



Structurally screening calixarenes as peptide transport activators†

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Calixarenes are reportedly excellent activators that can remarkably improve the transport efficiencies of cell penetrating peptides. We employed eight calixarenes to systematically study the influence of structure on activation efficiency, which revealed that the scaffold, head group, and alkyl chain are all significant factors for activation efficiency by affecting affinities with the peptide and membrane.

Cellular membranes protect cells from harmful exogenous molecules while preventing some useful biologically active molecules from entering cells; hence transporting biologically active molecules into cells in a safe and effective manner remains a significant topic of interest.^{1,2} Cell penetrating peptides (CPPs) are short peptides rich in arginine and very membrane permeable.^{3–6} Benefiting from their low cytotoxicities, low immunogenicities, and high efficiencies,^{1,3,7,8} these peptides have been widely used to transport proteins, peptides, nucleotides, RNA, and DNA across membranes.^{1,4,8–12} Some counteranions (activators) improve CPP transport efficiency.^{13–18} Hence, designing and developing excellent activators have been scientific interests for many years; an ideal activator should recognize the CPP and interact with the membrane.

Calixarenes, which are third-generation supramolecular hosts, have pre-organized structures and tunable recognition and assembly abilities;¹⁹ hence, they are potentially excellent activators.²⁰ As early as 2005, Matile *et al.* reported calixarene-based activators,²¹ and our group recently reported that *p*-sulfonatocalix[4]arene tetrapentyl ether acts as an extremely effective activator, with EC₅₀ values more than three orders of magnitude smaller than those of classic activators.²² Our group also reported that amphiphilic sulfonatocalix[5]arene (sCx5-6C)

activates lysine-rich peptide and protein transport, which cannot be achieved using other established counterion activators.¹⁷ While calixarene has a tunable scaffold that is easily modified, leading to diverse structures,²⁰ the effect of calixarene structure on activation efficiency has not been explored. Investigating the calixarene structure–activity relationship is helpful for understanding their ultra-high activation efficiencies and for designing new activators.

In this work, we studied the effect of calixarene structure on activation efficiency by employing eight calixarenes (Fig. 2 and Fig. S1, ESI†) and dividing them into three groups: amphiphilic sulfonatocalix[4]arene (sCx4-6C), sCx5-6C, amphiphilic sulfonatocalix[6]arene (sCx6-6C), and amphiphilic sulfonatocalix[8]arene (sCx8-6C) were used to study the influence of the scaffold; sCx5-6C, amphiphilic carboxycalix[5]arene (cCx5-6C), and amphiphilic phosphinocalix[5]arene (pCx5-6C) were used to study the influence of the head group, while sulfonatocalix[4]arene (sCx4), sCx4-6C, and amphiphilic sulfonatocalix[4]arene (sCx4-12C) were used to study the influence of the alkyl chain length.

The carboxyfluorescein (CF) assay (Fig. 1) (see ESI† for details) was used to study the activation efficiencies of the chosen calixarenes,^{16,17,22,24,25} and nonaarginine (Arg9) as well as α -poly-L-lysine (polyLys) were used in this study (Fig. 2). The time-dependent fluorescence of CF-encapsulated large unilamellar egg-yolk phosphatidylcholine vesicles (EYPC-LUVs \supset CF) (Fig. S5, ESI†) was measured in response to varying concentrations of calixarene and appropriate concentrations of Arg9 or polyLys (0.350 μ M for Arg9 or 0.125 μ M for polyLys; Fig. S6 and S7, ESI†). No CF efflux was observed prior to the addition of Arg9 or polyLys, which indicates that the calixarenes do not cause transient pores or vesicle lysis (Fig. S6–S9, ESI†).

