

# Adenovirus Endocytosis

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## Abbreviations:

AAV, adeno-associated virus; AP2, adaptor protein complex 2; Ad, Adenovirus; C3b, classical complement cascade product; CAR, Coxsackie virus B Adenovirus receptor; CED-1, cell death gene for scavenger receptor which serves the removal of apoptotic cells; CHO, chinese hamster ovary; CR, complement receptor; ECM, extracellular matrix; EEA1, early endosome antigen 1; EH, Eps15 homology; EGF, epidermal growth factor; EIPA, 5-(N-ethyl-N-isopropyl) amiloride; EM, electron microscopy; GAG, glycosaminoglycan; iC3b, alternative cleavage product of the complement cascade; IgG, Immunoglobulin G; Fc $\gamma$ -R, immunoglobulin receptor; ITAM, immunoreceptor tyrosine-based activation motif; M-CSF, macrophage colony stimulating factor; MDCK, Madin-Darby canine kidney; PA, phosphatidic acid; Pb, penton base; PDGF, platelet derived growth factor; PH, pleckstrin homology; PI, phosphatidylinositol; PI3K, phosphatidylinositol 3-kinase; PKC, protein kinase C; PLC $\gamma$ , phospholipase C $\gamma$ ; RTK, receptor tyrosine kinase; SH3, Src homology 3; src, non-receptor tyrosine kinase identified in sarcoma caused by Rous sarcoma virus; SYK, tyrosine kinase; UIM, ubiquitin-based intracellular network motif

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## Summary

Pathogen entry into cells occurs by direct penetration of the plasma membrane, clathrin-mediated endocytosis, caveolar endocytosis, pinocytosis or macropinocytosis. For a particular agent, the infectious pathways are typically restricted, reflecting a tight relationship with the host. Here, we survey the uptake process of human Adenovirus (Ad) type 2 and 5 and integrate it into the cell biology of endocytosis. Ad2 and Ad5 naturally infect respiratory epithelial cells. They bind to a primary receptor, the Coxsackie virus B Ad receptor (CAR). The CAR-docked particles activate integrin coreceptors and this triggers a variety of cell responses, including endocytosis. Ad2/Ad5 endocytosis is clathrin-mediated and involves the large GTPase dynamin and the adaptor protein 2. A second endocytic process is induced simultaneously with viral uptake, macropinocytosis. Together, these pathways are associated with viral infection. Macropinocytosis requires integrins, F-actin, protein kinase C and small G-proteins of the Rho family but not dynamin. Macropinocytosis *per se* is not required for viral uptake into epithelial cells, but it appears to be a productive entry pathway of Ad artificially targeted to the high affinity Fc $\gamma$  receptor CD64 of hematopoietic cells lacking CAR. In epithelial and hematopoietic cells, the macropinosomal contents are released to the cytosol. This requires viral signalling from the surface and coincides with particle escape from endosomes and infection. It emerges that incoming Ad2 and Ad5 distinctly modulate the endocytic trafficking and disrupt selective cellular compartments. These features can be exploited for effective artificial targeting of Ad vectors to cell types of interest.

## **Adenoviruses**

Adenoviruses (Ads) are nonenveloped icosahedral DNA viruses of about 90 nm diameter. They produce progeny virions within the nucleus of infected cells and release the newly synthesized virions upon cell lysis [reviewed in 1]. The species C Ads, such as the human Ad2 and Ad5, are nononcogenic to humans and cause infections of the respiratory tracts. They are widely used as gene transfer vehicles or oncolytic agents in diseased somatic cells [2, 3]. The Ad particle is composed of an outer capsid and an inner DNA-associated core with a 36 kbp linear DNA, two terminal proteins and condensing proteins V and VII [4]. The chromosome also contains about ten copies of the cysteine protease p23 and is linked to the outer capsid by protein VI. The capsid consists mainly of hexon and is stabilized by proteins IIIa, VIII and IX. The vertices are made up of penton base (Pb) containing an exposed arginine-glycine-aspartate (RGD) motive and the protruding fiber protein which attaches the particle to target cells [5].

