

## Reviews

### Transgenic mice carrying chimeric or mutated type III intermediate filament (IF) genes

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**Abstract.** Mice carrying chimeric, truncated or mutated genes encoding intermediate filament (IF) proteins type III do not show any detectable severe pathology. However, upon (over)expression of the transgene in the eye lens all animals develop lens opacification (cataract). At the cellular level the loss of visual acuity is preceded by interference with the terminal differentiation of lens fibre cells, plasma membrane damage, distorted assembly of the IF cytoskeleton and perturbation of the cytoskeleton-membrane complex. The degree of expression is paralleled by the extent of the damages.

**Key words.** Transgenic mice; chimeric IF-genes; cataract.

#### Introduction

The skeleton of cells (cytoskeleton) consists of three major fibrillar networks: microtubules, intermediate filaments (IFs) and microfilaments. These filamentous structures can be distinguished on the basis of morphological features, composition and immunological properties (compare tables 1 and 2).

A wealth of data led to the notion that IF subunits consist of three distinct domains: (1) a central 'rod' of conserved length [approximately 310 amino acid residues (365 in case of type V IFs)], conserved secondary structure and partly conserved sequence; (2) a nonhelical amino-terminal domain ('head'); (3) a nonhelical carboxyl-terminal domain ('tail').

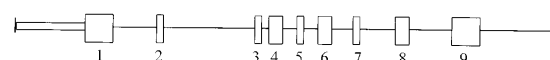
The two nonhelical domains flank the conserved central rod that has a helix structure, interrupted only by three short nonhelical linkers. The resulting four  $\alpha$ -helical subdomains are named 1A (N-terminal end of the rod); 1B, 2A and 2B (C-terminal end of the rod) (fig. 1).



Figure 1. Schematic representation of the tripartite structural organization of an IF protein subunit. Single lines = non-helical flanking regions; white boxes = helices; black boxes = nonhelical linkers.

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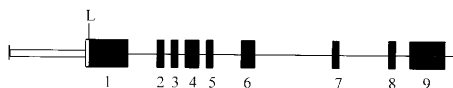
#### Vimentin



#### Desmin



#### pVDes



#### pV Vim<sub>1-6</sub> Des<sub>7-9</sub>



#### pV Des<sub>1-5</sub> Vim<sub>9</sub>



#### pD Des<sub>1-5</sub> Vim<sub>9</sub>



Figure 2. Schematic representation of the vimentin and desmin genes and the constructs derived thereof.

The amino acid sequence in the central helix follows the heptad rule, which means that this sequence comprises sets of seven amino acids with hydrophobic amino acid

Table 1. Major constituents of the cytoskeleton.

Filament	Protein/subunit(s)	Diameter (nm)
Microtubuli	tubulin	25
Intermediate filaments	IF-protein*	10
Micro-filaments	actin	5

\*See table 2.

residues at positions 1 and 4. This typical structure enables the formation of coiled-coil dimers.

It has been established that in the living cell microtubules and microfilaments are involved in important roles in processes such as mitosis, locomotion, anchorage and protoplasmic streaming. In contrast, the function of intermediate filaments is not yet fully understood. Apparently cytoplasmic IFs do not fulfil a typical 'household' function, as some cell lines and a few cell types in vivo do not contain an IF network. Recently, so-called knock-out experiments, in which a particular type-III IF gene has been removed from the genome in transgenic animals, led to controversial conclusions concerning the evidence of an abnormal phenotype [1–3]. Furthermore, various studies with cell lines or transgenic animals show that disruption of the IF networks does not affect mitosis, cell morphology or motility. However, there is a strong belief among workers in the field that IFs play a role in the cellular organization as a whole and that the networks provide mechanical strength and maintenance of the architecture of the living cell. Evidence for such a structural role stems from studies upon the expression of domi-

nant negative, truncated human keratin-14 subunits in transgenic mice [4–7] that exhibit features of a serious autosomal human genetic skin disease called epidermolysis bullosa simplex (EBS). Expression of another truncated human gene, encoding keratin-10, causes a skin disease comparable to human epidermolytic hyperkeratosis [6]. Indeed patients suffering from those diseases reveal point mutations in the highly conserved amino carboxyl ends of the rod domain of a particular keratin.

A number of reviews dealing with expression, assembly and interaction of IFs with other cell components, particularly the plasma membrane, and putative functions have been published previously [8–34].

