

Receptor FGFR1 acts as a tumor suppressor in nude mice when overexpressed in HEK 293 Tet-On cells

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Abstract. Fibroblast growth factor receptor-like 1 (FGFR1) is a transmembrane receptor that interacts with heparin and FGF ligands. In contrast to the classical FGF receptors, FGFR1 to FGFR4, it does not appear to affect cell growth and proliferation. In the present study, an inducible gene expression system was utilized in combination with a xenograft tumor model to investigate the effects of FGFR1 on cell adhesion and tumor formation. It was determined that recombinant FGFR1 promotes the adhesion of HEK 293 Tet-On® cells *in vitro*. Moreover, when such cells are induced to express FGFR1ΔC they aggregate into huge clusters. If injected into nude mice, the cells form large tumors. Notably, this tumor growth is completely inhibited when the expression of FGFR1 is induced. The forced expression of FGFR1 in the tumor tissue may restore contact inhibition, thereby preventing growth of the cells in nude mice. The results of the present study demonstrate that FGFR1 acts as a tumor suppressor similar to numerous other cell adhesion proteins. It is therefore likely that FGFR1 functions as a regular cell-cell adhesion protein.

Introduction

Cell adhesion proteins form connections between individual cells and mediate the interactions of cells with the surrounding extracellular matrix (ECM) (1). In this way, they participate in various cellular functions, including signal transduction, communication, embryogenesis, inflammation and apoptosis.

The majority of cell adhesion proteins can be grouped into one of five families: The immunoglobulin family, the

integrins, the cadherins, the selectins and the syndecans (1-3). Members of the immunoglobulin family promote cell adhesion in a calcium-independent manner. These proteins contain an extracellular section with a variable number of immunoglobulin (Ig)-like domains, a relatively short intracellular domain and a single transmembrane domain. For example, cell adhesion molecule-1 (CADM1; also known as nectin-like protein 2) contains three extracellular Ig-like domains and binds with its intracellular domain to the adapter protein DAL1, which in turn anchors CADM1 to the actin cytoskeleton (1-3). Integrins are involved in the integration of the ECM with the cytoskeleton. These proteins represent noncovalently linked heterodimers composed of an α and a β subunit. The majority of integrins acts as receptors for ECM proteins, including fibronectin, vitronectin, collagens and laminin, and recognizes the Arg-Gly-Asp sequence within target proteins (1-3). Cadherins are calcium-dependent cell adhesion molecules. E-cadherin, for example, contains an extracellular domain with 5 cadherin repeats, an intracellular domain and a transmembrane domain. The intracellular domain interacts with catenin and binds to the actin cytoskeleton (1-3). Selectins mediate the interactions between leukocytes and endothelial cells. They are composed of a single transmembrane domain, a short intracellular domain and an extracellular domain with a variable number of sushi motifs. Typically, selectins assist with the homing of lymphocytes to the lymph node (1-3). Finally, syndecans are proteoglycans located on the cell membrane. They contain a variable number of glycosaminoglycan chains attached to strategic serine residues of the polypeptide chain. In addition to mediating cell-matrix and cell-cell interactions, syndecans contribute to modulating the activity of heparin binding growth factors (1-3).

Evidence from previous studies indicates that cell adhesion proteins participate in tumor formation and metastasis (2,3). Indeed, the progression of cancer is often associated with the loss of at least one cell adhesion protein. However, cell adhesion proteins can also function as tumor suppressors when overexpressed in tumor tissues (2,3). It appears that their forced expression restores contact inhibition, a phenomenon observed in normal cells, but not in the majority of malignant cells. E-cadherin, for example, is often lost in tumors of epithelial origin and it has been demonstrated that the disruption of cell adhesion mediated by E-cadherin is associated with the development and progression of cancer (4). The

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Abbreviations: FGF, fibroblast growth factor; FGFR, fibroblast growth factor receptor; FGFR1, FGFR-like protein 1

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forced expression of E-cadherin in tumor cell lines slows down cell proliferation and reduces cell invasiveness. By contrast, inhibiting E-cadherin using antibodies or antisense RNA restores the invasiveness of the cells (4). Likewise, integrin $\alpha 7$ is frequently mutated in human malignancies and such mutations are associated with cancer recurrence (5). It has been demonstrated that the increased expression of integrin $\alpha 7$ in leiomyosarcoma cells resulted in the reduction of colony formation. Moreover, increasing the expression of $\alpha 7$ integrin in mice with xenografted tumors inhibited tumor growth (5). Likewise, CADM1 has been implicated in cancer progression. Downregulation of CADM1 synthesis has been observed in a variety of human tumors, including breast cancer and esophageal squamous cell carcinoma (6,7). However, it has been demonstrated that restoring CADM1 expression suppresses cell growth and slows down tumor invasion (7). It has therefore been concluded that the majority of the cell adhesion proteins can function as tumor suppressors (2,3).