## **Access to the plasma membrane**

The plasma membrane of epithelial cells is protected by the extracellular matrix (ECM), a dense meshwork of proteins and sugars. The ECM has crucial functions in shaping the tissue architecture, controlling cell migration and surface receptors, and it regulates the activity of growth factors and hormones [6]. The ECM and the extracellular surface contain glycosaminoglycans (GAGs), i.e. highly sulfated, heterogeneous oligosaccharides, including heparin, heparan sulfate, chondroitin sulfate, dermatan sulfate and hyaluronic acid. GAGs can store chemokines and growth factors and also viral particles, such as herpesviruses [7], respiratory syncytial virus [8], human immunodeficiency virus [HIV, 9], adeno-associated virus [10] and foot-and-mouth disease virus [11]. Virus trapping in the ECM has been associated with a reduction [12] and in some cases with an enhancement of infection [13, 14]. The key to the plasma membrane is the viral docking to a selected cell surface receptor. This can trigger viral penetration of the plasma membrane or endocytic uptake, and thus not only overcomes the ECM but also provides opportunities to directly interfere with cell signalling [reviewed in 15].

Some Ad serotypes have been reported to bind to GAGs. Binding of the species C Ad2 and Ad5 to epithelial cells is partially inhibited by high concentrations of heparin [16]. The species D serotype Ad37 binds sialic acid linked to an oligosaccharide in an alpha 2-3 glycosidic bond [17, 18]. This residue is found on glycoproteins and glycolipids and could, in part, account for the ocular tropism of Ad37. Ad37 has recently been reported to also bind to a protein component of unknown identity [19]. Infections of polarized epithelial cells from the apical side with a variety of Ad serotypes of the species A, C, D, E and F appear to be inefficient [20]. These viruses are known to bind the Coxsackie virus B Adenovirus receptor CAR [21]. The same is true for amphotropic retroviruses [22], lentiviruses [23], adeno-associated virus (AAV) type 2 [24] and measles virus [MV, Edmonston vaccine strain, 25]. In the case of MV, the limiting factor does not seem to be the apically localized viral receptor CD46, but viral access to the plasma membrane through the glycocalyx or similar structures seems to be blocked [26, 27]. In contrast, AAV5 is reported to transduce airway cells from the apical side and it binds to alpha 2-3 linked sialic acid [24, 28]. Further experiments in HeLa cells suggested that AAV5 bound to apical microvilli and basolateral membranes with different kinetics and it was internalized from both membranes in clathrin-coated and non-coated vesicles [29]. Interestingly, a fraction of it trafficked to the Golgi complex, but the infectious pathway of AAV is presently unknown.

### **The Coxsackie virus B Adenovirus receptor (CAR)**

Fibers of all Ad species except species B have been reported to bind the Coxsackie virus B Ad receptor [21], termed CAR [30-32]. Intriguingly, the enteric Ad41 of species F encodes a second fiber gene in the late region 5 which is shorter (20 nm) than its long fiber of 34 nm [33, 34]. While the long fiber binds to CAR, the receptor(s) of the short fiber, present in six trimeric copies per virion is unknown. It has been speculated that the short fibers may be responsible for the exceptionally high selectivity of Ad41 for the digestive tract [35, 36].

CAR is an important receptor for Ad entry into a large variety of cultured cells, including respiratory endothelial and epithelial cells [20], although CAR mRNAs in human lung tissue are relatively low [37]. The mature human CAR is a type 1 transmembrane protein

of 346 amino acids. Based on sequence similarities, CAR has been placed into a conserved subfamily of the Ig superfamily of proteins, together with the A33 protein of colon carcinoma cells and the CTX protein of *X. laevis* cortical thymocytes [reviewed in 37]. More recently, a new member of the Ig superfamily has been identified, termed BT-IgSF [brain- and testis-specific Ig superfamily, 38]. Its DNA sequence predicts it to be a type-1 transmembrane protein with extracellular V and C2-type Ig domains and significant homology to CAR and endothelial cell-selective adhesion molecule [ESAM, 39]. In humans, CAR is abundantly expressed in heart, pancreas, the central and peripheral nervous system, prostate, testis and also in lung, liver and intestine. Little or no CAR is found in B and T-cells, adult muscle and on many malignant cells. Notably, the expression of CAR is developmentally regulated [40] and seems to be induced by mediators of inflammation [reviewed in 37]. There is also an intriguing inverse correlation of CAR expression and growth of prostate carcinoma cells [41]. The large cytoplasmic domain of CAR contains basolateral targeting information and might be involved in homotypic signalling, although the latter has not been directly demonstrated [42, 43]. The extracellular portion comprises 216 amino acids and is folded into a V-like immunoglobulin (Ig) fold comprising amino acid 1 to 120 and a C2-like fold of amino acids 124-210. The amino-terminal V-like Ig domain 1 (D1) functions as the Ad binding domain by binding the protruding fiber knob domain. This has been demonstrated upon expression cloning of the human and the mouse CAR in chinese hamster ovary (CHO) cells and NIH 3T3 mouse fibroblasts, respectively [30, 31]. X-ray crystallography studies subsequently showed that viral docking to D1 involves the same CAR interface as D1 dimerization [44-47]. This indicates that Ad may subvert the intrinsic property of CAR to oligomerize. Recently, it was shown that Ad2 fiber proteins that are known to be released from infected cells together with Ad particles interfere with CAR oligomerization in the infected respiratory epithelium [48]. It was suggested that fibers disrupt the adherens junctions of the epithelial barrier and promote viral release to the airway lumen, implying that CAR is also an exit receptor. Intriguingly, another study recently suggested that Ad2 Pb was exported from the infected cell prior to the release of Ad particles and, by doing so, Pb abduced the fiber protein to the surrounding cells of the monolayer [Fig. 1, and 49]. If this occurs in the infected respiratory epithelium, viral spreading is expected to be even more effective due to CAR blocking by free fiber-Pb complexes prior to the release of viral particles.

