

Summary and Conclusion

It is widely recognized that endometrial carcinoma(EC) represents the most frequent type of genital malignancy in women. Endometrial hyperplasia(EH) has been linked with endometrial carcinoma for many years and it has since long been proposed to be a predecessor to endometrial carcinoma.

The existence of a high risk subgroup among EH patients is a well established and accepted fact today. The crucial problem is the definition of this high risk population of EH patients.

The aim of our study was to evaluate recent molecular markers: DNA ploidy status, DNA index (DI), S-phase (SPF) and c-erbB-2 oncoprotein; also biological marker CEA to identify their value in classifying endometrial hyperplasia into low and high risk groups and their possible role in malignant transformation.

The study group consisted of 93 women and the endometrial specimens were histopathologically classified into normal group (n = 36) (16 secretory type and 20 proliferative type), hyperplastic group (n = 50) (32 simple type and 18 complex one) and malignant group (n = 7).

The mean \pm SD of age for the normal group was 41.6 ± 8.3 years for the hyperplastic 46 ± 6.1 years and for malignant cases 57 ± 9.1 years (the difference between the groups was statistically significant P < 0.001).

DNA analysis was made on fresh endometrial samples by FCM. All the normal endometrial tissues (secretory or proliferative) were DNA diploid (DI = 0.9-1.1), also all cases of simple hyperplastic group showed no abnormality of DNA ploidy pattern, whereas in the complex hyperplastic group one case (6%) was DNA aneuploid (DI = 1.7) which may represent a high risk case of an increased malignant potential. For the malignant cases 1 case (14%) was diploid while the majority (86%) were DNA aneuploid (DI > 1.1).

All the seven non-diploid cases were above 40 years old and six of them were postmenopausal.

The mean \pm SD SPF for the malignant cases (13.73 \pm 6.44) was significantly higher than the normal (5.54 \pm 1.73) and the hyperplastic (5.63 \pm 2.50) groups (P < 0.01), while that of hyperplastic was not significantly different from normal (P>0.05). Also non significant difference was obtained between mean SPF of normal secretory (5.28 \pm 1.37) and normal proliferative (5.76 \pm 1.9) or simple hyperplastic (5.23 \pm 2.28) and complex hyperplastic (6.32 \pm 2.66) subgroups P>0.05).

The highest value for SPF in normal cases was 9, i.e. all normal cases ≤ 9 , whereas 12 % of hyperplastic group were > 9; 2 % > 10 and no cases > 15. In the malignant group 71 % were > 9; also 71 % > 10

while 29 % > 15 which was statistically significant as revealed by X^2 test.

Regarding c-erbB-2 oncoprotein expression measured in the endometrial tissue by EIA technique, the mean \pm SD (HNU/µg protein) in the malignant group (3.74 \pm 2.45) was significantly higher than normal group (1.47 \pm 0.84) ,P < 0.05; but not significantly different from the hyperplastic group (2.4 \pm 2.46) P > 0.05.

The highest value for *c-erbB-2* oncoprotein in the normal cases was 3.46 HNU / μ g protein, 8 hyperplastic cases (16%) were > 3.46 HNU / μ g protein and 3 malignant cases (43%) were > 3.46 HNU / μ g protein (P<0.01). Using a cut off value for *c-erbB-2* oncoprotein = 3.15 HNU/ μ g protein (mean + 2 SD) 1 (3%) normal case was > 3.15 while 10 (20%) hyperplastic cases and 4 (57%) malignant cases were > 3.15 HNU / μ g protein, a statistically significant difference (P < 0.01). From the 15 cases > 3.15 HNU / μ g protein 9 (60%) premenopausal and 6 (40%) postmenopausal; also one (7%) was \leq 40 years and 14 (93%) were > 40 years.

Concerning serum CEA the mean \pm SD (ng/ml) of normal group (3.72 \pm 0.54), hyperplastic (3.75 \pm 0.60) and malignant (3.83 \pm 0.17) which was a non significant difference.

Non significant correlation was found between age, parity, c-erbB-2, CEA, and SPF in the normal group while in the hyperplastic group only a positive significant correlation between parity and SPF was detected.

Whereas in the malignant group significant positive correlation between tumor grade with both DI and SPF also between DI and SPF.

From the data of the present study we could conclude that: FCM of cellular DNA content is a rapid, objective, quantitative and sensitive method to determine a highly specific and stable tumor cell marker. The utility of this technology in the evaluation of premalignant and malignant tissues (e.g. EH and EC) is becoming increasingly recognized. DNA analysis data by FCM may be useful for selecting a subset of EH patients with high risk of developing EC.

FCM DNA measurements (ploidy status, DI, and SPF) might add information independent of clinical and histopathological examination and might help the clinician about the decision and way of management of some cases of EH.

A major advantage of FCM measurements is that they can be done on fresh or stored (paraffin embedded) tissues as well as they are much less time consuming than other methods of DNA analysis.

To ascertain whether aneuploid EH cases represent definitely a premalignant lesions, the DNA results should be correlated with the final histologic and clinical follow up of patients.

In this era of rapid acceleration of knowledge of molecular biology in the oncologic sciences, we can expect growth factors and oncogenes to take a prominent place in molecular epidemiology and the assay of tissue and tumor virulence. These virulence factors should reduce our

complacency about the ease of curability of EH and EC and sharpen our interest in individualizing treatment of these diseases.

Measurements of c-erbB-2 oncoprotein might help to detect more risky cases of EH since alteration in the expression of c-erbB-2 oncoprotein might play an important role in malignant transformation.

Whether *c-erbB-2* proto-oncogene activation is actually involved in the transformation from normal to neoplastic endometrium remains to be proved. However, the presence of the *c-erbB-2* product in the hyperplastic endometrial tissues in amount higher than normal endometrial tissue and near to that of malignant tissue support this hypothesis in a subset of EH populations.

A much larger numbers of patients must be studied to determine the role and prognostic significance of *c-erbB-2* (and other) oncogene(s) amplification and/or overexpression in normal, hyperplastic and malignant endometrial tissues.

Clinically, this study indicates that molecular markers and biologic data must be taken into account in the management of patients with EH and EC. The more we understand about the molecular biology and markers, not only in malignant tissues, but also in normal and premalignant tissues, the closer we will come to using molecular markers to their potential in diagnosis, prognosis and therapeutic management of cancer patients.

Ultimately, our future treatment might include a form of gene therapy or be directed at inhibitors of growth factors and oncoproteins that might have some role in malignant transformation.