The data were fitted to the Hill equation (ESI,† eqn (S2)), which provided maximum transmembrane activity (Y_{\max}), EC₅₀ values and Hill coefficient (N). Transport efficiency (E), which is defined in eqn (S3) in the ESI,† was introduced to simultaneously reflect Y_{\max} and EC₅₀,²¹ a large E value corresponds to a high Y_{\max} and low EC₅₀ (*i.e.*, high transport efficiency).

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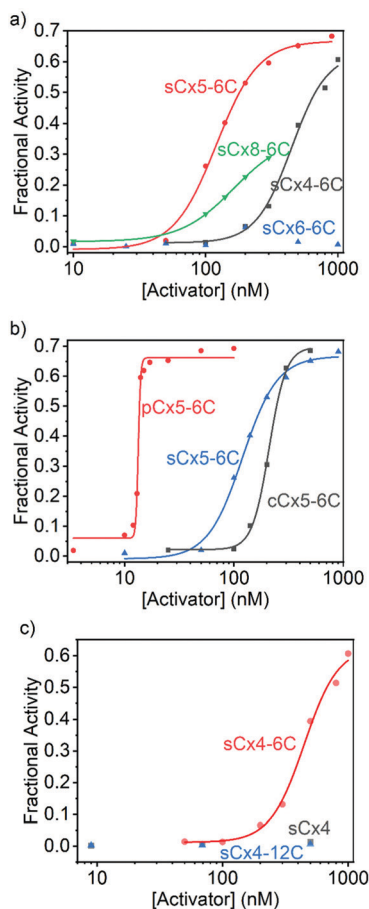


Fig. 4 Hill plots for (a) sCx4-6C, sCx5-6C, sCx6-6C, and sCx8-6C, (b) sCx5-6C, cCx5-6C, and pCx5-6C, and (c) sCx4, sCx4-6C, and sCx4-12C for polyLys transport activation. With the exception of sCx6-6C, data for sCx4, and sCx4-12C, were fitted to the Hill equation.

Table 2 Y_{\max} , EC_{50} , N , and E values of calixarenes for polyLys transport. Experiments were performed at 25 °C in HEPES buffer (10 mM HEPES, 107 mM NaCl, pH = 7.4)

Activator	Y_{\max}	EC_{50} (nM)	N	E
sCx4	—	—	—	—
sCx4-6C	0.628 ± 0.049	446 ± 36	3.22 ± 0.70	10.2 ± 0.9
sCx4-12C	—	—	—	—
sCx5-6C	0.667 ± 0.018	120 ± 6	2.72 ± 0.35	12.7 ± 0.4
sCx6-6C	—	—	—	—
sCx8-6C	0.362 ± 0.005	164 ± 3	2.15 ± 0.05	6.65 ± 0.11
cCx5-6C	0.698 ± 0.018	210 ± 5	5.60 ± 0.67	12.5 ± 0.4
pCx5-6C	0.662 ± 0.015	13.3 ± 0.1	40.1 ± 8.0	15.7 ± 0.4

and skeleton flexibility are important for Arg9 activation. As the negative charge increases, the interaction between calixarene and Arg9 will become stronger. In addition, a flexible skeleton can adapt well to the Arg9 sites and interact with more sites. Compared to arginine-rich peptides that exhibit “arginine magic”,^{21,27,28} lysine-rich peptides are more challenging to translocate^{29,30} because the ammonium group in lysine is kosmotropic, while the guanidinium in arginine is

