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**Communication between the plasma membrane and tonoplast
is an emergent property of ion transport**

Running head: Emergent membrane transport interactions

Dear Editor,

A very substantial body of data now exists for ion transport at both the plasma membrane and the tonoplast of several plant cell models, notably for guard cells, root hairs and epidermal cells of several species to name a few (Grierson et al., 2014; Jezek and Blatt, 2017; Wang et al., 2019). This knowledge has its foundation in detailed electrophysiological and flux studies that have provided quantitative biophysical and kinetic information. Our understanding has expanded through molecular biology to include the genetic identities of many transporters that operate at both membranes. Among these studies, it is common to deconstruct the mechanics and genetics of transport and to characterize each transporter in isolation. In many cases, this work has included the cloning and heterologous expression of specific transport gene products, their analysis under voltage clamp and by radiotracer flux measurements.

Where such studies often struggle is in phenotypic analysis in vivo to associate a genetic lesion with a function for the transport gene of interest. We suggest that, in focusing on single gene products, the deconstructionist approach can lose sight of the unique feature of transport, namely its physiological integration and apparent communication with other transport processes in situ, what is often referred to as 'emergent' properties arising from transport interactions. This communication is particularly evident when the transporter in question moves charge across the membrane. Although less often appreciated, similar considerations apply to transport across serial membranes when these operate on common pools of

34 solutes within an enclosed and finite compartment. Both situations are common to
35 plants. Communication is especially important between serial membranes, for
36 example between the plasma membrane and tonoplast, when the consequences of
37 manipulating transport at one membrane affects the cytosolic contents and, hence,
38 transport across the other membrane. In these circumstances quantitative
39 mathematical modeling is frequently essential to gain true insight into the
40 communication between membranes and their transport processes.

41 Within a single membrane there exist connections between different
42 transporters that ensure their fundamental interdependency. Notably, voltage exerts
43 a dominant control on charge-carrying transport, acting both as a driving force for
44 transport and as a product of charge flux across the membrane (Jezek and Blatt,
45 2017). Because physical laws require that net charge flux across a membrane is
46 zero in the steady state, transport of each ionic species is necessarily joined to the
47 transport of all other ions that affect voltage across the same membrane. Only by
48 imposing the circuit of a voltage clamp is this interconnection between transporters
49 bypassed. In short, manipulations affecting charge flux through any one transporter
50 will necessarily impact on all other charge-carrying transporters in the same
51 membrane, often with unforeseen consequences. As one example, the
52 interdependency between H^+ -coupled K^+ transport and the primary H^+ -ATPase (Blatt
53 and Slayman, 1987; Gibrat et al., 1990; Maathuis and Sanders, 1994) was ultimately
54 a key factor in explaining why more than a decade of research failed to uncover a
55 hypothetical H^+/K^+ exchange ATPase in plants (Leonard and Hotchkiss, 1976),
56 proposed to operate in a manner analogous to the mammalian Na^+/K^+ -ATPase.

57 Similar considerations apply to communication between the plant plasma
58 membrane and tonoplast. Because these serial membranes operate through a
59 common and enclosed compartment - the cytosol - the transport activities of both are
60 connected through this single pool of solutes. Analogies may be drawn here to
61 transport interactions across the apoplast between cells, for example as highlighted
62 in the plant vasculature (Gajdanowicz et al., 2011) and fungal symbiosis (Dreyer et
63 al., 2019). Some caution in drawing such analogies is advisable however, as the
64 nature of apoplast, if semi-open, may moderate interactions between membranes
65 and thus precludes a direct comparison with the enclosed cytosolic compartment
66 within a cell.

67 We focus here on communication between the serial membranes of the
68 plasma membrane and tonoplast and how alterations in transport at one are a
69 predictable consequence of manipulations affecting transport across the other
70 membrane. The examples are for guard cells and the ensuing stomatal phenotypes,
71 but apply equally to transport in other plant cell types. We use two studies (Wang et
72 al., 2012; Wang et al., 2017) to address the mechanics of how Cl⁻ flux mediated by
73 the SLAC1 channel at the plasma membrane affects transport across the tonoplast.
74 The physiology of the SLAC1 Cl⁻ channel in guard cells ensures that it mediates Cl⁻
75 efflux for stomatal closure and, as expected, the *slac1* null mutation suppresses
76 stomatal closure and greatly slows stomatal kinetics. Surprisingly, the mutant also
77 greatly slows the kinetics of stomatal opening (Vahisalu et al., 2008; Wang et al.,
78 2012; Wang et al., 2017), an effect that we now know arises because the mutation
79 indirectly suppresses the activity of the K⁺ channels that mediate K⁺ uptake (Wang et
80 al., 2012; Wang et al., 2017).