## **Adenovirus infection of the respiratory epithelium**

But how do Ads infect the respiratory epithelium? This is a burning question for essentially every respiratory agent infecting the apical membrane of polarized epithelial cells. In polarized respiratory epithelial cells and in airway cultures, the Ad receptor CAR is localized in the tight junctions near the top of the cells and along the basolateral membranes [Fig. 1, and 42, 50]. The amount of tight junction association of CAR seems to depend on the particular cell type and the culture conditions and, accordingly, CAR has been found also in adherens junctions along the lateral sides of airway cells [48]. The absence of CAR in the apical membrane is thought to preclude Ad infection of polarized airway epithelia from the luminal side. Interestingly, the expression of CAR mutants lacking the transmembrane or the cytoplasmic domains gave rise to both apically and basolaterally distributed CAR in polarized Madin-Darby canine kidney (MDCK) cells [42]. This constellation did not allow significant Ad5 transduction, unless the apical glycocalyx was experimentally removed. Another report, however, stated that in transfected airway cells apical CAR was sufficient for infection [51]. In any case, it is unlikely that incoming Ad binds to CAR in tight junctions of polarized cells. Rather, airway infections may start in specialized nonpolarized cells that express CAR on the luminal membrane, or that lesions of the epithelium prior to infection expose basolateral membranes to allow viral docking to CAR or other receptors (Fig. 1). Such models would also be compatible with the basolateral localization of integrins, an Ad coreceptor involved in viral endocytosis. However, other types of infection cannot be excluded. One interesting concept of epithelial infection has recently been discussed again, transcytosis. *Streptococcus pneumoniae* are able to co-opt polymeric Ig receptors, which normally transport antibodies across mucosal epithelial cells as part of the host defense, and thus the bacteria gain entry into airway epithelial cells [52].

## **Endocytic uptake**

Endocytosis serves to sample the extracellular environment and regulate the activity of cell surface receptors [for reviews, see e.g., 53, 54, 55]. Notably, all types of endocytosis can be abused by pathogens as gates into target cells [56]. Over the last decades, one particular type of receptor-mediated endocytosis was intensively studied, clathrin-

dependent endocytosis. It originates at specialized plasma membrane regions where the cytosolic adaptor protein complex 2 (AP2) binds the endocytic receptors and phosphoinositides and together with other cytosolic proteins mediates the formation of clathrin coats [57]. Coats function to deform the donor membrane to produce a vesicle, and they also assist in sorting of transmembrane receptors and their cargo. In addition to receptor-mediated nutrient uptake, clathrin-mediated endocytosis contributes to the initiation, propagation and downregulation of signalling [recently reviewed by 58, 59]. Less is known, however, about other receptor-mediated uptake processes, such as caveolar endocytosis and constitutive non-clathrin uptake [for recent reviews, see 60, 61]. Caveolae are flask-shaped invaginations that are formed and stabilized by the coat like protein caveolin-1. The recruitment of receptors is thought to occur by the cholesterol and sphingolipid-rich nature of caveolar membranes and this serves as a signalling platform at the cell surface. Similarly, clathrin and caveolin-independent uptake can be mediated by specialized cholesterol and sphingolipid domains, called lipid rafts.

In contrast, phagosomes and macropinosomes are designed to remove large particles, such as invading microorganisms. Phagocytosis often occurs upon activation of Fc-receptors [Fc-R, 62] and in extreme cases even appears to involve the recruitment of membranes from the endoplasmic reticulum to the plasma membrane [63]. Macropinocytosis is related to phagocytosis but is less specific, as first discovered by Lewis in 1931 [64]. It is a major endocytic pathway, found in epithelial cells, fibroblasts, neutrophils and macrophages [60, 65]. Typically, macropinocytosis is triggered by growth factor stimulation or downstream activated signalling molecules. It plays a key role in the entry of *Salmonella* and *Shigella* bacteria [66], and it is crucial for antigen presentation of dendritic cells and viral clearance [67-69]. Macropinosomes are dynamic structures formed by the closure of lamellipodia at ruffling membranes and can reach several  $\mu\text{m}$  in diameter. They assure the endocytic removal of large membrane domains, alter the adhesive and communicative properties of the cell and are involved in cell contraction and migration.