chaotropic.³¹ We conclude that the translocation of lysine-rich peptides requires special complexation rather than electrostatic interactions to provide the strong affinity required to promote the desolvation of the ammonium groups. To verify this, we performed the ¹H NMR experiments. We previously reported that lysine was encapsulated into the cavity of sCx5-6C in a threading manner,¹⁷ and the NMR results obtained in this work show that the binding mode has not changed when lysine becomes part of the peptide (Ac-GKG-NH₂). As shown in Fig. S11 and S12 (ESI[†]), the complexation-induced shifts become larger and larger from lysine protons α to ϵ , indicating increasing proximity to the cavity center, which is consistent with the threading mode.^{32,33} As control, the complexation-induced shifts were also observed in the case of Ac-GKG-NH₂ with sCx4-6C, Ac-GRG-NH₂ with sCx4-6C or sCx5-6C (Fig. S13–S15, ESI[†]), but not as remarkable as Ac-GKG-NH₂ with sCx5-6C. This is because the cavity of calix[4]arene is too small to form the threading complex. The guanidinium residue is too large, regardless of whether for sCx4-6C or sCx5-6C, so it cannot form the threading complex either. Therefore, calix[5]arene, which has the most appropriate cavity size among the calix[*n*]arene ($n = 4, 5, 6, 8$) family for recognizing lysine, exhibits the highest activation efficiency.

Calixarenes with phosphonate groups exhibit the best activation efficiencies among the calix[5]arenes, regardless of whether for Arg9 or polyLys. There may be two factors contributing to the good efficiency of pCx5-6C. On one hand, it is well-established that phosphonates had strong affinities for guanidinium and ammonium.^{34–36} Moreover, phosphonate/guanidinium binds at vesicle surfaces reportedly stronger than in water.^{37–39} On the other hand, the solubility of pCx5-6C is much poorer than that of sCx5-6C, which indicates that pCx5-6C is less hydrophilic and may therefore be able to cross the phospholipid membrane more easily. It should be noted that the activator needs to bind moderately strongly (rather than strongly) to peptides, as an affinity that is too strong impedes peptide release, thereby reducing transport efficiency.

The sCx4-6C calixarene showed the best activation efficiency among sCx4, sCx4-6C, and sCx4-12C. The length of the alkyl chain appears to mainly affect interactions between the calixarene and the membrane. We used fluorescence polarization experiments to study the abilities of the calixarenes to embed in the membrane (see ESI[†] for details).⁴⁰ Fig. S16 (ESI[†]) shows that the addition of sCx4-6C or sCx4-12C increases the fluorescence anisotropy of membrane-bound 1,6-diphenyl-1,3,5-hexatriene in EYPC-LUVs, while the addition of sCx4 resulted in little change, which indicates that sCx4-6C and sCx4-12C are embedded in the membrane, whereas sCx4 hardly interacts with the membrane. Moreover, the addition of sCx4-12C resulted in a stronger change in fluorescence anisotropy than the addition of sCx4-6C, which suggests that sCx4-12C binds more-strongly to the membrane (Fig. S16, ESI[†]). The above results suggest that the activator needs to be amphiphilic and interact with the membrane; however, excessive membrane interaction restricts the ability of the activator-peptide complex to move, which reduces activation efficiency.

According to above experiments, pCx5-6C has the best activation efficiency among calixarenes we used. In order to verify whether this conclusion is still valid for peptides/proteins other than Arg9 and polyLys, we further employed histones. Histones are the main components of chromatin, which are rich in both arginine and lysine.⁴¹ pCx5-6C still has the best activation efficiency among all calixarenes (Fig. S10, ESI[†]), consisting with the conclusion we got when using Arg9 and polyLys.

In summary, we studied and compared the activation efficiencies of eight calixarenes for the transmembrane transportation of arginine- and lysine-rich peptides, and discussed the influence of activator structure on activation efficiency. The results show that the influence of scaffold is related to the recognition mode. The negative charge and flexible scaffold are conducive to the translocation activation of arginine-rich peptides, whereas cavity size is more important for lysine-rich peptides. The head group facilitates calixarene/peptide complexation, thereby improving activation efficiency. Alkyl chains of appropriate length that provide moderate interactions with the membrane are also necessary; a weak affinity will result in activation failure, while an affinity that is too strong restricts the ability of the activator-peptide complex to move, which reduces activation efficiency. This study provides meaningful information for the design of new peptides and (even) protein transport activators.

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Conflicts of interest

The authors declare no conflicts of interest.

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