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FLESHING OUT THE REGULATORY NETWORKS CONTROLLING HUMAN PANCREAS DEVELOPMENT

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SUMMARY

The transcription factor gene *Pancreatic and Duodenal Homeobox 1 (PDX1)* encodes an evolutionarily conserved transcription factor that is expressed during the earliest stages of human pancreatic development. The loss of *PDX1* results in complete pancreatic agenesis, a dramatic phenotype that emphasizes the critical role *PDX1* plays in coordinating the morphogenesis of this indispensable metabolic organ. My laboratory routinely models human pancreatic development *in vitro* by differentiating human pluripotent stem cells into early pancreatic progenitors (ePP). Using a protocol that tightly adheres to developmental logic, abundant *PDX1*+ ePP cells emerge after two weeks of *in vitro* culture. These ePP cells display a molecular signature that significantly overlaps with the developing human pancreatic primordium. In my talk, I will describe how we have used a variety of “omics” and bioinformatic approaches to unveil the gene regulatory network operating directly downstream of *PDX1*. Interestingly, several *PDX1* target genes currently under investigation in the lab are not only essential for pancreatic development but are also redeployed in insulin-secreting beta cells that ensure glucose homeostasis and whose dysfunction leads to diabetes.