## Clathrin-mediated endocytosis

Recent studies have revealed intricate details of the regulation underlying clathrin-dependent endocytosis. Initially, clathrin is recruited by the heterodimer Hip1/Hip1R which leads to an actin-dependent formation of flattened coated-pit zones at the plasma membrane [70]. Hip1 is termed according to its interaction with the Huntingtin protein and binds the adaptor protein AP2. AP2 is a heterotetramer comprising two large subunits,  $\alpha$  and  $\beta$ 2, and two smaller chains,  $\mu$ 2 and  $\sigma$ 2 [57]. The  $\mu$ 2 subunit of AP2 recognizes a tyrosine (Y)-based signal (YxxØ) with a large hydrophobic residue (Ø) or a dileucine motif on cytosolic domains of plasma membrane receptors [71]. While the  $\beta$ 2 subunit of AP2 binds the clathrin-triskelions, the  $\alpha$  subunit is responsible for the interaction with a complex network of accessory proteins. These proteins interact with each other by various protein-protein recognition modules, including Eps15 homology (EH) domain that bind to NPF-motives on AP180/CALM, epsin and synaptojanin, and Src homology 3 (SH3) domains recognizing proline rich regions. The  $\alpha$ -subunit of AP2 preferentially binds to membranes rich in phosphatidylinositol (4,5)-bisphosphate (PI(4,5)P<sub>2</sub>) and PI(3,4,5)P<sub>3</sub>. Oligomers of the transmembrane protein synaptotagmin are thought to cooperatively tighten AP2 membrane association [72]. In many cases, the coat contains the additional monomeric protein AP180/CALM that later in the process appears to define the size of the coated vesicle [73]. The membrane curvature of clathrin-coated pits is driven by epsin binding to PI(4,5)P<sub>2</sub> [74]. PI(4,5)P<sub>2</sub>, in turn, is generated by PI(5)P-4-kinase and PI(4)P-5-kinase and stimulated by phosphatidic acid (PA). PA is produced by endophilin or generated by phospholipase D-mediated hydrolysis of phosphatidyl-choline. The enzymatic activity of PLD is enhanced by the small GTPase Arf6 and PI(4,5)P<sub>2</sub>, thus generating a positive-feedback loop. This cycle can be inhibited by the phosphoinositide phosphatase synaptojanin and by amphiphysin [for a review, see 72]. PI(4,5)P<sub>2</sub> is also recognized by epsin amino-terminal homology domains, present in epsin, Ap180/CALM and Hip1R.

Like the SH3 domain of amphiphysin, Eps15 is an important diagnostic tool to discriminate clathrin-mediated endocytosis from other types of endocytosis. It was originally identified as epidermal growth factor (EGF) receptor pathway substrate clone 15 [75]. Benmerah and colleagues showed that overexpression of an EH-domain deleted GFP-fusion protein inhibited transferrin uptake, whereas the deletion of the AP2 binding domain had no effect [76, 77]. Eps15 also plays a role in the recruitment of monoubiquitinated receptors [78].



Ubiquitin is a 76 amino acid peptide that becomes covalently attached to other proteins through an isopeptide bond between its C-terminus and a lysine residue of the target protein [reviewed by 79]. Monoubiquitination of receptors at the cytoplasmic tail serves as a signal for clathrin-mediated internalization [reviewed in 80]. In contrast, proteasomal degradation is mediated through polyubiquitination at the ubiquitin lysine residue 48 [for a review, see 81]. In the case of the alternative ubiquitination, a single ubiquitin-based intracellular network motif (UIM) is responsible for ubiquitin recognition and monoubiquitination. Two of these UIM's allow Eps15 to act as an adaptor between ubiquitinated membrane cargo and endocytic coats and their dual functionality leads to a amplification of this network in the endocytic system [78].