A previous study by our group described a novel transmembrane protein that resembles CADM1 (8,9). This protein contains three extracellular Ig-like domains, a relatively short intracellular domain and a single transmembrane domain. It interacts with heparin and fibroblast growth factors (FGFs) in a manner similar to the classical FGF receptors (FRFRs), FGFR1-FGFR4 (10,11). For this reason, the novel protein was termed FGFR-like 1 protein (FGFRL1). However, FGFRL1 does not possess the intracellular tyrosine kinase domain required for signal transduction by transphosphorylation and consequently cannot mediate FGF signaling by itself.

The function of the novel receptor is currently unclear. Knockout mice with a targeted disruption of the FGFRL1 gene present a striking phenotype; they lack metanephric kidneys (12) and die at birth due to a weak, malformed diaphragm that cannot inflate the lungs after birth (13,14). Cell culture experiments have demonstrated that FGFRL1 can act as a typical cell adhesion protein when coated on plastic dishes. FGFRL1 forms heterophilic interactions with heparan sulfate proteoglycans, such as glypican, at the cell surface of neighboring cells (15). A tetracycline-inducible system has been utilized, in which the expression of FGFRL1 could be controlled by the addition of the inducer doxycycline (16). It was observed that in the presence of doxycycline, cells aggregated and formed huge clusters, whereas in its absence, they tended to remain as individual clones (17).

In the present study, one of the tetracycline-inducible cell clones was characterized in more detail. The primary aim of the study was to determine whether FGFRL1 functions as a tumor suppressor in a manner similar to other cell adhesion proteins.

Materials and methods

Cell culture. The generation of tetracycline-inducible cell clones has been described in detail in a recent publication (17). The stable clone K13 Δ C was produced by transfection of a cDNA for truncated FGFRL1 (corresponding to amino acid residues 1-417) into HEK 293 Tet-On[®] cells (Clontech Laboratories, Takara Bio Europe SAS, Saint-Germain-en-Laye, France). The cells were cultivated in an atmosphere of 5% CO₂ in Dulbecco's modified Eagle's medium (DMEM)

supplemented with 10% fetal bovine serum, 100 U/ml penicillin and 100 μ g/ml streptomycin (all Sigma-Aldrich, Buchs, Switzerland). To maintain the selective pressure for stable transfection, 100 μ g/ml Hygromycin B (InvivoGen, San Diego, CA, USA) was used. Expression of FGFRL1 from the tetracycline responsive promoter was induced by the addition of 1 μ g/ml doxycycline.

Adhesion experiments. Recombinant FGFRL1 protein was isolated from the conditioned media of HEK 293 cells that had been stably transfected with a cDNA for human FGFRL1 (corresponding to amino acid residues 1-357), as previously described (18). The recombinant protein was purified by chromatography on a column of Heparin Sepharose[®] 6 Fast Flow (GE Healthcare Bio-Sciences, Pittsburgh, PA, USA) as previously described (18). FGFRL1 protein or bovine serum albumin (BSA; Sigma-Aldrich), which served as a control, was diluted to 20 μ g/ml in phosphate-buffered saline (PBS) and droplets of the solution (15 μ l) were spotted onto 35-mm petri dishes (non-tissue culture; catalog no. 82.1135; Sarstedt Co., Nümbrecht, Germany). Following incubation in a humidified chamber at 4°C for 16 h, the solution was carefully aspirated and all residual sites of the petri dish were blocked with 1% BSA in PBS. Cells were seeded onto pre-coated petri dishes in serum-free medium (2x10⁶ cells/plate) and incubated for 1 h at 37°C. Non-adherent cells were carefully removed by washing with PBS. Adherent cells were fixed with 4% paraformaldehyde (Sigma-Aldrich) and inspected under a microscope (Nikon Eclipse E800; Nikon AG, Zurich, Switzerland).

