

9 Reprogramming

To date, several different approaches can be applied to the reprogramming of somatic cells to a pluripotent state [144, 145]. Briefly, a somatic cell nucleus transferred to an unfertilized egg acquires pluripotency and develops into a blastocyst, allowing cloned ES cells to be established. Alternatively, somatic cells can be reprogrammed by cell–cell hybridization (cell fusion). The disadvantages of these techniques are the destruction of the embryo (i.e. ethical problem) and the tetraploid karyotype of the fused cells (i.e. risk problem in transplantation), respectively. Because these problems can be avoided using iPS cells, most recent reprogramming studies have utilized iPS cell systems.

In iPS induction, *Oct3/4*, *Sox2*, *Klf4*, and *Myc* (4 factors) are simultaneously introduced into somatic cells, and the reprogrammed cells are selected by marker gene expression and/or morphology [146]. Retroviruses were initially used to introduce the four factors, but the resulting iPS cells showed numerous insertions in the chromosome, raising safety issues. Currently, iPS cells can be established by a DNA-free method, in which the proteins of the four factors are tagged with an arginine stretch that confers membrane permeability [147, 148]. The cell types in which successful reprogramming have been reported include fibroblasts, primary hepatocytes, and completely differentiated B lymphocytes, suggesting that all the cell types in the body can be reprogrammed using the iPS technique [149–151]. The progression of reprogramming in iPS induction is relatively slow when compared with other techniques. In cell fusion, the upregulation of *Oct3/4* in completely differentiated cells is observed within 2 days [152], whereas in the iPS process this upregulation is first detectable 16 days after induction [153].

The molecular mechanism of this reprogramming is not clear at present. The efficiency of iPS establishment is low, less than a few percent of cells treated [154], indicating that in most cells the reprogramming is aborted even in the presence of the four factors. The function of *Myc* for iPS induction is dispensable, although it enhances the efficiency of the iPS establishment, probably through repressing the expression of differentiated cell-specific genes while promoting binding of *Oct3/4*, *Sox2*, and *Klf4* (OSK) to their target genes [75, 155]. In fact, in partially reprogrammed cells OSK does not bind to the target genes (which are thus not expressed), suggesting that the cellular environment ensuring access of these factors to the target genes is rate limiting in reprogramming [155]. In addition, administration of chemical inhibitors targeting epigenetic factors that are associated with transcriptional repression is effective for enhancing iPS cell induction. These inhibitors include BIX-01294 (G9a inhibitor) [156], AZA (5-aza-cytidine,

Dnmts inhibitor) [154, 157], VPA (valproic acid, Hdac inhibitor) [158], and TSA (trichostatin A, Hdac inhibitor) [158], and such findings suggest that target accessibility accompanying the global transcriptional activation seen in ES cells is critical to the initiation of the pluripotency transcriptional network. During iPS induction, OSK activity is enhanced by the transcription factors known to co-regulate with OSK, which include *Esrrb*, *Sall4* and *Tcf3* (via Wnt signaling), since forced expression of those factors can enhance the efficiency of iPS induction [80, 142, 159–161].

Collectively, the mechanism of reprogramming in iPS induction can be hypothesized as follows. Upon introduction of the four factors, the endogenous genes for the transcription factors necessary to pluripotency are primed and gradually induced to express through the regulatory region targeted by OSK. Subsequently, the transcriptional circuit begins to self-stabilize via increasing expression of the endogenous core transcription factors, through a positive-feedback loop, while repressing the developmentally regulated genes through recruitment of epigenetic factors and various protein complexes such as Paf1C and NODE. Once the stable transcriptional network is established/self-stabilized, exogenous cDNA expression is no longer necessary, and iPS cells indistinguishable from ES cells can be obtained after selection based on the expression of endogenous *Oct3/4* and/or *Nanog*. This outcome is the result of concerted action by a group of molecules that play a central role in pluripotency.

10 Conclusion

Pluripotency of ES cells is externally regulated through several molecules, including Wnt and Lif, whose signaling pathway activates transcription factor genes such as *Klf4* and *Nanog* in the nucleus. The core transcription factors, including *Oct3/4*, *Sox2*, and *Klf4* positively self-regulate while also repressing developmentally regulated genes by co-occupation with a variety of protein complexes. Introduction of *Oct3/4*, *Sox2*, and *Klf4* into the somatic cells gradually reconstitutes the above transcriptional network with the aid of *Myc* and epigenetic modifiers, which might allow the regulatory regions of the target genes more access to these transcription factors.

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