Another way to test if a ligand enters by endocytosis is to express dynamin mutants, such K44A-dynamin, defective of GTP loading and hydrolysis [82]. K44A-dynamin inhibits clathrin-mediated endocytosis, caveolar endocytosis and phagocytosis [reviewed by 83]. Dynamin is targeted to flattened clathrin lattices of the plasma membrane and is also found at the neck of emerging clathrin-coated vesicles by binding of its poly proline rich domain to the SH3 domain of amphiphysin and the pleckstrin homology (PH) domain recognizing PI(4,5)P<sub>2</sub> [84]. In the presence of GTP dynamins form oligomers and thus may act as a constrictase upon GDP-GTP exchange and GTP hydrolysis, thus leading to vesicle detachment. It was, however, also recognized that dynamin is a signal transducing molecule that activates downstream effectors rather than just being the constrictase. One of the dynamin effectors is endophilin which appears to exhibit a lysophosphatidic acid acyl transferase activity and may facilitate the formation of a strong membrane curvature [85]. Additional regulation is provided by the adaptor associated kinase 1 which phosphorylates the  $\mu$ 2 subunit of AP2 thus allowing initial concentration of receptors [86, 87]. Following dephosphorylation by an unknown phosphatase receptors are then rapidly internalized. Yet another regulatory mechanism can be activated by the cargo itself. Binding of EGF to the EGF receptor appears to lead to clathrin phosphorylation and facilitates EGF receptor uptake [88]. Furthermore, EGF activates the small GTPase Rab5, and provides a link between signalling and endocytic trafficking [89].

## Clathrin-mediated endocytosis of Adenovirus

Initial electron microscopy (EM) studies of Ad entry had suggested that the virus penetrated the plasma membrane [90, 91]. Subsequent EM analyses detected incoming Ad2 and Ad5 in clathrin-coated pits and clathrin-coated vesicles [92-95]. It was also noted that endocytic uptake of Ad2 from noncoated membranes could occur at very high multiplicities of infection [94]. Several functional assays subsequently showed that the infectious entry pathway of Ad2 and Ad5 occurs either in clathrin-coated or non-coated membranes [96-99]. More recently, the use of dominant-negative expression constructs, including Eps15, clathrin fragments and K44A-dynamin underlined that the infectious entry pathway of Ad2 and Ad5 into epithelial cells is clathrin-mediated [100]. This and additional requirements of the infectious pathway in epithelial cells are depicted in Fig. 2. For additional information, the reader is referred to a recent review [15].

It had been noted earlier that viral endocytosis depends on the activation of  $\alpha_v$  integrins [101], and several other integrins are also implicated in Ad endocytosis [for a review, see 15]. This suggests that Ad endocytosis is regulated. Most likely, Ad particles are endocytosed together with their integrin coreceptors, possibly by an NPXY motif present in the  $\beta_3$  and  $\beta_5$  subunits [102, 103]. NPXY motives have been implicated in the localization of certain receptors to coated pits [104]. That integrins bind to Ad particles has been directly shown by cryo-EM reconstructions and biochemical experiments using soluble  $\alpha_v\beta_5$  integrin heterodimers attaching to the RGD domains of Ad2 and Ad12 pentons in a five-fold symmetrical manner [105]. The binding of RGD peptides to the extracellular domain of  $\alpha_v\beta_3$  integrins leads to discrete conformational changes in the integrins [recently reviewed in 106]. This suggests that the incoming Ad particle activates integrins in a spatially controlled manner and triggers cell signalling. The internalization of Ad5 has been associated with an integrin-dependent activation of p85/p110 PI(3) kinase [PI3K, 107]. The products of p85/p110 activation, PI(3,4)P<sub>2</sub> and PI(3,4,5)P<sub>3</sub>, have been proposed to act as second messengers for cell cycle progression and cytoskeletal changes underlying the cell plasma membrane and for regulating vesicular trafficking [108]. For instance, PI(3,4,5)P<sub>3</sub> can bind to and activate various isoforms of protein kinase C (PKC). Another target of PI(3,4)P<sub>2</sub> and PI(3,4,5)P<sub>3</sub> is the small GTP-binding protein Rab5, involved in coordinating viral endocytosis and homotypic vesicular fusion with early endosomes [109]. It was reported that the GTP-binding defective S34N-Rab5 mutant slightly reduced

Ad5 uptake and infection. Interestingly, members of the Rho family of small GTPase, Rac1 and Cdc42 acting downstream of PI3K induced actin polymerization and promoted Ad endocytosis [110], and viral endocytosis was recently found to be complemented by macropinocytosis [100].

