Syrian hamster embryo cell lines useful for detecting transforming genes in mouse tumours: detection of transforming genes in X-ray-related mouse tumours

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Summary  The Syrian hamster embryo cell lines, SHOK and MC-1, were used as recipient cells for DNA transfection assay to detect transforming genes in experimental mouse tumours. A mouse repeat sequence was utilised to check whether each transformed focus included mouse genomic DNA in the Hamster background. We investigated five mouse tumours that are related to X-ray radiation, and detected activated c-K-ras, c-mos, and c-cot oncogenes which induced foci of hamster cells. These results show that SHOK and MC-1 cells have unique properties for detecting transforming genes in experimental mouse tumours.

Development of the mouse NIH3T3 transfection assay has led to the identification of several oncogenes in human tumours and animal cells transformed by chemical carcinogens (Parada et al., 1982; Shimizu et al., 1983a; Sukumar et al., 1983). Although the roles of such oncogenes in multi-stage carcinogenesis are not fully understood, we are interested in detecting transforming genes involved in the mouse tumours which are related to X-ray-radiation, to obtain some insight about the role of murine oncogenes. In this experimental system, the descendants of the irradiated mouse N5 strain tend to develop tumours and their susceptibility to tumour formation is inherited as an autosomal dominant trait as if it were induced by germ-line mutation (Nomura, 1982; 1983; 1986). It remains unknown, however, what kind of genetic changes are involved in developing tumours in these mice and whether activated oncogenes are actually present in these tumours or not.

NIH3T3 cells are commonly used for detecting oncogenes because of their high efficiency of transformation by oncogenes. Since mouse-specific repeated DNA sequences do not serve as molecular markers of foreign DNA in NIH3T3 cells, we have tried to find recipient cells of non-mouse origin. Although there are other transfection systems using non-mouse origin cells such as 4DH2 cells (Shiner et al., 1988), their sensitivities against oncogenes are not as good as NIH3T3. We developed the hamster SHOK and MC-1 cell lines whose sensitivity against oncogenes is comparable to NIH3T3 cells.

Here we report the novel properties of these hamster cells as indicator cells for the focus-forming assay and detection of the activated oncogenes in tumours formed in N5 mice exposed to X-rays and cancer prone descendants whose parents’ germ line were irradiated with X-rays.

Materials and methods

Cell culture

Syrian hamster embryo fibroblast cell lines, SHOK and MC-1 cells, were grown in Dulbecco’s modified minimum essential medium (D-MEM) supplemented with 10% foetal calf serum. NIH3T3 cells were grown in D-MEM supplemented with 10% calf serum.

DNA transfection assay

High molecular weight DNAs extracted from mouse tumours were transfected into NIH3T3, SHOK and MC-1 cells by the calcium phosphate coprecipitation method (Wigler, 1978). After transfection, NIH3T3 cells were maintained in D-MEM supplemented with 5% calf serum, SHOK and MC-1 cells were maintained in D-MEM supplemented with 3% foetal calf serum. The foci were scored 21 days after transfection. DNA samples examined were as follows. Genomic DNA: T24 for c-H-ras (Sukumar et al., 1983), HL-60 for N-ras (Murray et al., 1983), CC-013 for c-K-ras (a NIH3T3 transfromant from colon cancer tissue; Sasaki, unpublished work). Cloned plasmid (Tsuda et al., 1981), pEJRas for c-H-ras (Shih et al., 1982), pSVneo-fgr for GF-FeSV (Miyoshi et al., 1989), pZLP62B for AEV (Aoki, unpublished work), pSR2 for RSV (DeLorbe et al., 1980), AFPS2 for FSV (Shibuya et al., 1982), AAB3 for Abelson MLV (Goff et al., 1980), pSVSL-1L for Moloney-MSV (Beveren et al., 1981) and pC60 for SVV (Gelmann et al., 1981).