The present paper gives an overview of our previous and more recent work aimed to contribute to a better understanding of the function of vimentin and desmin IFs and the role of the membrane-cytoskeleton complex in cells and in the developing organism. With this goal in mind, several chimeric and truncated IF protein-encoding genes (fig. 2) were either transfected into cells in culture or microinjected into the murine germ line. Some of the resulting mutant IF subunits appeared to be assembly-incompetent and capable of disrupting pre-existing vimentin or desmin filaments in a dominant negative fashion. In all cases the expression of the gene constructs was tissue-specific in transgenic mice.

Strikingly, the disruption of endogenous vimentin or desmin IFs resulted in aberrations at the ultrastructural level but did not lead to obvious developmental or functional abnormalities with the exception of cataract formation in cases where the gene product was expressed in the eye lens.

Table 2. Classification of IF subunits.

Cell type	IF protein	Sequence type	Size (kD)
Epithelial cells	acidic keratins	I	40–60
Epithelial cells	basic keratins	II	50–70
Mesenchymal cells	vimentin	III	55
Lens cells*			
Muscle cells	desmin	III	53
Peripheral nervous system	peripherin	III	57
Glial cells and astrocytes	GFAP (glial fibrillary acidic protein)	III	53
Retinal ganglion cells (goldfish)	plastin	III	64
Most neurons	neurofilament proteins (NF-L, NF-M, NF-H)	IV	68, 145, 200
Neuroepithelial stem cells	nestin	IV	200
Central nervous system	$\alpha$ -internexin	IV	66
Nucleated cells	lamins (A/C and B)	V	60, 67, 70

\*A 49-kDa and a 115-kDa lens protein termed *phakinin* and *filensin*, respectively, have been reported to be related to the family of IF proteins. These two proteins have a cytoskeletal structure which is distinct from classical 8–11 nm intermediate filaments, thus possibly comprising a novel form of intermediate filaments [49]. Evidence that the 49-kDa protein belongs to the keratin family has been provided by Orii et al. [52].

In the following sections the effects of various constructs are described briefly.

### pVDes

pVDes comprising the vimentin promoter linked to the complete desmin coding region was microinjected into fertilized mouse eggs [35]. Immunofluorescence studies of eye lens cells derived from the resulting transgenic animals revealed a mosaic expression pattern of the transgene product [36]. This means that, in contrast to the endogenous vimentin, desmin cannot be detected in all cells of mesenchymal origin or in the eye lens (fig. 3). We observed that the number of cells expressing the transgene in the embryo is higher (fig. 3A) than in adult lens cells (fig. 3B). In cells where both intermediate filament proteins are coexpressed, hybrid IFs as well as amorphous desmin deposits are found (not shown). The deviating assembly of IF networks induced by genetic manipulation is in some way correlated with abnormal lens cell differentiation. Immunofluorescence

experiments show that lenticular cell nuclei, which are normally extruded from the elongating fibre cells, are now detectable in a number of inner fibres. The result of the distorted terminal differentiation process can also be demonstrated by electron microscopy (fig. 4).

Furthermore, marked aberrations of membranous and junctional domains can be observed that frequently appear as internalized structures, whereas a number of cells are fused as a consequence of membrane breakdown (fig. 5). The level of expression is paralleled by the extent of plasma membrane damage.

In lens fibre ghost preparations from transgenic mice impressive conchoidal whorls of IFs can be visualized on thin sections. These abnormal structures are frequently found near and in contact with the nuclear envelope (fig. 6). Eventually, the alterations of the architecture of the plasma membrane-cytoskeleton complex lead to opacification of the adult eye lens (cataract).

It appears that lens opacification is not the only aberration caused by the transgene. Alterations of the incisor tooth organ were seen in immunochemical and ultrastructural studies [37]. Apparently the expression of the