In a further experiment, the cells were cultivated in complete growth medium (as aforementioned) on uncoated petri dishes (non-tissue culture) for 1-3 days, and cell adhesion and cell-cell clustering were documented with a Zeiss Axiovert 10 microscope (Carl Zeiss AG, Oberkochen, Germany).

Immunocytochemistry. Cells grown on coverslips were washed with ice-cold PBS, fixed with 4% paraformaldehyde and permeabilized with 0.2% Triton X-100 (Sigma-Aldrich) in PBS. Non-specific sites were blocked with 3% BSA in PBS. The fixed cells were incubated for 3 h at room temperature with a humanized monoclonal antibody (1 μ g/ml) against FGFRL1 that had been prepared by our group in a previous study (19). Following three steps of washing with PBS, bound antibodies were detected with Cy2-labeled secondary antibodies (catalog no. 109-225-097; dilution 1:200; Jackson ImmunoResearch Laboratories, West Grove, PA, USA). The nuclei of the cells were stained with 1 μ g/ml 4',6-diamidino-2-phenylindole (Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA). To detect filamentous actin, fixed cells were treated in a similar way with tetramethylrhodamine-labeled phalloidin (Sigma-Aldrich).

Electron microscopy. Cells grown on cover slips were fixed with 2.5% glutaraldehyde in 30 mM potassium phosphate buffer. The fixed cells were dehydrated in ethanol, critical point dried and sputter-coated with gold, as previously described (20). Finally, the specimens were inspected with a Philips XL 30 FEG scanning electron microscope operated at 10 kV (Philips, Amsterdam, The Netherlands).

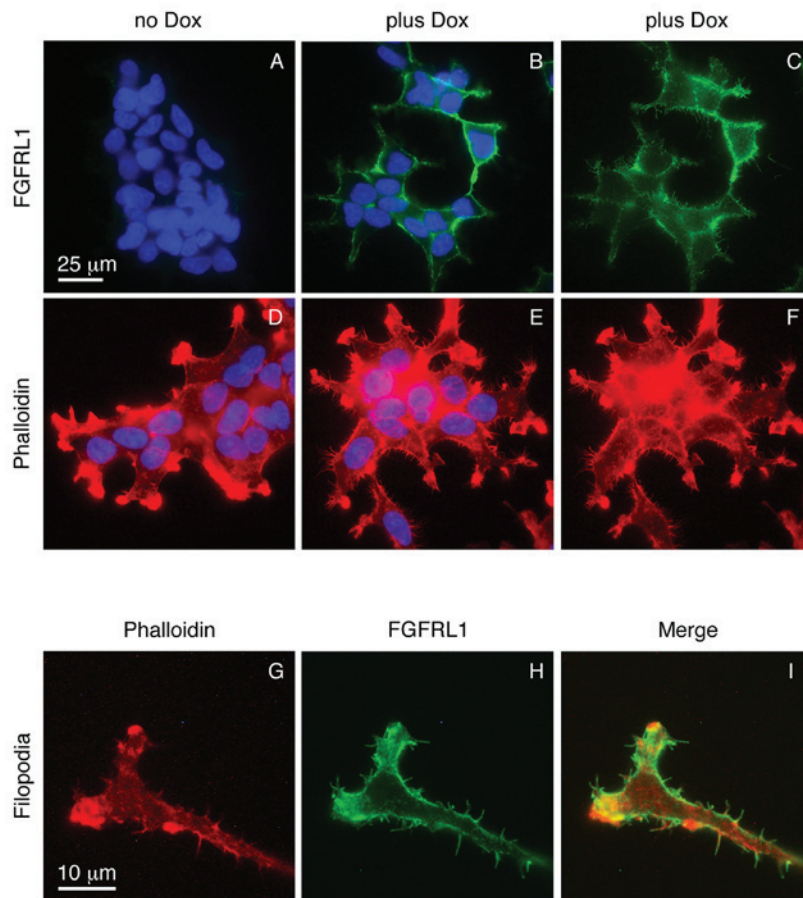


Figure 1. Expression of FGFR1 in HEK 293 Tet-On-FGFR1ΔC cells induces the formation of filopodia-like spikes. FGFR1-inducible cells (clone K13ΔC) were cultivated in the absence (no Dox) or presence (plus Dox) of the inducer doxycycline. (A-C) Expression of FGFR1ΔC was visualized with a monoclonal antibody against human FGFR1, followed by Cy2-labeled secondary antibodies (green). FGFR1ΔC was observed primarily at the cell membrane and in microspikes that emerged from the cell membrane. (D-F) Filamentous actin was stained with tetramethylrhodamine-labeled phalloidin (red). (G-I) Merging of the pictures stained for FGFR1 and for actin indicated that the filopodia-like spikes partially colocalized with filamentous actin. In the absence of doxycycline, no FGFR1ΔC expression was observed and the number of filopodia-like spikes was much lower. For easy reference, cell nuclei stained with 4',6-diamidino-2-phenylindole are included in panels A, B, D, and E. FGFR1, fibroblast growth factor receptor-like 1; Dox, doxycycline.

