

Cellular, extracellular and extracellular vesicular miRNA profiles of pre-ovulatory follicles indicate signaling disturbances in polycystic ovaries

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International journal of molecular sciences 2020 / art. 9550, 23 p. : ill <https://doi.org/10.3390/ijms21249550> [Journal metrics at Scopus](#)
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Heterogeneity of preovulatory follicle cell types between normo- and hyporesponders

Roos, Kristine; **Rooda, Ilmatar;** **Keif, Robyn-Stefany;** **Liivrand, Maria;** Salumets, A.; **Smolander, Olli-Pekka Aukusti;** **Velthut-Meikas, Agne;** **Liivrand, Maria** Human Reproduction : ESHRE 2022 : abstracts 2022 / art. P-595

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Hsa-mir-548 family expression in human reproductive tissues

Rooda, Ilmatar; **Kaselt, Birgitta;** **Liivrand, Maria;** **Smolander, Olli-Pekka;** Salumets, Andres; **Velthut-Meikas, Agne** BMC

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In-depth analysis of transcriptomes in ovarian cortical follicles from children and adults reveals interfollicular heterogeneity

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Intercellular signalling in the human pre-ovulatory follicle : microRNA expression in granulosa cells and detection in the follicular fluid = Rakkudevaheline kommunikatsioon inimese munasarja preovulatoorses folliikulis : mikroRNA-de ekspressioon granuloosrakkudes ja follikulaarvedelikus

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MicroRNAs expressed from FSHR and aromatase genes target important ovarian functions [Online resource]

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Single-cell RNA-seq analysis and cell-cluster deconvolution of the human preovulatory follicular fluid cells provide insights into the pathophysiology of ovarian hyporesponse

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Target prediction and validation of microRNAs expressed from FSHR and aromatase genes in human ovarian granulosa cells

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Uterine fluid proteins for minimally invasive assessment of endometrial receptivity

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