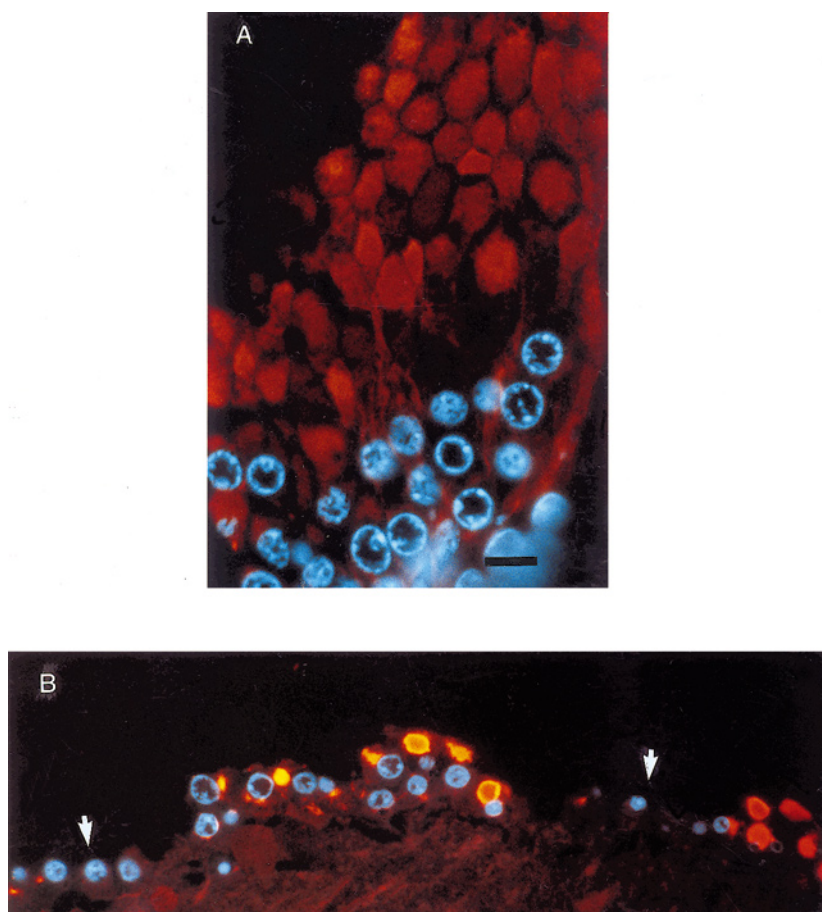


Figure 3. (A) Cryosection of an embryo lens (E<sub>14</sub>) carrying the pVVim<sub>1-6</sub>Des<sub>7-9</sub> transgene; staining with antidesmin and DAPI. The primary fibres appear virtually uniformly labelled. (B) Cryosection of the adult transgenic pVVim<sub>1-6</sub>Des<sub>7-9</sub> lens. The mosaic mode of desmin expression is clearly shown; staining with antidesmin and DAPI. White arrows point to cells devoid of desmin expression.

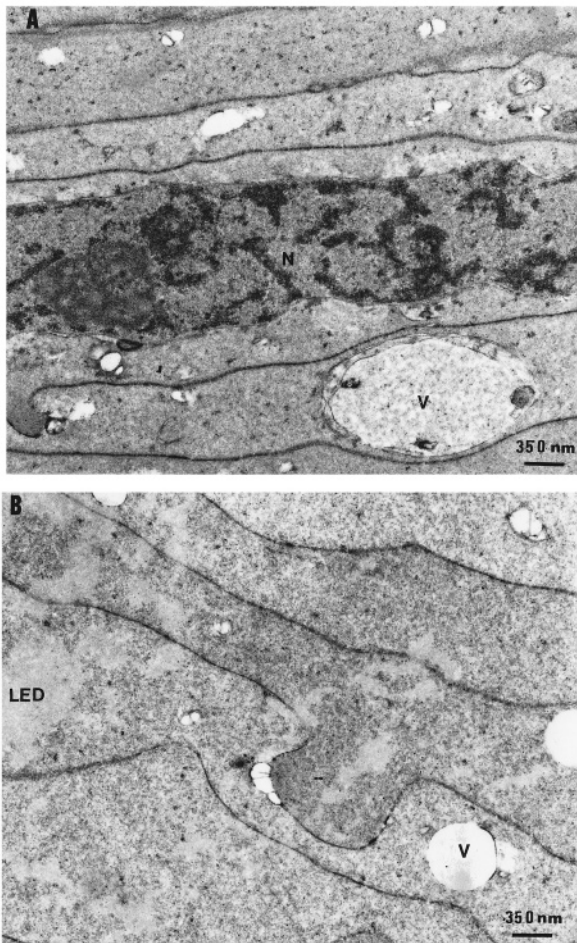


Figure 4. Thin sections of intact transgenic lens stained with uranyl acetate and lead citrate. (A) The lens fibres differ considerably in shape. Indentations and projections of the plasma membrane can be observed. The cytoplasm is occupied by remnants of endoplasmic reticulum and vacuoles (V) of different sizes wrapped by membrane profiles. Note the nucleus in a fibre cell. Bar = 350 nm. (B) The irregular framework comprising homogenous material of low electron density (LED). N = nucleus. (Reprinted with permission from the European Journal of Cell Biology)

transgenic desmin in the odontoblasts modifies the composition and assembly of the endogenous IF network. Furthermore, our results show that alterations of odontoblast differentiation has a negative effect on the induction capacity of these cells in regulating the function of ameloblasts. Presumably odontoblast-ameloblast interactions are mediated by cytoskeleton-dependent polarization and by specialized domains of the plasma membrane generating specific signals between the two cell types.

