

VIEWPOINT

Could vesicular transport of Na⁺ and Cl⁻ be a feature of salt tolerance in halophytes?

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- **Background** Halophytes tolerate external salt concentrations of 200 mM and more, accumulating salt concentrations of 500 mM and more in their shoots; some, recretahalophytes, excrete salt through glands on their leaves. Ions are accumulated in central vacuoles, but the pathway taken by these ions from the outside of the roots to the vacuoles inside the cells is poorly understood. Do the ions cross membranes through ion channels and transporters or move in vesicles, or both? Vesicular transport from the plasma membrane to the vacuole would explain how halophytes avoid the toxicity of high salt concentrations on metabolism. There is also a role for vesicles in the export of ions via salt glands.
- **Scope and Methods** We have collected data on the fluxes of sodium and chloride ions in halophytes, based on the weight of the transporting organs and on the membrane area across which the flux occurs; the latter range from 17 nmol m⁻² s⁻¹ to 4.2 μmol m⁻² s⁻¹ and values up to 1 μmol m⁻² s⁻¹ need to be consistent with whatever transport system is in operation. We have summarized the sizes and rates of turnover of vesicles in plants, where clathrin-independent vesicles are 100 nm or more in diameter and can merge with the plasma membrane at rates of 100 s⁻¹. We gathered evidence for vesicular transport of ions in halophytes and evaluated whether vesicular transport could account for the observable fluxes.
- **Conclusions** There is strong evidence in favour of vesicular transport in plants and circumstantial evidence in favour of the movement of ions in vesicles. Estimated rates of vesicle turnover could account for ion transport at the lower reported fluxes (around 20 nmol m⁻² s⁻¹), but the higher fluxes may require vesicles of the order of 1 μm or more in diameter. The very high fluxes reported in some salt glands might be an artefact of the way they were measured.

Key words: Salt tolerance, endocytosis, exocytosis, pinocytosis, Na⁺ flux, ion transport, vesicle

INTRODUCTION

All plants can tolerate some NaCl in the medium in which they grow, but the concentration varies considerably, from under 50 mM to over 1 M (see [Flowers *et al.*, 2010](#); [Moir-Barnetson *et al.*, 2016](#)). While there is a continuous spectrum of salt tolerance, it is useful to separate the less tolerant glycophytes from the more tolerant halophytes and there have been various suggestions (see [Huchzermeyer and Flowers, 2013](#)) of where to draw a rather arbitrary dividing line between the two groups – from about 90 mM ([Aronson, 1989](#)) to 200 mM ([Flowers and Colmer, 2008](#)). Here we adopt the higher concentration, the arguments for which were put forward in [Flowers and Colmer \(2008\)](#). Halophytes are remarkable in that not only do they grow in external salt concentrations lethal to most plants, but they also accumulate high concentrations of salts in their leaves (in excess of 500 mM in some species; see [Flowers,](#)

[1985](#)): some species (recretahalophytes) secrete salt from their leaves. However, despite much research, the molecular mechanisms by which halophytes transport and sequester salts within tissues and secrete salts from leaves are still largely unknown.

This Viewpoint explores the idea that vesicles might be involved in salt transport. Vesicular transport of ‘particles’ was suggested in the 1950s (see below) and remains an attractive but controversial topic for discussion and research into transport processes in plants. Historically, the concept of vesicular ion transport was based on microscopic observations and preceded ideas about the existence of ion channels and transporters ([Behrends, 2012](#)). However, the rapid rise of research into membrane transport proteins and the intensive accumulation of knowledge of ion channels pushed ideas on vesicular ion transport into the shadows, virtually suppressing and abolishing them. Nevertheless, MacRobbie, in a series of

papers published between 1966 and 1977 (see MacRobbie, 1999), drew attention to the fact that the kinetics of Cl^- movement between the cytoplasm and vacuole in algae with giant cells suggested transport in ‘salt-filled vesicles’. That ions may be moved in vesicles gains support from the fact that vesicular transport is widespread throughout eukaryotic organisms, with the mechanisms of endo- and exocytosis being conserved (Cucu *et al.*, 2017). The idea we examine is that, in addition to ion channels and transporters, halophytes use vesicles to move salts through the cytoplasm (endocytosis and exocytosis) and to expel excess salts through salt glands (exocytosis). A consequence of vesicular transport is that Na^+ and Cl^- , which are toxic at the concentrations found in the cells of halophytes, would not come into direct contact with the proteins that they have been shown to disrupt *in vitro*. Using the recent literature on ion transport and vesicular trafficking in plants, we test the feasibility of vesicular transport as an integral part of salt tolerance in halophytes. Baral *et al.* (2015) have previously argued that the stress of salinity modulates endo- and exocytosis in glycophytes.

Na⁺ and Cl⁻ toxicity in the cytosol

Plants growing in saline soils take up large quantities of salt for osmotic adjustment. Perhaps the leading theory to account for this uptake of Na^+ and Cl^- is that it occurs passively according to thermodynamic gradients or using secondary active transport, since no pumps for Na^+ (Pedersen and Palmgren, 2017) or Cl^- are known in higher plants. In halophytes, Na^+ and Cl^- are sequestered in central vacuoles of cells (Flowers *et al.*, 1977, 1986) and it has been presumed that cytosolic concentrations of Na^+ are kept within a reasonable range through the activity of Na^+/H^+ antiporters, other transporters and ion channels in the vacuolar and plasma membranes in both root and leaf cells (Fig. 1; see also Volkov, 2015a, cf. fig. 2 in Maathuis, 2014). Consequently, it has been argued that enzymes in the cytosol (which is not a simple aqueous solution of proteins; Cheeseman, 2013; Yu *et al.*, 2016) are not exposed to concentrations of salt higher than 100–200 mM (Flowers *et al.*, 2015). However, vesicular transport of Na^+ and Cl^- would allow large fluxes of Na^+ and Cl^- through the cytosol while avoiding exposing enzymes to toxic concentrations of transiting ions.

A further argument in favour of vesicular transport of ions is that the mechanisms are completely different from those involved in transport by ion channels and transporters. Vesicular ion transport depends more on intracellular machinery, cell metabolism and cytoskeletal activity than on trans-membrane electric potential and ion concentrations, and hence could be a safety valve for directing intracellular ion fluxes. Another important point is that while the selectivity of ion transport via ion channels and transporters is determined by the molecular protein structure of the ion channels and transporters (e.g. Volkov *et al.*, 2004), for vesicular ion transport the selectivity of trapped ion cargo would depend on local ion concentrations near forming vesicles and on electric charges of vesicular membranes. Here is another mechanism to generate selective (or semi-selective) ion fluxes.

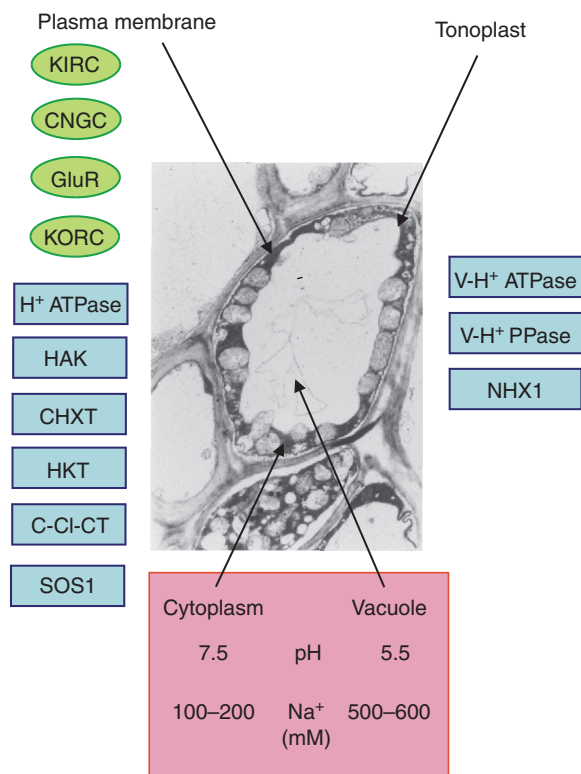


Fig. 1. The transporters that might be found in the membranes of a halophyte cell (after Volkov, 2015a). The estimated concentration of Na^+ is that of a mesophyll cell of *Suaeda maritima* and the pH values are taken from the literature: abbreviations enclosed by ovals are ion channels, and those within rectangles are transporters. Plasma membrane: KIRC, inward rectifying potassium channels; CNGC, cyclic-nucleotide gated ion channels; GluR, glutamate receptors; KORC, outward rectifying potassium channels; H^+ ATPase, plasma membrane proton pump; HAK, high affinity potassium transporter; CHXT, cation H^+ exchange transporters; HKT, high affinity K^+ transporters; C-Cl-CT, cation-chloride co-transporters; SOS1, sodium-proton antiporter. Tonoplast: V- H^+ ATPase, tonoplast proton pump; V- H^+ PPase, tonoplast pyrophosphatase; and NHX1, tonoplast sodium (cation)/proton antiporter.