Molecular hybridisation of transformed work

High molecular weight DNAs were digested with restriction endonuclease and fractionated by electrophoresis in 0.7% agarose gel (Maniatis et al., 1982). Ten micrograms of denatured total RNA were fractionated by electrophoresis in 1.0% agarose formaldehyde gels (Perbal, 1988). DNA and RNA samples were transferred to a nylon membrane filter and hybridised to the following probes: p 014 (MIF/Bam/R/B1 superfamily sequence) for the mouse repetitive sequence (Fujimoto et al., 1985), HindIII fragment (880 bp) of Ha-MSV for Ha-ras (Hager et al., 1979), R fragment (1 kb) for N-ras (Shimizu et al., 1983b), HIIHI 380 fragment for K-ras (Ellis et al., 1981), BalI-HindIII fragment (790 bp) of Mo-MSV-HT-1 for mos (Yande Woude et al., 1979) and EcoRI-HaeIII fragment (170 bp) of cot c-DNA (Miyoshi et al., 1991). The DNA probes were labelled by the multiprimer extension method to give 10 c.p.m. µg⁻¹ DNA. The filters were hybridised at 65°C over night in hybridisation buffer containing 6 x SSC, 5 x Denhardt, 0.1% SDS, 20 µg ml⁻¹ salmon sperm DNA. The filters were washed in 2 x SSC, 0.1% SDS at 65°C for 30 min, followed by washing with 0.2 x SSC and 0.1% SDS for 10 min. They were exposed to an X-ray film at -70°C with an intensifying screen.

Results

New cell lines for detecting murine oncogenes

We have established a hamster cell line SHOK (Syrian Hamster Osaka, Kanazawa) from a GHE-L strain of Syrian hamster embryo cells (Higashi et al., 1990). The GHE-L
strain was shown to be transformed by an activated c-H-ras oncogene by Suzuki et al., but it was found to be transformed less efficiently than NIH3T3 cells and to have a tendency to generate spontaneous foci due to overgrowth. SHOK cells were isolated as a subclone, which do not exhibit the disadvantages observed with GHE-L cells and were used for focus-forming assay satisfactorily. The MC-1 (Myc-Clone-1) cell line is one of the transfectedants of the SHOK cell line obtained with pSV2gpt-c-myc containing the second and third exons of the mouse c-myc (Land et al., 1983) which was expected to have higher transforming ability than the SHOK cell line. Table I summarises the focus-forming ability of various oncogenes in SHOK, MC-1 and NIH3T3 cells. All recipient cells were transformed by tumour cellular DNAs containing activated H-, N- and K-ras oncogenes. Typical examples presented in Figure 1 indicate that spontaneous overgrowth is a rare event in SHOK and MC-1 cell lines. Both hamster cells were more sensitive than NIH3T3 cells to N- and K-ras. However, both SHOK and MC-1 cells showed quite different properties from those of NIH3T3 cells with regard to transformation by other viral oncogenes. Whereas hamster cells were transformed less efficiently than NIH3T3 cells by v-fgr, v-erbB, v-src, v-fps and v-abl which are all classified into tyrosine-specific protein kinases, they showed about ten times higher efficiency than NIH3T3 cells to transformation by v-mos, a serine/threonine-specific protein kinase. Since the ability of incorporating foreign DNAs and promoter/enhancer activity of viral LTRs were almost at equal levels in all these cell lines (data not shown), these differences in protein kinase-mediated transformation efficiency between mouse and hamster cells might be attributed to species-specific cellular factors which respond to oncogenic function of the protein kinase family.

