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<tr>
<td><strong>Citation</strong></td>
<td>Journal Of Clinical Pathology, 1993, v. 46 n. 3, p. 204-207</td>
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<tr>
<td><strong>Issued Date</strong></td>
<td>1993</td>
</tr>
<tr>
<td><strong>URL</strong></td>
<td><a href="http://hdl.handle.net/10722/45490">http://hdl.handle.net/10722/45490</a></td>
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Expression of c-myc and c-fms oncogenes in trophoblastic cells in hydatidiform mole and normal human placenta

A N Y Cheung, G Srivastava, S Pittaluga, T K Man, H Ngan, R J Collins

Abstract

Aims: To compare the expression of c-myc and c-fms proto-oncogenes in the placenta and hydatidiform mole. Methods: Twelve hydatidiform moles and six induced abortion cases were collected. c-myc and c-fms proto-oncogene expression was analysed by northern blot hybridisation and immunohistochemical staining. Results: The results of northern blot hybridisation analysis showed that c-fms was expressed more strongly in hydatidiform moles compared with normal placenta of similar gestational age. Moreover, c-fms mRNA concentrations increased with more advanced gestational age in moles but not in normal placentas. c-myc expression was very low in hydatidiform moles and normal placentas. Both oncogenes, however, had no direct correlation with the clinical course of the molar pregnancies. Conclusion: The difference in c-fms expression between hydatidiform moles and normal placentas suggests that c-fms may have a role in the development of molar pregnancies.

(J Clin Pathol 1993;46:204–207)

The trophoblast of early placenta can proliferate rapidly and infiltrate the decidua and myometrium at the placental site.1 Expression of certain cellular proto-oncogenes may be associated with rapid cell growth in these early trophoblast. A strong correlation between c-myc transcript concentrations and trophoblastic proliferation has been reported.2 It has also been suggested that colony stimulating factor-1 (CSF-1), a glycoprotein growth factor required for the proliferation and differentiation of mononuclear phagocytic cells, may have a role in placental development and function under hormonal influence.3 The c-fms proto-oncogene product is related to the receptor for the CSF-1,4 but in situ hybridisation studies indicate that c-fms mRNA is associated with trophoblast in human placenta.5 The pathogenesis of hydatidiform moles has remained a mystery for decades. The difference between physiological oncogene expression in early placenta and that seen in molar pregnancies is unknown. It would also be useful to ascertain whether there is a particular pattern of oncogene expression in moles which will later progress into invasive moles or chorionic carcinoma. This study aimed to investigate the expression of c-myc and c-fms proto-oncogenes in hydatidiform moles compared with that in normal placenta.

Methods

Twelve clinically and ultrasonographically suspected cases of hydatidiform mole and six legally induced abortions were suction evacuated. Fresh chorionic villi and molar vesicles were selected and snap frozen in liquid nitrogen and stored at −70°C. The rest of the tissue was fixed in formalin for histological examination.

Human HL-60 promyelocytic leukaemia cell line6 and human HUT78 T cell lymphoma cell line were maintained in logarithmic growth phase; total RNA was extracted and used as positive controls for c-myc and c-fms expression, respectively.

The probes used were c-myc-exonIII (1-8 kilobase insert) from Oncor (Gaithersburg, USA) and c-fms cDNA clone pCfms 104 in PUC12 (1-23 kilobase insert) from American Type Culture Collection (Maryland, USA). The cDNA probes were labelled by an oligolabelling procedure (Bresatec, Australia).