Transport of Na⁺ and Cl⁻ in plants

Much more is known about ion transport in glycophytes than in halophytes. For example, in *Arabidopsis thaliana* about 5 % of around 25 000 genes are likely to encode transport proteins in membranes, with >150 predicted from their nucleotide sequences to be cation channels or transporters (see Mäser *et al.*, 2001; <http://aramemnon.uni-koeln.de/>). Of these, non-selective cation channels (NSCCs), including cyclic nucleotide gated ion channels and probably glutamate receptors, have been assumed to be the major pathway for Na^+ entry into root cells of *A. thaliana* (Davenport, 2002; Demidchik and Tester, 2002; Demidchik *et al.*, 2002; Essah *et al.*, 2003) and in some other plants (reviewed in Maathuis *et al.*, 2014). However, in a critical review of the transport of Na^+ into plants, Kronzucker and Britto (2011) concluded that the involvement of NSCCs was promulgated prematurely and that evidence for their involvement was highly speculative. There is, however, strong evidence in support of the role of $\text{Na}^+/\text{K}^+/\text{H}^+$ exchangers (NHXs; see Pilar Rodriguez-Rosales *et al.*, 2009 for information on NHX transporters in plants) in the transport of Na^+ , both across the plasma membrane (SOS1; see Ji *et al.*, 2013)

and into vacuoles (Volkov, 2015a). Interestingly, the aquaporin AtPIP2 also has non-selective cation channel activity and could provide a link between water and Na⁺ transport (Byrt et al., 2017). Chloride channels are also known for plants (see Ward et al., 2009) and inhibitor studies with bumetanide suggest that cation–chloride cotransporters may be involved in the transport of Na⁺ and Cl[−] (Zhang et al., 2010). For halophytes, 18 or more transporters may potentially be involved in the transport of Na⁺, K⁺ and Cl[−] into cells (Flowers and Colmer, 2008; Fig. 1), but, in spite of >80 years of research (see Huchzermeyer and Flowers, 2013), we still do not know the precise identity of the proteins involved. Low-affinity cation transporter 1 (LCT1), AKT-type ion channels, K⁺ transporters from the KUP/HAK/KT family as well as members of the HKT1 and HKT2 classes of transporters possibly participate in Na⁺ transport across the plasma membrane (see, for example, Wang et al., 2007; Almeida et al., 2013), but the exact means by which Na⁺ and Cl[−] enter the cells of halophytes remains uncertain. Before reviewing evidence for vesicular transport of ions, first suggested in the 1950s and developed for ion transport in the 1970s, we evaluate the magnitude of the fluxes involved.

ION FLUXES BASED ON ROOT WEIGHT

In attempting to ascertain the feasibility of vesicular transport, as opposed to transport through channels or transporters, we next assess the magnitude of fluxes of Na⁺ and Cl[−] so that we can calculate required rates of membrane turnover to deliver these measured fluxes. The effects of external concentration of K⁺ and its flux into roots have been analysed over many years and often viewed as separable into a high-affinity system saturating at external concentrations of around 1 mM and a low-affinity system where the flux increases with external concentration > 1 mM (recently reviewed in Nieves-Cordones et al., 2016). The magnitude of influxes has generally been determined from measurements of the uptake of isotopes, which it has been suggested may be difficult to separate from concurrent efflux (Coskun et al., 2016), so the values may need to be viewed with caution.

High- and low-affinity systems are presumed to occur for Na⁺ as well as K⁺ fluxes through roots (see Nieves-Cordones et al., 2016). Halophytes are normally exposed to high concentrations of NaCl in the medium in which they grow, certainly in the low-affinity range as, for example, seawater contains about 480 mM Na⁺ and 560 mM Cl[−]. Salt concentrations around the roots could be higher than in any bulk solution, as salts tend to accumulate in the rhizosphere following water uptake by the roots, when salts are left behind in the soil. For example, plants of the halophyte *Tamarix ramosissima* had soil-solution salinities of 280–620 mM NaCl in the capillary fringe above the water table from which they extracted water, when aquifer salinities were 54–172 mM NaCl (Glenn et al., 2013); *Salicornia bigelovii* irrigated daily with 670 mM seawater at 2.25 times the potential evaporation rate had soil-solution salinities in the range 1228–1315 mM (mostly NaCl) while still maintaining high productivity of biomass and seeds (Glenn et al., 1997). To maintain such high growth rates requires high, but controlled, fluxes of ions from roots to shoots (see, for example, Yeo and Flowers, 1986).

For halophytes, we have collected the values of Na⁺ fluxes (Table 1; see also Flowers and Colmer, 2008; Kronzucker and Britto, 2011), where Na⁺ concentrations were in the low-affinity range (>25 mM and often >200 mM). Influx values were determined from short-time (a few minutes) uptake of labelled isotope (²²Na⁺ or ²⁴Na⁺) and steady-state values generally from changes in chemical content over days. Influx values range from 4 to 110 nmol g^{−1} f. wt root s^{−1} and steady-state (net) fluxes from about 3 to 11 nmol g^{−1} f. wt root s^{−1}; net fluxes are generally lower than influxes since part of the initial uptake could be expelled by, for example, Na⁺/H⁺ antiporters (see, for example, Wang et al., 2006). For the halophyte *Suaeda maritima*, Zhang et al. (2013) recorded a short-term influx of 28 nmol g^{−1} f. wt root s^{−1} in 200 mM NaCl (this is equivalent to 295 nmol g^{−1} d. wt root s^{−1}, calculated from the fresh weight figure assuming a f. wt:d. wt ratio of 10.4; Yeo and Flowers, 1980) and a very similar net flux of 25 nmol g^{−1} f. wt root s^{−1} over 12 h (equivalent to 260 nmol g^{−1} d. wt root s^{−1}). While the high influx of 28 nmol g^{−1} f. wt root s^{−1} (Zhang et al., 2013) and the value of 110 nmol g^{−1} f. wt root s^{−1} recorded for *Spergularia marina* (Lazof and Cheeseman, 1986, 1988) may be subject to the objections raised by Coskun et al. (2016), a sustained (over 7 d) net flux of around 11 nmol g^{−1} f. wt root s^{−1} (110 nmol g^{−1} d. wt root s^{−1}; Yeo and Flowers, 1986) has been measured through the roots of *S. maritima* plants growing in 340 mM NaCl (calculated from the f. wt:d. wt ratio in roots of *S. maritima* of 10.4; Yeo and Flowers, 1980); this is a net flux of ions from roots to shoots determined from changes in chemical content (Table 1) and so not subject to objections raised by Coskun et al. (2016). An additional feature of this net flux (measured over 5 d) through the roots of *S. maritima* to the shoots is that it does not saturate as the external concentration of Na⁺ rises from 20 to 200 and 400 mM; Al-Zahrani, 1990; Table 1). We will return below to the very high influx value calculated for *S. marina* (Lazof and Cheeseman, 1986).

Conclusion

Influx of Na⁺ through the roots of halophytes is about 25 nmol g^{−1} f. wt root s^{−1}; net fluxes measured over days can reach values close to 10 nmol g^{−1} f. wt root s^{−1} or >100 nmol g^{−1} d. wt root s^{−1}, setting values that have to be accounted for by the transport systems that operate through membranes in these plants.

FLUXES OF IONS BASED ON MEMBRANE AREA

While fluxes calculated on the basis of fresh or dry weight are often the best data available for higher plants, they do not provide values for the fluxes of ions across membranes *per se* – although they do provide a context by which to judge such membrane fluxes as are available. A flux based on membrane area is a crucial parameter for determining whether vesicular transport of salts could be feasible and, as so few are available (note fluxes based on the use of vibrating microelectrodes external to a root provide estimates based on the area of the plane in which the electrode vibrates, not on a membrane area; see, for example, Bose et al., 2015), we have collected data from both freshwater and saltwater species.

TABLE 1. A summary of the magnitude of Na⁺ fluxes in plants, reported on the basis of fresh (nmol g⁻¹ f. wt s⁻¹) or dry (nmol g⁻¹ d. wt s⁻¹) weight

Summary or species	NaCl (mM)	Period of measurement and direction	Flux	Reference
Reported on the basis of f. wt				
<i>Spergularia marina</i>	25	Influx 0 min	13	Lazof and Cheeseman (1988)
<i>Spergularia marina</i>	90	Influx 0 min	110	Lazof and Cheeseman (1986)
<i>Suaeda maritima</i>	25	Influx 2 min	3.8	Wang et al. (2007)
<i>Suaeda maritima</i>	200	Influx 2 min	30	Wang et al. (2007)
<i>Suaeda maritima</i>	200	Influx 2 min	25	Zhang et al. (2013)
<i>Suaeda maritima</i>	200	Influx 2 min	60	Zhang et al. (2013)
<i>Hordeum vulgare</i>	80	Influx 5 min	15–39*	Chen et al. (2007)
<i>Arabidopsis thaliana</i>	50	Influx 2 min	31	Essah et al. (2003)
<i>Arabidopsis thaliana</i>	100	Influx 3 min	11	Wang et al. (2006)
<i>Arabidopsis thaliana</i>	100	Efflux 10 min	7.3–8.5	Wang et al. (2006)
<i>Eutrema halophilum</i>	100	Influx 3 min	5.2	Wang et al. (2006)
<i>Eutrema halophilum</i>	100	Efflux 10 min	1.4–4.0	Wang et al. (2006)
<i>Spergularia marina</i>	25	Steady state 10–15 min	2.9	Lazof and Cheeseman (1988)
<i>Spergularia marina</i>	90	Linear influx 1–20 min	1.2	Lazof and Cheeseman (1986)
<i>Suaeda maritima</i>	25	Net	3.3	Wang et al. (2007)
<i>Suaeda maritima</i>	150	Net	9.3	Wang et al. (2007)
<i>Arabidopsis thaliana</i>	50	Net 21 d	0.5	Wang et al. (2007)
Median influx			Approx. 25	
Reported on the basis of d.wt				
<i>Arabidopsis thaliana</i>	100	6 h	28.2	Wang et al. (2006)
<i>Eutrema halophilum</i>	100	6 h	4.3–9.5	Wang et al. (2006)
<i>Suaeda maritima</i>	100	24 h	31	Yeo and Flowers (1986)
<i>Suaeda maritima</i>	300	24 h	62	Yeo and Flowers (1986)
<i>Arabidopsis thaliana</i>	100	25 h	56	Volkov et al. (2004)
<i>Arabidopsis thaliana</i>	100	25 h	16.2–20.2	Wang et al. (2006)
<i>Eutrema halophilum</i>	100	25 h	16.0	Volkov et al. (2004)
<i>Eutrema halophilum</i>	100	25 h	5.3	Wang et al. (2006)
<i>Suaeda maritima</i>	340	Net 7 d	110	Yeo and Flowers (1980)
<i>Suaeda maritima</i>	20	Net 5 d	36	Al-Zahrani (1990)
<i>Suaeda maritima</i>	200	Net 5 d	67	Al-Zahrani (1990)
<i>Suaeda maritima</i>	400	Net 5 d	80	Al-Zahrani (1990)
<i>Tamarix ramosissima</i>	242	Net 30 d	4.6	Glenn et al. (2012)
<i>Arabidopsis thaliana</i>	50	42 d	2.0	Wang et al. (2006)
<i>Elymus × oliveri</i>	75	33 d	5.5	Ansari (1982)
<i>Elymus × oliveri</i>	150	33 d	5.6	Ansari (1982)
<i>Eutrema halophilum</i>	100	42 d	0.63	Wang et al. (2006)
<i>Puccinellia maritima</i>	100	63 d	3.3	Ansari (1982)
<i>Puccinellia maritima</i>	200	63 d	3.5	Ansari (1982)

Influx data represent uptake into the roots (estimated from flux analysis or isotope uptake over minutes); net fluxes were gathered over longer periods (hours, days or weeks) and are fluxes into the whole plant.

