

Vascular permeability in a human tumour xenograft: molecular charge dependence

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Summary Molecular charge is one of the main determinants of transvascular transport. There are, however, no data available on the effect of molecular charge on microvascular permeability of macromolecules in solid tumours. To this end, we measured tumour microvascular permeability to different proteins having similar size but different charge. Measurements were performed in the human colon adenocarcinoma LS174T transplanted in transparent dorsal skinfold chambers in severe combined immunodeficient (SCID) mice. Bovine serum albumin (BSA) and IgG were fluorescently labelled and were either cationized by conjugation with hexamethylenediamine or anionized by succinylation. The molecules were injected i.v. and the fluorescence in tumour tissue was quantified by intravital fluorescence microscopy. The fluorescence intensity and pharmacokinetic data were used to calculate the microvascular permeability. We found that tumour vascular permeability of cationized BSA (pI-range: 8.6–9.1) and IgG (pI: 8.6–9.3) was more than two-fold higher (4.25 and 4.65×10^{-7} cm s⁻¹) than that of the anionized BSA (pI \approx 2.0) and IgG (pI: 3.0–3.9; 1.11 and 1.93×10^{-7} cm s⁻¹, respectively). Our results indicate that positively charged molecules extravasate faster in solid tumours compared to the similar-sized compounds with neutral or negative charges. However, the plasma clearance of cationic molecules was $\sim 2 \times$ faster than that of anionic ones, indicating that the modification of proteins enhances drug delivery to normal organs as well. Therefore, caution should be exercised when such a strategy is used to improve drug and gene delivery to solid tumours. © 2000 Cancer Research Campaign

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Inadequate delivery of macromolecules limits the efficacy of systemic therapies of tumours with cytokines, monoclonal antibodies, and oligonucleotides (Presant et al, 1994; Jain, 1994; 1998). One of the major barriers to the delivery is the tumour microvessel wall. The permeability of the wall to macromolecules is heterogeneous in tumours: there exist regions where tumour vessels are nearly impermeable to these molecules, while other regions are much more leaky than normal vessels (Yuan et al, 1994). The leakiness of tumour vessels is presumably caused by the large pores (\sim several hundred microns) in the vascular endothelium (Yuan et al, 1995), and discontinuity in the basement membrane (Bosman et al, 1985). Therefore, electrically neutral macromolecules and nanoparticles circulating in the blood accumulate preferentially in tumour tissues.

In addition to the size selectivity, the transvascular transport is influenced by the charge of molecules. The charge-dependence of the transvascular transport is especially important for gene delivery to tumours, using polycationic liposomes (Gao and Huang, 1995) or amino polymers (Goldman et al, 1997). These non-viral vectors have shown promising results in gene delivery across the cell membrane (Gao and Huang, 1995; Goldman et al, 1997). Recently, Thurston et al (1998) demonstrated that cationic liposomes are preferentially taken up by angiogenic tumour endothelium, mostly through vesicular organelles.

In normal tissues, the luminal endothelial membrane is negatively charged (Curry et al, 1987; Turner et al, 1983; Vehaskari et al, 1984; Deen et al, 1980; Baldwin et al, 1991; Dermietzel et al, 1983). Thus, it restricts the extravasation of anionic macromolecules, as has been demonstrated in cultured endothelial cell monolayer (Sahagun et al, 1990) and various normal tissues (Jain, 1997; Curry et al, 1987; Turner et al, 1983; Vehaskari et al, 1984; Deen et al, 1980; Baldwin et al, 1991; Dermietzel et al, 1983; Adamson et al, 1988; Barrowcliffe et al, 1990; Gandhi and Bell, 1992; Gilchrist and Parker, 1985; Haraldsson et al, 1983; Khaw et al, 1991; Leypoldt and Henderson, 1993; Michel and Phillips, 1985; Öjteg et al, 1987; Parker et al, 1985; Rasio and Goresky, 1985; Triguero et al, 1989). Adamson et al (1988) have demonstrated that the microvascular permeability to α -lactalbumin ($M_r = 14\,176$, net charge -10) is approximately 50% of that to ribonuclease ($M_r = 13\,683$, net charge $+4$), suggesting that the microvascular permeability to the positively charged molecules is higher than the permeability to the negative ones. The transport restriction of anionic macromolecules is crucial for maintaining a fluid homeostasis in the body, due to the osmotic effect (Curry, 1984). However, tumour vessels are significantly different from normal vessels. The role of molecular charge in the transport across tumour vessel wall is still unknown. To address this issue, we quantified tumour microvascular permeability of bovine serum albumin (BSA) and IgG, as well as their derivatives, on which different charge groups were covalently attached. The quantification was based on the intravital fluorescence microscopy, in combination with the human colon adenocarcinoma LS174T model implanted in the transparent dorsal skinfold chambers in

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severe combined immunodeficient (SCID) mice. The results of the present study may have important implications in designing vehicles for delivering genes and drugs to solid tumours.