Nude mice xenograft experiments. To investigate tumorigenicity, immunodeficient CD-1[®] NU/NU-Foxn1 nude mice (Charles River Wiga GmbH, Sulzfeld, Germany) were utilized. Approximately 1×10^7 K13ΔC cells in PBS were subcutaneously injected at two ventral sites into the animals. Half of the mice (4 mice per group, randomly selected) were treated with 100 μg/ml doxycycline, which was directly added to the drinking water of the animals; the other half received regular water. After 5 weeks, the mice were sacrificed and images of the tumors were captured. All animal experiments had been approved by the Ethics committee of the County of Bern.

Statistical analysis. The significance of the results from the xenograft experiments was analyzed with the exact Fisher test utilizing an online calculation tool (<http://www.quantitativeskills.com>). $P \leq 0.05$ was considered to indicate a statistically significant difference.

Results

FGFR1 overexpression in cell culture. All the following experiments were performed with clone K13ΔC from the Tet-On-FGFR1ΔC cell line (17). This clone had been

prepared with cDNA for FGFR1, which covered the extracellular domain and the transmembrane helix of the protein, but lacked the intracellular domain. Hence, all effects observed with this clone may be attributed to the extracellular and the transmembrane domains of FGFR1.

In the absence of doxycycline, the K13ΔC clone did not express any detectable FGFR1ΔC protein, as verified with our monoclonal antibodies (Fig. 1A). Following induction with doxycycline, a strong signal for FGFR1 was observed at the cell membrane (Fig. 1B). Thus, K13ΔC cells tightly controlled the expression of FGFR1ΔC depending on the presence or absence of doxycycline. When the focus of the microscope lens was changed to the plane of the glass slide (rather than the site of strongest fluorescence), some fluorescent signal was observed at numerous protrusions that emerged from the plasma membrane of the cells in a spike-like fashion (Fig. 1C). To investigate whether these microspikes represented normal filopodia, the cells were stained with fluorescently labeled phalloidin, which is known to interact with filamentous actin. Again, a number of spikes was observed that protruded in a perpendicular manner from the plasma membrane (Fig. 1D-F). At higher magnification (objective lens, 100X), the FGFR1ΔC signal (green) partially

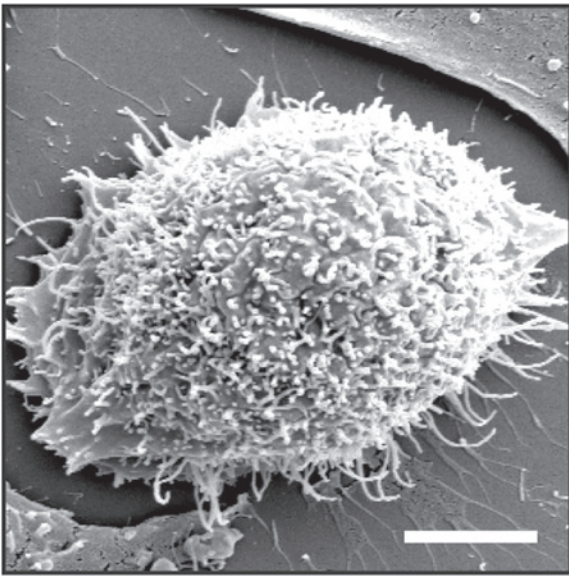
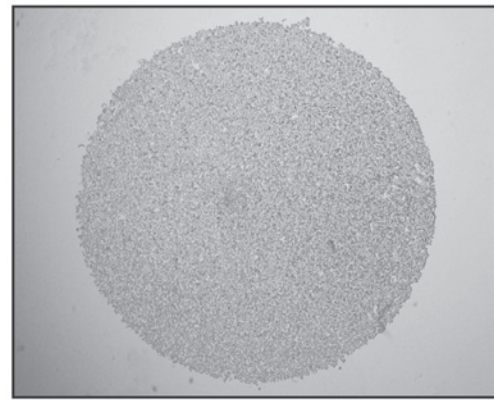


Figure 2. Electron microscopy of HEK 293 Tet-On-FGFR1 Δ C cells. Clone K13 Δ C was cultivated in the presence of doxycycline, fixed with glutaraldehyde and prepared for SEM. Inspection by SEM at 10 kV detected numerous filopodia-like spikes with a diameter of ~ 200 nm. Bar, 5 μ m. SEM, scanning electron microscopy.