*Dependent on the external calcium concentration.

Fluxes in algae

Fluxes have been estimated for freshwater algae, where cell dimensions can be calculated; for those growing in 'pond' water (0.1 mM K, 1.0 mM Na, 0.1 mM Ca and 1.3 mM Cl). MacRobbie (1971a) concluded that fluxes ranged from 10 to 20 nmol m⁻² s⁻¹ (fluxes: K⁺, 10 nmol m⁻² s⁻¹; Na⁺, 5 nmol m⁻² s⁻¹; Cl⁻, 10–20 nmol m⁻² s⁻¹), similar to the values of tracer fluxes of 2–35 nmol m⁻² s⁻¹ for freshwater algae summarized by Raven (1976; external K⁺ from 0.1 to 0.2 mM and fluxes from 2 to 20 nmol m⁻² s⁻¹; external Na⁺ from 1 to 2 mM, fluxes 15–30 nmol m⁻² s⁻¹; external Cl⁻ from 1 to 2 mM, flux 30–35 nmol m⁻² s⁻¹). The tracer fluxes

for marine algae (Raven, 1976), where the external concentrations of monovalent ions were in the low-affinity range (K⁺, 10 mM; Na⁺ and Cl⁻, 550 mM) were significantly higher: K⁺, flux up to 2.1 μmol m⁻² s⁻¹; Na⁺ flux up to 1.1 μmol m⁻² s⁻¹; and Cl⁻ flux up to 10 μmol m⁻² s⁻¹ (Table 2).

Membrane fluxes in salt glands

In order to effect osmotic adjustment, ions are transported across roots and then via the xylem to the shoots where they are

TABLE 2. A summary of the magnitude of Na⁺ fluxes in algae and plants, based on area of the membrane

	NaCl (mM)	Na flux (nmol m ⁻² s ⁻¹)	
Freshwater algae	0.1	5	MacRobbie (1971a)
Freshwater algae	1–2	1–20*	Raven (1976)
Marine algae	550	170	Raven (1976)
<i>Suaeda maritima</i> roots	340	262	Yeo and Flowers (1986)
<i>Suaeda maritima</i> leaves	66 [†]	17	Flowers and Yeo (1986)
<i>Arabidopsis thaliana</i> root protoplasts [‡]	100	700	Volkov and Amtmann (2006)
<i>Eutrema halophilum</i> root protoplasts [‡]	100	150	Volkov and Amtmann (2006)
<i>Cynodon dactylon</i> gland	75 [§]	408	Oross <i>et al.</i> (1985)
<i>Limonium perezii</i> gland	300	4200 [‡]	Faraday and Thomson (1986a)

*Flux across the plasma membrane in the light.

[†]Estimated xylem concentration (Clipson and Flowers, 1987).

[‡]Efflux was, in the words of the authors (Faraday and Thomson 1986a), calculated as though ‘evenly distributed across the non-transfusion zone plasma membrane of the secreting gland’.

[§]Fed to the cut shoots.

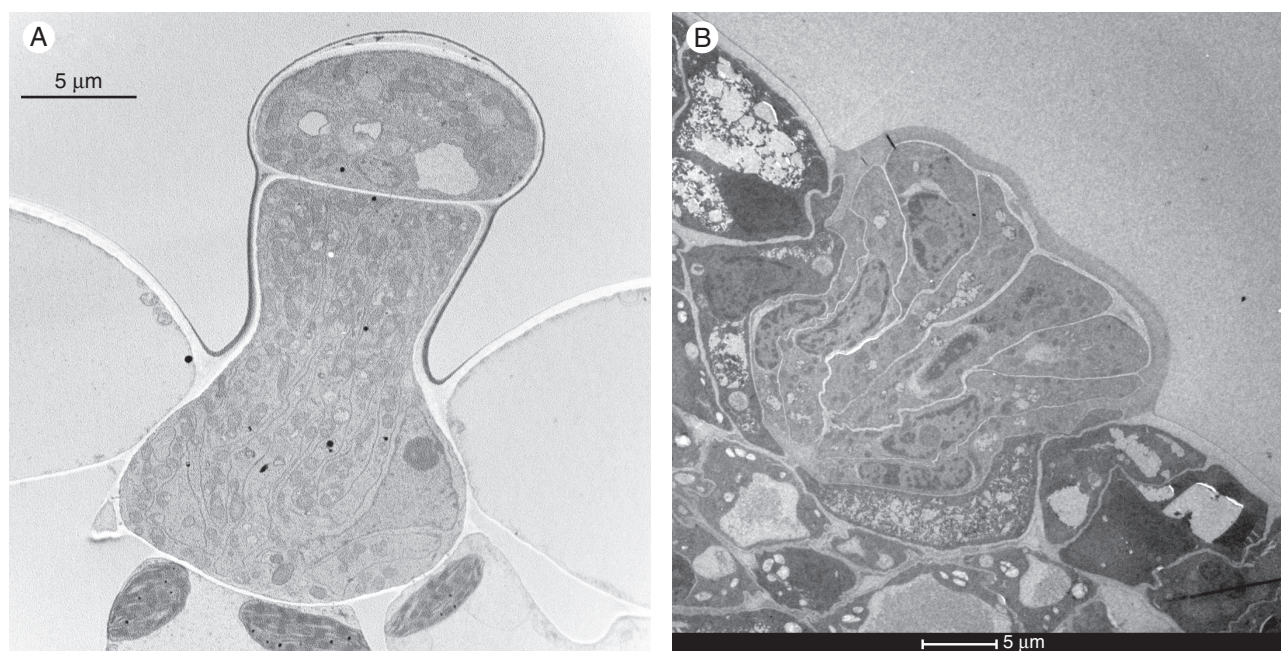


FIG. 2. Examples of transverse sections of salt glands seen by transmission electron microscopy. (A) A bicellular gland from the abaxial leaf surface of Rhodes grass (*Chloris gayana*) and (B) a multicellular gland on a leaf of sea lavender (*Limonium bicolor*). (A) Courtesy of Takao Oi of Nagoya University; more detail can be found in Oi *et al.* (2012). (B) Courtesy of Bao-Shan Wang, Shandong Normal University, China; detail can be found in Feng *et al.* (2014).

accumulated. However, since continued import into the transpiration stream follows from the need for open stomata necessary for continued carbon fixation, halophytes require some mechanism to avoid excess levels of Na⁺ and Cl⁻ building up in their shoots. Halophytes generally reduce transpiration over time (Flowers *et al.*, 1986; Koyro *et al.*, 2013; Rakhmankulova *et al.*, 2014), thus reducing excessive ion import and balancing import with growth. Salts can be removed by leaf shedding (Albert, 1975; Cram *et al.*, 2002; Ibrahim, 2013), accumulation in external bladders (Thomson and Healey, 1984; Shabala *et al.*, 2014) or excretion via glandular structures (Thomson and Healey, 1984; Flowers and Colmer, 2008; Shabala and Mackay, 2011; Roy and Chakraborty, 2014; Yuan *et al.*, 2016b), an attribute that has evolved >10 times (Dassanayake and Larkin, 2017). Not all halophytes excrete salts, but, in those that do, the

mass of excreted salt can exceed those retained in the plant for osmotic adjustment. For example, Rozema and Gude (1981) found that excreted salts accounted for 60 % of total uptake by *Spartina anglica*, while Glenn *et al.* (2013) found that 80 % of salt uptake was excreted by *Tamarix ramosissima* growing in salinized soil.

Glands and bladders have been reported in 117 halophytes (eHALOPH; <https://www.sussex.ac.uk/affiliates/halophytes/index.php>) and are mainly found within the Amaranthaceae (29 species, largely salt bladders in the genus *Atriplex*), the Plumbaginaceae (29 species) and the Poaceae (29 species): their structures and function have been the subject of a number of reviews (Lüttge, 1975; Thomson, 1975; Hill and Hill, 1976; Fahn, 1988; Thomson *et al.*, 1988; Salama *et al.*, 1999; Shabala and Mackay, 2011; Tyerman, 2013; Shabala *et al.*,

2014; Yuan *et al.*, 2016b; Dassanayake and Larkin, 2017). Salt glands are two- to multicelled epidermal structures (Fig. 2; a unique exception seems to be the single-celled glands of *Porteresia coarctata*; Flowers *et al.*, 1990) that are responsible for the movement of salt to an external aerial surface: salt bladders are epidermal cells that accumulate salt, and may burst, releasing the salt solution contained in a large vacuole. Here we are concerned not with the mechanism of secretion *per se* (see Yuan *et al.*, 2016b; Dassanayake and Larkin, 2017 for recent discussions of salt glands and the mechanism of salt secretion), but with any evidence that vesicles and exocytosis might be involved in a process where Na⁺ and Cl⁻ fluxes are high (Table 1).

The simplest of glands are the bicellular trichomes (Fig. 2A) found on the leaves of most grasses (with the exception of the Pooideae; Amarasinghe and Watson, 1988). Some of these trichomes (in the subfamily Chloridoideae) are able to secrete salt (Amarasinghe and Watson, 1989; Ceccoli *et al.*, 2015) and, in one case, fluxes based on membrane area have been estimated. For the glands of a hybrid, *Cynodon dactylon* × *C. transvalensis*, Oross *et al.* (1985) were able to calculate a flux of Na⁺ of 408 nmol m⁻² s⁻¹ (see Table 2).