## Macropinocytosis

Macropinocytosis is the best studied type of receptor and coat-independent endocytosis, closely related to the Fc-R mediated phagocytosis [65, 111]. Depending on the trafficking of macropinosomes and the kind of receptor, macropinocytosis can be divided into a recycling type and a processive type (Fig. 3). Recycling macropinosomes are generated upon binding of a ligand such as EGF, platelet-derived growth factor or nerve growth factor to receptor tyrosine kinases and they can be induced by the activation of downstream activators, including GTP-Rac1 and GTP-Cdc42 [112-115]. In contrast, macrophage colony stimulating factor-triggered phagocytic cells generate processive macropinosomes that shrink in size and readily mix with late endosomes and lysosomes [116]. Processive macropinocytosis is mediated by a diverse group of receptors, including non-complement-receptor integrins, like  $\alpha_v\beta_3/\beta_5$  and  $\alpha_5\beta_1$ , lectins such as mannose receptor, the lipopolysaccharide receptor CD14 and the *Caenorhabditis elegans* scavenger receptor CED-1, and it is involved in the uptake of particles, bacteria and apoptotic cells [62, 117]. Clearly, processive macropinocytosis has several features in common with recycling macropinocytosis, including F-actin-dependent surface ruffling, formation of large endocytic vesicles and a strong dependence on PKC. It is not sensitive to K44A-dynamin, as shown for macropinosomal uptake of *Chlamydia* bacteria [118]. PI3K is acting downstream of receptor tyrosine kinases and of tether/stimulating receptors, such as  $\alpha_v$  integrins, and it is required for the completion of ruffles and the formation of macropinosomes [111, 119]. In addition, Rac1 and Cdc42 are needed for promoting actin polymerization.

## Phagocytosis

In mammalian cells, the zippering phagocytosis is mechanistically similar to processive macropinocytosis (see Fig. 3). Both processes involve the sequential activation of

phospholipase C $\gamma$  and PKC, the activation of Ras and Src and the activation of PI3K to promote F-actin dynamics via Rac1 and Cdc42. Four steps can be distinguished during zippering phagocytosis [62, 120]. First, opsonized particles bind via the conserved Fc domains of Igs to the Fc $\gamma$ -R, including the high affinity Fc $\gamma$ -RI (CD64) and the low affinity Fc $\gamma$ -RIIA (CD32) and IIIA (CD16). The interaction of Ig and the Fc $\gamma$ -R triggers the phosphorylation of the ITAM (immunoreceptor tyrosine-based activation motif) which in turn provides docking sites for SYK, a tyrosine kinase, which activates PI3K. Interestingly, ligand induced monoubiquitination of the  $\mu$  subunit of Fc $\gamma$ -RI is thought to lead to clathrin-mediated endocytosis of soluble immune complexes, as shown for Fc $\gamma$ -RIIA, whereas the nonubiquitinated Fc $\gamma$ -RI appears to be involved in phagocytosis of opsonized particles [121]. The decision whether the Fc $\gamma$ -RI receptor is internalized by the one or the other pathway may be taken according to a thresholding concentration of activated kinases, such as SYK, recruited towards the Fc $\gamma$ -RI. Second, zippering drives the initial extension of membrane around the particle and this does not require actin polymerisation or signalling from the cytoplasmic domain of the receptors. Third, active signalling from the receptor leads to the recruitment of numerous cytoskeletal proteins, including the Arp2/3 complex, which nucleates actin filaments beneath the particle and thus pushes the plasma membrane further around the target in a process called pseudopod extension. Fourth, class I PI3K (p85 $\alpha$ /p110 $\beta$ ) activity leads to the generation of PI(3,4,5)P $_3$  and the closure of phagosomes. It is likely that PI(3,4,5)P $_3$  recruits proteins that control the actin cytoskeleton. Candidates are the PH-domain carrying Vav and ARNO, which bind PI(3,4,5)P $_3$  and act as GDP/GTP exchange factors for the small GTPases Rac1 and Arf6. By contrast, the class III PI3K (Vps34) product PI(3)P is required for phagosome maturation as it allows the recruitment of the Rab5 effector protein early endosome antigen 1 (EEA1) via its FYVE zinc finger [109].

One striking difference between macropinocytosis and zippering phagocytosis is the recruitment of dynamin to early phagosomes which requires PI3K and amphiphysin II $m$  [122]. Although dynamin is involved in the actin-based motility of macropinosomes [123], it does not participate in the initial internalization of macropinosomes [100, 118]. Phagocytosis differentiates also on the level of molecular scaffolds beneath the respective receptors which are responsible for actin remodelling. Integrins  $\alpha_v\beta_5$  recruit the p130<sup>cas</sup>-CrkII-Dock180 complex, which in turn triggers Rac1 activation and spacious phagocytosis of apoptotic cells in both professional and nonprofessional phagocytes [124]. Likewise,