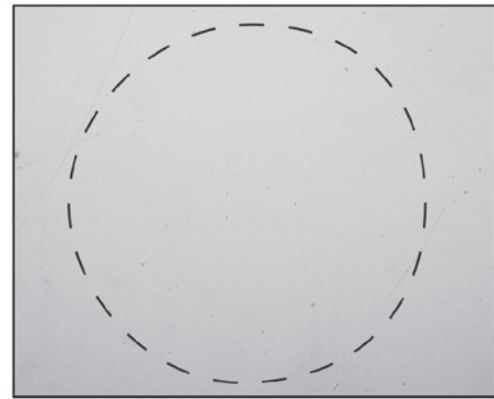
co-localized with the phalloidin signal (red) (Fig. 1G-I). Additionally, the spikes were visualized under a scanning electron microscope (Fig. 2). Here, numerous protrusions were detected that had an average diameter of 180 ± 28 nm ($n=6$). It is therefore likely that the spikes represent regular filopodia. Notably, a much larger number of filopodia was observed in the presence of doxycycline compared with the number observed in its absence (Fig. 1D-F). Thus, forced expression of FGFR1 Δ C appears to stimulate the formation of filopodia.

Next, it was investigated whether K13 Δ C cells would bind to purified FGFR1 protein, as has previously been documented in Chinese hamster ovary (CHO) cells (15). The surface of a plastic plate (non-tissue culture) was coated with droplets of recombinant FGFR1 solution comprising the Ig1-Ig3 domains, or with droplets of BSA that served as a control. Within 1 h, the K13 Δ C cells attached to the recombinant FGFR1 protein, however, they did not attach to the control BSA (Fig. 3). It is likely that this interaction was accomplished by the binding of FGFR1 to cell surface heparan sulfate proteoglycans, since the binding could be blocked with soluble heparin (data not shown), as previously reported in CHO cells (15).

In the following experiment, the effect of doxycycline (and therefore the effect of FGFR1 Δ C) on cell-cell adhesion and cell clustering was recapitulated (Fig. 4). K13 Δ C cells formed large patches of 10-30 cells following 9 h of incubation on bacterial plates in the presence of doxycycline. In the absence of the inducer, the cells also started to form clusters, however, these clusters were much smaller. After 3 days in culture, the K13 Δ C cells had merged to form a continuous network of cells. Notably, the borders between the individual cells were no longer distinguishable in the presence of doxycycline, whereas they were clearly detectable in its



FGFR1



BSA

Figure 3. FGFR1 induces cell adhesion. Droplets of a solution of recombinant FGFR1 (or of BSA, which acted as a control) were spotted onto bacterial petri dishes and incubated overnight at 4°C in a humidified chamber. Residual binding sites of the plastic surface were blocked with BSA. HEK 293 Tet-On-FGFR1 Δ C cells (clone K13 Δ C) were seeded onto the pre-coated dishes and allowed to attach for 1 h. Non-adherent cells were washed off; adherent cells were fixed. FGFR1 promoted cell attachment, whereas BSA (indicated by the dotted circle) did not. FGFR1, fibroblast growth factor receptor-like 1; BSA, bovine serum albumin.

absence. Thus, the presence of FGFR1 Δ C promotes tight, intimate interactions between cells. Taken together, the three experiments suggested that FGFR1 represents a typical cell adhesion protein.

Xenograft tumor model. The majority of cell adhesion proteins can function as tumor suppressors (2,3); therefore, the effects of FGFR1 on tumor formation were investigated. K13 Δ C cells were injected subcutaneously into immunocompromised nude mice at two ventral sites. Half of the mice received doxycycline in their drinking water in order to induce the expression of FGFR1 Δ C. The other half served as controls and received regular drinking water. Within 5 weeks, the control animals had developed large tumors (diameter ≥ 12 mm) at five out of the eight injection sites (Fig. 5). In sharp contrast, none of the doxycycline-treated mice had produced any tumors ($P=0.0257$). A control experiment confirmed that doxycycline itself had no effect on tumor growth (data not shown). Thus, FGFR1 may function as a typical tumor suppressor that effectively inhibits the outgrowth of xenografted tumors *in vivo*.