More complex salt-secreting glands are found in five halophytic species of the family Plumbaginaceae (*Aegialitis*, *Armeria*, *Limoniastrum*, *Limonium* and *Plumbago*; Fig. 2B). The glands consist of 16 secretory cells surrounded by a cuticle that isolates most of the apoplast of the gland from the rest of the leaf (there is some apoplastic continuity between what are

termed sub-basal cells lying next to the mesophyll, the transfusion zone and the innermost gland cells; Thomson *et al.*, 1988). There is also symplastic continuity via plasmodesmata between the mesophyll cells and the gland cells. For the glands of *Limonium perezii*, detailed analysis of membrane areas has been carried out, allowing fluxes of Na⁺ across the plasma membrane that lines wall protuberances to be calculated (Faraday and Thomson, 1986a; see also Table 3). Values were dependent upon the area over which efflux was deemed to occur, but the maximum fluxes of Na⁺ and Cl⁻ were 70 and 85 μmol m⁻² s⁻¹, respectively, if the flux occurred across the plasma membrane of the transfusion zone. Even the lowest fluxes estimated – for fluxes across the plasma membrane lining wall protuberances – were 13 μmol m⁻² s⁻¹ for Na⁺ and 16 μmol m⁻² s⁻¹ for Cl⁻; the flux for Na⁺ is about 32 times the value reported for the glands of *C. dactylon*.

Membrane fluxes through roots of halophytes

Membrane fluxes have been estimated for the roots of one halophyte, *S. maritima*, an annual species within the Amaranthaceae (a family with 24 % of known halophytes; <https://www.sussex.ac.uk/affiliates/halophytes/index.php?content=plantStats>). Plants of *S. maritima* grow optimally in 200–300 mM NaCl, when the sustained net flux of Na⁺ through the roots is about 100 nmol g⁻¹ d. wt s⁻¹ over a period of 7 d (Yeo and Flowers, 1986; Table 1). Using estimates of the membrane area per unit

TABLE 3. The presence of numerous small vesicles has been described in some but not all ultrastructural studies of glands

Genus	Species	Micro-vesicles	Wall protuberances	Reference
(a)				
<i>Armeria</i>	<i>maritima</i>	Yes		Heumann (2002) – for the uptake of Zn
<i>Atriplex</i>	<i>halimus</i>	Yes	Yes	Smaoui <i>et al.</i> (2011)
<i>Atriplex</i>	<i>spongiosa</i>	Yes		Osmond <i>et al.</i> (1969)
<i>Avicennia</i>	<i>marina</i>	Yes		Drennan <i>et al.</i> (1987)
<i>Limonium</i>	<i>bicolor</i>	Yes		Feng <i>et al.</i> (2015); Yuan <i>et al.</i> (2015)
<i>Lysimachia</i>	<i>maritima</i>	Yes	Yes	Rozema <i>et al.</i> (1977)
<i>Odyssea</i>	<i>paucinervis</i>	Yes	Yes	Somaru <i>et al.</i> (2002)
<i>Porteresia</i>	<i>coarctata</i>	No*		Flowers <i>et al.</i> (1990)
<i>Spartina</i>	<i>foliosa</i>	Yes	Yes	Levering and Thomson (1971)
<i>Tamarix</i>	<i>aphylla</i>	Yes		Thomson <i>et al.</i> (1969)
(b)				
<i>Aegiceras</i>	<i>corniculatum</i>			Cardale and Field (1971)
<i>Avicennia</i>	<i>germinans</i>			Balsamo and Thomson (1993)
<i>Atriplex</i>	<i>semibaccata</i>			Campbell <i>et al.</i> (1974)
<i>Cynodon</i>	<i>dactylon</i>			Oross and Thomson (1982)
<i>Distichlis</i>	<i>spicata</i>			Oross and Thomson (1982)
<i>Frankenia</i>	<i>pauciflora</i>			Olesen, 1979
<i>Frankenia</i>	<i>salina</i>			Balsamo and Thomson (1993)
<i>Frankenia</i>	<i>grandifolia</i>			Campbell and Thomson (1976a, b)
<i>Limonium</i>	<i>platyphyllum</i>			Vassilyev and Stepanova (1990); Campbell <i>et al.</i> (1974)
<i>Limonium</i>	<i>perezii</i>			Faraday and Thomson (1986a); Faraday <i>et al.</i> (1986)
<i>Limonium</i>	<i>sinuatum</i>			Faraday and Thomson (1986b)
<i>Limonium</i>	<i>vulgare</i>			Ziegler and Lüttge (1967)
<i>Plumbago</i>	<i>auriculata</i>			Faraday and Thomson (1986b)
<i>Tamarix</i>	<i>aphylla</i>			Thomson and Liu (1967); Campbell <i>et al.</i> (1974)
<i>Tamarix</i>	<i>usneoides</i>			Wilson <i>et al.</i> (2017a)

The data in (a) include only cases where plants growing in saline media were investigated and exclude those where cut stems were exposed to higher salt concentrations than might be expected in the xylem (see text). These excluded papers are listed in (b).

*The glands are unlike the bicellular glands of other grasses (Type 4 in Dassanayake and Larkin, 2017).

dry weight of roots and their component cells from the data of Hajibagheri *et al.* (1985), Yeo and Flowers (1986) calculated fluxes of Na⁺ to be 763 nmol m⁻² s⁻¹ across the epidermal plasmalemma, 262 nmol m⁻² s⁻¹ across the combined area of epidermal and cortical plasmalemma, and 6.25 μmol m⁻² s⁻¹ across the endodermal surface area. They (Yeo and Flowers, 1986) argued a flux across epidermal and cortical cells to be the most likely, based on its size: loading of the xylem, however, required a flux of 1.18 μmol m⁻² s⁻¹.

Ions passing through the roots of *S. maritima* are transferred to the xylem, where the concentration of Na⁺ ions in these plants growing in 200 mM NaCl (around the optimum salinity for growth) is about 66 mM (Clipson and Flowers, 1987); this results in the delivery of 74 mol Na⁺ m⁻³ shoots d⁻¹ (Flowers and Yeo, 1986). Using the stereological data of Hajibagheri *et al.* (1984), Flowers and Yeo (1986b) calculated that the supply of Na⁺ to the leaf would average about 17 nmol m⁻² plasma membrane s⁻¹, similar to values in the range reported for freshwater algae (see Table 2).

Fluxes have also been calculated from electrophysiological data. Volkov and Amtmann (2006) obtained values of 700 and 150 nmol m⁻² s⁻¹ (with an external concentration of 100 mM NaCl and -80 mV applied to the interior of the plasma membrane) of protoplasts of the glycophyte *A. thaliana* and the halophyte *Eutrema halophilum* (syn. *Thellungiella halophila*), respectively (values could vary with the NaCl concentration and applied voltages).

Fluxes in other systems

There have been other estimates of the magnitude of solute fluxes in plants – of K⁺ and of sugars. For example, K⁺ influx into protoplasts isolated from roots of *Zea mays* from a 1 mM K⁺ concentration was 99 nmol m⁻² s⁻¹ (Gronwald and Leonard, 1982), while the flux of K⁺ into guard cells of *Commelina communis* was estimated to be up to 1.9 μmol m⁻² s⁻¹ (Penny and Bowling, 1974). For the roots of rye, K⁺ flux across the epidermal cells (White and Lemtiri-Chlieh, 1995) was calculated to lie between 14 and 27 nmol m⁻² s⁻¹. By way of comparison, oocytes of the African clawed toad *Xenopus laevis*, which are used for heterologous expression and overexpression of membrane ion channels and transporters from different organisms, can provide the range of maximal potential fluxes via these membrane proteins. These artificial systems may give ion currents of around 10 μA (sometimes slightly higher) per oocyte for high-level expression, which is about 30 μmol m⁻² s⁻¹ (e.g. Volkov, 2015a).

Turning to non-electrolytes, estimated fluxes of sugars for plants range between 3 nmol m⁻² s⁻¹ and 157 μmol m⁻² s⁻¹. Although not charged and so different from Na⁺ or Cl⁻, the magnitude of the fluxes of sucrose does illustrate the capacity of vesicular transport in plants. Phloem transport of sugars ranges from 20 to 200 nmol m⁻² s⁻¹ (MacRobbie, 1971b), while a flux of sucrose through plasmodesmata in the nectaries of *Abutilon megapotamicum* var. *anviogatum* was calculated to be 157 μmol m⁻² s⁻¹, about three orders of magnitude greater than for other fluxes (Gunning and Hughes, 1976). Using data in Etxeberria *et al.* (2005b), we calculate that the transport rate of

sugar in *Citrus reticulata* is about 90 nmol m⁻² plasma membrane s⁻¹ over 24 h. This transport process, which was sufficient to support the rapid growth of citrus fruits from fruit set to ripening at a sucrose import rate of 2.5 nmol g⁻¹ d. wt s⁻¹, is within the range of Na⁺ uptake rates listed for halophytes in Table 1.

Conclusions

Membrane fluxes of Na⁺ in halophytes are commonly in the range 150–1000 nmol m⁻² s⁻¹. There are, however, instances of higher fluxes, such as that of K⁺ across stomata (1.9 μmol m⁻² s⁻¹), Na⁺ in *Limonium* salt glands (4 μmol m⁻² s⁻¹), Na⁺ and Cl⁻ fluxes in marine algae (up to 1.1 and 10 μmol m⁻² s⁻¹, respectively), sucrose fluxes in *A. megapotamicum* (157 μmol m⁻² s⁻¹) and fluxes across oocyte membranes under artificial conditions (up to 30 μmol m⁻² s⁻¹). The high value of 4 μmol m⁻² s⁻¹ that has been estimated for *Limonium* salt glands depends on an assessment of their morphology and judgement of the area of available membranes or contact zones (via membranes or plasmodesmata) between cells in the gland. A concerning feature of the *Limonium* estimates, which used cut shoots and leaf disks to measure fluxes, is the difference in secretion rates from those calculated from intact shoots of another species in the same genus. The flux of Na⁺ estimated by sequential rinsing of leaf tissues of *L. vulgare* growing in 200 mM NaCl by Rozema *et al.* (1981) was around 1 pmol per gland per hour, dramatically lower than the 230–390 pmol per gland per hour reported for glands on disks punched from detached leaves of *L. perezii* floated on 300 mM NaCl or detached whole leaves of *L. perezii* with their petioles in 300 mM NaCl (Faraday *et al.*, 1986). This suggests to us that ions were reaching the apoplast at higher concentrations than might be expected in an intact transpiring leaf and that this influenced secretion rates, raising doubts over the very high fluxes estimated and requiring additional assessments and checks. While we do not have an explanation for the very high symplastic fluxes via plasmodesmata in the *Abutilon* gland, the high fluxes across oocyte membranes are artificially created with a very high density of heterologously expressed membrane proteins. We conclude that fluxes of the order of 150–1000 nmol m⁻² s⁻¹ need to be accounted for in halophytes. The question is, could such fluxes occur through vesicular transport?

