SM, Ludwinski FE, Gnanalingham KK, Atkinson RA, Freemont AJ et al. (2017). Notochordal and nucleus pulposus marker expression is maintained by subpopulations of adult human nucleus pulposus cells through aging and degeneration. *Scientific Reports* 7: 1356. DOI 10.1038/s41598-017-01567-w.
- Risbud MV, Schoepflin ZR, Mwale F, Kandel RA, Grad S et al. (2015). Defining the phenotype of young healthy nucleus pulposus cells: Recommendations of the Spine Research Interest Group at the 2014 annual ORS meeting. *Journal of Orthopaedic Research* 33: 283–293. DOI 10.1002/jor.22789.
- Rodrigues-Pinto R, Richardson SM, Hoyland JA (2014). An understanding of intervertebral disc development, maturation and cell phenotype provides clues to direct cellbased tissue regeneration therapies for disc degeneration. *European Spine Journal* 23: 1803–1814. DOI 10.1007/ s00586-014-3305-z.
- Shapiro IM, Risbud MV (2010). Transcriptional profiling of the nucleus pulposus: Say yes to notochord. Arthritis Research & Therapy 12: 117. DOI 10.1186/ar3003.
- Sheyn D, Ben-David S, Tawackoli W, Zhou Z, Salehi K et al. (2019). Human iPSCs can be differentiated into notochordal cells that reduce intervertebral disc degeneration in a porcine model. *Theranostics* 9: 7506–7524. DOI 10.7150/thno.34898.
- Tang R, Jing L, Willard VP, Wu CL, Guilak F et al. (2018). Differentiation of human induced pluripotent stem cells into nucleus pulposus-like cells. *Stem Cell Research & Therapy* 9: 79. DOI 10.1186/s13287-018-0797-1.
- Tang S, Richards J, Khan S, Hoyland J, Gallego-Perez D et al. (2019). Nonviral transfection with Brachyury reprograms human intervertebral disc cells to a pro-anabolic anti-catabolic/ inflammatory phenotype: A proof of concept study. *Journal of Orthopaedic Research* **37**: 2389–2400. DOI 10.1002/ jor.24408.
- Tang S, Salazar-Puerta A, Richards J, Khan S, Hoyland J et al. (2021). Non-viral reprogramming of human nucleus pulposus cells with FOXF1 via extracellular vesicle delivery: An *in vitro* and *in vivo* study. *European Cells and Materials* **41**: 90– 107. DOI 10.22203/eCM.v041a07.
- Tang X, Jing L, Richardson WJ, Isaacs RE, Fitch RD et al. (2016). Identifying molecular phenotype of nucleus pulposus cells in human intervertebral disc with aging and degeneration. *Journal of Orthopaedic Research* 34: 1316–1326. DOI 10.1002/jor.23244.
- Xu J, Chen M, Wu Y, Zhang H, Zhou J et al. (2020). The role of transcriptional factor Brachyury on cell cycle regulation in non-small cell lung cancer. *Frontiers in Oncology* **10**: 7066. DOI 10.3389/fonc.2020.01078.