MATERIALS AND METHODS

Animals and tumour model

Dorsal skinfold chambers bearing the LS174T tumour were prepared in male severe combined immunodeficient (SCID) mice as described earlier (Leunig et al, 1992). In brief, titanium chambers were implanted in the dorsal skin of mice (male, 6–8 weeks old, 25–35 g) under anaesthesia (75 mg ketamine hydrochloride and 25 mg xylazine kg⁻¹ body weight, subcutaneously). Two days later, 2 µl of a dense suspension of human colon carcinoma cells (approx. 2×10^5 cells in phosphate buffered saline) was inoculated onto the striated muscle layer of the subcutaneous tissue in the chambers. Experiments for measurement of permeability were performed 2 weeks after tumour cell implantation.

Preparation of charge-modified BSA

Bovine serum albumin (A7030; Sigma Chemical Co, St. Louis, MO, USA) was first fluorescently labelled by conjugation with carboxytetramethylrhodamine succinimidyl ester (C-1171; Molecular Probes, Eugene, OR, USA). Free dye was removed on a gel filtration column (Econo-Pac 10DG; Bio-Rad Laboratories, Hercules, CA, USA) equilibrated with 50 mM phosphate buffered saline (PBS; Sigma) containing 0.002% sodium azide (Sigma). This procedure yielded a molar dye/protein ratio of 1.3. Subsequently, aliquots of the BSA solution were anionized by succinylation, or cationized by conjugation with hexamethylenediamine. For succinylation (Klotz, 1967; Rennke and Venkatchalam, 1978), 10 mg of succinic anhydride (Sigma) in 50 ml of dimethylsulfoxide (DMSO, Sigma) was added dropwise in small increments to the solution of 10 mg protein in 1 ml of 0.2 M bicarbonate buffer, pH 8.0. During the incubation time (30 min at room temperature) pH was maintained at 8.0–8.5. For cationization (Triguero et al, 1989; Kumagai et al, 1987; Hoare and Koshland, 1967), free carboxylic groups of the protein (10 mg in 1 ml of 5 mM MES, pH 5.3, if necessary pH was maintained at 5.3 with 1 M HCl) were activated with 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide (CDI, Sigma). For this purpose, two portions of CDI (10 mg in 100 ml water) were added to the protein solution with a 10 min interval. Immediately after the addition of the second portion of CDI, the activated protein was added upon stirring into 2 ml of 2 M hexamethylenediamine (Sigma) in water. pH of the solution was adjusted to 6.8 and the mixture was incubated for 5 h at room temperature. After conjugation, the products were purified by overnight dialysis, and high molecular weight aggregates were removed by elution through a Sepharose CL-6B column (Pharmacia). The major protein peak was pooled and stored at 4°C.

Preparation of charge modified IgG

The solvent of mouse monoclonal IgG antibody (MOPC21; M-9269, Sigma) was exchanged to 0.2 M sodium bicarbonate buffer on a gel filtration column (Econo-Pac 10DG; Bio-Rad) and the solution concentrated to 1 mg protein ml⁻¹ by centrifugal filtration (Ultrafree-CL; Millipore, Bedford, MA, USA). pH was

adjusted to 9.3. For fluorescent labelling, Cyanine 3 monofunctional dye (Cy3-Mono-OSu) (PA13104; Amersham Life Science, Arlington Heights, IL, USA) was used to yield high fluorescence intensity, which allowed in vivo experiments with small amounts of IgG (0.5 mg IgG animal⁻¹). Cy3-Mono-OSu was added at a ratio of 2 mg dye 10 mg protein⁻¹ and the solution stirred at room temperature for 60 min. Free dye was removed on a gel filtration column (Econo-Pac 10DG; Bio-Rad) equilibrated with 50 mM PBS containing 0.002% sodium azide (Sigma), and the solution concentrated to 1.8 mg protein ml⁻¹ by centrifugal filtration. Subsequently, anionic and cationic derivatives of IgG were prepared as described above for BSA.