Although we obtained no significant differences between the SHOK and MC-1 cell lines, we used MC-1 as well as SHOK cells for further experiments as indicator cells in transfection assays, because MC-1 has a lower background of overgrowth than SHOK cells.

| Genomic DNA* | T24 (c-H-ras) | 16.0* | 20.0 | 36.0 |
| Cloned DNA* | NIH3T3 | SHOK | MC-1 |
|---|---|---|---|
| c-H-ras (pT24-ras) | 65.5 | 200+ | 200+ |
| v-fps (FSV) | 23.5 | 2.5 | 16.5 |
| v-abl (Ab-MLV) | 21.5 | 0 | 0 |
| v-mos (Mo-MSV) | 4.0 | 51.0 | 47.0 |
| v-sis (SSV) | 6.0 | 0 | 0 |
| control (carrier only) | 0 | 0 | 0 |

*Ten μg cellular DNA was transfected into 100 mm dish seeded with 5 × 10^5 cells. Fifty ng plasmid or phage DNA with 10 μg carrier DNA was transfected into 60 mm dish seeded with 2 × 10^5 cells. The values are the means for duplicate culture dishes (the number of foci/dish)

| Tumour strain | Generation* | Histopathology |
|---|---|---|
| N5-2057 | male parent | Osteosarcoma |
| N5-BMX-2710 | F3 | Multiple undifferentiated tumour* |
| N5-BMX-3013 | F3 | Rhabdomyosarcoma |
| N5-BMX-2596 | F2 | Multiple undifferentiated tumour* |
| N5-BMX-2305 | F1 | Fibrosarcoma |

*Generation in which tumours developed. F1 to F3 are descendants of the irradiated male mouse. *Difficult to specify the histological type because of the undifferentiated morphology of the tumour cells.

Figure 1 Giemsa staining of SHOK, MC-1 and NIH3T3 cells transfected by the activated c-H-ras gene (T24-ras). a and d, MC-1 cells; b and e, SHOK cells; c and f, NIH3T3 cells. The upper dishes a, b and c contain several transformed foci and the lower dishes d, e and f are mock-transfected negative controls. Cells were fixed in ethanol and stained with Giemsa solution 21 days after transfection.
Transfection of mouse tumour DNAs into hamster cells

As shown in Table II, cellular DNAs were obtained from transplantable tumours derived from a non-heritable tumour formed in the male N5 mouse irradiated with 504 rad of X-rays (Nomura, 1986) and four heritable tumours formed in the F1 to F3 cancer prone descendants. We transfected tumour DNAs into three recipient cells and scored the number of transformed foci (Table III). In primary transfection, DNAs extracted from a non-heritable tumour N5-2057 and two heritable tumours, N5-BMX-2710 and N5-BMX-3013, showed transforming ability in SHOK and MC-1 cells (Figure 2). The N5-BMX-2710 tumour DNA scored one focus in NIH3T3 cells as well. DNAs extracted from the primary foci were confirmed to induce transformation of all the recipient cells examined in the second cycle of transfection. The other two heritable tumours, N5-BMX-2305 and N5-BMX-2596, were both negative in primary and secondary transfections.

Detection of transforming genes

DNA extracted from hamster cells transformed by transfection of mouse tumour DNAs were analysed for the presence of the mouse repetitive sequence using Southern blots probed with p 014. All primary transformants except for SHOK 2710 f-1 were found to have mouse repetitive sequences (Figure 3). Although SHOK 2710 f-1 derived from the heritable tumour did not show a clear hybridisation pattern, its DNA was confirmed to induce transformed foci in the secondary and tertiary transfection. These results suggest that mouse repetitive DNA sequences were not closely associated with the location of the responsible transforming gene in SHOK 2710 f-1.

We then analysed whether DNAs of these three transformants contained additional bands of mouse origin which might be homologous to several known oncoregene sequences under the background of hamster cellular DNA. We found that SHOK 2057 f-1, the transformant from the X-ray-induced osteosarcoma showed an extra band (about 12 kb EcoRI fragment) which hybridised to mos (Figure 4a). In contrast, other transformants induced by the same tumour DNA, MC-1 2057 f-11 and 12 clearly had 16 kb, 3.7 kb and 1.6 kb HindIII fragments which hybridised to the K-ras probe (Figure 4b). These additional bands homologous to the mouse c-K-ras gene were also found in the other three MC-1 transformants derived from N5-2057 (data not shown).