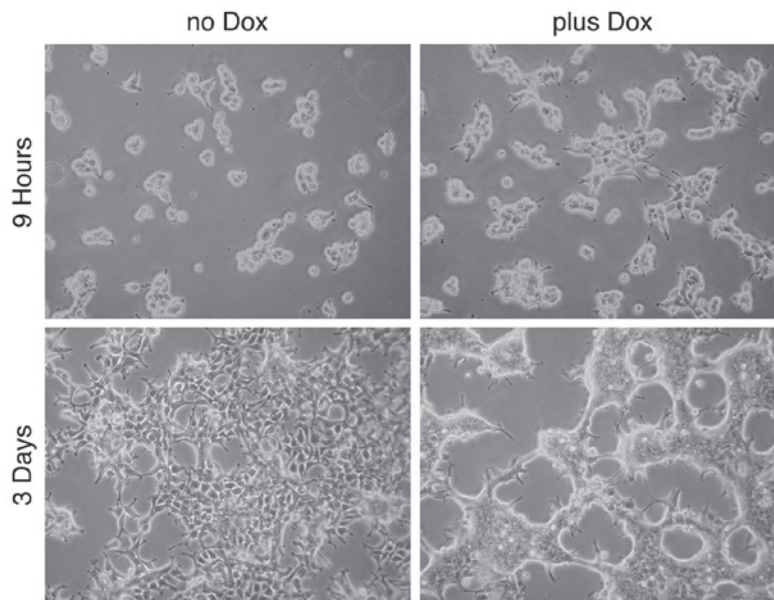


Figure 4. FGFR1 induces cell-cell adhesion and cell clustering. FGFR1-inducible HEK 293 Tet-On cells (clone K13ΔC) were seeded into bacterial petri dishes and cultivated for up to 3 days in the absence (no Dox) or presence (plus Dox) of doxycycline. Expression of FGFR1ΔC induced cell-cell adhesion and cell clustering. After 3 days, the boundaries between cells were barely detectable in the presence of doxycycline, whereas they were clearly detectable in its absence. FGFR1, fibroblast growth factor receptor-like 1; Dox, doxycycline.



Figure 5. FGFR1 acts as a tumor suppressor *in vivo*. Inducible HEK 293 Tet-On-FGFR1ΔC cells (clone K13ΔC) were injected into nude mice at two ventral positions. Expression of FGFR1 was induced in half of the animals by doxycycline, which was added to their drinking water. The other half received normal drinking water. Tumor formation was inspected at the injection sites after 5 weeks. Two representative examples of each group are depicted. The injected cells were able to generate tumors only in those mice that did not receive any doxycycline. FGFR1, fibroblast growth factor receptor-like 1; Dox, doxycycline.

Discussion

The current study, taken together with results from the literature, demonstrates that the novel receptor FGFR1 represents a regular cell-cell adhesion protein. This conclusion is based on the following facts: i) The domain structure of FGFR1, which has a single transmembrane domain, three extracellular Ig-like repeats and a short intracellular domain (21), resembles the structure of other cell adhesion proteins from the Ig superfamily, namely the nectins and nectin-like molecules (22,23). ii) Recombinant polypeptides corresponding to the extracellular domain of FGFR1 promote cell adhesion *in vitro*. If a mutation is introduced into the polypeptide chain, the activity is completely lost (15). iii) When overexpressed in different cell lines, FGFR1 protein accumulates at intersections where two cells touch each other (15). Overexpression

in HEK 293 Tet-On cells leads to the aggregation of the cells and to the formation of large clusters (17). iv) Finally, the results of the current study indicate that FGFR1, like other typical cell adhesion proteins, acts as a tumor suppressor in a xenograft tumor model. It was determined that the forced expression of FGFR1 in HEK 293 Tet-On cells completely inhibited the outgrowth of tumors in immunocompromised mice.

