Morphological and molecular events during H9c2 differentiation: role for pPKCδ/SC35 interaction

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H9c2 cells are rat embryonic myoblasts with skeletal muscle properties which reach differentiation when cultured in HS low concentration (1%). Their terminal differentiation consists of fusing and forming multinucleated myotubes. At ultrastructural level the organization of differentiated cells undergoes modifications in terms of morphological features: they appear thin and elongated, often fused to form multinucleated syncytia (myotubes) and well organized actin filaments are recognizable and detected by immunogold labeling. The decreased expression of cyclin A, required cell cycle regulator for the onset of DNA replication, and the increased expression of myogenin, marker of skeletal muscle differentiation, accompany such morphological modifications.

Among the molecules involved in a cascade of reactions controlling the functioning of nuclear complexes and leading to cell differentiation, a role has been assigned to Protein Kinases C. In our experimental model a ten fold greater signal of pPKC δ in 1%HS cells, with respect to 10%FBS cells, is shown. In order to investigate a potential activation of SC35 splicing factor by pPKC δ , pPKC δ -SC35 co-immunoprecipitation has been performed and revealed only in differentiated cells, supporting a possible interaction between the two molecules. Immunogold co-localization of SC35 factor with pPKC δ , carried out by FEI-SEM and TEM, evidences in 1% HS cells an increased level of pPKC δ and its accumulation in the nucleus, which represents reasonable evidence of protein activation along with an increased expression of SC35. In order to check the specificity of PKC δ involvement in H9c2 differentiation, PKC δ silencing has been performed and both PKC δ and myogenin expression have been detected by immunofluorescence analysis, evidencing a decrease of myogenin expression in 1% HS PKC δ siRNA- transfected H9c2 cells in parallel to the reduced percentage of PKC δ positive cells.

Thus these results suggest that $pPKC\delta/SC35$ interaction represents a crucial event resulting in downstream changes in transcription of components which determine the morphological modifications related to cardiomyocyte differentiated phenotype.