#### pVVim<sub>1-6</sub>Des<sub>7-9</sub>

We also designed a construct that comprises the 5' region of the hamster vimentin gene including the head and rod domain, and the 3' region of the desmin gene,

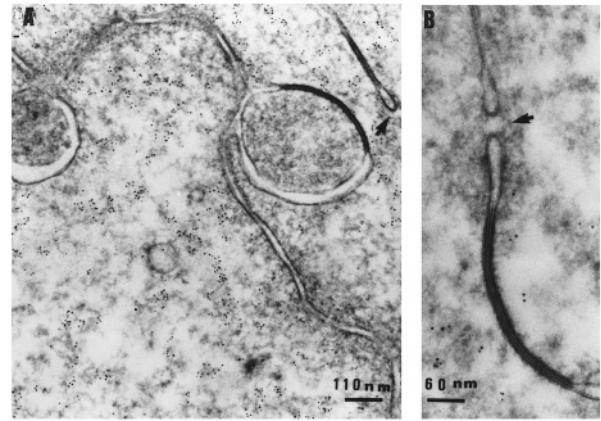


Figure 5. Ultrathin section of transgenic mouse lens showing the cortical region immunolabelled with antidesmin antibodies. Note the internalizations of the general plasma membranes and intercellular junctions. The arrows point to interruptions of the plasma membrane of adjacent fibre cells connected by a cytoplasmic bridge. The gold particles are associated with amorphous desmin. Bars = 110 nm (A), 60 nm (B). (Reprinted with permission from the European Journal of Cell Biology)

comprising the tail domain. This chimeric gene has been transfected into C<sub>2</sub>C<sub>12</sub> cells or microinjected into fertilized mouse eggs [38]. Several tissues of the transgenic animals were analysed by immunofluorescence in order to demonstrate the presence of the corresponding expression product (fig. 7).

Strong staining could be observed in all tissues of mesenchymal origin that normally express only vimentin. The fact that the fluorescence patterns often show the filamentous feature of the IF network strongly suggests that the chimeric Vim-Des protein assembles into pre-existing normal vimentin IF networks.

Since in addition to its occurrence in tissues of mesenchymal origin, vimentin is also expressed in the eye lens, it was not surprising that the chimeric protein could also be demonstrated in the water-insoluble protein fraction derived from the transgenic mouse lens. Obviously the extent of expression was high enough to allow detection even on a Coomassie Brilliant Blue-stained one-dimensional SDS-containing polyacrylamide gel (fig. 8).

From our experiments it may be concluded that the presence of appreciable amounts of pVVim<sub>1-6</sub>Des<sub>7-9</sub> protein in all vimentin-expressing tissues does not interfere with normal development of the animal. However, the expression of this protein hampers normal lens cell differentiation by blocking the cell denucleation process and leads to cataract.

#### pVDes<sub>1-5</sub>Vim<sub>9</sub>

To further investigate the putative function of vimentin IFs in intact tissues and the developing organism, a

construct under direction of the vimentin promoter and encoding a truncated desmin subunit was injected into fertilized mouse eggs [39]. The expression product, a mutant desmin protein, appeared to be assembly-incompetent and capable of disrupting pre-existing IFs, both in transgenic mouse tissue and in cells derived thereof. Tissue specificity and levels of expression of the construct were analysed by Northern blotting in different tissues, including lens, heart, skeletal muscle, spleen, esophagus, tongue, brain, testis and ovary. Three independent transgenic mouse strains expressed the gene product to virtually the same extent (fig. 9). Western blots displaying the corresponding protein were in agreement with the Northern blotting patterns (not shown). Expression of the truncated protein caused disruption of the endogenous vimentin filament network, as can be visualized by indirect immunofluorescence of tail sections and blood smears derived from founders and offspring of the the transgenic mice (fig. 10). Instead of

the normal filamentous structure, strongly fluorescent dots and clumps were seen in stromal cells. The disruption of the vimentin IF network has also been demonstrated on several tissues and cultured cells at the ultrastructural level. Electron microscopy was performed on thin sections of lens fibre ghosts from transgenic mice and compared with similar material from wild-type mice (fig. 11). The ultrastructural study demonstrated the presence of a disrupted IF network and the accumulation of amorphous transgene product.

