

<b>Supplemental Table 1: Examples of genes that promote lineage plasticity and cancer</b>		
<b>Gene name</b>	<b>Roles in homeostasis and cancer</b>	<b>Selected references</b>
<i>Homeobox transcriptional regulatory genes</i>		
<i>CDX2</i>	Expressed in intestinal epithelium and promotes its differentiation; dysregulation leads to colon cancer; principal driver of Barrett’s metaplasia	( <a href="#">Strumpf et al. 2005</a> , <a href="#">Colleypriest et al. 2010</a> )
<i>NKX2.1</i>	Expressed in lung epithelium and promotes its differentiation; dysregulation leads to lung cancer; represses a gastric phenotype in lung epithelium	( <a href="#">Winslow et al. 2011</a> , <a href="#">Snyder et al. 2013</a> )
<i>NKX3.1</i>	Expressed in luminal prostatic epithelial cells; forced expression in non-prostatic cells leads to transdifferentiation to prostate; master regulator for reprogramming to prostate; dysregulation leads to prostate cancer	( <a href="#">Dutta et al. 2016</a> , <a href="#">Taloz et al. 2017</a> , <a href="#">Abate-Shen et al. 2008</a> )
<i>PDX1</i>	Expressed in endodermal cells that give rise to pancreas; forced expression in non-pancreatic cells leads to transdifferentiation to pancreatic cells; reprograms pancreatic exocrine cells to the endocrine cells; dysregulated in a stage-specific manner in pancreatic cancer	( <a href="#">Horb et al. 2003</a> , <a href="#">Zhou et al. 2008</a> , <a href="#">Roy et al. 2016</a> )
<i>PAX4</i>	Expressed in pancreas during development and essential for specification of insulin-producing $\beta$ cells; forced expression in pancreatic progenitor cells promotes their conversion to $\beta$ cells	( <a href="#">Collombat et al. 2009</a> , <a href="#">Sosa-Pineda et al. 1997</a> )
<i>OCT4</i>	Expressed in pluripotent cells in the developing embryo; promotes pluripotency and reprogramming; ectopic expression leads to dysplasia	( <a href="#">Takahashi et al. 2007</a> , <a href="#">Takahashi &amp; Yamanaka 2006</a> , <a href="#">Hochedlinger et al. 2005</a> )
<i>SOX transcriptional regulatory genes</i>		
<i>SOX2</i>	Expressed in pluripotent cells in the developing embryo; promotes pluripotency and reprogramming; forced expression in lung promotes cancer; driver of lineage plasticity in prostate cancers arising from loss-of-function of <i>RB</i> and <i>TP53</i> , particularly in contexts of drug resistance	( <a href="#">Sarkar &amp; Hochedlinger 2013</a> , <a href="#">Takahashi et al. 2007</a> , <a href="#">Takahashi &amp; Yamanaka 2006</a> , <a href="#">Ferone et al. 2016</a> , <a href="#">Mu et al. 2017</a> )

<i>SOX9</i>	Expressed in pancreatic progenitors during development; promotes pancreatic metaplasia; acts coordinately with <i>SOX2</i> to promote EMT in lung cancer	( <a href="#">Kopp et al. 2011</a> , <a href="#">Lin et al. 2016</a> )
<i>SOX11</i>	Expressed during neuronal development; promotes lineage plasticity in prostate cancers arising from loss-of-function of <i>PTEN</i> and <i>TP53</i> , particularly in contexts of drug resistance	( <a href="#">Bergsland et al. 2011</a> , <a href="#">Zou et al. 2017</a> )
Other transcriptional regulatory genes		
<i>KIF4</i>	A member of the Krüppel-like transcription factor family; promotes pluripotency and reprogramming; collaborates with <i>Kras</i> to promote cellular reprogramming in pancreatic cancer initiation	( <a href="#">Takahashi et al. 2007</a> , <a href="#">Takahashi &amp; Yamanaka 2006</a> , <a href="#">Wei et al. 2016</a> )
<i>MITF</i>	Member of the basic helix-loop-helix leucine zipper transcription factor family; required for survival and specification of melanocytes; driver of “phenotype switching” and drug resistance in metastatic melanoma	( <a href="#">Hoek et al. 2008</a> , <a href="#">Konieczkowski et al. 2014</a> )
<i>AR</i>	Essential regulator of prostate differentiation; master regulator for reprogramming to prostate; among the most frequently dysregulated gene in prostate cancer	( <a href="#">Talos et al. 2017</a> , <a href="#">Watson et al. 2015</a> )
<i>Oncogenes</i>		
<i>MYC</i>	Promotes pluripotency and reprogramming; collaborates with <i>Kras</i> to reprogram primary (non-tumorigenic) cells to metastatic cells	( <a href="#">Takahashi et al. 2007</a> , <a href="#">Takahashi &amp; Yamanaka 2006</a> , <a href="#">Ischenko et al. 2013</a> )
<i>KRAS</i>	Cooperates with <i>SOX9</i> in metaplasia of pancreatic adenocarcinoma; cooperates with <i>Myc</i> in reprogramming to metastatic cells	( <a href="#">Kopp et al. 2011</a> , <a href="#">Ischenko et al. 2013</a> )
<i>Tumor suppressor genes</i>		
<i>TP53</i>	Inhibits induced pluripotency and reprogramming; collaborates with <i>RB</i> and <i>PTEN</i> in promoting plasticity in several cancer types, including prostate cancer in contexts of drug resistance	( <a href="#">Marion et al. 2009</a> , <a href="#">Utikal et al. 2009</a> , <a href="#">Yi et al. 2012</a> , <a href="#">Kawamura et al. 2009</a> , <a href="#">Zou et al. 2017</a> , <a href="#">Mu et al. 2017</a> )

<i>RB</i>	Collaborates with <i>TP53</i> in conversion of adenocarcinoma to neuroendocrine differentiation in lung and prostate cancer, particularly in contexts of drug resistance	<a href="#">(Mu et al. 2017, Ku et al. 2017)</a>
<i>PTEN</i>	Collaborates with <i>TP53</i> in transdifferentiation of adenocarcinoma to neuroendocrine differentiation in prostate cancer, particularly in contexts of drug resistance	<a href="#">(Zou et al. 2017)</a>