Measurement of molecular charge and weight

We did not quantify the net charge per molecule, since it depends on local tissue environment, especially extracellular pH which is heterogeneous in solid tumours (Helmlinger et al, 1997). Therefore, only the fraction of free lysine amino-groups in proteins that were modified with charge-bearing groups was quantified, using trinitrobenzenesulphonic acid titration method (Fields, 1971). We found that approximately 25–35% of all amino-groups accessible for modification was modified with charge-bearing groups. In addition, isoelectric points of the proteins were determined by isoelectric focusing using polyacrylamide gel slabs on a vertical electrofocusing apparatus. pI was quantified by comparison with protein standards (Bio-Rad) after Coomassie Blue staining of the gels. The proteins' molecular weight was analysed by SDS-PAGE (Mini-Protean II; Bio-Rad) with and without reducing agent β-mercaptoethanol in the sample buffer. Molecular size of native and charge-modified proteins was estimated by HPLC on analytical gel filtration column (Shodex Protein KW 804; Waters, Milford, USA) with precolumn (Shodex Protein WS 800P) using Waters 600E system controller and 996 photodiode array detector. Mobile phase was 0.05 M phosphate buffer at pH 7.4 and flow rate was 0.7 ml min⁻¹. We used three standard molecules: chymotrypsin (MW 23 kDa), amylase (MW 200 kDa), and thyroglobulin (MW 669 kDa), and found that positive, negative and native BSA and IgG gave indistinguishable peaks. All BSAs showed the same peak not very far from chymotrypsin, while all IgGs showed the same peaks close to amylase. These data indicate that charge modification did not significantly change the size of molecules.

Measurement of tumour microvascular permeability

Effective microvascular permeability of the different proteins was measured using fluorescence microscopy as reported previously (Yuan et al, 1993; 1995). In brief, animals were anaesthetized as described above and immobilized in a polycarbonate tube on the microscope stage. The fluorescently labelled protein was dissolved in PBS and injected into the tail vein as a bolus (0.1 ml 25 g body weight⁻¹). Fluorescence intensity of tumour tissue was monitored by a 20× objective of a fluorescence microscope (Axioplan, Zeiss), and quantified over a period of 20 min by a photomultiplier. We have quantified S/V in different tumours of the same cell line (LS174T) and found that $S/V = 239 \pm 82 \text{ cm}^2 \text{ cm}^{-3}$ (Yuan et al, 1993). Therefore, we used the mean value of S/V in the estimation of microvascular permeability. In separate experiments using three animals per protein, the time constant of plasma clearance (K) was quantified through collecting arterial blood

Table 1 Characteristics of tracer molecules and their microvascular permeability in tumours. Anionized and cationized BSA and IgG were prepared from the native proteins as described in Materials and methods. * $P < 0.01$ vs the corresponding anionized compound (Mann-Whitney U test)

Name	M_r	pI ^a (range)	Median K ^a (range) (100s)	Median P_v ^a (range) (10^{-7} cm s ⁻¹)
Native BSA ^b	66 000	4.5	80.5 (77–130)	1.61 (0.65–1.93)
Anionized BSA	64 000	≈2.0	40.8 (37–45)	1.11 (0.95–2.38)
Cationized BSA	64 000	8.6–9.1	12.7 (12–13)	4.25* (3.57–5.34)
Native IgG ^b	160 000	6.0	50.0 (34–82)	2.82 (1.47–4.07)
Anionized IgG	155 000	3.0–3.9	39.3 (37–44)	1.93 (1.11–2.53)
Cationized IgG	155 000	8.6–9.3	11.0 (5–15)	4.65* (4.25–5.27)

^apI, isoelectric point; K, time constant of concentration decay in the plasma; P_v , microvascular permeability. ^bData from Yuan et al (1995).

samples at $t = 0, 1, 3, 5, 10$ and 30 min following injection of the proteins, and fitting these data to an exponential function. Tumour microvascular permeability from experiments with 6–7 individual tumours for each of the different proteins, was calculated from these data according to Yuan et al (1993). Results for microvascular permeability are given as range or median \pm standard error of median. Mann-Whitney U test was used to compare the difference in tumour microvascular permeability between the altered proteins.