#### **pDDes<sub>1-5</sub>Vim<sub>9</sub>**

We also constructed a similar mutant desmin gene as described in the foregoing section, but driven by the desmin promoter [40]. Again, the first five exons of the desmin gene were linked to exon 9 of the vimentin gene. Transgenic mice were generated by pronuclear micro-injection. Like the previous mutant gene product, the expression product of the desmin promoter-driven gene was assembly-incompetent and capable of disrupting pre-existing vimentin and desmin filaments in a dominant negative fashion. Since the construct comprised the desmin promoter, expression occurred exclusively in transgenic mouse muscle tissue and cultured muscle cells. Visual inspection of the transgenic animals did not reveal gross morphological defects; however, severe damage was detected at the ultrastructural level of transgenic mice carrying the desmin promoter-driven truncated gene. The alterations were observed in sartorius muscle. Expression of the latter truncated gene induced radical changes of the myofibrillar organization, notably the disruption of the T system [transverse tubular system] (fig. 12). These findings suggest that alterations of the desmin IF network lead to collapse and damage not only of the T system but also to fragmentation of specialized membrane domains of the sarcoplasmic reticulum [40].

#### **pVMDR3**

As we observed considerable damage of the cytoskeleton-membrane complex in all cases of genetic manipulation described afore, we wondered whether or not a specific membrane protein gene, linked to the vimentin promoter, might cause similar effects in transgenic animals. We chose the human multidrug-resistant phosphoglycoprotein MDR3 Pgp [41]. This protein belongs to a class of highly conserved membrane proteins that can extrude hydrophobic drugs from mammalian cells [42–45]. The tissue distribution of this non-lens-specific protein has been described previously [46].

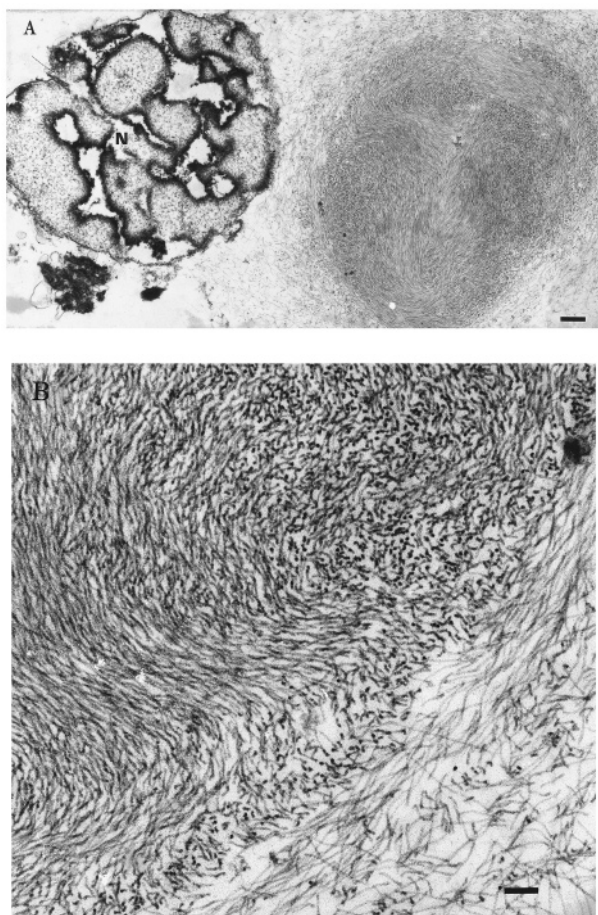


Figure 6. (A) Thin section of fibre ghosts prepared from the lens cortex of pVDes carrying transgenic mice. Note the giant conchoidal whorl of collapsed IFs still associated with the nucleus (N). Staining with uranyl acetate and lead citrate. Bar = 600 nm. (B) High magnification of a region of the conchoidal whorl showing irregular loops and compact IF bundles running in various directions. Bar = 100 nm. (Reprinted with permission from the European Journal of Cell Biology)