the stimulation of Fc $\gamma$ -RIIA leads to the recruitment of Crkl to the Cbl-Nck-Grb2 complex [125]. In this scaffold the downstream effector of Rac and Cdc42 GTPases, Pak1, binds to Nck and leads to actin remodelling in part through Grp2 mediated WASP/N-WASP interaction followed by Arp2/3 activation [120]. In contrast, the complement receptor (CR) 3 is coupled to the focal adhesion scaffold, including talin,  $\alpha$ -actinin, vinculin and paxillin. Accordingly, CR-mediated phagocytosis is morphologically distinct from Fc-R mediated phagocytosis. Upon binding of iC3b-complement opsonized particles to CR3 ( $\alpha_m\beta_2$  integrin) the actin cytoskeleton is reorganized into stress fibers as controlled by the Rho GTPase, and this triggers the sinking of the particle into the phagocyte. This type of endocytosis is exploited, e.g., by *Mycobacterium tuberculosis* for macrophage entry [62]. The signalling events involved in sinking phagocytosis seem to be different from triggering or zippering phagocytosis, as they include, e.g., inside out signalling of R-Ras on integrins via the small GTPase Rap1 [126]. Notably, R-Ras has a high degree of sequence homology to Ras, yet it does not activate the ERK/MAP kinase pathway nor does it activate the JNK or p38/MAP kinase pathway.

### **Macropinocytosis in Adenovirus entry**

Earlier studies had noted that endocytosed macromolecules were apparently released from pinocytic vesicles into the cytosol in the presence of Ad2 or Ad5 [127, 128]. It had been widely assumed that these macromolecules, including enzymes, dextrans and DNA, were codelivered with the incoming viral particles from receptosomes, i.e., endosomes that were bearing both the viral particle and the macromolecules [129-135]. Additional studies found that the delivery of the macromolecules was sensitive to lysosomotropic agents, such as chloroquin and ammonium chloride [136, 137]. Remarkably, the efficiency of fluid phase stimulation was high, i.e., in the range of 2 to 15 fold over the noninfected cells, depending on the cell type tested. This implied that Ad controlled an endocytic pathway. The nature of this pathway was recently shown to be macropinocytosis [Fig. 4, and 100]. Viral activation of macropinocytosis required signalling through  $\alpha_v$  integrins, F-actin, the small Rho family GTPases and also PKC, and it was sensitive to amiloride, an inhibitor of the sodium / proton exchanger implicated in recycling and processive macropinocytosis [Fig. 3, and 138, 139]. In contrast to processive macropinocytosis, the Ad-induced macropinosomes were triggered to efficiently release their contents, i.e., more than 50% of

the cytoplasmic endosomes that were generated during the entry phase of Ad2 were destroyed [100]. These endosomes also included Ad-bearing endosomes, since Ad particles were released to the cytosol when macropinocytosis and pinosomal release occurred. Expression of the dominant-negative K44A dynamin further showed that the presence of Ad particles in macropinosomal membranes was not sufficient for rupturing the macropinosomes, but that discrete signals from the cell surface were involved in breaking the pinosomal membranes. These results imply that appropriate endocytic sorting of viral and cellular vesicles cooperates with cell signalling to mediate endosome rupture and viral escape.

### **Adenovirus escape to the cytosol**

Viral escape is one of the most crucial steps of entry but also one of the least understood ones. It is clear that the escape of Ad2 and Ad5 happens rapidly after endocytosis. It is also known that the delivery of the Ad particles into the cytosol is inhibited by lysosomotropic agents [95, 97] and that the Ad-mediated disruption of endosomes *in vitro* requires an acidic pH [140, 141], although a couple of studies reported rather weak effects of lysosomotropic amines on Ad infection [142, 143]. This implies then that low endosomal pH and the presence of Ad particles in endosomes are not sufficient for endosome disruption and additional factors must be involved. One of these factors has been identified as the  $\alpha_v\beta_5$  integrin [144, 145]. The attachment of the Ad penton base protein to integrin  $\alpha_v\beta_5$  at reduced extracellular pH promoted the permeabilization of the plasma membrane as measured by the release of small molecules into the extracellular medium. Another factor could be a spatially controlled activation of integrins, as suggested to be important for proper cell signalling through integrins [reviewd by 146]. While many signals emerge from the binding of wild type Ad2 or Ad5 to integrins [for a review, see 15], an Ad2 mutant called ts1 [for temperature sensitive 1, isolated by 147] elicits an incomplete spectrum of signals upon entry. Ts1 lacks the viral protease and contains a capsid which is not proteolytically processed and ts1 does not dissociate the fibers upon entry [reviewed in 148]. It binds to CAR, is endocytosed into the cells with indistinguishable efficiency and kinetics as the wild type virus, but it is unable to penetrate the endosomal membrane and ends up in late endosomes and lysosomes where it is degraded [149]. Interestingly, ts1 activates PI3K [data not shown in 107] but not the MAPK pathways ERK and p38 [150]

