



Expression of oestrogen receptor α and β in uterine endometrial and ovarian cancers

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Oestrogen, as well as the progestins [1], is recognised as a significant modifier of the growth, development, invasion and metastasis of uterine endometrial cancers and ovarian cancers. Most uterine endometrial cancers in their early stages conserve sex steroid hormone dependency. Together with angiogenic factors [2,3] the expression patterns of sex steroid receptors such as the ER (oestrogen receptor) and PR (progesterone receptor) are extremely important [4,5] and are related to patient prognosis in other gynaecological cancers. Novel human *ER β* was identified in cDNA libraries from human testis in 1996 [6]. *ER β* has a different expression pattern to classical *ER α* . *ER β* may not have the same physiological functions as *ER α* . Therefore, *ER α* and *ER β* mRNA expression patterns in oestrogen-dependent cancers, uterine endometrial cancers and ovarian cancers were investigated by competitive reverse transcription-polymerase chain reaction (RT-PCR) with recombinant *ER α* and *ER β* mRNA. The level of *ER β* mRNA was significantly lower than those of *ER α* mRNA in uterine endometrial cancers, ovarian cancers and the corresponding normal tissues. The level of *ER β* was regulated, like those of *ER α* in normal uterine endometria, with a peak in the periovulatory phase during the menstrual cycle ($n=20$) and the ratio of *ER β* mRNA to *ER α* mRNA was stable during this cycle [7]. The ratio in most primary uterine endometrial cancers ($n=60$) was similar to that in normal uterine endometria ($n=20$). In contrast, in 14 of the 20 lymph node metastasis-positive cases of uterine endometrial cancers, the ratio in the metastatic lesion was significantly higher than that in the primary lesion of the corresponding case, and the patient's prognosis in these cases was extremely poor [8]. Immunohistochemical staining for *ER α* and *ER β* revealed that *ER α* and *ER β* were expressed predominantly in the nuclei of the cancer cells

of the primary and metastatic lesions of uterine endometrial cancers and in the glandular cells of normal uterine endometria. Semi-quantitative analysis by immunohistochemical staining showed that there was no discrepancy between *ER α* and *ER β* mRNA and protein levels [8]. The ratio of *ER β* mRNA to *ER α* mRNA in ovarian cancers ($n=42$) had a wider range than ratios in the normal ovaries ($n=21$). The patient's prognosis in ovarian cancers with a low or a high ratio of *ER β* mRNA to *ER α* mRNA was worse than that in ovarian cancers with a medium ratio [9]. Therefore, it is suggested that the intact synchronised expression of *ER β* interacting with *ER α* might be disrupted in most metastases of uterine endometrial cancers and some ovarian cancers, leading to advancement of the cancer and a poor patient prognosis, especially in uterine endometrial cancers that are refractory to hormonal treatments.

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