Identification of the 14-3-3 zeta and epsilon isoforms in mouse estrous uterine fluid

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摘要

關鍵字: endometrial damage、14-3-3 epsilon、14-3-3 zeta、LC/MC/MS、生物指標

Abstract

During the mouse estrous cycle, the endometrium prominently proliferates in proestrus and dramatically degenerates in estrus. Proteins released from the degenerating endometrial cells may not only serve as surrogate markers for endometrial damage potentially measurable in accessible luminal fluids, but the profiles of protein efflux may reflect the underlying mechanisms for exclusion of endometrial cells. To determine the potential biomarker of degenerating endometrial cells, we studied the differential proteins of uterine luminal fluid (ULF) between estrus and proestrus. In comparison of two patterns of ULF constituents resolved by SDS-PAGE, an approximately 30 kDa proteins was predominately shown in estrous ULF distinct from proestrous ULF. The protein band from SDS-PAGE was subjected to tryptic digestion, followed by LC/MS/MS sequences analysis and identified as 14-3-3 epsilon and 14-3-3 zeta from peptide fragments analysis. Immunoblot analysis revealed that zeta forms were specifically increased in ULF at estrus distinct from proestrus. After separation of mouse estrous ULF from Sephacryl S-200 chromatography, two 14-3-3 isoforms were predicted to be a dimmer form, but not co-fractionated in the gel filtration fractions. In addition, the 14-3-3 epsilon was not co-immunoprecipitated with anti-14-3-3 zeta antibody from the same fraction. Taken together, this study has determined that the analysis of 14-3-3 zeta levels from the estrous ULF could be useful as a biomarker for mouse endometrial damage.

Keywords: endometrial damage · 14-3-3 epsilon · 14-3-3 zeta · LC/MC/MS · biomarker