HDAC4 is necessary for satellite cell differentiation and muscle regeneration.

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In response to injury, skeletal muscle exhibits high capacity to regenerate and epigenetics controls multiple steps of this process (Giordani et al., 2013). It has been demonstrated *in vitro* that completion of muscle differentiation requires shuttling of histone deacetylase 4 (HDAC4), a member of class IIa HDACs, from the nucleus to the cytoplasm and consequent activation of MEF2-dependent differentiation genes (McKinsey et al., 2000). *In vivo*, HDAC4 expression is up-regulated in skeletal muscle upon injury, suggesting a role for this protein in muscle regeneration. With the aim to elucidate the role of HDAC4 in skeletal muscle regeneration, we generate mice lacking HDAC4 in the satellite cells (HDAC4^{1//fl};Pax7^{CE} Cre). Lack of HDAC4 inhibits satellite cells proliferate less and have less pax7 than controls. Importantly, muscle regeneration in vivo is impaired in HDAC4^{1//fl};Pax7^{CE} Cre mice. These results are confirmed by molecular analyses of the expression of myogenic markers. All together, these data delineate the importance of HDAC4 in muscle regeneration and suggest a protective role in response to muscle damage.

References

(1) Giordani L, Puri PL. Epigenetic control of skeletal muscle regeneration: Integrating genetic determinants and environmental changes. FEBS J. 2013 Sep;280(17):4014-25.

(2) McKinsey TA, Zhang CL, Lu J, Olson EN. Signal-dependent nuclear export of a histone deacetylase regulates muscle differentiation. Nature. 2000 Nov 2;408(6808):106-11.

Keywords

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