Total cellular RNA was extracted by the guanidine hydrochloride procedure and degraded RNA was obtained from eight moles and six abortion cases.7 Haematoxylin and eosin stained sections of each frozen tissue block were examined to exclude contamination by decidua and to ensure roughly equal numbers of molar villi in the tissue block. Aliquots of total RNA (20 μg) from each sample were denatured and subjected to electrophoresis on 1-0% agarose gels containing 1-1M formaldehyde. Equivalent amounts of RNA were loaded into each lane as judged by ethidium bromide staining. RNA was then transferred to nylon membranes (Gene Screen Plus, NEN, Du Pont, Boston, Massachusetts, USA). The membranes were prehybridised in a solution containing 50% formamide, 1M sodium chlo-
ride, 10% dextran sulphate, 1% sodium dodecyl sulphate (SDS), 0.2% polyvinylpyrrolidone, 0.2% ficoll, 0.2% bovine serum albumin, 50 mM TRIS-HCl (pH 7.5), 0.1% sodium pyrophosphate, 0.1 mM EDTA, and denatured salmon sperm DNA (200 μg/ml) at 42°C for overnight. The membranes were hybridised to the denatured 32P-oligolabelled c-myc and c-fms DNA probes (0.5–2 ng/ml) in the same solution for 24 hours. The final wash of the membranes was in a solution containing 0.1 x SSC (30 mM sodium chloride, 3 mM sodium citrate), 1% SDS, 0.1% sodium pyrophosphate, and 0.1 mM EDTA at 65°C for 60 minutes. The membranes were also stripped and reprobed with 28S ribosomal RNA oligomeric probe to ensure that equal amounts of RNA were loaded in each lane and that the transfer was quantitative. Autoradiography was carried out using X-OMAT AR x-ray film (Kodak) with intensifying screens at -70°C.

Signals were quantitated by densitometric scanning of the autoradiograms.

Frozen sections (6 μm) of the stored frozen samples were cut, air-dried, and fixed in 4% paraformaldehyde. The sections were stained with mouse monoclonal antibody to c-myc oncoprotein and rabbit polyclonal antibody to c-fms oncoprotein (Cambridge Research Biochemicals, England) and monoclonal mouse antibody to human Ki-67 antigen (Dako, Copenhagen, Denmark). The alkaline-phosphatase anti-alkaline-phosphatase method, as described by Cordell, was used. Negative controls were used by omitting the primary antibody in each case. Positive controls were obtained by cytoxin of the human HL-60 and HUT-78 cell lines as well as by including a lymphoma case with a high proliferative rate.

Results
Histological examination showed that all the moles collected were complete moles. There were no detectable differences in the accumulation of macrophages in the moles and placentas examined. The gestational ages of the hydatidiform moles thus selected for analysis ranged from six to 16 weeks while the abortion cases ranged from five to 16 weeks. In the cases studied two hydatidiform moles had persistently raised serum concentrations of human chorionic gonadotrophin (hCG) during follow up and single agent chemotherapy was given.

The results of northern blot hybridisation analysis showed that higher mRNA concentrations of c-fms were found in hydatidiform moles compared with those in normal placenta of comparable gestational age (fig 1). Moreover, c-fms expression increased with more advanced gestational age in moles, but in normal placentas the c-fms concentration reduced with increased gestational age. Densitometric measurements of the relative c-fms RNA expression in normal placentas and hydatidiform moles is shown in fig 2. Immunohistochemical staining carried out with the anti-c-fms antibody clearly localised c-fms antigen expression to the syncytiotrophoblast with occasional positive staining in the Hofbauer cells in the molar villi (fig 3). There was no detectable accumulation of macrophages in the sections. No correlation between c-fms expression and serum concentrations of hCG during follow up and subsequent chemotherapy in the molar pregnancies studied could be demonstrated.

Contrary to previous reports, c-myc expression was very low in the cases of moles and normal placentas studied except in one case of induced abortion at five weeks' gestation (fig 4). However, the total RNA extracted from the HL60 cell line showed high concentrations of c-myc mRNA. The Ki67 indices were less than 5% in all the placentas and molar pregnancies studied.