Northern blot analysis of the mos transformant (SHOK 2057 f-1) shows that mos is expressed at high levels (Figure 5) and further analysis of genomic structure of mos gene in this transformant shows that the small part of N-terminal coding region of this gene was replaced with a hamster genomic

| Donor DNA          | NIH3T3 | Recipient cells | SHOK | MC-1 |
|--------------------|--------|-----------------|------|------|
| Primary transf.⁴   |        |                 |      |      |
| N5-2057            | 0/340 µg | 1/310 µg        | 5/90 µg |      |
| N5-BMX-2710        | 1/280 µg | 1/420 µg        | 0/90 µg |      |
| N5-BMX-3013        | 0/220 µg | 1/220 µg        | 0/90 µg |      |
| N5-BMX-2596        | 3/280 µg | 0/380 µg        | 0/90 µg |      |
| N5-BMX-2305        | 0/310 µg | 0/160 µg        | ND²   |      |
| Secondary transf.⁵ |        |                 |      |      |
| N5-2057            | 111/45 µg | 129/45 µg       | ND    |      |
| MC-1 2057 f-11     | ND      | 57/100 µg       | ND    |      |
| MC-1 2057 f-12     | ND      | 79/100 µg       | ND    |      |
| N5-BMX-2710        | 8/430 µg | 2/90 µg         | ND    |      |
| NIH 2710 f-1       | ND      | ND              | ND    |      |
| SHOK 2710 f-1      | ND      | 4/60 µg         | ND    |      |
| N5-BMX-3013        | 6/90 µg | 43/60 µg        | ND    |      |
| SHOK 3013 f-1      | ND      | 43/60 µg        | ND    |      |
| N5-BMX-2596        | ND      | 0/190 µg        | ND    |      |
| NIH 2596 f-1       | 0/60 µg | 0/300 µg        | ND    |      |
| NIH 2596 f-2       | 0/60 µg | 0/175 µg        | ND    |      |
| NIH 2596 f-3       | 0/40 µg | 0/175 µg        | ND    |      |
| Tertiary transf.⁶  |        |                 |      |      |
| N5-2057            | 9/60 µg | >200/60 µg       | >200/60 µg |      |
| SHOK 2057 f-1-1    | 12/90 µg | ND              | ND    |      |
| SHOK 2710 f-1-4    | ND      | 8/60 µg         | 8/60 µg |      |

⁴Total cellular DNAs extracted from tumours of the N5 mouse and its progeny were transfected and scored by number of foci per µg donor DNA. ²Not determined. ³DNAs were extracted from primary transformants and transfected into mouse or hamster recipient cells.
sequence which has strong promoter activity (Ouchi et al., 1992). Therefore this activation of the mos gene was probably generated artificially during the step of transfection.

As for the two hamster transformants obtained from heritable mouse tumour DNAs, probes of the ras family failed to hybridise to any exogenous restriction fragments of both SHOK 2710 f-1 and SHOK 3013 f-1. While we did not perform further characterisation of the 'non ras' transforming gene in SHOK 2710 f-1, the other transformant SHOK 3013 f-1 was found by molecular hybridisation to contain the cot oncogene, a member of serine-specific protein kinases, which we have recently isolated from human thyroid cancer using the SHOK transfection assay (Miyoshi et al., 1991). Figure 4c shows that the approximately 4 kb EcoRI fragment, which hybridised to the cot probe and corresponded in size to the endogenous mouse cot restriction fragment, was present in the SHOK 3013 f-1 transformant (lane 3).