and also fails to trigger macropinocytosis [100]. These data support the notion that Ad precisely activates integrins preparing the host for its arrival. Precise activations seem to be also crucial for successfully retargeting of Ad particles to new receptors and cell types normally not affected by Ad. For example, Ad targeted to the transferrin receptor by an insertion into the HI loop of the fiber knob was able to transduce brain microcapillary cells to low levels but the virus was difficult to propagate and overall ineffective [151]. A more successful retargeting example is the delivery of Ad5 to the high affinity Fc $\gamma$ -RI (CD64) of hematopoietic cells by virtue of a soluble adaptor consisting of the extracellular domains D1 and D2 of CAR and the constant region of a human Ig [152]. These hematopoietic cells are CAR-negative and express only very low levels of  $\alpha_v$  integrins. Virus internalization and cytosolic delivery required a phagocytic mechanism and the transduction was effectively inhibited by anti-Fc $\gamma$ -RI antibodies, particularly at low adaptor to particle ratios, i.e., when large virus aggregates had been formed. It remains to be tested if viral transduction via CD64 requires also a clathrin-dependent pathway or if it solely relies on Fc $\gamma$ -RI mediated phagosome formation.

## Conclusions

There are many ways to get into a cell, and pathogens essentially exploit every one of the entry pathways with astounding efficiency. However, under normal conditions, a particular pathogen usually takes only one defined entry pathway. This allows the exploitation of discrete signalling pathways at the cell surface and distinct intracellular sites. In the case of epithelial cells, it is clear that clathrin-mediated uptake of Ad2 or Ad5 together with signal stimulated macropinocytosis leads to infection. This does not preclude the uptake of viral particles by alternative routes, but these routes alone are considered to be noninfectious, unless the infectious route is used at the same time and the appropriate signalling switches are turned on. The lessons that pathogens teach us on their natural entry are crucial to further develop viral vectors into useful therapeutics.

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## Figure Legends

### **Fig. 1: A hypothetical scheme of events leading to epithelial infection with Ad2 or Ad5.**

Ad is thought to access the epithelial cells by virtue of cells displaying CAR receptors in the apical membrane. This first leads to a local infection and the spreading of the particles to basolateral membranes of epithelial cells. Viral spreading is aided by the release of fiber-penton base complexes which are thought to disrupt the tight junctions of epithelial cells prior to virus release by virtue of binding the CAR protein localized to cell-cell contacts. For further explanations, see main text.

### **Fig. 2: The infectious entry pathway of Ad2 and Ad5 into epithelial cells.**

Ad fibers bind to CAR and locally activate  $\alpha_v$  integrins which triggers clathrin-mediated viral endocytosis. This requires the large GTPase dynamin, PI3K, the small GTPases Rac1 and Cdc42 and also Rab5. Ad is then delivered to a slightly acidic intracellular compartment and escapes to the cytosol upon cell signalling. Additional signalling, including the activation of protein kinase A and the p38/MAPK cascade boost microtubule-dependent and dynein/dynactin-dependent viral transport towards the nucleus [150, 153-155]. Virus then docks to the nuclear pore complex receptor CAN/Nup214 and it disassembles by recruiting the nuclear histone H1 and the H1 import factors importin  $\beta$  and importin 7 [156].

### **Fig. 3: Mechanisms of macropinocytosis and phagocytosis.**

While macropinocytosis is a spacious process, i.e., engulfing substantial amounts of extracellular fluid, phagocytosis tightly wraps plasma membrane regions around an extracellular particle, e.g., an opsonized bacterium. Recycling and processive macropinocytosis differ with respect to the trafficking of macropinosomes, while zippering and sinking phagocytosis differ with respect to the Rho GTPases involved. Common

signalling molecules to all four types of endocytosis include Ras, PI3K and PLC $\gamma$ . Abbreviations see above.

**Fig. 4: Ad2 or Ad5 triggered macropinocytosis.**

Simultaneous with viral uptake by clathrin-coated pits and vesicles, the incoming Ad triggers macropinocytosis, independent of dynamin (Dyn), but depending on protein kinase C (PKC) and the sodium / proton exchange inhibitor 5-(N-ethyl-N-isopropyl) amiloride (EIPA). It appears that macropinosomes and virus-bearing endosomes simultaneously release their contents into the cytosol.