Discussion
Using northern blot and in situ hybridisation, Pfeifer-Ohlsson et al, have demonstrated the
expression of human c-myc proto-oncogenes in early human villous cytotrophoblast with a peak expression at five weeks after conception and a deline by the end of the first trimester of pregnancy related to differentiation of cytotrophoblasts into syncytiotrophoblasts. The low expression of c-myc in the normal placentas analysed in our study corresponds to the advanced gestational age of the studied cases—five to 16 weeks. On the other hand, the low expression of c-myc in the moles we studied is contrary to the findings of previous studies. Sarkar et al have demonstrated c-myc and c-ras proto-oncogene expression by in situ hybridisation in cytotrophoblasts of six hydatidiform moles. The gestational age of the moles studied was not specified. Yokoyama et al have also reported the expression of c-myc, c-fms, and c-sis proto-oncogenes in one mole at the 10th week of gestation by northern blot analysis and in situ hybridisation.

While such low c-myc expression in our study may have been related to the comparatively advanced gestational age of the moles studied—six to 16 weeks—it may be important to the pathogenesis of hydatidiform mole: this has always been controversial. Park attributed the primary disturbance to abnormalities in the trophoblast, resulting indirectly in death of the fetus, but Hertig and Edmonds proposed early death of the embryo as the initiating process. Fox also proposed that moles were simply a form of abortion and not neoplastic. Suressh et al, from their comparative study of silver stained nucleolar organiser region counts on complete and incomplete hydatidiform moles as well as on missed abortions, supported the latter view. Flow cytometric studies also suggested that the correlation of the clinical course with proliferative index was not significant while the correlation with ploidy was positive. Thus the low c-myc expression in the moles we studied favours the hypothesis that most hydatidiform moles are a form of abortion due to abnormal conceptions, rather than a true neoplasia as c-myc is usually high in actively proliferating cells. This is further supported by the low growth fraction determined by anti-Ki67 antibody.

Figure 3  Immunohistochemical staining with anti-fms antibody in hydatidiform mole to localise c-fms antigen expression in syncytiotrophoblast.

Figure 4  Representative northern blot hybridised with 32P labelled c-myc cDNA probe. Total RNA was isolated from normal placenta at five, eight, nine, 10, and 16 weeks of pregnancy and from hydatidiform moles at six, nine, 10, 12, and 16 weeks' gestational age. The transcript of c-myc was detected in the range of 2-4 kilobases in the placenta at five weeks' gestation. The human HL-60 leukaemia cell line RNA was used as the positive control. The positions of 18S and 28S ribosomal RNA are indicated. The results of the probing of 32P-labelled 28S ribosomal RNA is shown in the lower panel B.
Pregnancy induces a 1000-fold increase in the murine uterine CSF-1 concentration, which appeared to be regulated by chorionic gonadotrophin,18 and the synergistic action of the female sex hormones, oestradiol and progesterone. c-myc expression in murine placenta peaks early throughout gestation to plateau on the 18th day.16 Moreover, the expression of the CSF-1 receptor and the c-fms proto-oncogene product has been demonstrated in previous studies.19 20 Simultaneous expression of α-subunits of hCG was also observed in the rat placenta. The c-fms transcript has been localised to the syncytiotrophoblast of hydatidiform moles and chorionic carcinoma, where hCG is produced.13 Hence, c-fms may have a role in placental development and may be related to hCG metabolism. In our study the highest c-fms expression found in the normal placentas was at the 5th week of gestation; there was a steady decline in its expression in placentas of more advanced pregnancies. On the other hand, a parallel increase in c-fms mRNA concentration with advancing gestational age of moles was observed. However, there is no demonstrable relation between the hCG activity at diagnosis and the intensity of c-fms expression in this study. The patients with hydatidiform moles were followed up with an estimation of hCG activities and two of them received treatment for persistently high hCG concentrations. Both oncogenes did not seem to have a direct effect on the clinical course of molar pregnancies. This difference in the intensity of c-fms expression between normal placentas and moles suggests that c-fms may have a role in the development of hydatidiform moles.

This study was supported by grants from the Committee on Research and Conference Grant (335/046/0035) and the Medical Faculty Fund (362/030/3410).