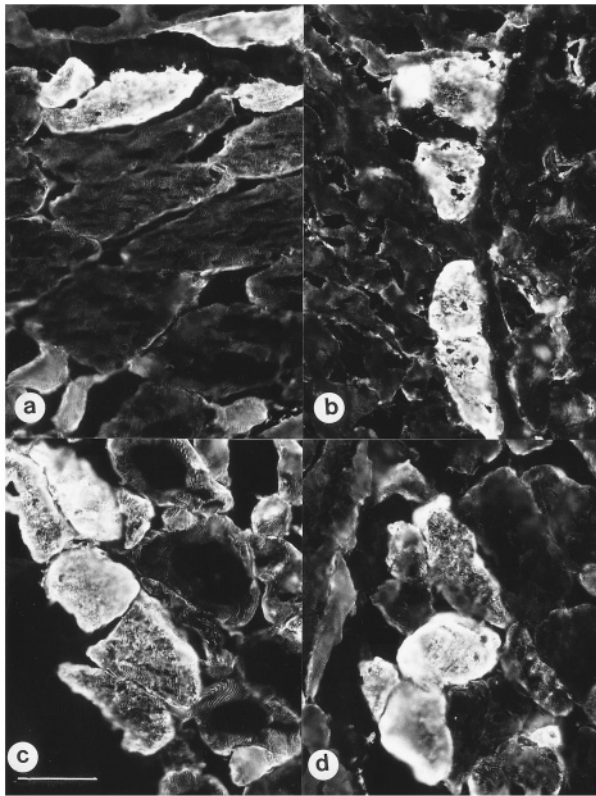


Figure 10. Indirect immunofluorescence of skeletal muscle (*a, b*) and tongue tissue sections (*c, d*) from pVDes<sub>1-5</sub>Vim<sub>9</sub> transgenic mice strains 47 (*a, c*) and 50 (*b, d*), using poly-des. Note strongly fluorescent, diffuse staining pattern in a minority of muscle fibres. The striated pattern normally observed upon desmin staining also seemed to be present in the strongly fluorescent muscle fibres (not shown). Bar = 4  $\mu$ m. (Reprinted with permission from the European Journal of Cell Biology)

transgenic mouse lens. Transgenes with a vimentin promoter linked to a heterologous IF coding region (complete, truncated or mutated) cause aberrations of the IF assembly process. Such abnormality is paralleled by impairment of membrane formation and membrane domain integrity. As far as the eye lens is concerned, the distortions lead to lenticular opacification independent of the type of IF coding sequence. Apparently the stringent prerequisite is the presence of a lens-specific promoter. It can be anticipated that strong viral promoters which drive the coupled gene into a variety of tissues in an aspecific way will act in a similar way as described here for IF or membrane protein-encoding genes. Actually, it appears from the literature that one of the obvious effects of viral promoter-driven transgenes is cataract, sometimes in addition to microphthalmia or other eye pathologies (table 3).

In our opinion earlier statements that (over)expression of IF protein-encoding sequences in transgenic mice does not lead to a defined phenotype are hardly ten-

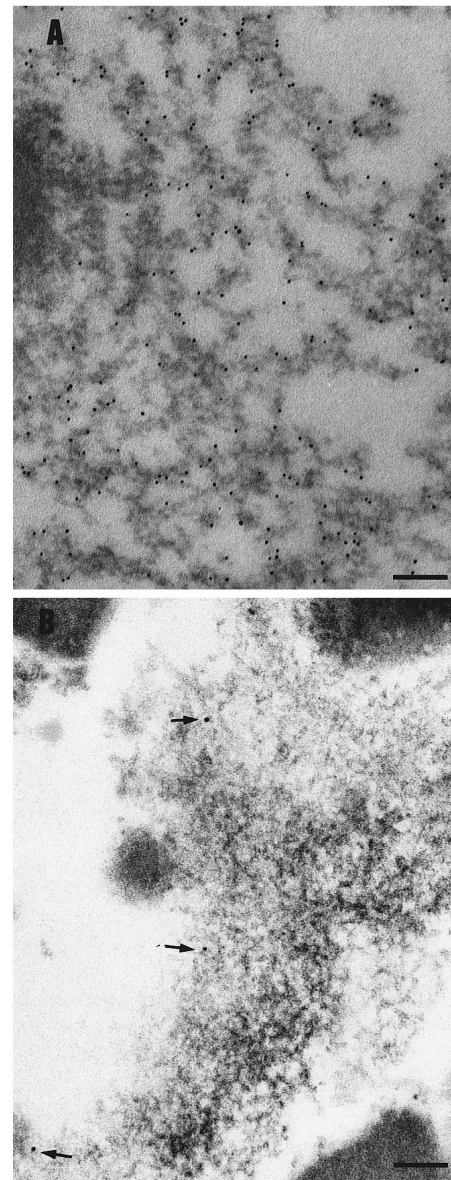


Figure 11. Thin sections of lens fibre ghosts from the cortical region of lens from transgenic mouse strain 55. Immunogold labelling with mono-vim (*A*) and poly-des (*B*). Note disruption of vimentin network (*A*) and poor accessibility of accumulated mutant desmin to the antibody (*B*). Gold particles (arrows) are 5 nm. Bars = 100 nm. (Reprinted with permission from the European Journal of Cell Biology)

able, as one may argue that the phenomenon cataract is a phenotypic feature.