During the xenograft experiments, it was noted that the cells were not extremely tumorigenic. A large number of HEK 293 Tet-On-FGFR1ΔC cells had to be injected into the mice to initiate any tumor growth. Other studies have also noted that HEK 293 cells exhibit particularly low tumorigenicity. Shen *et al* (24) demonstrated that HEK 293 cell tumorigenicity increased with increasing passage number, and finally reached 100% when the passage number was >65; however, the original

isolates of the cells did not form tumors at all in nude mice. In the present study, an outgrowth of tumors was detected in the absence of doxycycline in 5 cases (63%). Following induction of FGFR1 synthesis, tumor growth was observed in no cases (0%). Doxycycline itself had no effect on tumor growth, as demonstrated in a control experiment and as previously published in the literature (25). Thus, it is FGFR1 that can act as a tumor suppressor.

Previous studies have reported that alterations in the synthesis of FGFR1 occur in tumor cells (26-31). The screening of 241 different human tumor samples with a cancer-profiling array suggested that major changes in the relative expression of FGFR1 occur in ovarian tumors (26). In several samples, a significant decrease in FGFR1 expression was observed in the tumor tissue relative to the matched control tissue. However, in one ovarian tumor sample there was a 25-fold increase (26). Furthermore, the overexpression of FGFR1 in certain ovarian tumor samples was confirmed in a study aiming to identify novel tumor-specific marker genes (27). An association of FGFR1 expression with tumor growth and metastasis was also suggested by a study of patients suffering from head and neck tumors (28,29). In this case, FGFR1 overexpression appeared to correlate with tumor growth. Another study noted a significant decrease of FGFR1 protein expression in bladder tumors and bladder cancer cell lines (30). This decrease was explained by heterozygous deletions at the chromosomal region 4p16.3, which includes the locus of the FGFR1 gene. Finally, mutations affecting the reading frame of FGFR1 have been observed relatively frequently in colorectal cancer cell lines (31). The majority of these mutations occurred in the intracellular domain of the protein.

The molecular mechanism governing how FGFR1 may inhibit tumor growth in a xenograft model is complex. With other cell adhesion proteins it has been concluded that forced expression partially restores contact inhibition of the tumor cells. In the case of CADM1, it was demonstrated that homophilic interactions of CADM1 at the surface of two adjacent cells activated the phosphatidylinositol 3-kinase (PI3K) pathway and led to the reorganization of the actin cytoskeleton (32). The intracellular domain of CADM1 formed a complex with membrane-associated guanylate kinase homologues, including MPP3 and Dlg, thus linking CADM1 and PI3K. In the case of FGFR1, the effect may be exerted by heterophilic interactions of the extracellular domain of FGFR1 with another transmembrane protein, since the protein expressed by K13ΔC cells lacks the intracellular domain of FGFR1. The extracellular domain has the ability to interact with target proteins of neighboring cells containing heparan sulfate chains, including syndecans and glypicans (15). Two target proteins, glypican-4 and glypican-6, which specifically interact with FGFR1, have previously been identified by our group using tandem LC mass spectrometry (16). Therefore, it is hypothesized that FGFR1 inhibited tumor growth in the current study in the following manner: HEK 293 Tet-On cells lost their normal contact inhibition and grew in an unrestricted way, as they were subcloned multiple times during the generation of the HEK 293 Tet-On-FGFR1ΔC clones. When injected into nude mice, these cells gave rise to large tumors due to a defect in contact inhibition. Enforcing the expression of FGFR1ΔC in

the injected cells by adding doxycycline induced the adhesion of the cells with each other and restored contact inhibition, thereby inhibiting tumor growth.

It remains to be demonstrated whether the identification of FGFR1 as a tumor suppressor may be exploited to improve the diagnosis and therapy of cancer patients. FGFR1 may serve as a tumor marker to identify aggressive tumors that have lost FGFR1 expression (27,31). Furthermore, there may be the possibility of enhancing FGFR1 synthesis in the tumor tissue. Levels of FGFR1 mRNA are regulated by microRNA-120 (28,33), which specifically interacts with the 3' end of the FGFR1 mRNA and leads to its degradation. Therefore, if it were possible to downregulate microRNA-120 expression in the tumor tissue, the endogenous levels of FGFR1 would increase and contact inhibition of the cells in the tumor tissue would be restored, thus suppressing tumor growth. Future studies are necessary to determine whether this could be developed as a novel therapeutic strategy for patients with cancer.

In conclusion, the current study taken together with results from the literature, demonstrates that FGFR1 is a cell-cell adhesion protein that acts as a tumor suppressor similar to numerous other cell adhesion proteins.

Acknowledgements

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