Insulin Sensitivity and NO Availability (p 784)

Viswambharan et al investigate the effects of enhancing endothelial cell insulin sensitivity.

Stimulation of endothelial nitric oxide synthase (eNOS) by insulin increases the production of nitric oxide (NO), which is a suppressor of atherosclerosis. Indeed, insulin resistance in type 2 diabetes is associated with both reduced production of NO and increased risk of atherosclerosis. To find out whether boosting insulin sensitivity in ECs could increase NO production and reduce atherogenesis, Viswambharan and colleagues genetically engineered mice to overexpress the human insulin receptor in ECs. Surprisingly, they found that even though insulin stimulation of the engineered ECs prompted Akt phosphorylation (a downstream mediator of insulin signaling), NO availability was reduced. Instead, insulin stimulation increased the activity of Nox2 NADPH oxidase, which in turn ramped-up production of damaging superoxide. The team then examined the EC-specific insulin receptor over-expression in atherosclerosis-prone mice, and found that the animals were even more prone to atherosclerosis than usual, with significantly accelerated plaque development in their aortas. The study suggests that upregulating insulin signaling may not be necessarily a desirable goal for diabetes treatment.

ER71/ETV2-Mediated Reprogramming of HDF to Ecs (p 848)

Lee et al describe a method for generating endothelial cells for therapy, disease analysis, and more.

Because endothelial cells (ECs) are critical for the repair and growth of blood vessels, researchers have been investigating possible ways to generate ECs for treating vascular disorders and injury. However, adult stem and progenitor cells are poor at endothelial cell conversion, carry a risk of tumorigenesis, and are not an easy source of patient-specific ECs. Lee and colleagues, therefore, investigated an alternative idea of converting differentiated somatic cells into endothelial cells directly. Using human postnatal fibroblast cells, the team overexpressed a combination of 7 transcription factors, including eNOS, to drive their conversion into cells resembling endothelial cells and then, by a process of elimination, discovered that just one of these transcription factors—ETV2—was sufficient for the conversion. The reprogrammed endothelial cells were capable of contributing to vessel formation and recovery from hind limb ischemia in mice. Based on these findings, the authors suggest that direct reprogramming of differentiated fibroblasts into endothelial cells could be used for autologous cell therapies as well as personalized disease studies and drug testing.

ATF6 Links ER Stress and Oxidative Stress (p 862)

Jin et al. report that acute stimulation of ATF6 may be beneficial in protecting against ischemia–reperfusion damage. To investigate the mechanism behind this protective effect, Jin and colleagues examined cardiomyocytes lacking ATF6. They showed that on induction of ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced usually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase. The team then showed that when mouse hearts were subjected to ischemia–reperfusion, the cells produced unusually high levels of reactive oxygen species. ATF6 overexpression relieved this effect. Gene expression analysis revealed that ATF6 activated not only ER stress related genes, but also genes encoding antioxidant proteins, including catalase.
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Ruth Williams

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