Moreover, one should take into consideration that lens opacification is always preceded by subtle alterations which already are typical for an abnormal phenotype. The finding of the distorted differentiation process preceding incisor tooth formation demonstrates that introduction of the transgene may interfere with normal differentiation in any tissue where it is expressed.



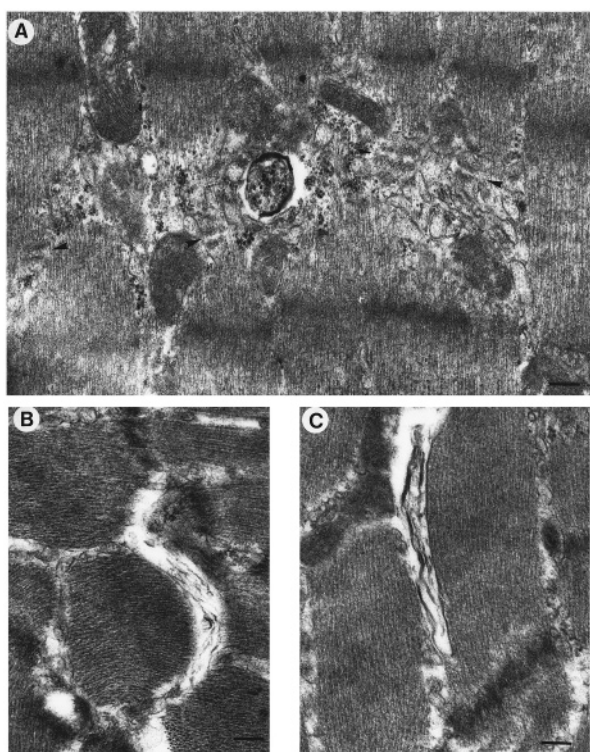


Figure 12. Longitudinal thin sections of sartorius muscle from transgenic mice stained with uranyl acetate and lead citrate. A. The T system at the A-I band junction is disrupted (arrowheads). Note the sequestration of sarcoplasmic areas wrapped by multi-layered membrane profiles and disruption of the membrane vesicles of the sarcoplasmic reticulum longitudinal tubular system. (B, C) Transverse (B) and longitudinal (C) sections of myofibrils which are wrapped by fibrillar and membranous material. Bars = 200 nm. (Reprinted with permission from the European Journal of Cell Biology)

Finally, a phenomenon discovered several years ago [36] and which has to be clarified unequivocally is the mosaic expression of the transgene (fig. 3B). A straightforward interpretation might be that all lens cells actually do carry the transgene but do not have the capability of expressing the gene to a similar extent. Apparently the expression in some cells may be so low that it cannot be detected. From our results we may anticipate that this local phenotypic expression of the transgene indicated as 'mosaic' mode is not dependent on the level of transgene expression. An interesting feature is that the majority of cells expressing the transgene are actively dividing cells (fig. 3A). It cannot be excluded that, at least in a number of cells, the promoters of our constructs are downregulated as soon as the cells become post-mitotic (fig. 3B).

Our understanding of the structural and functional properties of IFs has been broadened by using transgenic mice expressing mutated IF proteins. With the