Mathematical modeling of tumour uptake

Electrical charge of molecules affects both transvascular transport and plasma clearance of drugs. Therefore, the amount of drug accumulation in solid tumours depends on the balance between these two processes. In this study, a two-compartmental model was used to investigate the effect of charge manipulation on the total accumulation of drugs in tumours. The governing equation of the average concentration of drugs in tumours (C) is as follows:

$$(1) \quad \frac{dC}{dt} = \frac{PS}{V} (C_p - C) \\ C_p = C_{p0} e^{-\frac{t}{K}}$$

where P is the vascular permeability, S/V is the vascular surface area per unit tissue volume, t is the time, C_{p0} is the plasma concentration (C_p) immediately after intravenous injection, and K is the time constant of plasma clearance. We assumed that the initial concentration in tumours was zero, thus the solution of equation 1 is:

$$(2) \quad \frac{C}{C_{p0}} = \frac{(PS/V)K}{1 - (PS/V)K} (e^{-\frac{t}{V}} - e^{-\frac{t}{K}})$$

We further assumed that $S/V = 239 \text{ cm}^2 \text{ cm}^{-3}$ (Yuan et al, 1993). The values of K and P used in the simulation were measured directly in this study (see Table 1).

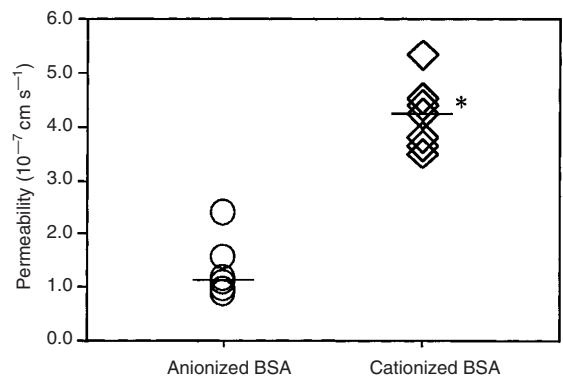


Figure 1 Microvascular permeability of tumours to anionized ($n = 6$ tumours) and cationized BSA ($n = 7$ tumours). Bars depict median values of the groups. * $P < 0.01$ (Mann-Whitney U test)

RESULTS

Fluorescence microscopy revealed that the different proteins extravasated homogeneously along the vessel walls. Tumour microvascular permeability (P_v) to anionized and cationized BSA is shown in Figure 1. Permeability values were low for negatively charged BSA ($P_v = 1.11 \pm 0.44 \times 10^{-7} \text{ cm s}^{-1}$; median \pm standard error of median), and almost fourfold higher to positively charged BSA ($P_v = 4.25 \pm 0.26 \times 10^{-7} \text{ cm s}^{-1}$). Characteristics of tracer molecules are given in Table 1.

The microvascular permeability in tumours of anionized and cationized IgG was slightly above values measured for corresponding BSA samples (Figure 2, Table 1). Similar to the data for charge-modified BSA, cationized IgG revealed highest permeability ($P_v = 4.65 \pm 0.29 \times 10^{-7} \text{ cm s}^{-1}$), and permeability of anionized IgG ($P_v = 1.93 \pm 0.41 \times 10^{-7} \text{ cm s}^{-1}$) was below the value measured for the native protein.

The plasma clearance rate revealed a similar behaviour for both proteins (Table 1). The time constant of the plasma clearance (K) was highest for the native proteins, slightly lower for the anionized compounds, and significantly lower for cationized BSA and IgG. These results are consistent with the data in the literature (Deen *et al*, 1980).

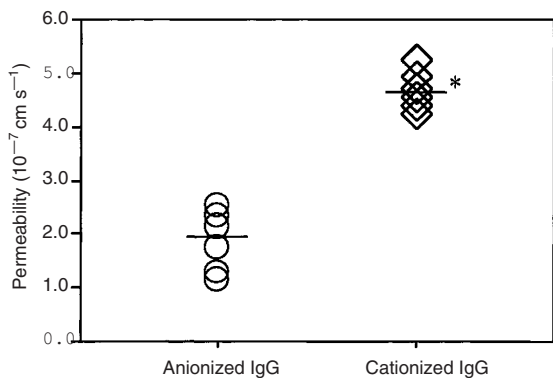


Figure 2 Tumour microvascular permeability to anionized ($n = 6$ tumours) and cationized IgG ($n = 6$ tumours). Bars depict median values of the groups. * $P < 0.01$ (Mann-Whitney U test)

The effect of charge on the accumulation of molecules in tumours was investigated, based on the mathematical simulation of average concentrations. We found that charge modification of BSA and IgG decreased the total accumulation of drugs (Figure 3A), through either reduction in microvascular permeability or increase in the plasma clearance. The total accumulation in tumours could be increased through cationization of drugs, if the plasma concentration was maintained at a constant level, i.e. the time constant of plasma clearance, $K \rightarrow \infty$ (Figure 3B). In this case, the accumulation was affected only by the microvascular permeability.