81 How does the *slac1* mutation influence transport at the tonoplast? Direct
82 access to the vacuolar membrane in vivo is not practicable. However, it is possible to
83 assess the consequences for solute contents and to examine the underlying
84 mechanisms that can explain these phenomena through simulation. We used the
85 OnGuard platform (Chen et al., 2012; Hills et al., 2012; Wang et al., 2012) to explore
86 the connections between these membranes. OnGuard2 (freely available at
87 www.psrq.org.uk) incorporates all of the quantitative detail for transport and the
88 relevant metabolic activities in guard cells to reproduce the characteristics of solute
89 flux, stomatal aperture and conductance known in the literature, and it has yielded a
90 number of unexpected predictions, many now validated experimentally. For
91 comparison with guard cells of wild-type Arabidopsis in OnGuard2, simulations of the
92 *slac1* mutant were generated by setting to zero the ohmic (voltage-independent) Cl⁻
93 conductance and the dominant fraction of the voltage-gated Cl⁻ conductance which,
94 combined, normally represent the characteristics of SLAC1 (Wang et al., 2017). Negi
95 et al (2008) reported that the *slac1* mutant accumulates osmotically-active solutes,
96 not only K⁺ and Cl⁻ but also substantial amounts of organic anions. The simulations
97 carried out by Wang et al (2012; 2017) similarly yielded accumulations of K⁺, Cl⁻ and
98 organic anions, the latter subsumed as malate in the OnGuard platform.

99 It is not surprising that eliminating a major pathway for anion efflux should

100 result in its accumulation in the cytosol and vacuole through its build-up in the
101 cytosol and trans-inhibition of efflux across the tonoplast. However, a comparison of
102 the fluxes through each of the major tonoplast transporters is instructive. These data
103 are available in the Supplemental Figures of Wang et al (2012). The simulations
104 predict an early increase in daytime K^+ flux through the tonoplast TPK1 and FV K^+
105 channels (see Figure S7 of Wang et al (2012)), thereby accounting for the overall
106 accumulation of this cation in the guard cell as reported previously (Negi et al.,
107 2008); they also shows a daytime reversal of Cl^- flux through the tonoplast Cl^-
108 channels (VCl). Additional to these outputs, a detailed analysis here throws up a
109 number of other predictions highlighting the apparent communication between
110 plasma membrane and tonoplast. It predicts, counterintuitively, roughly 3-fold
111 increases in net Ca^{2+} transport by the tonoplast Ca^{2+} -ATPases (V Ca^{2+} -ATPase) and
112 the Ca^{2+} channels (V Ca_{in}), as is clearly evident in the Supplemental Figure S6 of
113 Wang et al (2012) and a substantial increase in the activity of both sets of
114 transporters (Figure 1). These changes arise from the enhanced Ca^{2+} influx with
115 plasma membrane hyperpolarization in the *slac1* mutant and stimulation of
116 endomembrane Ca^{2+} release and recycling (Wang et al., 2012). The juxtaposition of
117 these two Ca^{2+} fluxes accounts for the overall lower total Ca^{2+} levels in the vacuole
118 as well as the elevated cytosolic-free $[Ca^{2+}]_i$ ([Ca^{2+}]_i) reported both in simulation and
119 as validated through experimental measurements (Wang et al., 2012; Wang et al.,
120 2017).

121 OnGuard2 yields a number of other predictions which, although still to be
122 tested experimentally, gain credibility from the accuracy of simulations to date in
123 predicting experimental observations. Notable among these, in simulation the effect
124 of the *slac1* mutant is to increase the activity of the tonoplast FV, TPK and VCl
125 channels as would be resolved under voltage clamp (Figure 1). The effect on all
126 three currents is substantially greater than might be expected for the overall K^+ and
127 Cl^- fluxes [Supplemental Figure S7 of Wang et al (2012)] and is seemingly
128 counterintuitive. However, the effects on these transporters is a natural consequence
129 of changes in $[Ca^{2+}]_i$ and cytosolic pH: all three channels are $[Ca^{2+}]_i$ -sensitive and the
130 K^+ channels are also subject to the elevated cytosolic pH that is characteristic of the
131 *slac1* mutant (Allen and Sanders, 1996; Wang et al., 2012; Wang et al., 2017). In
132 short, the simulation predicts, in the *slac1* mutant, an enhanced *capacity* for K^+ and

133 Cl⁻ flux, even if this capacity is kinetically restricted by charge balance and the free-
134 running voltage across the tonoplast.