Figure 3 Detection of mouse repetitive sequences in SHOK and MC-1 cells transformed by X-ray-induced mouse tumour DNAs. Total cellular DNAs extracted from transformed cells were digested with EcoRI, electrophoresed in 0.7% agarose gel, transferred to a nylon membrane and hybridised with 4.8 kb HindIII fragment of p 014 containing the mouse repetitive sequence. Lane 1, SHOK; lane 2 to 8, SHOK and MC-1 primary transformants. (lane 2, SHOK 2057 f-1; lane 3 to 6, MC-1 2057 f-11 to f-14; lane 7, SHOK 2710 f-1; lane 8 SHOK 3013 f-1).

Figure 4 Analysis of mos, K-ras and cot sequence in the DNAs of SHOK and MC-1 cells transformed by X-ray-induced tumour DNAs. Ten micrograms of each DNA was digested with EcoRI a and c or HindIII b, electrophoresed in 0.7% agarose gel, transferred to a nylon membrane and hybridised with the following probes: a, the 0.79 kb BalI-HindIII fragment of v-mos; b, the HiHi 380 fragment of v-K-ras, and c, the EcoRI-HaeIII fragment of cot c-DNA. a, Lane 1, mouse NIH3T3 DNA; lane 2, hamster SHOK DNA; and lane 3, SHOK 2057 f-1 DNA. b, Lane 1, mouse NIH3T3 DNA; lane 2, hamster SHOK DNA; and lane 3 to 4, MC-1 2057 f-11 to 12 DNAs. c, Lane 1, mouse NIH3T3 DNA; lane 2, hamster SHOK DNA; and lane 3, SHOK 3013 f-1 DNA.
they are more useful tools than NIH3T3 cells to analyse mouse tumour DNAs since the mouse repetitive DNA sequence serves as a molecular marker to detect and isolate transforming DNA sequences. Although mice and hamsters are closely related rodents, we have shown that the mouse tumour DNAs are clearly distinguished from hamster cellular DNA by Southern hybridisation probing the cloned mouse repetitive sequence. Second, their differential sensitivities to transformation by a variety of oncogenes open up the possibility to detect new classes of oncogenes which have not been identified by the NIH3T3 transfection assay. The novel cot oncogene encoding a serine-specific protein kinase was actually detected in human thyroid carcinoma (Miyoshi et al., 1991) and colon cancer tissue DNA (Sasai et al., in preparation) using SHOK cells. The murine cot and mos were also identified in this study. These findings indicate that SHOK and MC-1 cells are highly susceptible to oncogenes of the serine/threonine-specific protein kinase family. Third, their growth behaviour and low activity in developing spontaneous overgrowth were ideal properties for screening oncogenes. We have estimated that their transformation efficiency was much greater than that of Rat 1, Rat 2 of Swiss 3T3, and comparable to that of NIH3T3 and C3H10T1/2 cells (data not shown).

In the osteosarcoma N5-2057 formed in the irradiated parent mouse, the c-K-ras oncogene was detected. We presume the activation of c-K-ras would be correlated with X-ray-induced carcinogenesis since a similar case has been reported in a X-ray-induced mouse thymic lymphoma (Guerrero et al., 1984).

The activation of the mos oncogene in SHOK 2057 f-1, which is due to promoter insertion during the step of transfection, has similarity with the case of a murine plasticocytoma which has the insertion of intercisternal A-particle serving as a strong promoter in mos gene (Canaani et al., 1984).

Although we have detected cot and an unidentified 'non-ras' oncogene in two of four heritable tumours, it is presently unknown whether they have been activated by germ-line mutations. Since we have only one primary transformant in each of these cases, it is possible that activation of these oncogene are due to the artifact of transfection. The characterisation of the transformant (SHOK 3013 f-1) should be done using the mouse c-DNA of c-cot gene which has recently been cloned.

Negative results in detecting the ras gene family during the survey using SHOK, MC-1 and NIH3T3 cells suggest that the ras oncogenes may not be frequently involved in development of these X-ray-induced heritable tumours.

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