DISCUSSION

Molecular charge-dependence of vascular permeability

In comparison to tumour microvascular permeability of native BSA and native IgG (Table 1, Yuan et al, 1995), the permeability values were lower for negatively charged BSA and IgG, and more than twofold higher for the positively charged proteins. The charge-dependence in vascular permeability in tumours is consistent with observations in normal tissues (Jain, 1997; Curry et al, 1987; Turner et al, 1983; Vehaskari et al, 1984; Deen et al, 1980; Baldwin et al, 1991; Dermietzel et al, 1983; Adamson et al, 1988; Barrowcliffe et al, 1990; Gandhi and Bell, 1992; Gilchrist and Parker, 1985; Haraldsson et al, 1983; Khaw et al, 1991; Leyboldt and Henderson, 1993; Michel and Phillips, 1985; Öjteg et al, 1987; Parker et al, 1985; Rasio and Goresky, 1985; Triguero et al, 1989), suggesting the luminal surface of tumour vascular endothelium is negatively charged as well.

The effect of molecular charge on microvascular permeability has been studied in several organs including the mesentery (Adamson et al, 1988), the kidney (Rennke and Venkatchalam, 1978), the lung (Gilchrist and Parker, 1985) and the brain (Kumagai et al, 1987). On the one hand, negatively charged glycoproteins, coated on the luminal side of the endothelium known as glycocalyx in various normal organs (Turner et al, 1983; Vink and Duling, 1996) form a physiological barrier for the transcapillary movement of negatively charged molecules. Vehaskari et al (1984) have demonstrated that intravenous injection of polycations causes a significant increase in haematocrit and decrease in plasma albumin in nephrectomized rats. These results suggest that the

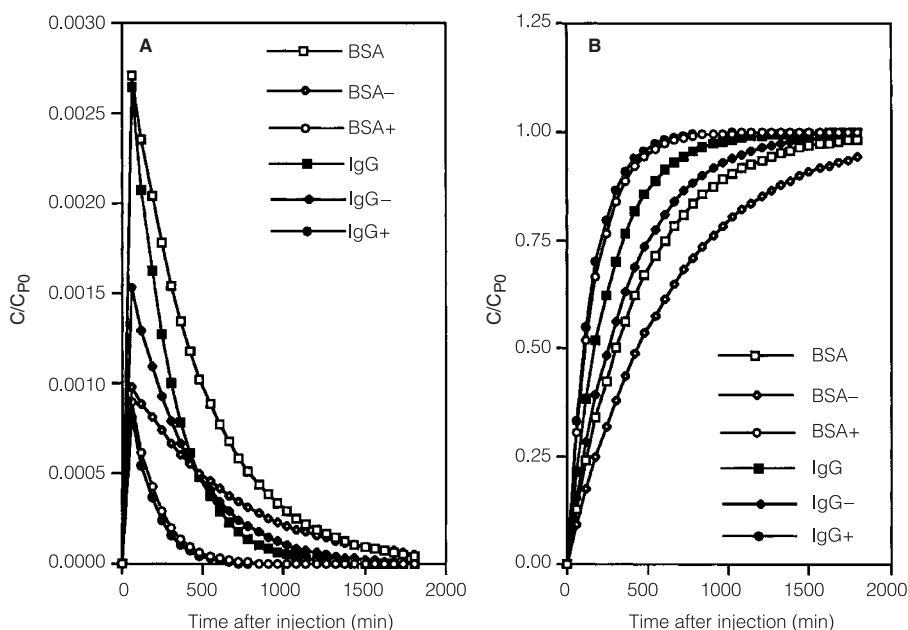


Figure 3 Mathematical simulation of the average concentration (C) of BSA and IgG in a solid tumour as a function of time. (A) Simulation of the relation between tumour concentration and plasma concentration (C/C_{p0}) for the different proteins based on our experimental results following bolus injection. (B) Simulation of the relation between tumour concentration and plasma concentration (C/C_{p0}) for the different proteins, if the plasma concentration was maintained at a constant level: '+' and '-' indicate anionic or cationic modification of molecules