VESICULAR TRANSPORT IN PLANTS

Vesicle trafficking in cells involves both exo- and endocytosis (sometimes called pinocytosis), the first being the fusion of vesicles produced by the Golgi apparatus with the plasmalemma and the latter being the recovery of membrane material from the plasma membrane (Fig. 3). There is now strong evidence that both endo- and exocytosis occur in plants (Battey *et al.*, 1999). Early evidence of endocytosis came from the uptake of the membrane-impermeable dye, Lucifer Yellow, and of electron-dense tracers by plant protoplasts (see Gradmann and Robinson, 1989; Low and Chandra, 1994). More recently, the use of patch clamp to measure electric capacitance has allowed detection of vesicle fusions (see Homann and Tester, 1997; Thiel and Battey, 1998) and the uptake of nanoparticles

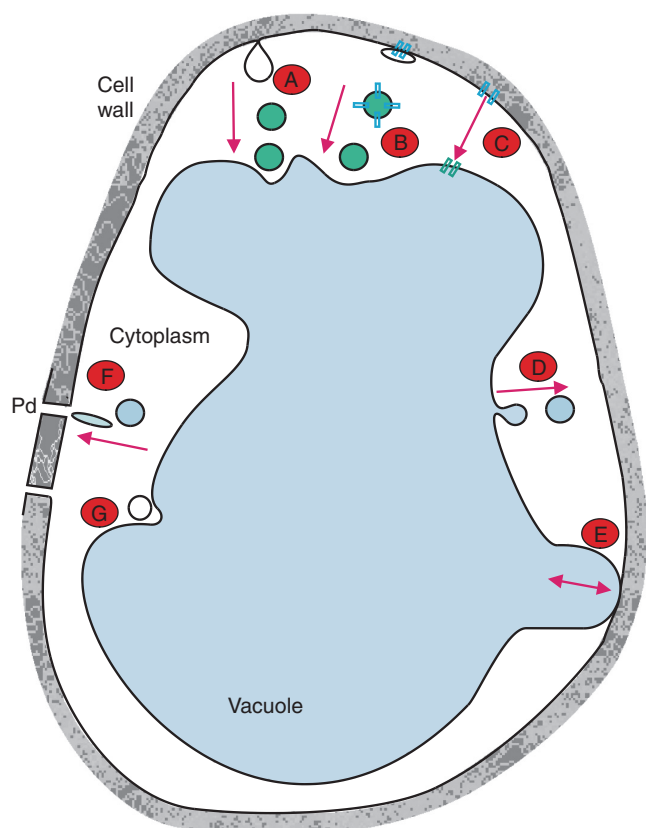


FIG. 3. Hypothetical vesicular transport of Na^+ ions in halophytes. The figure illustrates a hypothetical cell from a halophyte. The average Na^+ concentrations in the cytoplasm and vacuole are about 150–200 mM (Flowers *et al.*, 2015) and 500 mM (Flowers, 1985), respectively; the turgor pressure is low for a plant cell at about 0.2 MPa (Flowers *et al.*, 2015), due to high apoplastic ion concentrations (Harvey *et al.*, 1981). In the figure, we hypothesize six possible transport scenarios, none of which excludes combination with other possibilities. A, clathrin-dependent or clathrin-independent (more likely due to the size of the vesicles) endocytosis with vesicles either fusing with the vacuole or moving from cell to cell via plasmodesmata (F). B, a microdomain of the plasma membrane rich in transporters invaginates to form a vesicle that is loaded with Na^+ from the cell wall or cytoplasm; the fate of the vesicle is as in A. C, non-vesicular ion transport through transporters and channels. D, exocytosis of vesicles originating from the tonoplast allows export of ions (Echeverria, 2000; see also Mellander *et al.*, 2014 for possible mechanisms). E, junctional complexes (Vassililyev and Stepanova, 1990); in the words of Echeverria (2000), ‘Juxtaposition of membrane-bound carriers located at the tonoplast and the plasmalemma. These allow the direct transfer of metabolites to the apoplast’. F, vesicular transport through plasmodesmata (Wilson *et al.*, 2017). G, formation of a multivesicular body (MVB; see Balnokin *et al.*, 2007 and text above). pd, plasmodesma.

by plants has been established (reviewed in Anjum *et al.*, 2016), including gold nanoparticles in tomato and rice (Li *et al.*, 2016). Evidence for exocytosis has been gathered from studies of tissues such as pollen tubes and nectaries (Battay *et al.*, 1999; Grebnev *et al.*, 2017).

Although there is now strong evidence in favour of endo- and exocytosis in plants (see Echeverria, 2000), in 1980 Cram wrote in the abstract of his review on ‘Pinocytosis in plants’: ‘Thus it appears not only that pinocytosis does not occur, but also that it could not occur in plant cells.’ He (Cram, 1980) raised objections on the grounds of selectivity of uptake, depolarization of the membrane, turgor, energetics, the rate

of endocytosis in relation to measured rates of uptake and the density of putative binding sites. However, the energy required for plasmalemma recycling has been estimated to be a very small (0.06 %) fraction of the total energy required for biochemical synthetic processes in cells (Raven, 1987, appendix II) and sufficient for receptor-mediated endocytosis (RME), since the vesicles are small (Saxton and Breidenbach, 1988). As far as turgor is concerned, Gradmann and Robinson (1989) concluded that endocytotic vesicles could only form where turgor was less than about 0.1 MPa. More recently, however, a thermodynamic analysis has shown that endocytosis is feasible if the tension in the membrane is low and vesicles are flat with little volume as they detach from the plasma membrane (Fricke *et al.*, 2000). The thermodynamic approach sets a biophysical framework for the processes and has been applied assuming membranes have a homogeneous structure and surface tension. However, adjustments may be necessary for real biological membranes, where, for example, micro-domains with localized ion channels have been described for plasma membranes (e.g. Mongrand *et al.*, 2004; Sutter *et al.*, 2006; reviewed in Malinsky *et al.*, 2013) demonstrating their complex structure in plant cells. More recent modelling suggests that vesiculation can occur even under high membrane tensions (Hassingier *et al.*, 2017). Furthermore, it is reasonable to consider that all the membrane enzymes and ion channels in plants operate at the pressures found in their cells (approx. 0.5 MPa), so there could be a variety of unknown effects at the molecular level with extra weak molecular forces involved (e.g. Van der Waals forces and London forces), which may outweigh the turgor pressure. The forces may act non-linearly and could even be negative depending on the distance between interacting molecules (e.g. Israelachvili, 2011). So, while the thermodynamic approach remains valid for the whole cells and larger systems, it may need refining for individual parts of systems and for their membrane domains. A further complication lies in the fact that the physical properties of membranes and their behaviour are influenced by the underlying cytoskeleton: myosin-I controls cell membrane tension for living animal epithelial cells and fibroblasts (e.g. Nambiar *et al.*, 2009), while for fungal cells of *Aspergillus nidulans* (which has cell walls and high turgor pressures), a mutated version of *myoA* (the same class I myosin gene) constitutively activated endocytosis (Yamashita and May, 1998). Myosin from *Chara corallina* moves actin filaments an order of magnitude faster than myosin from skeletal muscle (Ito *et al.*, 2007). Cytoplasmic streaming, which is driven by myosin motors (Kurth *et al.*, 2017), was thought to be a possible limiting factor in the movement of Na^+ (and Cl^-) across the roots of *S. maritima* (Yeo and Flowers, 1986).

In spite of early scepticism, there is now good evidence for endocytosis in plants and its role in solute transport (Echeverria, 2000). For example, a sucrose-induced endocytosis of sucrose into the vacuoles of cultured cells of sycamore and juice cells of *Citrus* (Etcheberria *et al.*, 2005a) has been clearly demonstrated. Where sucrose or other sugars are transported, they are delivered in high concentration, typically 250 mM or more, to the apoplast of the receiving cells via unloading from the phloem. For citrus juice cells, uptake of sucrose into the cells does occur via transporters as well as by relatively large endocytotic vesicles (up to 7 μm in diameter) that form at the plasma membrane (Etcheberria *et al.*, 2005b, 2007, 2009). Transport via

an endocytic system allows uptake to bypass both the plasma-membrane and tonoplast transporters and to incorporate sucrose directly from the apoplast to the vacuole (Etxeberria *et al.*, 2005a, b). The process has been followed by labelling the outside lamella of the plasma membrane with vital fluorescent stains, transport of which equates quantitatively with the formation and transport of sucrose into vacuoles. In protoplasts of *Citrus aurantifolia* (sweet lime), fluid-phase endocytosis transports sucrose from the medium to the vacuoles in vesicle populations of two sizes, one about 1 μm in diameter and the other 1–7 μm in diameter (Etxeberria *et al.*, 2009). Labelling with fluorescent dyes suggested that the larger vesicles were the primary transport system for fluid-phase endocytosis (Etxeberria *et al.*, 2012) and independent of the endosome.

