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Forkhead Transcription Factors Coordinate Expression of Myocardial KATP Channel Subunits and Energy Metabolism

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Alex J. Baertschi **Web Site Feature**★e20–e35

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On the cover: Bioinformatics suggests a major role for Fox transcription factors in the expression of SUR2-KIR6.1 gene. Over- and under-expression of Fox in atrial cells confirm the hypothesis (green: FoxO1-EGFP; red: KIR6.1; blue: Dapi). See related article, pages e20–e35.