neutralization of the negative charge sites on the vessel wall can abolish the barrier effect of the glycocalyx layer on the transport of anionic macromolecules. On the other hand, the negatively charged endothelial surface may facilitate interactions between cationic molecules and the vascular endothelium. Kumagai et al (1987) observed a rapid binding and endocytosis of cationized albumin to isolated brain capillaries, suggesting that the anionic barrier facilitates the endothelial absorption and endocytosis of cationic molecules.

Cationic liposomes have been demonstrated to be taken up by endothelial cells in an organ-specific pattern with highest accumulation in the lung (McLean et al, 1997). However, angiogenic endothelial cells in tumours and in chronic inflammation revealed a preferential uptake of cationic liposomes, with a high proportion being associated with endothelial fenestrae (Thurston et al, 1998). Endothelial fenestrae are very frequently found on tumour endothelium (Roberts and Palade, 1997; Hobbs et al, 1998), and may thus be the site of extravasation of cationic proteins.

The absorption of cationic molecules to the endothelium and their subsequent internalization may potentially affect the vascular permeability measurement, since most techniques used are based on the quantification of the total accumulation of tracers in the vicinity of vessels. Therefore, the apparent vascular permeability data of cationic molecules accounts for the combined effect of the endothelial uptake (bound plus internalized) and the extravasation. The exact contributions of individual mechanisms to the apparent vascular permeability cannot be determined separately. However, we did not observe a significant staining of tumour vessels with the fluorescent tracers, indicating that the amount of bound proteins was insignificant compared to the extravasated molecules. The ratio of the bound versus extravasated macromolecules should be much lower in tumours than normal tissues, since tumour vascular endothelium is leaky (Yuan et al, 1993; 1995; Wu et al, 1993; Gerlowski and Jain, 1986; Dvorak et al, 1995) and the basement membrane is incomplete (Hobbs et al, 1998). Therefore, the effect of the endothelial uptake on the permeability measurement, if any, may be less important in tumours.

Plasma clearance of charged macromolecules

The mechanism of the plasma clearance of macromolecules is different from that of small molecules. The cut-off size of renally filterable proteins and peptides is approximately 6.3 nm ($M_r \sim 60\,000$) (Behr et al, 1995), which is slightly smaller than the size of albumin. For dextran, this size is increased to approximately 8 nm (Deen et al, 1980), presumably due to the difference in the molecular configuration. Consequently, the excretion of proteins larger than the renally filterable size may initially require catabolism in the liver; and smaller metabolites can then be filtered and excreted through the kidney (Behr et al, 1995). In addition to liver and kidney, organs such as the lung (McLean et al, 1997) and the immune system may be involved in the clearance of charge modified molecules. Bass et al (1990) have shown that cationization of albumin alters its molecular conformation. The conformation change may increase the immunogenicity of exogenous molecules, in part due to an increased uptake by antigen-presenting cells (Apple et al, 1988). The immune response can cause vascular leakiness and significantly reduce the plasma half-life of molecules (Yamamoto et al, 1986; Adamski et al, 1987). However, such a response was unlikely in our experiments, since the permeability was quantified within 1 h after the intravenous injection of

proteins. Taken together, we hypothesize that the charge-dependence of the plasma clearance soon after the injection depends on three effects: microvascular permeability in various normal tissues, catabolism of the macromolecules in the liver, and the clearance of their metabolites in the plasma through the kidney.

Implications for drug and gene delivery to solid tumours

In conclusion, positively charged macromolecules extravasate faster in solid tumours compared to similar sized compounds with neutral or negative charges, suggesting that cationization may enhance delivery of therapeutic agents to solid tumours. However, the rapid clearance of cationic molecules from the plasma indicates that the charge modification enhances the drug delivery to normal organs as well. Therefore, caution should be exercised when such a strategy is used to improve the drug and gene delivery to solid tumours. This issue is especially important for gene therapy where polycationic liposomes and amino polymers are used as the delivering vectors. In general, the tumour specificity versus the enhanced delivery of drugs and genes has to be balanced in the development and evaluation of charge-modified drugs or vectors.

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