Although lagging behind information available for animals, much is now known about the endosome in plants and about the mechanism of endocytosis (Valencia *et al.*, 2016). During endocytosis, parts of the plasma membrane enclose extracellular substances in vesicles that then fuse with membranes originating from the Golgi apparatus, the endosome. This endosome is comprised of the *trans*-Golgi network (TGN), which as an early endosome receives the contents of vesicles containing a cargo; these early endosomes mature to multivesicular endosomes that can fuse with lysosomes and vacuoles (Valencia *et al.*, 2016). The process of trafficking (most often investigated in terms of protein as the cargo; see, for example, Uemura, 2016) involves a protein coat (Hwang and Robinson, 2009), which prevents fusion of the vesicle membrane with other membranes: coated pits have long been known to be present in plant cells (see Robinson and Hillmer, 1990). A key feature of the transport process is the presence of micro-domains in the plasma membrane surface that are able to recognize and concentrate the cargo. It is the cargo that is postulated to bind to proteins that in turn binds to other proteins that cause the membrane to form a pit and then, in a complex process, a vesicle. These pits may be coated by a protein, clathrin (clathrin-mediated endocytosis, CME; see also Hassinger *et al.*, 2017), or vesiculation may be clathrin independent (Fan *et al.*, 2015). In plants, there is evidence that flotillins (membrane proteins that function in endocytosis; see Otto and Nichols, 2011; Danek *et al.*, 2016) are involved in clathrin-independent endocytosis (CIE). These vesicles involved in CIE have a diameter of 65–200 nm in yeast (median value 115 nm; Cucu *et al.*, 2017) and 100 nm in *A. thaliana*; clathrin-coated vesicles appear to be of similar size (Beevers, 1996; Li *et al.*, 2012; Baral *et al.*, 2015). Danek *et al.* (2016) speculated that for the same cargo, CME may be constitutive while CIE may be induced by stress or signalling. The large (up to 7 μm in diameter) clathrin-independent vesicles seen in sweet lime are probably the primary pathway for fluid-phase endocytosis (Etxeberria *et al.*, 2012). Interestingly, it is CIE, some kinds of which involve flotillins, whose function is associated with membrane sterols (Danek *et al.*, 2016). Membrane sterols are normally a small component of plant membranes, but are relatively enriched in the halophyte *S. maritima* (Leach *et al.*, 1990). Chloroplast detergent-resistant membrane fractions from the halophyte *Salicornia perennans* contained a higher concentration of sterols than the less tolerant *Artemisia santonicum* (Nesterov *et al.*, 2017).

Endocytosis requires exocytosis to recycle the membrane material and water that accompanies the solutes (Robinson

and Hillmer, 1990; Thiel and Battey, 1998; Battey *et al.*, 1999; Etxeberria *et al.*, 2013). In beet storage cells, rapid endocytosis was balanced by the formation of vacuole-derived vesicles which transiently counter-balanced the addition of surplus endocytic membrane (Etxeberria *et al.*, 2013). However, as far as we are aware, little is known of how the lipid composition is modified or of the role of flippases (see, for example, Sahu and Gummadi, 2008; Lopez-Marques *et al.*, 2012; Underwood *et al.*, 2017) and how they work under turgor pressure. Exocytosis is also the mechanism by which plants secrete a range of substance – from nectar, lipophilic materials, resins, gum and the cellulose and other polysaccharides for cell wall synthesis (e.g. Yoneda and Doering, 2006) to salts, for which there is, at least, circumstantial evidence for the role of vesicular transport (Fahn, 1988). Mechanical energy is released when vesicles at high pressure in the cytoplasm fuse with the membrane at the lower pressure of the cell wall, so that turgor favours exocytosis (but not endocytosis; Fricke *et al.*, 2000).

Might ions be transported in vesicles?

Although theoretically possible, the question remains as to whether plants and particularly halophytes use vesicular transport for the movement of Na^+ and Cl^- into and out of cells. Bennett (1956) hypothesized that ‘membrane flow may be an important part of a type of active transport mechanism carrying particles, including ions, along, within, into and out of cells’, a concept promulgated for plants by Sutcliffe (1962). Evidence for and against the presence of micro-vesicles and their involvement in ion uptake as well as the radial transport of ions was reviewed by Baker and Hall (1973), at a time when kinetic analysis of influx and efflux in plants was much in vogue. They (Baker and Hall, 1973) emphasized the notion of a dynamic cell, where endocytotic vesicles, which incorporated membrane-bound carriers, formed at the plasma membrane. Once formed, the vesicles would release ions either in the cytoplasm by breakdown of the membrane or in a larger vacuole by fusion with the tonoplast. An immediate question that arose was identifying any vesicle involved in transport from other small vacuoles. Hall (1970) had earlier provided evidence of the presence of small (average diameter of 130 nm) vesicles in the meristematic and cortical cells of maize roots that stained heavily for ATPase and were associated with the plasma membrane. Treating the roots with KCl increased the proportion of cell sections with vesicles. These data were taken as providing significant evidence in favour of the role of endocytosis in the uptake of K^+ into roots. Evidence in favour of vesicular transport of ions now comes from a variety of experiments. (see Echeverria, 2000).

Endocytosis has been implicated in the uptake of toxic elements (uranyl and lead) into oats, barley and maize roots (Wheeler and Hanchey, 1971; Wheeler *et al.*, 1972; Malone *et al.*, 1974) and in the glycophyte *Raphanus sativus*, where treatment with uranyl acetate induced vesiculation and uranyl phosphate was visible in both the apoplast and vesicles (Romanenko *et al.*, 1986). NaCl-induced vesiculation was seen in the halophyte *Salicornia europaea* and the glycophytes *Hordeum vulgare* and *Phaseolus vulgaris* (Nassery and Jones, 1976a). Treatment of samples containing Cl^- with

a soluble salt of Ag leads to precipitation of AgCl, which has very low solubility and is electron dense, so visible under an electron microscope. Electron-dense deposits of Cl as its Ag salt have been demonstrated in the halophytes *Limonium vulgare*, *Salicornia pacifica*, *Tamarix aphylla*, *Aegiceras corniculata* and *Petrosimonia triandra* (Ziegler and Lüttge, 1967; Hess et al., 1975; Campbell and Thomson, 1975; Van Steveninck et al., 1976b; Kurkova et al., 1992) and the glycophytes *Pinus radiata*, *Phaseolus vulgaris* and *Hordeum vulgare* (Nassery and Jones, 1976b; Van Steveninck et al., 1976a, b; Foster and Sands, 1977). Support has also come from the localization of La^{3+} in the bladders of *Atriplex halimus*, where, after submergence of excised leaves in a solution of $\text{La}(\text{NO}_3)_3$, La^{3+} is present in small vesicles (Smaoui et al., 2011). While the presence of electron-dense deposits of AgCl argues in favour of the role of vesicles in the transport of Cl^- , it was clear from the early work of Harvey et al. (1976) that there is a significant loss of Cl^- during preparation of tissue for electron microscopy. So, localization, especially in small vesicles, could be compromised and remains a technical challenge, especially as side-scattering of electrons means the minimal area that can be resolved is $>100 \text{ nm}^2$ (see Ralle and Lutsenko, 2009). Losses of Cl^- can be reduced by the use of freeze substitution, and precipitates of AgCl have been seen in the cytoplasm of the halophyte *Suaeda maritima* prepared using this procedure (Harvey et al., 1979), although with this technical improvement the resolution in unstained tissues was insufficient to localize Cl to small vesicles. Overall, however, the presence of Ag deposits in vesicles is supportive of their role in ion transport.

In studies of the halophyte *Suaeda altissima*, Balnokin and colleagues (Kurkova et al., 1992; Kurkova and Balnokin, 1994) suggested that vesicles they could see in electron micrographs were composed of more than one membrane and led to the view (see Balnokin et al., 2007) that these structures were composed of plasmalemma and tonoplast. AgCl was identifiable in Ag-treated material, but could not be unequivocally located to the large vesicles seen in glutaraldehyde-fixed material. These vesicles, which were 2–3 μm in diameter, are not consistent with the endo- and exocytotic vesicles described above, but could be examples of multivesicular bodies, (MVBs) about which much has been discovered over recent years (Cui et al., 2016; Schoneberg et al., 2017; Wang et al., 2017), including a role in response to salt stress (Wang et al., 2015a). The evidence from *S. altissima* suggests that these MVBs are involved in Cl^- transport, although the evidence is again compromised by the potential movement of Cl^- during fixation. However, Shuvalov et al. (2015) have provided evidence that a Cl^-/H^+ antiporter is associated with a Golgi membrane fraction prepared from the roots of *S. altissima*.

Attempts have also been made to use the electron opacity of Rb^+ in order to identify its localization within cells as it is accumulated by plants and has been used as a tracer for K^+ . Following exposure of roots of *S. maritima* to RbCl , electron-dense deposits can be seen in the central vacuoles (Hall et al., 1974). However, significant quantities of Rb are lost during conventional fixation for microscopy (Hall et al., 1974), raising questions about the quality of the evidence for endocytosis in K^+ (Rb^+) uptake. Nevertheless, the presence of Rb-containing vesicles is suggestive of a role in ion transport. Further evidence in favour of vesicular transport in halophytes has been obtained from the study of salt glands.

The salt glands of species of the genus *Tamarix* consist of eight cells: six secretory cells and two basal cells (Thomson and Liu, 1967; Bosabalidis and Thomson, 1984; Fig. 2). Secretion, which takes place through four secretory pores (Feng et al., 2014), is greater during the day than at night (Imada et al., 2012) and reflects the salt concentration around the roots (Thomson et al., 1969; Sookbirsingh et al., 2010). Circumstantial evidence in favour of the involvement of vesicles in transport (Table 3) comes from electron microscopic investigations of *Tamarix aphylla* that found micro-vesicles associated with the plasma membrane (Thomson and Liu, 1967; Thomson et al., 1988; Wilson et al., 2017). After exposure of cut twigs to RbSO_4 , these vesicles contained electron-dense material (Thomson et al., 1969). Faraday and Thomson (1986b) looked at the anatomy and ultrastructure of 11 species and six genera of Plumbaginaceae. They noted that ‘While microvacuoles and vesicles were often present, they were not prominent, and there was no apparent correlation between their presence or absence and whether the glands were secreting or non-secreting’. They (Faraday and Thomson, 1986b) went on to conclude that the absence of numerous vesicles in the glands of *Limonium perezii* was evidence of a trans-membrane rather than a vesicle-mediated pathway. Nevertheless, there are genes associated with vesicle formation that are upregulated in *L. bicolor* when plants are exposed to salinity (Yuan et al., 2015).