135 Each of these predictions, and other outputs of OnGuard2, highlight the
136 emergent properties of transport communication within and, especially, between the
137 membranes of the guard cells. This communication is natural consequence of a
138 system of non-linear biological processes that share substrates and products across
139 each membrane and within cellular compartments. It arises from the membrane
140 voltage that is shared between all charge-carrying transporters on any one
141 membrane, as well as the common pool of ionic substrates shared between the
142 membranes and enclosed by them. Between the plasma membrane and tonoplast, it
143 arises from the common pool of cytosolic solutes that contribute to transport across
144 both membranes.

145 We stress that there is nothing unusual to this network of interactive
146 communication or the component transport processes. However, the intrinsic non-
147 linearities in flux behavior of each transporter ensures that the consequences of
148 experimental manipulations are beyond intuitive understanding. Thus, in vivo the
149 consequence of manipulating a single transporter at a membrane is rarely (if ever)
150 restricted to this one process, the distributions of the transported species alone or, in
151 plants, solely to the target membrane. Distinguishing between the primary effects of
152 a mutation and 'off target' effects clearly benefits in these circumstances from
153 quantitative mathematical modelling.

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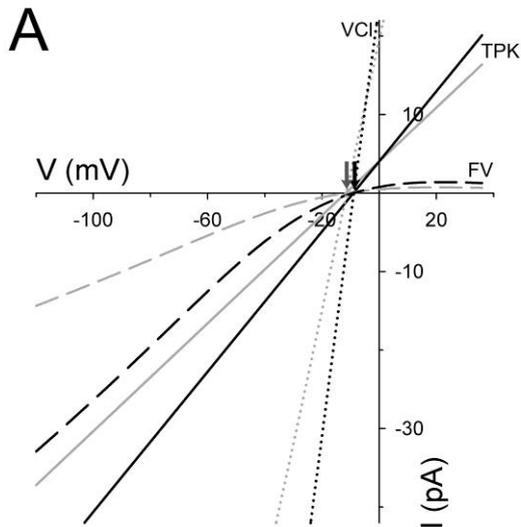
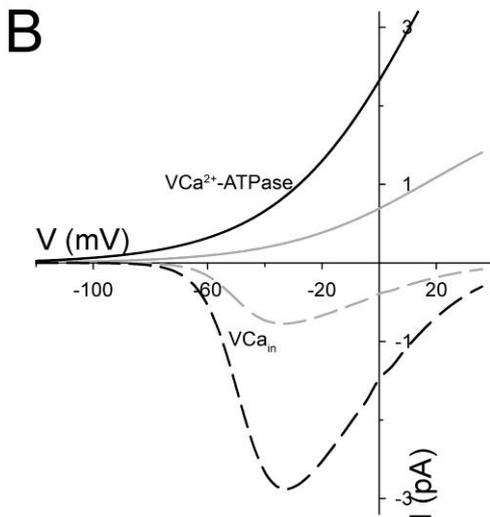


Figure 1. Current-voltage (IV) curves predicted for the major tonoplast K^+ and Cl^- channels and for the tonoplast Ca^{2+} ATPases and channels in guard cells of wild-type (grey lines) and the *slac1* mutant (black lines) Arabidopsis. Data were extracted from OnGuard2 simulations, as described, at 8 h into the daylight period, this timepoint corresponding to the maximum diurnal stomatal conductance in each case. The curves are plotted separately here for clarity.



(A) IV curves predicted for the TPK1 (solid lines) and FV (dashed lines) K^+ channels and for the VCl^- Cl^- channel (dotted lines). The free-running tonoplast voltages are indicated by the arrows (wild-type, grey; *slac1*, black). Note the substantial increase in the conductances (slopes) for the current of each channel type in the *slac1* mutant.

192 (B) IV curves predicted for the VCa_{in} Ca^{2+} channels (dashed lines) and for the VCa^{2+} -
 193 ATPase (solid lines). The identity of the VCa_{in} remains unknown and, in the OnGuard
 194 platform, the VCa^{2+} -ATPase subsumes the characteristics of all endomembrane
 195 Ca^{2+} -ATPases (Chen et al., 2012; Hills et al., 2012; Wang et al., 2012; Wang et al.,
 196 2017). Again, the analysis yields a substantial increase in the conductances and
 197 amplitudes of both currents in the *slac1* mutant.

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