

Deep-learning based blood cells classification and initial edge device implementation

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Research resource : small RNA-seq of human granulosa cells reveals miRNAs in FSHR and aromatase genes

Velthut, Agne; **Simm, Jaak**; Tuuri, Timo; Tapanainen, Juha S.; **Metsis, Madis**; Salumets, Andres Molecular endocrinology 2013 / p. 1128-1141 : ill