As well as the species listed in Table 3a, there have been many other species (Table 3b) where vesicles have been described, but where plants were grown on garden soil and in some cases where the ends of cut shoots were inserted into salt solutions much higher than might be expected in the xylem. For example, Campbell and Thomson (1976b) induced salt secretion in glands of *Frankenia grandifolia* by immersing cut stems into NaCl (513 mM). Secretion began 4 h after treatment, and after 15 h the authors noted micro-vesicles scattered through the cytoplasm, which they speculated arose from the endoplasmic reticulum (ER). They (Campbell and Thomson, 1976b) suggested ‘transport components, perhaps synthesized in the ER, might feasibly be functional while being transferred to the plasma membrane in microvesicular form’. The treatment, however, may have supplied the leaf apoplast with a much higher salt concentration ($>500 \text{ mM}$) than that normally experienced, as all plants, including halophytes, prevent a significant proportion of the ions present in the external solution from reaching the xylem (Munns, 2005). For example, in the halophyte *S. maritima*, the mean Na^+ concentration in the xylem of plants growing in 200 mM NaCl was just 66 mM (Clipson and Flowers, 1987). The data from the experiments listed in Table 3b have not been included in our analysis.

In addition to the microscopic evidence for the presence of vesicles in plant cells, there is also physiological evidence consistent with vesicular transport. MacRobbie (1971a) reviewed analyses of influx and efflux kinetics of ions in algae and concluded that for *Nitella translucens*, an alga with giant cells, there were within the cytoplasmic and vacuolar phases rapid entry components that were consistent with movement of ions in vesicles. Subsequently, Lazof and Cheeseman (1986) provided kinetic evidence in the roots of the halophyte *Spergularia marina* for a small (1 %) rapidly exchanging compartment of the cytoplasm, which they could not identify, but would be consistent with a vesicular pool (evidence in the literature which

they discussed in some detail). In a later review, MacRobbie (1999) suggested it would be timely to reinvestigate the transfer of Cl⁻ and Br⁻ between the cytoplasm and vacuole of giant algal cells using methods that would test whether vesicular transport is involved in this transport process and in the rapid movement of K⁺ and sucrose in stomatal closure as well as the rapid movement of water in the leaf of species of *Mimosa*. There is now evidence for endocytosis of K⁺ in guard cells of *Vicia faba*, where turgor pressures are high, obtained using a styryl dye, FM4-64 (Meckel et al., 2004; see also below). In the future, the use of FM dyes (lipophilic styryl dyes that bind to membranes; see, for example, Bolte et al., 2004; Malinska et al., 2014; Rigal et al., 2015) and other fluorescent probes may aid localization of vesicular transport of ions; for example, the use of Di-4-ANEPPDHQ (Zhao et al., 2015) and of filipin (a sterol-specific marker; Klima and Foissner, 2008) to label membrane domains.

Evidence from molecular biology

There is molecular evidence pointing towards a role for vesicular transport in the tolerance of plants to salt (reviewed in Baral et al., 2015).

- (1) In *A. thaliana*, overexpression of a gene, *AtRabG3e*, that regulates vesicle trafficking increased vesicular endocytosis and raised Na⁺ contents and salt tolerance in transgenic plants compared with control plants (Mazel et al., 2004): the authors suggested that salinity tolerance was associated with an increase in vacuolar volume and the ‘process requires delivery of membrane material to the vacuole by an intracellular vesicle trafficking system, which is in line with the predicted function of *AtRabG3e*’. Salt stress (150 mM NaCl) induced the production of reactive oxygen species (ROS) and increased endocytosis (monitored with FM dyes) in the root tips of *A. thaliana*, leading to the suggestion that ROS act as a signal in the response to salt (Leshem et al., 2007). In separate experiments, treatment of roots of *A. thaliana* with high (200 mM) salt concentrations resulted in accumulation of Na⁺ in small vesicles (Hamaji et al., 2009). However, whether this is an adaptive or a pathological response is not clear, as the treatment with 200 mM NaCl is likely to be lethal for *A. thaliana* (Inan et al., 2004, Attia et al., 2008).
- (2) In habanero pepper exposed to NaCl, Na⁺ was associated with vacuoles and small intracellular compartment in a relatively tolerant genotype, whereas this was not so in a relatively salt-sensitive genotype (Bojorquez-Quintal et al., 2014).
- (3) The NHX antiporter genes *AtNHX5* and *AtNHX6* are located in the endosome (see Wang et al., 2015b) and expressing *AtNHX5* in transgenic soybean raised its tolerance to salt compared with wild-type plants and increased the concentration of Na⁺ in leaves and roots, suggesting a role for endosomal transport in the efficient movement of Na⁺ (Wu et al., 2016).
- (4) Annexins (widely conserved Ca²⁺- and phospholipid-binding proteins) are thought to have a role in membrane trafficking as well as in exocytosis due to their ability to bind to membranes and position membrane structures in a Ca-dependent way (see the review by Konopka-Postupolska and Clark, 2017). Since annexins also appear to play a role in stress tolerance in plants through ROS-induced Ca²⁺ signalling (see Shabala et al., 2015; Wilkins et al., 2016), there is a possible link between salt stress, the generation of ROS in the apoplast by NADPH oxidases and vesicular transport, perhaps involving MVBs, where annexins may also play a role (Konopka-Postupolska and Clark, 2017). Such a link may prove to be part of the explanation of the long-established importance of Ca²⁺ in salt tolerance (Greenway and Munns, 1980). There is also molecular evidence for the trafficking of plasma membrane transporters (Fan et al., 2015; see Valencia et al., 2016), of their clustering in membrane micro-domains (Fan et al., 2015) and regulation of alkali metal transporters through their recycling from the plasma membrane in yeasts, plants and mammals (Miguel Mulet et al., 2013).
- (5) The *A. thaliana* flotillin genes (*Aflot*) are induced under salt stress, and *Aflot2* binds to a cation hydrogen exchanger (CHX), CPA2, suggesting the flotillins may be involved in the uptake of monovalent cations (see Danek et al., 2016 for further discussion).
- (6) Evidence has been presented supporting the view that K⁺ channels are added and removed in clusters to and from the plasma membrane of isolated guard cell protoplasts of *V. faba* during pressure-driven volume changes (Hurst et al., 2004) and that the hormone abscisic acid initiates the endocytosis of the K⁺ channel KAT1 in *A. thaliana* (Sutter et al., 2007).
- (7) Other transporters have also been shown to be located in vesicular membranes. For example, the eight NHX transporters found in *A. thaliana* are divided between vacuolar, endosomal and plasma membranes (see Qiu, 2016), with *AtNHX5* and *AtNHX6* being localized in the Golgi and TGN, where they regulate endosomal pH (Bassil et al., 2011a, b; Reguera et al., 2015) as well as playing a role in K⁺ transport (see Wang et al., 2015b).
- (8) The cation–chloride co-transporters, from both *A. thaliana* and *Vitis vinifera* have also been shown located to the Golgi and TGN with a potential role in Na⁺ and Cl⁻ transport and salt tolerance (Henderson et al., 2015).

Membrane fluxes in halophytes

While molecular mechanisms for salt excretion through salt bladders can be postulated based on transporters alone (Shabala et al., 2014), recent reviews of multicellular salt glands have included a role for vesicular cell to cell transport followed by exocytosis onto leaves (Feng et al., 2014; Zouhaier et al., 2015; Yuan et al., 2016a; Wilson et al., 2017). Whether or not vesicular transport of ions is a feasible means of transport depends, at least in part, on the rate at which vesicles fuse with the tonoplast (for internal accumulation) or the plasma membrane (for secretion), on the concentration of ions in vesicles (see Table 4) and on movement of vesicles. Estimates for rates of movement of vesicles vary widely (three orders of magnitude) in different organisms. For yeasts, vesicles move at 150 nm s⁻¹ (Toshima et al., 2006), while in Characean algae, the rate of cytoplasmic streaming is 50 μm s⁻¹ (Tominaga et al., 2013). The rate of movement of clathrin-coated vesicles in yeast has

TABLE 4. Diameters of vesicles reported from eukaryotic cells

Vesicle type	Vesicle diameter (nm)	Characteristics		References
Endocytotic	80	Time to internalize an area equivalent to plasma membrane area (min)	20–115	See Steer (1988)
Exocytotic	70–400	Productivity/unit volume dictyosome ($\mu\text{m}^2 \mu\text{m}^{-3} \text{min}^{-1}$)	0.65–3.6	See Steer (1988)
Exocytotic	80–150	Arrival rate number s^{-1}	100	See Thiel and Battey (1998)
Exocytotic (yeast)	100	Fusions min^{-1} during growth	18	Cucu <i>et al.</i> (2017)
Endocytotic (Arabidopsis)	30–100	Clathrin mediated		Beevers (1996); Dhonukshe <i>et al.</i> (2007) Baral <i>et al.</i> (2015)
Endocytotic (glucose uptake)	80–220	Clathrin independent		Bandmann and Homann (2012)
Endocytotic (nano beads)	20–1000	Clathrin independent		Bandmann <i>et al.</i> (2012)
Endocytotic	100	Clathrin independent, flotillin mediated		Li <i>et al.</i> (2012)

been estimated at $5 \mu\text{m s}^{-1}$ (Ehrlich *et al.*, 2004). So, finally, we evaluate whether rates of membrane turnover match the fluxes – and whether the fluxes recorded in halophytes in Table 2 could feasibly occur via vesicles.

Although there are few data on membrane fluxes for halophytes, there are examples that cover movement of ions across roots and leaves and excretion of ions from salt glands. In order to model fluxes, we have used data on vesicular sizes taken from the literature (Table 4), i.e. vesicles with a diameter of 100 nm arriving at the plasma membrane at rates of up to 100 s^{-1} . Because of their size, this vesicular transport may be clathrin independent and probably involves flotillins (cf Table 4).