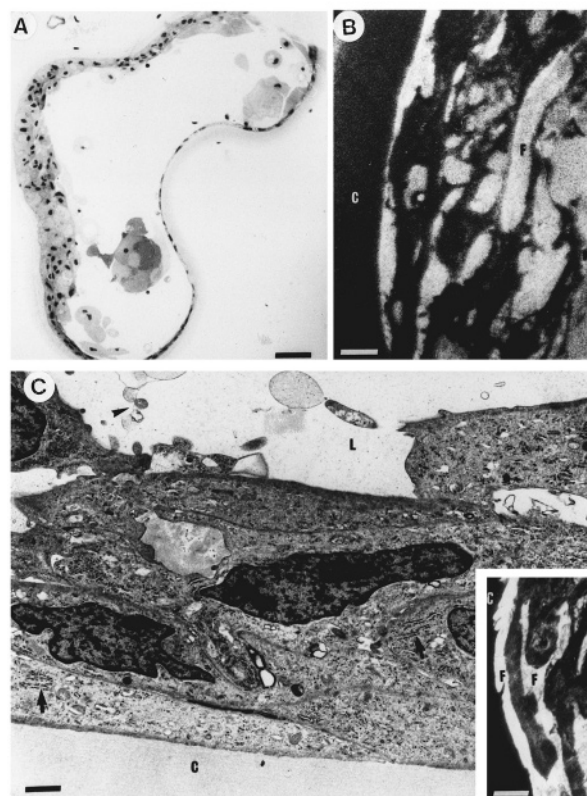


Figure 13. (A) Histological section of a cataractous adult transgenic mouse lens. The lens appears as a bladder surrounded by a thick capsule and formed by multilayered, round-shaped and elongated nucleated cells. The cavity is occupied by swollen cellular debris. Bar = 150  $\mu$ m. (B) Confocal laser scanning microscopy on a cryosection of a cataractous adult transgenic mouse lens incubated with the MDR3 antibody. Note the mosaic distribution of the transgene product highly expressed in elongated multilayered cells. The capsule (C) is not stained. Bar = 60  $\mu$ m. (C) Thin section of the wall of the lens bladder in an adult transgenic cataractous lens, stained with uranyl acetate and lead citrate. The cells are characterized by a well-developed rough endoplasmic reticulum (arrows). Note the presence of cytoplasmic vacuoles and shedded membranes (arrowheads). Bar = 2  $\mu$ m. L = lumen; C = capsule. Inset: Cryosection of the same material incubated with antivimentin antibody showing the mosaic distribution of this protein. F = fibres; C = fragment of capsule detached from the lens during sectioning. Bar inset = 80  $\mu$ m. (Reprinted with permission from the Journal of Cell Biology)

availability of knock-out mice for specific cytoskeletal or membrane genes new issues can be addressed. One of the problems that has to be firmly established is the possibility that deletion of a single cytoskeletal or membrane constituent could be compromised by an alternative component capable of maintaining stable assembly of the membrane-cytoskeleton complex. Also, the elucidation of the various aspects of the aberrant differentiation will be necessary. We believe that the results of such studies may have important implications for forthcoming efforts in the field of gene therapy.

Table 3. Cataract in transgenic mice.

Promoter <sup>1</sup>	Coding region	Expression in lens and consequences	Ref.
1 $\alpha A$ (m)	E6-E7-SV40 polyA	inhibition of lens fibre denucleation inhibition of elongation, cataract, lens tumours	53
2 $\alpha A$ (m)	CRABP-SV40 polyA	ablated secondary lens fibre differentiation	54
3 $\alpha A$ (m)	stop-SV40 TAg	cataract, lens tumours	55
4 $\alpha A$ (m)	RAR-LacZ-pA	cataract, microphthalmia	56
5 $\alpha A$ (m)	huAR-SV40 polyA	galactose and diabetic cataract microphthalmia	57
6 $\alpha A$ (m)	H-2D <sup>d</sup>	cataract, microphthalmia	58
7 $\alpha A$ (m)	HIV-1 protease	cataract, lens hydration	59
8 $\alpha A$ (m)	HIV-1 protease	cataract	60
9 $\gamma$ II (m)	tox 176	ablation of fibre cells, anophthalmia	51
10 vimentin (h)	desmin (h)	denucleation and elongation impaired, perturbed intermediate filament assembly, cataract	35, 62
11 vimentin (c)	vimentin (c) cataract	denucleation and elongation impaired,	63
12 vimentin (h)	P-glycoprotein	differentiation of primary fibres affected, terminal differentiation perturbed, microphthalmia, osmotic type cataract	41
13 MSV	murine NF-L	cataract	64
14 MSV-SV40	SV40 TAg	ablated secondary fibre differentiation, cataract	65
15 type I GT (hu)	rasT24 oncogene	cataract, microphthalmia	66
16 rho IF- $\gamma$	IFN- $\gamma$	cataract	67

$\alpha A$  (m) = mouse  $\alpha A$ -crystallin; E6, E7 = oncoproteins from human papilloma virus (HPV); SV40 = simian virus; polyA (PA) = polyadenylic acid; CRABP = cellular retinoic acid-binding protein; stop = regulatory region composed of the C terminal sequence of yeast His 3 gene, 825 bp of SV40 polyadenylation signal region and a synthetic oligonucleotide; TAg = large tumor antigen; RAR = retinoic acid receptor; LacZ =  $\beta$ -galactosidase;  $\gamma$ II =  $\gamma$ -crystallin II; tox176 = attenuated form of the diphtheria toxin-A gene; h = hamster; c = chicken; P-glycoprotein + ATP-dependent efflux pump; MSV = mouse sarcoma virus; NF-L = neurofilament low molecular weight subunit; IRBP = interstitial retinol-binding protein; rho = rhodopsin; IFN- $\gamma$  = interferon-gamma.

<sup>1</sup>Lens-specific promoters: 1–9; non-lens-specific: 10–16.

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