In our first example, we evaluate transport in the halophyte *S. maritima*. The flux of Na^+ across the root has been estimated to be between 262 and $6250 \text{ nmol m}^{-2} \text{ s}^{-1}$ (see above). A vesicle of 100 nm diameter containing a concentration of 500 mM NaCl would contain $2.62 \times 10^{-10} \text{ nmol}$ of Na^+ , so a flux of $262 \text{ nmol m}^{-2} \text{ s}^{-1}$ would require 10^{12} vesicles $\text{m}^{-2} \text{ s}^{-1}$. This number of vesicles would have a surface area of 0.03 m^2 , so 32×10^{12} vesicles merging per second would replace the membrane area of 1 m^2 . For an average cell (diameter $28 \mu\text{m}$ and area about $2500 \mu\text{m}^2$) this would mean around 2500 vesicles fusing per second. Larger vesicles of 400 nm diameter containing a similar concentration of ions would require 40 vesicles per cell per second and, if the diameter was $1 \mu\text{m}$ (Balnokin *et al.*, 2007; Klima and Foissner, 2008; Etxeberria *et al.*, 2009; Table 4), just three vesicles every second. A similar argument could be made for the flux of approx. $1000 \text{ nmol m}^{-2} \text{ s}^{-1}$ that crosses into the xylem (Yeo and Flowers, 1986; see also above) where fusing nine $1 \mu\text{m}$ diameter vesicles containing 500 mM Na^+ every second over the area of an average xylem parenchyma cell would deliver a flux of $1000 \text{ nmol m}^{-2} \text{ s}^{-1}$ or $100 \text{ nmol g}^{-1} \text{ d. wt roots s}^{-1}$.

The flux for leaves of *S. maritima* is $17 \text{ nmol m}^{-2} \text{ s}^{-1}$, where mesophyll cells have an average radius of $27 \mu\text{m}$ (volume of $80 \times 10^{-15} \text{ m}^3$; see Flowers and Yeo, 1986). With a vesicle diameter of 100 nm containing a solution of 500 mM Na^+ (close to that in the vacuoles of plants growing in 340 mM NaCl (Harvey *et al.*, 1981), a flux of $17 \text{ nmol m}^{-2} \text{ s}^{-1}$ could be delivered when the surface area is replaced every 16 min. It appears entirely feasible that the flux of ions from the xylem to cell vacuole could occur within vesicles. Flowers and Yeo (1986) also calculated fluxes for different leaves, which were expanding at different rates (between 2 and $10 \text{ m}^3 \text{ m}^{-3} \text{ s}^{-1}$) and for cells of different sizes. The flux required to maintain an internal salt

concentration of 350 mM (for plants growing in 200 mM NaCl, the measured range was between 420 and 380 mM as a leaf doubled in weight) was $<20 \text{ nmol m}^{-2} \text{ s}^{-1}$, again well within the range of membrane fluxes reported in Table 2 and feasible as a vesicular flux. Vesicles, however, carry water as well as ions, but water transport is hard to evaluate.

When endocytosis occurs the solution would be sampled from the cell wall, not the external medium. In *Suaeda maritima*, this would be between 100 and 200 mol m^{-3} cell wall volume for plants growing in 200 and 340 mM NaCl (Flowers, 1985; Hajibagheri and Flowers, 1989). If the cell wall has a water content of about 35 % solute available space (Flowers, 1985), then this would represent approx. $300\text{--}500 \text{ mM Na}^+$. However, the concentration of ions in solution and not bound to fixed charges in the cell wall is not known. In our calculations of fluxes, we used a figure of 500 mM as the ion concentration entering the vesicle. Water will enter along with ions, but we argue it will also leave during the balancing of membrane area via exocytosis. However, while we make assumptions of net water movement, we do not know of the detailed water relations between vesicles and the surrounding cytoplasm. These will be a function of reflection coefficients and osmotic potential differences, but complicated by the pressure, which in a droplet is determined by radius and surface tension – so, for a 300 nm vesicle, this might mean an internal pressure of 9.6 MPa depending on water interaction with lipid membrane, well above that in the cell as a whole. Ions remaining in the apoplast will serve to reduce cellular turgor (see Yeo and Flowers, 1986), which although unknown for root cells is low in the leaves of *S. maritima*, i.e. between 0.05 and 0.3 MPa (Clipson *et al.*, 1985). High elasticity in the cell wall will also mitigate against a turgor rise, as would a continuous slow increase in volume – as seen in the leaves of halophytes (Flowers, 1985).

Other important factors that are intrinsically related to vesicular transport are the speed of vesicles in the cytoplasm and the molecular motors for any vesicular transport (specific types of actin–myosins or dynein–kinesins). Several types of myosins participate in vesicular transport (e.g. Buss *et al.*, 2001; Peremyslov *et al.*, 2013; Kurth *et al.*, 2017); kinesins running on tubulin microtubules could also be involved (e.g. reviewed in Endow *et al.*, 2010). Here there are still many questions about how the size and type/molecular surface of the vesicles are linked to the speed of vesicular transport via the cytoplasm and the particular molecular motors involved.

Evidently, a thicker layer of cytoplasm requires more time and energy for the transport, while the estimates for the speed of transport differ by orders of magnitude and need more study. Developing methods that allow manipulation of the speed of cytoplasmic streaming and intracellular flows (Mittasch *et al.*, 2018) could be a way to estimate quantitatively the role of vesicular transport and the particular roles of specific vesicles in net salt transport in halophytes.

CONCLUSIONS

We have recorded a considerable range for the fluxes of Na⁺ and Cl⁻ into and out of the organs of halophytes, with many values lying between 150 and 1000 nmol m⁻² s⁻¹. Although the number of estimates is small, we take confidence that this range is reasonable from the fact that the values are derived from examples where fluxes calculated on a mass basis lie within what might be described as a ‘normal range’. We have then shown that such fluxes could be accounted for by rates of membrane turnover calculated for endo- and exocytosis, although there are too many uncertainties in the calculations to inspire real confidence in vesicles as a means of transport. In Fig. 3, we summarize possible pathways for the transport of ions into and out of cells. Endocytosis could transfer ions from the cell wall to the vacuole (A in Fig. 3), and ion transporters and channels may function as part of the vesicular membrane (B) or independently of vesicles (C). Vacuolar contents may be transferred to the apoplast via exocytosis (D), through so-called junctional complexes (E) or move to neighbouring cells through plasmodesmata (F); finally there is the possibility of movement through multivesicular bodies (G).

A key question is whether vesicular transport can support the range of fluxes needed for transport of NaCl into and out of halophytes as they grow. As we have shown, many assumptions are currently needed to attempt to answer this question definitively at the molecular, cellular and tissue levels of organization. What is now required for analysis of transport and membrane properties is the use of fluorescent dyes in the tissues of halophytes. At the molecular level, rates of exocytosis might be measured using fluorescence (Luo *et al.*, 2016), while the role of the myosin–MyoB complex in cytoplasmic streaming and its association with ion transport in halophytes is worthy of investigation (Kurth *et al.*, 2017).

It is not clear whether trans-membrane fluxes operate in parallel or in series with vesicular transport, nor how selectivity might be maintained during vesicular transport. However, as pointed out by Etzeberria *et al.* (2005a) for sucrose, transport through both membrane proteins and vesicles could occur simultaneously. As we mentioned earlier, vesicular transport is another mechanism to generate selective ion fluxes. While the selectivity of ion transport via ion channels and transporters is determined by the molecular protein structure of the ion channels and transporters, for vesicular ion transport the selectivity of trapped ion cargo would depend on local ion concentrations near forming vesicles and on electric charges of vesicular membranes. Variation of K⁺/Na⁺ selectivity amongst halophytes is worthy of further examination as are the energetics of the two processes, molecular vs. vesicular transport. Vesicular transport is likely to be energetically cheap (0.06 %; see above and

Raven, 1987), while the energy costs of ion transport are high (30–50 % of a cell's energy; Volkov, 2015b). However, plants may not be limited by energy due to that available through photosynthesis. For halophytes, the energetics of transport and compartmentation remain uncertain (see discussion in Flowers and Colmer, 2008).

The underlying issue in terms of large fluxes of ions across cells is how cytosolic ion concentrations compatible with metabolism are maintained – issues recently discussed by Flowers *et al.* (2015) and Munns *et al.* (2016). Constraining potentially toxic ions in vesicles would avoid interactions between damaging ion concentrations and the metabolic machinery of the cytosol. It is difficult to know if such vesicles would ‘leak’ but, if they had similar properties to the plasma and tonoplast membranes, then this is unlikely: ions do not leak through lipid bilayers (Hauser *et al.*, 1972) but will cross membranes through channel proteins (e.g. Demidchik and Tester, 2002; Demidchik *et al.*, 2002). Membranes of the halophyte *S. maritima* have been shown to have a relatively high cholesterol content with low leakage of Na⁺ (Leach *et al.*, 1990). ‘Leaks’ occur through channels or transporters and so are subject to gradients in electrochemical potentials with mechanisms to control transport through the protein (and there is evidence that Na⁺-permeable channels are closed in the tonoplast in *S. maritima* (Maathuis *et al.*, 1992)). In summary, ions could accumulate to high concentration in vesicles, creating protected compartments that may be moved through the cell to fuse with the vacuole. Movement from the vacuole to the apoplast could involve the reverse process – loading of a protected compartment that then moves to fuse with the plasma membrane, releasing ions to the apoplast. A high ion concentration in the apoplast would reduce turgor to a value that would not prevent vesicular transport (cf. Gradmann and Robinson, 1989). Low turgor pressure can, and does, occur in halophytes (Clipson *et al.*, 1985), where ions accumulate in the cell walls – and has been seen as a means to regulate cell turgor (James *et al.*, 2006; see also discussion in Flowers *et al.*, 2015). As far as the cell wall is concerned, its structure should not limit the movement of ions, although charge on the walls will alter the concentration of soluble ions through Donnan equilibria (Briggs *et al.*, 1961); only the movement of substances with dimensions greater than about 5–20 nm would be likely to be constrained by the wall structure (Carpita *et al.*, 1979; Anjum *et al.*, 2016).

So, what should physiologists and breeders pursue in defining the key features of salt tolerance? Certainly, as suggested almost 20 years ago (MacRobbie, 1999), the role of vesicular transport in euhalophytes requires further investigation and, if proven to be an important aspect of salt tolerance, this would influence the choices of genes that might be transformed into plants in the quest for salt-tolerant crops.

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